XyMTEX: A Macro Package Set for Typesetting Chemical Structural Formulas

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August 16, 1996 (Version 1.01)

History:

Version 1.00 (December 1, 1993)
"XMTEX: A Macro Package for Typesetting Chemical Structural Formulas" (for LATEX2.09)

Version 1.01 (August 16, 1996)
Rewritten for LATEX 2_ε and renamed to "XMTEX: A Macro Package Set for Typesetting Chemical Structural Formulas"

Version 1.01 (the same) (August 30, 2004)
Re-processed at the release of XMTEX Version 4.01.

Contents

Chapter 1

Introduction

1.1 Backgrounds

1.1.1 Backgrounds for version 1.00 (1993)

The text formatter TEX developed by Knuth [1] is widely used in preparing manuscripts of scientific papers and in the typesetting processes of several scientific journals and books (for a recent example, see [2]). In particular, LATEX, a TEX macro package that was released by Lamport [3], has expanded the society of TEX users because of plainness.

Since the beginning of its history, TEX (LATEX) places special emphasis on mathematics typesetting. Hence, it has been accepted by scientists who have to write mathematic equations. In contrast, the TEX/LATEX typesetting is less popular in chemistry than in mathematics and other fields. One of the reasons is that there are few TEX/LATEX utilities for typesetting chemical structural diagrams.

Although LATEX provides us with a picture enviroment for drawing simple figures, its original commands are so primitive as to be directly applied to the drawing of structural formulas. Hence, the commands should be combined to produce more convenient macros.

Pioneering works by Haas and O'Kane [4] and by Ramek [5] have provided such macros that allow us to typeset structural formulas. The macros of the former approach are available in the public domain, being named ChemTEX. Although they are easier to use than the original picture environment of LATEX, they still have some items to be improved. The most inconvenient item is the incapability of accommodating 10 or more substituents. It stems from the fact that one argument is used to assign one substituent (or one object) in each of the macros of Haas-O'kane's approach. Note that the direct usage of arguments enables us only to assign 9 or less substituents, because a macro in TEX/LATEX is capable of taking 9 or less arguments.

For example, the `\steroid` macro reported for typesetting a steroid skeleton takes 9 arguments [4]:

```
\steroid{A1}{A2}{A3}{A4}{A5}{A6}{A7}{A8}{A9}
```

where Argument 1 (`A1`) can take 'D' (a second bond between positions 1 and 2), 'Q' (no action), or 'R^{11}' (a substituent on position 11 and the corresponding double bond); Argument 2 (`A2`) can take 'D' (a second bond between positions 3 and 4), 'Q' (no action), or 'R^{3}' (a substituent on position 3 and the corresponding double bond); Argument 3 (`A3`) can take 'Q' (no action), or 'R^{3}' (a substituent on position 3 and the corresponding single bond); and so on. Through the total statement of arguments, only six substituents are specified, while the skeleton have 20 or more substitution positions to be considered.

Moreover, the specification of the arguments is not systematic, since so many functions are included into the macro within the restriction of the direct usage of arguments.

1. One argument (Argument 2) specifies objects of two different categories *e.g.,* inner double bonds and outer double bonds.

2. Arguments 2 and 3 specify a substituent attaching to the same position (position 3).

3. It is difficult without a reference manual to differentiate between one argument for specifying bonds and another argument for specifying substituents.

4. The argument 'Q' is selected to show no modification because this character is hardly ever found in a chemical structure formula. However, the use of this character may become necessary in future. Such explicit description of 'no action' should be avoided.

As a result, the formats and contents of arguments are different from one argument to another and from one macro to another such that a typical TEX user, a secretary or a chemist author, may give up to memorize such macros. Hence, more systematic and convenient macros are desirable in order to spread the typesetting of chemical structures with TEX/LATEX.

The present package set[1] XymTEX involves convenient macros for typesetting chemical structural formulas [6]. These macros are based on techniques in which inner bonds, substituents and hetero-atoms on a skeleton are separately assigned without such limitation of numbers. The package set XymTEX[2] will be a more versatile tool if it is coupled with the macros which the author has released in a book [8].

1.1.2 Backgrounds for version 1.01 (1996)

The package set XymTEX (version 1.00, 1993) described in the preceding subsection has been depositted to NIFTY-Serve archives (FPRINT library No. 7) by the author[9] and to the CTAN by volunteers[10]. Although the style files of XymTEX has originally aimed at using under the LATEX2.09 system, they also work effectively under the LATEX 2$_\varepsilon$ system[11, 12] without any changes. Thus, what you have to do is to rewrite a top statement for LATEX2.09 such as

```
\documentstyle[epic,carom,hetarom]{article}
```

into the counterpart for LATEX 2$_\varepsilon$, *e.g.*,

```
\documentclass{article}
\usepackage{epic,carom,hetarom}
```

The purpose of the present version is the updating of XymTEX to meet the LATEX 2$_\varepsilon$ way of preparing packages (option style files). The following items have been revised or added for encouraging the XymTEX users to write articles of chemical fields.

1. Each of the old sty files of XymTEX has been rewritten into a dtx file, from which we have prepared a new sty file by using the docstrip utility of LATEX 2$_\varepsilon$. If you want to obtain the document of each source file, you may apply LATEX 2$_\varepsilon$ to the corresponding drv file, which has also been prepared from the dtx file by using the docstrip utility.

2. Macros for drawing chair-form cyclohexanes and for drawing adamantanes of an alternative type have been added.

3. Macros for drawing polymers have been added.

4. The package chemist.sty, which was originally prepared for [8], has been rewritten into a dtx file and added to XymTEX as a new component. This package enables us to use various functions such as

   (a) the numbering and cross-reference of chemical compounds and derivatives,

[1] LATEX 2$_\varepsilon$ uses the term 'package' to designate a file with .sty extention, while XymTEX version 1.00 have used the same term to indicate a set of sty files. In order to prevent confusion, we now use the term 'package set' to indicate a set of sty files and the term 'package' to designate each sty file.

[2]

(b) various arrows of fixed and flexible length for chemical equations,

(c) 'chem' version and chemical environments for describing chemical equations, and

(d) various box-preparing macros for chemical or general use.

1.2 The Name of the Package

The word 'chemistry' stems from an Arabian root 'alchemy', which is, in turn, considered to come from Greek, $\chi\upsilon\mu\epsilon\acute{\iota}\bar{\alpha}$. The XYM of the name XYMTEX is an uppercase form of $\chi\upsilon\mu$. This conforms to a rule of coinage, because the name TEX is also a word of Greek origin ($\tau\epsilon\chi$).

The pronuncialtion of XYMTEX is recommended to be 'khýmtekh', in which the 'kh' sound may be a Russian 'kh' or more simply an English 'k' and the symbol 'y' is expected to be pronounced like a German 'ü'.

The logo XYMTEX is defined as being either of the following statements. The second one has been adopted throughout the present manual.[3]

```
%%%XyMTeX Logo: Definition 1%%%
\newcount\TestCount
\def\XyM{\ifnum\fam=-1\relax\fam=0\relax\fi\TestCount=\fam%
X\kern-.30em\smash{\raise.50ex\hbox{$\fam\TestCount\Upsilon$}}%
\kern-.30em{M}}
\def\XyMTeX{\XyM\kern-.1em\TeX}

%%%XyMTeX Logo: Definition 2%%%
\def\UPSILON{\char'7}
\def\XyM{X\kern-.30em\smash{\raise.50ex\hbox{\UPSILON}}\kern-.30em{M}}
\def\XyMTeX{\XyM\kern-.1em\TeX}
```

When such a raised Greek letter as the 'Υ' is not available, XYMTEX may be referred to by typing '`XyMTeX`'.

1.3 Requirements

The macro package set XYMTEX runs within LATEX, since it is based on the picture environment of LATEX. It also requires an package file '`epic.sty`' developed by Podar [13], because the `\dottedline` command of epic is used in the macros. Since the main package file `xymtex.sty` is prepared for convenience, a manuscript file should begin with such a statement as follows:

```
\documentclass{article}
\usepackage{xymtex}
```

by which all of the package files of XYMTEX as well aa epic.sty are input for processing.

1.4 Compatibility

Although we have used LATEX 2_ε commands in the dtx files of XYMTEX system, we have carefully excluded them from the resulting sty files. This is the tentative policy of XYMTEX system to assure the compatibility to LATEX2.09 (the native mode).

For example, one or more sty files are crossloaded if necessary during the process of loading a sty file. Such loading has been carried out by using `\input` and `\@ifundefined` within the command system of

[3] Definition 2 is adopted in the manual because of simplicity. The methodology used in Definition 1 is applicable to a wide variety of cases in which font sizes have to be changed in the LATEX2.09 system. However, it is not recommended under the LATEX 2_ε system, since lower-level commands are used to select fonts.

LaTeX2.09, though the combination of the commands may be replaced by the `\RequirePackage` command of LaTeX 2_ε.

This tentative policy is considered to be effective to prevent some confusion provided by the coexistence of LaTeX2.09 and LaTeX 2_ε.

Chapter 2

The Construction of XymTEX

2.1 Overview

XymTEX contains package files listed in Table 2.1 along with their documents.[1] The package file 'chemstr.sty' is the basic file that is automatically read within any other package file of XymTEX. It contains macros for internal use, *e.g.*, common commands for bond-setting and atom-setting. The other package files contain macros for users.

Table 2.1: Style Files of XymTEX

package name	included functions
aliphat.sty	macros for drawing aliphatic compounds
carom.sty	macros for drawing vertical and horizontal types of carbocyclic compounds
lowcycle.sty	macros for drawing five-or-less-membered carbocyles.
ccycle.sty	macros for drawing bicyclic compounds etc.
hetarom.sty	macros for drawing vertical types of heterocyclic compounds
hetaromh.sty	macros for drawing horizontal types of heterocyclic compounds
hcycle.sty	macros for drawing pyranose and furanose derivatives
chemstr.sty	basic commands for atom- and bond-typesetting
locant.sty	commands for printing locant numeres
polymers.sty	commands for drawing polymers
xymtex.sty	a package for calling all package files
chemist.sty	commands for using 'chem' version and chemical environments

These files are designed to be packages for LaTeX 2$_\varepsilon$ as well as option style files for LaTeX2.09 (native mode). The complete list of the XymTEX commands is shown in Appendix A.

[1] Each package file (.sty file) has been generated from the corresponding dtx file by the docstrip utility of LaTeX 2$_\varepsilon$. The source file of each package can be generated by the LaTeX 2$_\varepsilon$ processing of the corresponding drv file. The source file of the present reference manual is 'xymtex.tex', the LaTeX 2$_\varepsilon$ processsing of which will read the tex files involved in the same directory.

2.2 General Conventions

2.2.1 User Commands for Specified Use and for General Use

XymTEX user commands are classified into two types, *i.e.*, commands for specified use and those for general use.

Specified user commands of XymTEX are used to draw a narrow range of structures. More precisely speaking, they are short-cut commands of general user commands with a specific bond pattern for drawing carbocycles or with a specific pattern of skeletal hetero-atoms for drawing heterocycles. They take such general forms as follows:

`\Sformb[OPT]{SUBSLIST}`	for drawing carbocycles
`\Sformd[BONDLIST]{SUBSLIST}`	for drawing heterocycles

where `\Sformb` and `\Sformd` may be approriate command names. These are selected from chemical names that represent the compound-group names to be typeset. In accord with LaTeX conventions, an argument in brackets is an option.

The command `\Sformb` typesets a carbocyclic compound with a specific bond pattern which may be altered by the optional argument OPT. The command `\Sformd` prints a heterocyclic compound with a specific atom pattern on its skeleton.[2]

For example, `\bzdrv[OPT]{SUBSLIST}` is a command for the specified use of drawing benzene derivatives, where the stem '`\bzdr` without a suffix 'v' is an abbreviation of 'benzene derivative'. The command `pyridinev[BONDLIST]{SUBSLIST}` is a command for drawing pyridine derivatives, in which the nitrogen atom on the pyridine ring is automatically typeset.

On the other hand, more elaborate commands for general use can be used within XymTEX. They are designed to have a variable set of skeletal heteroatoms in accord with our designation so that they cover a wide range of structures. They have general formats as follows.

`\Sforma[BONDLIST]{SUBSLIST}`	for drawing carbocycles
`\Sformc[BONDLIST]{ATOMLIST}{SUBSLIST}`	for drawing heterocycles

where `\Sforma` and `\Sformc` may be approriate command names.

The command `\Sforma` for general use generates a carbocyclic structure, in which its individual bonds can be independently altered by means of BONDLIST. The command `\Sformd` prints a heterocyclic compound so that individual atoms on its skeleton can be independently typeset through ATOMLIST.

For example, `\cyclohexanev[BONDLIST]{SUBSLIST}` is a command for the general use of drawing cyclohexane derivatives, by which six-membered carbocyles of any unsaturation level can be typeset. The command `sixheterov[BONDLIST]{ATOMLIST}{SUBSLIST}` is a command for drawing six-membered heterocyclic compounds, which may have any set of skeletal hetero-atoms (ATOMLIST) and any set of unsaturation (BONDLIST).

2.2.2 Suffix and Arguments

Most user commands of XymTEX are suffixed with 'v', 'vi', 'h' and 'hi'. The suffix 'v' means that the command prints a structural formula of vertical form. The suffix 'h' means that the command typesets a structural formula of horizontal form. When alternative orientations are possible, XymTEX commands are differentiated by an additional suffix 'i'.

The specification of each argument in a XymTEXcommand is based on list-treating macros [6]. Thus, items to be specified are listed sequentially with or without appropriate delimeters.

The argument SUBSLIST lists substituents with bonds. The argument {1==Cl;3D==O;...}, for example, means that position 1 takes a chlorine atom (Cl) through a single bond, position 3 takes an

[2] If we take a strictly systematic approach, the `\Sformd` should be designed to take an option argument OPT instead of the BONDLIST. However, a simple format of the OPT cannot be designed because heterocyclic compounds take a wide variety of bond patterns.

oxygen atom (O) through a double bond, and so on. Thus, a character string before every semicolon represents a mode of substitution, where a locant number with a bond modifier is separated from a substituent by means of a double equality symbol (==). Each bond modifier consists of one or two characters listed in Table 2.2. The diagrams below Table 2.2 illustrate these bond modifiers by using a cyclohexane skeleton (`\cyclohexanev`).

The optional argument OPT of `\Sformb` contains a string of one or two characters for giving a pattern of double bonds (*e.g.*, 'r' for a right-hand set of aromatic double bonds 'l' for a left-hand set of aromatic double bonds, and 'c' for an aromatic circle for the macro `\bzdrv`). Since the argument OPT is an option, a default set of bonds is used when omitted.

The optional argument BONDLIST contains a character string, each character of which is used for assigning a specific inner double bond (*e.g.*, 'a', 'b', ... for the double bonds of a given bond-numbering). Since the argument BONDLIST is an option, a default is used when omitted: the commands `\Sformd` and `\Sformc` (for drawing heterocycles) typeset default sets of bonds, while most `Sforma` commands (for drawing carbocycles) print fully saturated skeletons.

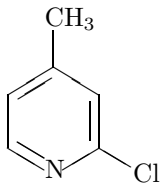
The argument ATOMLIST (*e.g.*, {1==O;4==O}) contains hetero atoms and their positions on the ring structure to be printed: this example argument produces a dioxane skeleton, when used in command `\sixheterov`.

2.2.3 Fonts

The character font used in each structral formula is `\normalfont` that is the default font of the LaTeX 2_ε text. For example, the statement

```
\pyridinevi{2==Cl;4==CH$_{3}$}
```

produces



Other fonts can be used by declaring the corresponding font selecting commands such as `\sffamily` and `\bfseries`. Thus, the code

```
{\sffamily \pyridinevi{2==Cl;4==CH$_{3}$}} \qquad
{\small\bfseries \pyridinevi{2==Cl;4==CH$_{3}$}}
```

produces the following structural formulas:

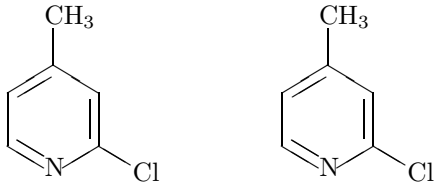
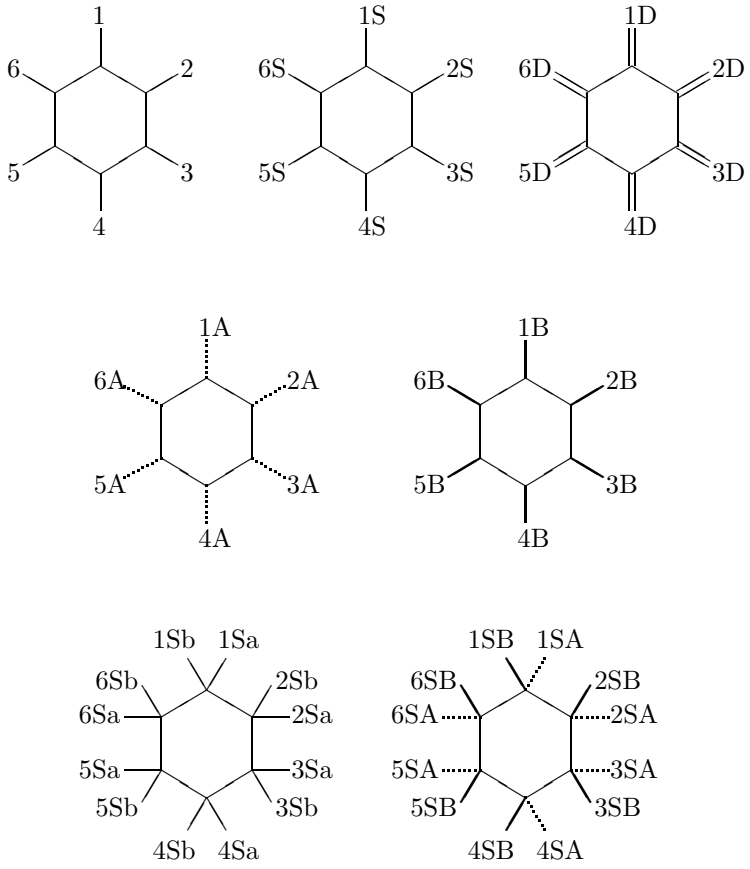


Table 2.2: Locant numbering and bond modifiers for SUBSLIST

Bond Modifiers	Printed structures
n or nS	exocyclic single bond at n-atom
nD	exocyclic double bond at n-atom
nA	alpha single bond at n-atom
nB	beta single bond at n-atom
nSa	alpha (not specified) single bond at n-atom
nSb	beta (not specified) single bond at n-atom
nSA	alpha single bond at n-atom (dotted line)
nSB	beta single bond at n-atom (boldface)



Chapter 3

Six-Membered Carbocycles

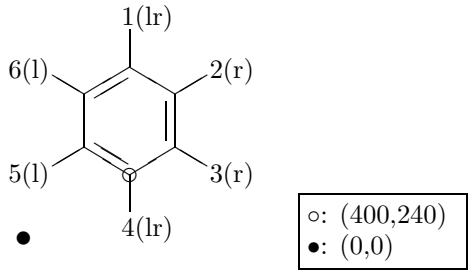
3.1 Drawing Benzene Derivatives

3.1.1 Vertical Forms of Benzene Derivatives

The macro `\bzdrv` is used to draw benzene derivatives of vertical type (carom.sty). The format of this command is as follows:

```
\bzdrv[OPT]{SUBSLIST}
```

The name and arguments of this command conform to the general conventions described in the preceding chapter. Thus, the suffix 'v' indicates that this macro produces a vertical-type structural formula. Locant numbers for designating substitution positions in the SUBSLIST are represented by the following diagram:



in which a character set in parentheses represent the handedness of each position. In accord with the default definitons of the macro `\bzdrv`, each of the right-handed positions (2 and 3) is designed to take only a right-handed substituent, while each of the left-handed positions (5 and 6) is to take only a left-handed substituent. Such positions (designated with the letter 'r' or 'l') are referred to as 'oriented' positions in this manual. In contrast, the top (and also the bottom) position of a benzene ring (designated with the string 'lr') can accommodate a substituent of both handedness. It is referred to as a 'double-sided' position in this manual. Although the default definition is to put a right-handed moiety, a left-handed substituent can be printed by means of the macro `\lmoiety`.

The symbols • and ○ in the diagram respectively represent the reference point and the inner origin of the macro. Since we select `\unitlength` to be equal to 0.1pt as a default value, the value 400, for example, corresponds to 40pt. These will be described in detail in Chapter 14.

The optional argument OPT specifying a bond pattern are shown in Table 3.1. Thereby, a wide variety of bond patterns (such as two patterns of benzene double bonds as well as an aromatic circle) can be depicted, as illustrated in Figure 3.1.

The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 6. For example, the statements,

Table 3.1: Argument OPT for commands \bzdrv and \bzdrh

Character	Printed structure
none or r	right-handed set of double bonds
l	left-handed set of double bonds
c	aromatic circle
p or pa	*p*-benzoquinone (A) (Oxygen atomes at 1,4-positions)
pb	*p*-benzoquinone (B) (Oxygen atomes at 2,5-positions)
pc	*p*-benzoquinone (C) (Oxygen atomes at 3,6-positions)
o or oa	*o*-benzoquinone (A) (Oxygen atomes at 1,2-positions)
ob	*o*-benzoquinone (B) (Oxygen atomes at 2,3-positions)
oc	*o*-benzoquinone (C) (Oxygen atomes at 3,4-positions)
od	*o*-benzoquinone (D) (Oxygen atomes at 4,5-positions)
oe	*o*-benzoquinone (E) (Oxygen atomes at 5,6-positions)
of	*o*-benzoquinone (F) (Oxygen atomes at 1,6-positions)

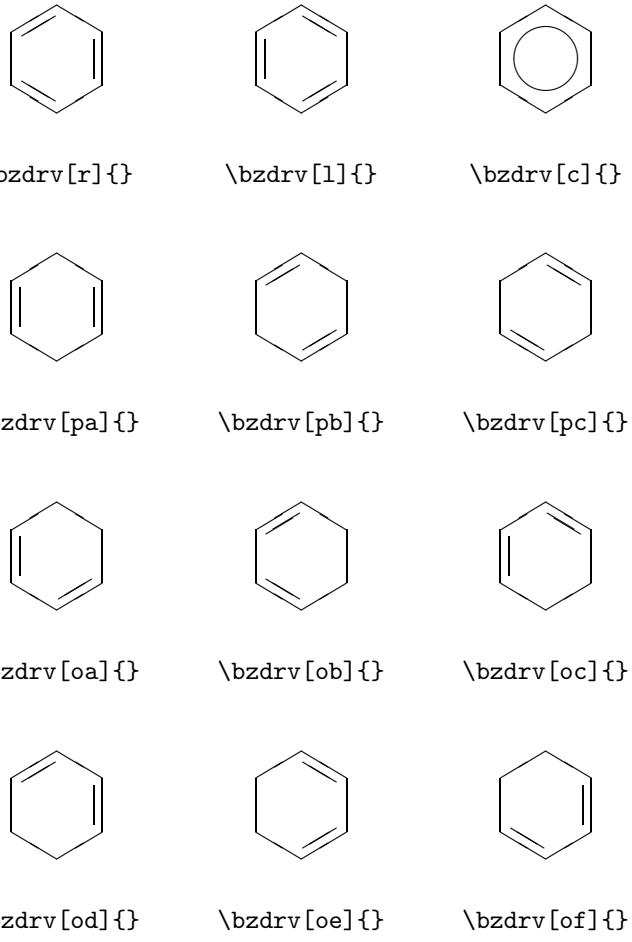
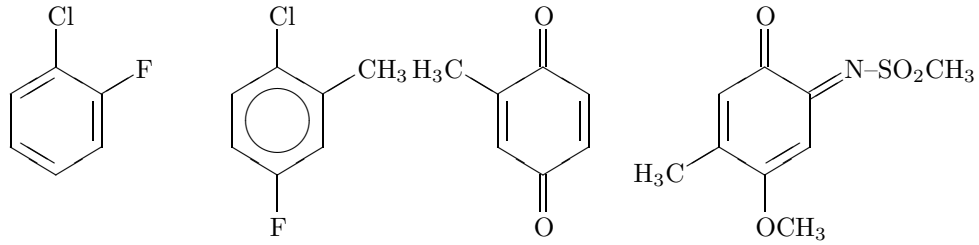


Figure 3.1: Endocylic bond patterns by the OPT argument

```
\bzdrv{1==Cl;2==F}
\bzdrv[c]{1==Cl;4==F;2==CH$_{3}$}\qquad
\bzdrv[pa]{1D==O;4D==O;6==H$_{3}$C}
\bzdrv[oa]{1D==O;2D==N--SO$_{2}$CH$_{3}$;4==OCH$_{3}$;5==H$_{3}$C}
```

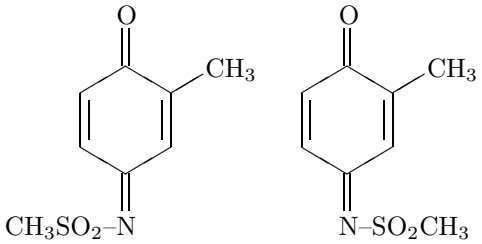
produce the following structures:



In order to designate the handedness of a substituent explicitly, you can use `\rmoiety` or `\lmoiety` commands. Thus, the statements,

```
\bzdrv[pa]{1D==O;4D==\lmoiety{CH$_{3}$SO$_{2}$--N};2==CH$_{3}$}
\bzdrv[pa]{1D==\rmoiety{O};4D==\rmoiety{N--SO$_{2}$CH$_{3}$};2==CH$_{3}$}
```

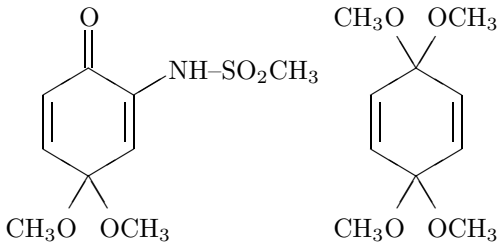
produce the following structures with left-handed and right-handed methanesulfonimido groups.



The macro `bzdrv` is used also to draw benzoquinone monoacetals and diacetals. The handedness of a substituent attached at such a tetrahedral position is determined in the light of chemical conventions. For example,

```
\bzdrv[pa]{1D==O;4Sb==CH$_{3}$O;4Sa==OCH$_{3}$;2==NH--SO$_{2}$CH$_{3}$}
\qquad \qquad
\bzdrv[pa]{1Sb==CH$_{3}$O;1Sa==OCH$_{3}$;4Sb==CH$_{3}$O;4Sa==OCH$_{3}$}
```

produce the following structures.

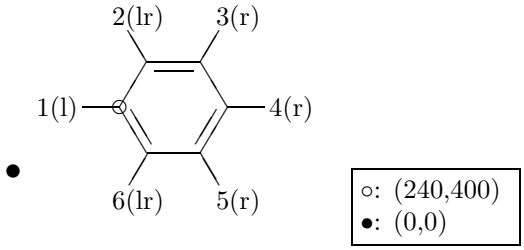


3.1.2 Horizontal Forms of Benzene Derivatives

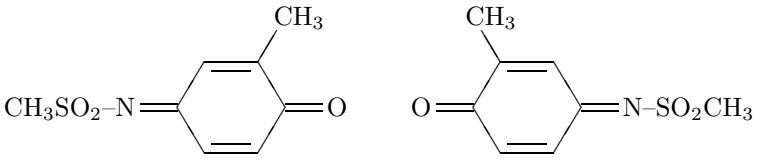
You can use the macro `\bzdrh` to draw benzene derivatives of horizontal type (carom.sty). The format of this command is as follows:

```
\bzdrh[OPT]{SUBSLIST}
```

The formats of the arguments are the same as those of `\bzdrv` (Tables 2.2 and 3.1). The locant numbering and the handedness of substitution are designed as follows:



For example, the diagrams:



are typeset by inputting the statements:

```
\bzdrh[pa]{4D==O;1D==CH$_{3}$SO$_{2}$--N;3==CH$_{3}$} \qquad
\bzdrh[pa]{1D==O;4D==N--SO$_{2}$CH$_{3}$;2==CH$_{3}$}
```

It should be noted the the commands `\bzdrv` and `\bzdrh` are based respectively on the commands `\cyclohexanev` and `\cyclohexaneh` that will be described in the next section. Hence, structures drawn with the former set of commands can be also drawn with the latter set of commands (see Figures 3.1 and 3.2).

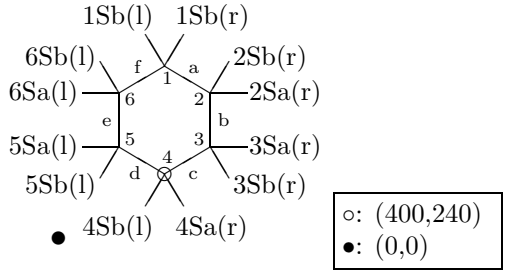
3.2 Drawing Cyclohexane Derivatives

3.2.1 Vertical Forms of Cyclohexane Derivatives

The macro `\cyclohexanev` is used to draw cyclohexane derivatives of vertical type (carom.sty). The format of this command is as follows:

```
\cyclohexanev[BONDLIST]{SUBSLIST}
```

Locant numbers (1–6) for designating substitution positions and characters (a–f) for showing bonds to be doubled are represented by the following diagram:



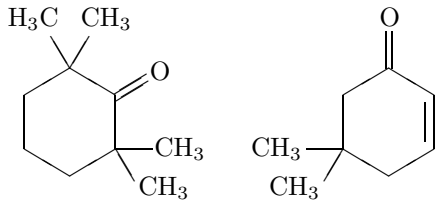
Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros.

The option argument BONDLIST is an character string in a pair of brackets, where each character indicates the presence of a double bond at the edge corresponding to the character. The bond-correspondence is rather arbitrary in some cases but conforms to chemical conventions as faithfully as possible if such conventions are presence (Table 3.2). Several examples for drawing endocyclic double bonds are listed in Figure 3.2. Note that Figure 3.2 provides alternative ways for designating endocyclic double bonds. Compare this with the results collected in Figure 3.1.

The argument SUBSLIST for this macro takes a general format, in which the modifiers listed in Table 2.2 are used. Suppose you input the commands:

```
\cyclohexanev{2D==O;1Sb==H$_{3}$C;1Sa==CH$_{3}$;%
3Sb==CH$_{3}$;3Sa==CH$_{3}$} \qquad\qquad
\cyclohexanev[b]{1D==O;5Sb==CH$_{3}$;5Sa==CH$_{3}$}
```

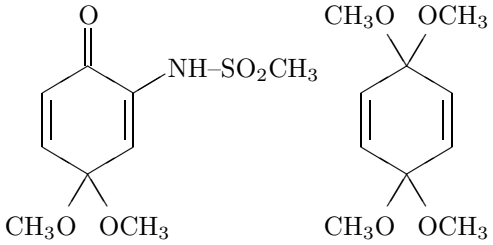
The first example illustrates the case that `\cyclohexanev` accompanies no optional argument. On the other hand, the second one take [b] as an optional BONDLIST, which prints an inner bond between 2 and 3 positions. Thus, you can obtain the following diagrams:



Since the macro `\cyclohexanev` is the basis of the macro `\bzdrv`, structural formulas depicted with the latter command can also be written by the former one. For example, the quinone acetals described above are also typeset by the following statements.

```
\cyclohexanev[be]{1D==O;4Sb==CH$_{3}$O;4Sa==OCH$_{3}$;2==NH--SO$_{2}$CH$_{3}$}
\qquad \qquad
\cyclohexanev[be]{1Sb==CH$_{3}$O;1Sa==OCH$_{3}$;4Sb==CH$_{3}$O;4Sa==OCH$_{3}$}
```

These commands are completely equivalent to those describe above and produce the following structures.



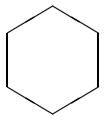
For the purpose of depicting the stereochemisty of a cyclohexane ring, input the following:

```
\cyclohexanev{2B==CH$_{3}$;3B==CH$_{3}$}\qquad\qquad
\cyclohexanev{2B==CH$_{3}$;3A==CH$_{3}$}
```

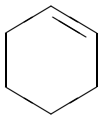
Thereby, you can obtain:

Table 3.2: Argument BONDLIST for commands \cyclohexanev and \cyclohexaneh

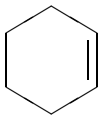
Character	Printed structure
none	cyclohexane
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,6-double bond
f	6,1-double bond
A	aromatic circle



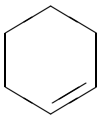
\cyclohexanev{}



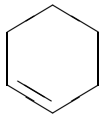
\cyclohexanev[a]{}



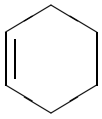
\cyclohexanev[b]{}



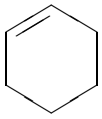
\cyclohexanev[c]{}



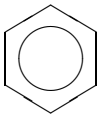
\cyclohexanev[d]{}



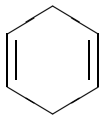
\cyclohexanev[e]{}



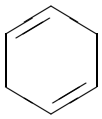
\cyclohexanev[f]{}



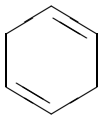
\cyclohexanev[A]{}



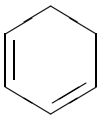
\cyclohexanev[be]{}



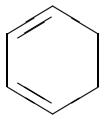
\cyclohexanev[cf]{}



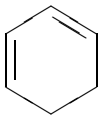
\cyclohexanev[ad]{}



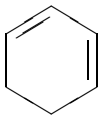
\cyclohexanev[ce]{}



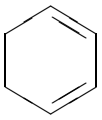
\cyclohexanev[df]{}



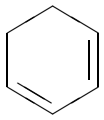
\cyclohexanev[ae]{}



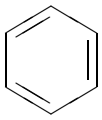
\cyclohexanev[bf]{}



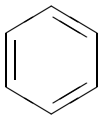
\cyclohexanev[ae]{}



\cyclohexanev[bd]{}



\cyclohexanev[bdf]{}



\cyclohexanev[ace]{}

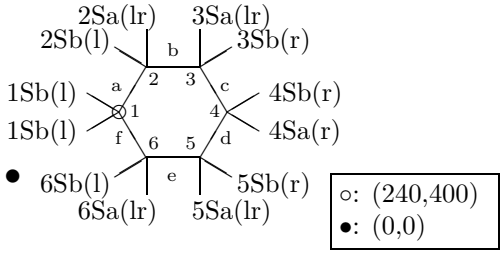
Figure 3.2: Endocylic bonds by the BONDLIST argument

3.2.2 Horizontal Forms of Cyclohexane Derivatives

The macro `\cyclohexaneh` is used to draw cyclohexane derivatives of horizontal type (carom.sty). The format of this command is as follows:

```
\cyclohexaneh[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros. The SUBSLIST and the BONDLIST format are shown in Table 2.2 and 3.2, respectively. Several examples for designating BONDLIST (Table 3.2) are collected in Figure 3.3. Note that this figure is obtained by the slight modification of Figure 3.2, where the suffix 'v' of the command `\cyclohexanev` is changed into 'h' to input the command `\cyclohexaneh`.

The following examples show the designation of SUBSLIST and of BONDLIST.

Example:

```
\cyclohexaneh{3D==O;5D==O;1Sb==CH$_{3}$;1Sa==CH$_{3}$;%
4==CH$_{2}$CO$_{2}$H}\qquad\qquad
\cyclohexaneh{4D==CH$_{2}$;3SB==CH$_{3}$;3SA==H}
```

These commands produce:

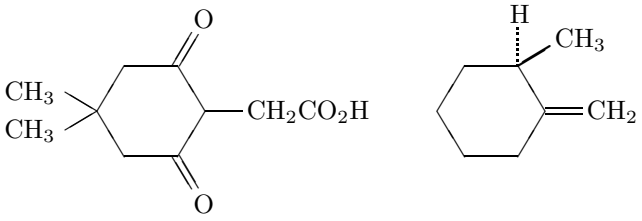


Figure 3.3: Endocylic bonds by the BONDLIST argument

Chapter 4

Carbocycles with Fused Six-Membered Rings

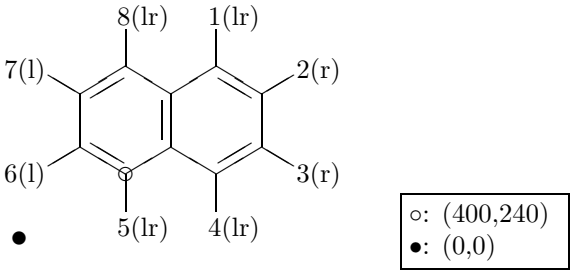
4.1 Drawing Naphthalene Derivatives

4.1.1 Vertical Forms of Naphthalene Derivatives

The macro `\naphdrv` is used to draw naphthalene derivatives of vertical type (carom.sty) as well as various naphthoquinone derivatives. The format of this command is as follows:

```
\naphdrv[OPT]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument OPT is used to specify a bond pattern as shown in Table 4.1.

Several endcyclic bond patterns typeset by the OPT argument of the `\naphdrv` command (Table 4.1) are shown in Figures 4.1 and 4.2.

The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 8.

Example:

```
\naphdrv{1==CH$_{2}$CH=CH$_{2}$;2==OH} \qquad
\naphdrv{6==H$_{3}$C;2==COCH$_{2}$CH$_{2}$COOH} \hskip1.5cm
\naphdrv[o]{1Sb==Cl;1Sa==Cl;2D==O}
```

These commands produce:

Table 4.1: Argument OPT for commands \naphdrv and \naphdrh

Character	Printed structure
none	naphthalene
A	aromatic circle
p or pa	1,4-quinone (A) left aromatic, right quinone
pb	1,4-quinone (B) right aromatic, left quinone
o or oa	*o*-quinone (A) (Oxygen atomes at 1,2-positions)
ob	*o*-quinone (B) (Oxygen atomes at 2,3-positions)
oc	*o*-quinone (C) (Oxygen atomes at 3,4-positions)
od	*o*-quinone (D) (Oxygen atomes at 4,5-positions)
oe	*o*-quinone (E) (Oxygen atomes at 5,6-positions)
of	*o*-quinone (F) (Oxygen atomes at 1,6-positions)
q or qa	2,6-quinone (A)
qb	2,6-quinone (B) (actually 3,7-positons)
qc	1,5-quinone (C)
qd	1,5-quinone (D) (actually 4,8-positions)
qe	1,7-quinone (E)
qf	1,7-quinone (F) (actually 2,8-positions)
qg	1,7-quinone (G) (actually 4,6-positions)
qh	1,7-quinone (H) (actually 3,5-positions)
P or Pa	: 1,4,5,8-quinone (A)
Pb	1,2,5,8-quinone (B)
Q	1,2,3,4-quinone
O or Oa	1,2,5,6-quinone (A)
Ob	1,2,7,8-quinone (B)
Oc	1,2,3,5-quinone (C)
Od	1,2,3,7-quinone (D)

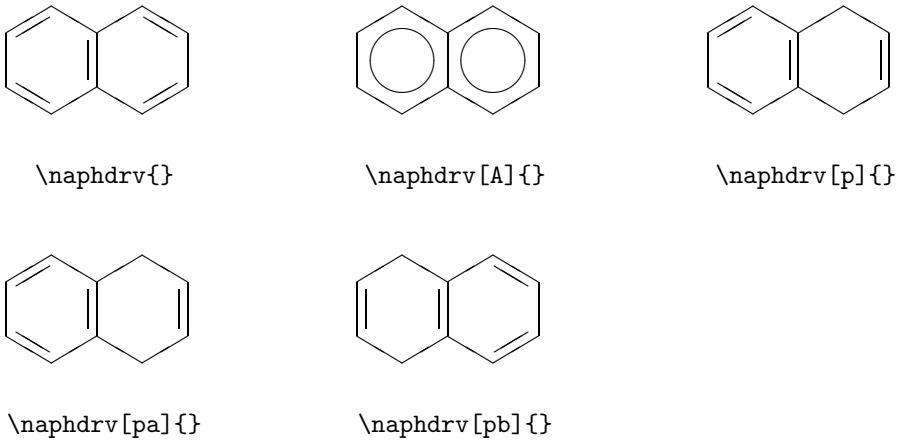


Figure 4.1: Endocylic bond patterns by the OPT argument

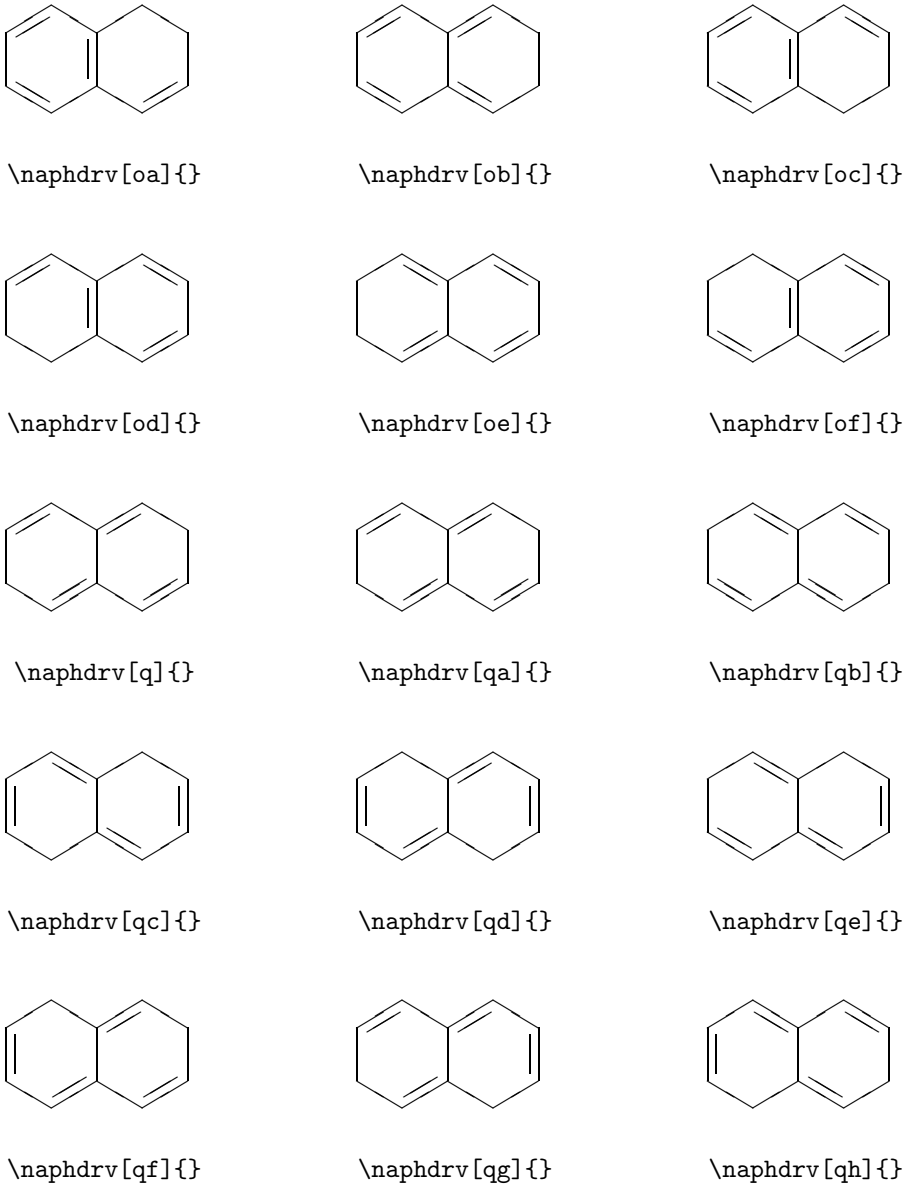
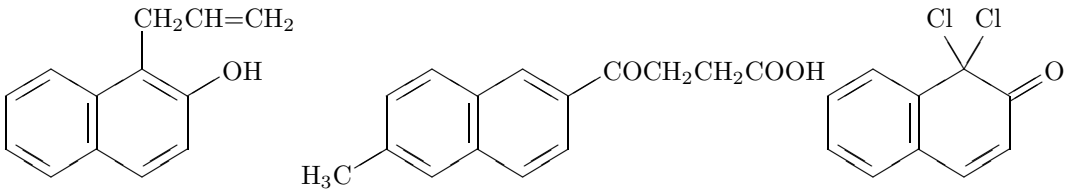


Figure 4.2: Endocylic bond patterns by the OPT argument (continued)

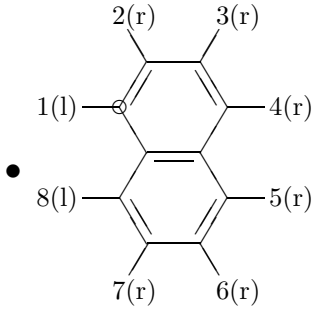


4.1.2 Horizontal Forms of Naphthalene Derivatives

The macro `\naphdrh` is used to draw naphthalene derivatives of horizontal type (carom.sty) as well as various naphthoquinone derivatives. The format of this command is as follows:

```
\naphdrh[OPT]{SUBSLIST}
```

The format of the argument OPT is the same as that of `\naphdrv` (Tables 4.1). The format of the argument SUBSLIST is the same as collected in Tables 2.2. The locant numbering and the handedness of substitution are designed as follows:

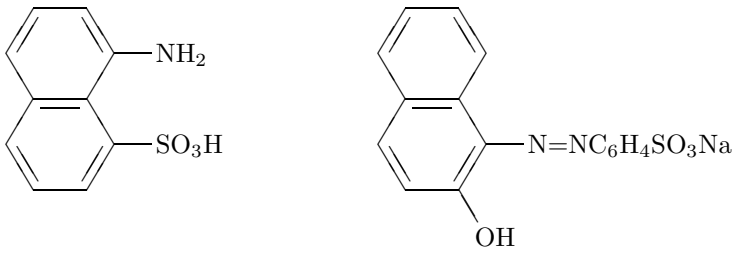


○: (240,400)
●: (0,0)

Example:

```
\naphdrh{4==NH$_{2}$;5==SO$_{3}$H}\qquad
\naphdrh{5==N=NC$_{6}$H$_{4}$SO$_{3}$Na;6==OH}
```

These commands produce:



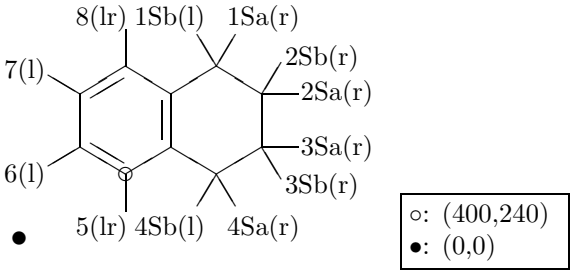
4.2 Drawing Tetraline Derivatives

4.2.1 Vertical Forms of Tetraline Derivatives

The macro `\tetralinev` is used to draw tetraline derivatives of vertical type (carom.sty) as well as various naphthoquinone derivatives. The format of this command is as follows:

```
\tetralinev[OPT]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument OPT is used to specify a bond pattern as shown in Table 4.2.

Table 4.2: Argument OPT for commands `\tetralinev` and `\tetralineh`

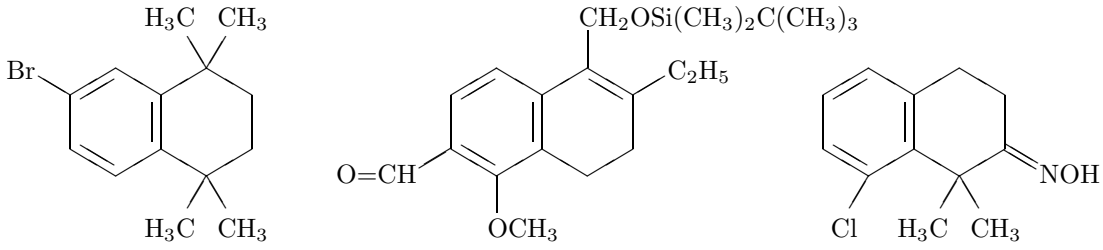
Character	Printed structure
none	tetraline
A	aromatic circle
e or ea	1,2-double bond
eb	2,3-double bond
ec	3,4-double bond

A bond modifier in the argument SUBSLIST for $n = 1$ to 4 can be one of the bond modifiers shown in Table 2.2, which allows α- or β-orientation. On the other hand a bond modifier in the argument SUBSLIST for $n = 5$ to 8 should be vacant. If there appears the overcrowding between 1- and 8-substituent or between 4- and 5-substituent, the bond modifier 5Sb or 8Sb is allowed to avoid such overcrowding.

Example:

```
\tetralinev{1Sb==H$_{3}$C;1Sa==CH$_{3}$;%
4Sb==H$_{3}$C;4Sa==CH$_{3}$;7==Br}\qquad
\tetralinev[ea]{1==CH$_{2}$OSi(CH$_{3}$)$_{2}$C(CH$_{3}$)$_{3}$;
2==C$_{2}$H$_{5}$;5==OCH$_{3}$;6==O=CH}\qquad
\tetralinev{3D==NOH;4Sb==H$_{3}$C;4Sa==CH$_{3}$;5Sb==Cl}
```

These commands produce:

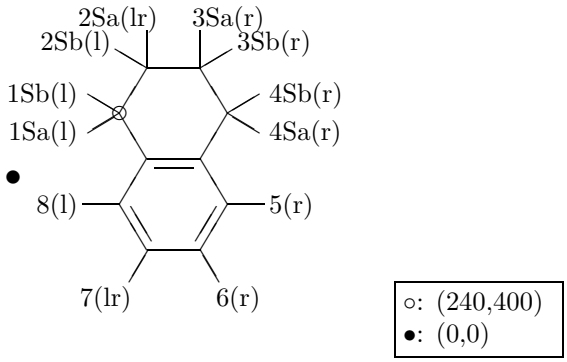


4.2.2 Horizontal Forms of Tetraline Derivatives

The `\tetralineh` is the horizontal counterpart of the command `\tetralinev`:

```
\tetralineh[OPT]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:

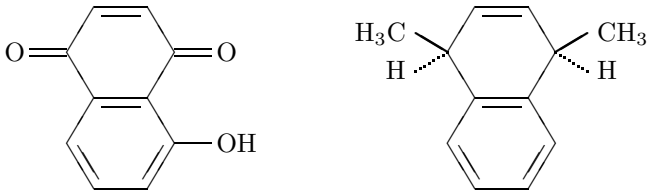


The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument OPT is used to specify a bond pattern as shown in Table 4.2. The argument SUBSLIST is the same as that of `\tetralinev`.

Example:

```
\tetralineh[eb]{1D==O;4D==O;5==OH} \qquad
\tetralineh[eb]{1SB==H$_{3}$C;1SA==H;4SB==CH$_{3}$;4SA==H}
```

These commands produce:



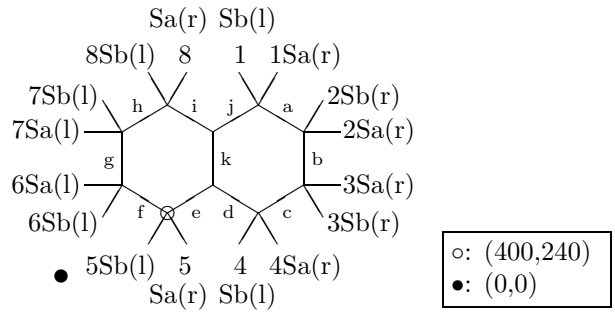
4.3 Drawing Decaline Derivatives

4.3.1 Vertical Forms of Decaline Derivatives

The macro `\decalinev` is used to draw decaline derivatives of vertical type (carom.sty). The format of this command is as follows:

```
\decalinev[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions and characters for showing bonds to be doubled are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The option argument BONDLIST is based on the assignment of characters (a–k) to respective bonds as shown in the above diagram. A bond modifier in the argument SUBSLIST for n = 1–8 can be one of bond modifiers shown in Table 2.2. The substitution at the bridgehead positions is designated as shown in Table 4.3.

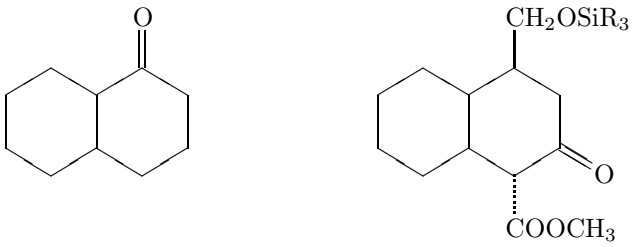
Table 4.3: SUBSLIST for bridgehead positions in \decalinev and \decalineh

Character	Printed structure
0FA	alpha single bond at 8a
0FB	beta single bond at 8a
0FU	unspecified single bond at 8a
0GA	alpha single bond at 4a
0GB	beta single bond at 4a
0GU	unspecified single bond at 4a

Example:

```
\decalinev{1D==O;0FB==H;0GA==H} \qquad
\decalinev{1B==CH$_{2}$OSiR$_{3}$;3D==O;4A==COOCH$_{3}$;%
0FB==CH$_{3}$;0GA==H}
```

These commands produce:

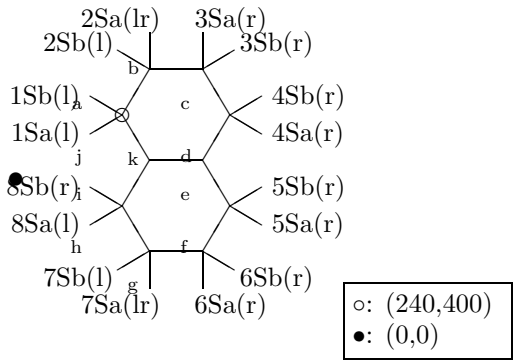


4.3.2 Horizontal Forms of Decaline Derivatives

The macro \decalineh (carom.sty) is the horizotal counterpart of \decalinev. The format and the assignment of BONDLIST and SUBSLIST of the former macro are the same as the latter (see Tables 2.2 and 4.3).

```
\decalineh[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:

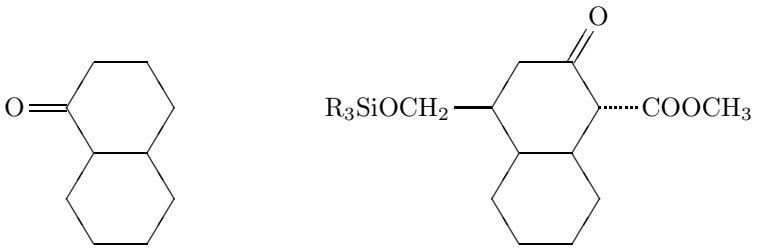


The handedness for each oriented or double-sided position is shown with a character set in parentheses.

Example:

```
\decalineh{1D==O;0FA==H;0GB==H} \qquad\qquad
\decalineh{1B==R$_{3}$SiOCH$_{2}$;3D==O;4A==COOCH$_{3}$;%
0FB==CH$_{3}$;0GA==H}
```

These commands produce:



Chapter 5

Fused Tricyclic Carbocycles and Steroids

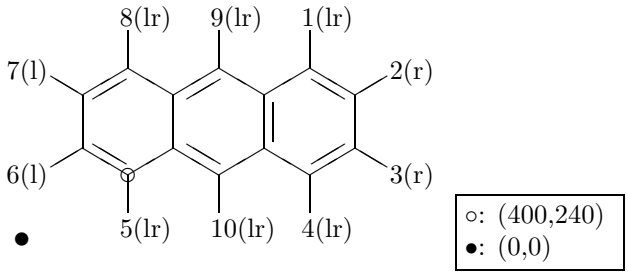
5.1 Drawing Anthracene Derivatives

5.1.1 Command for Specified Use

The macro `\anthracenev` is used to draw anthracene derivatives of vertical type (carom.sty) as well as various quinone derivatives. The format of this command is as follows:

```
\anthracenev[OPT]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument OPT is used to specify a bond pattern as shown in Table 5.1. The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 10.

Example:

```
\anthracenev[pa]{9D==O;{{10}D}==O;2==COOH}\hskip1.5cm
\anthracenev[pA]{9D==O;{{10}D}==O;2==COOH}
```

These commands produce:

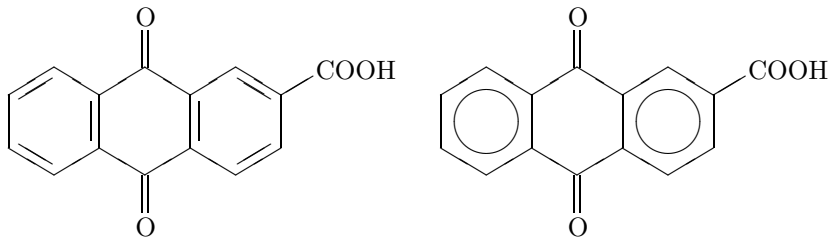


Table 5.1: Argument OPT for commands \anthracenev

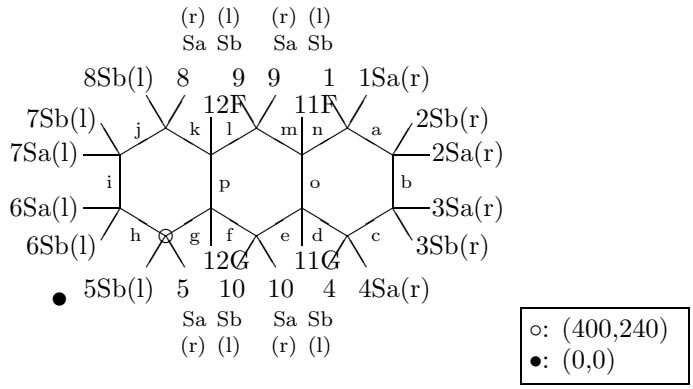
Character	Printed structure
none or r	right-handed double bonds
l	left-handed double bonds
A	aromatic circle
p or pa	9,10-anthraquinone (A)
pA	9,10-anthraquinone (circle type)
o	1,2-anthraquinone (A)
oa	1,2-anthraquinone (A')
oA	1,2-anthraquinone (circle type)
ob	2,3-antharquinone (B)
oc	1,2-anthraquinone (C)
q	1,4-anthraquinone (A)
qa	1,4-anthraquinone (A')
qA	1,4-anthraquinone (circle type)

5.1.2 Command for General Use

The macro \hanthracenev (carom.sty) is a more general macro than anthracenev, where the latter is actually a short-cut command of the former. The \hanthracenev command takes the following format:

```
\hanthracenev[BONDLIST]{SUBSLIST}
```

Locant numbers (1–12) for designating substitution positions and bond descriptors (a–p) are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The option argument BONDLIST is based on the assignment of characters (a–p) to respective bonds as shown in the above diagram and Table 5.2. A bond modifier in the argument SUBSLIST for n = 1–10 is selected from those shown in Table 2.2. The substitution at the bridgehead positions is designated as shown in Table 5.3.

Example:

```
\hanthracenev[C]{5==\lmoiety{CH$_{3}$O};%
8==\lmoiety{CH$_{3}$O};9==CN;{{10}D}==O}\qquad
\hanthracenev[hjp]{{{11}FA}==H;{{11}GA}==H;1A==OBz;4B==OH;2D==O}
```

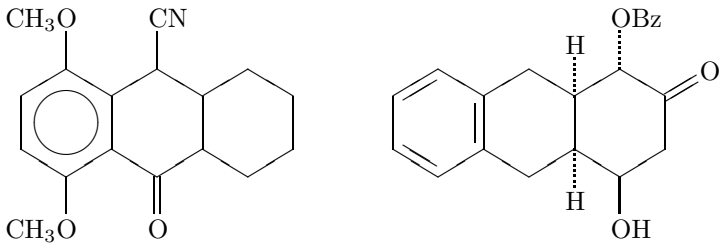
These commands produce:

Table 5.2: Argument BONDLIST for commands \hanthracenev

Character	Printed structure
none	perhydro-anthracene
a	1,2-double bond
b	2,3-double bond
c	3,4-double bond
d	4,4a-double bond
e	10,4a-double bond
f	10,10a-double bond
g	5,10a-double bond
h	5,6-double bond
i	6,7-double bond
j	8,7-double bond
k	8,8a-double bond
l	9,8a-double bond
m	9,9a-double bond
n	1,9a-double bond
o	4a,9a-double bond
p	10a,8a-double bond
A	right aromatic circle
B	central aromatic circle
C	left aromatic circle

Table 5.3: SUBSLIST for bridgehead positions in \hanthracenev

Character	Printed structure
11FA	alpha single bond at 9a
11FB	beta single bond at 9a
11FU	unspecified single bond at 9a
11GA	alpha single bond at 4a
11GB	beta single bond at 4a
11GU	unspecified single bond at 4a
12FA	alpha single bond at 8a
12FB	beta single bond at 8a
12FU	unspecified single bond at 8a
12GA	alpha single bond at 10a
12GB	beta single bond at 10a
12GU	unspecified single bond at 10a



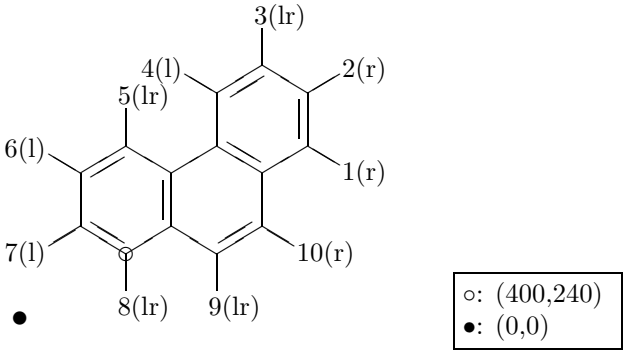
5.2 Drawing Phenanthrene Derivatives

5.2.1 Command for Specified Use

The macro `\phenanthrenev` is used to draw phenanthrene derivatives of vertical type (carom.sty) as well as various quinone derivatives. The format of this command is as follows:

```
\phenanthrenev[OPT]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument OPT is used to specify a bond pattern as shown in Table 5.4.

Table 5.4: Argument OPT for commands `\phenanthrenev`

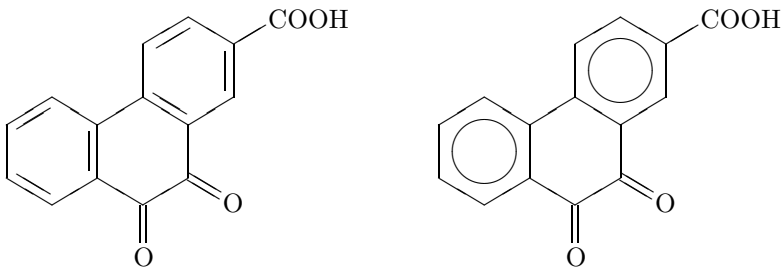
Character	Printed structure
none or r	right-handed double bonds
A	aromatic circle
p or pa	1,4-quinone (A)
pA	1,4-quinone (circle type)
o or oa	1,2-quinone (A)
oA	1,2-quinone (circle type)
ob	2,3-quinone (B)
oc	3,4-anthraquinone (C)
q or qa	9,10-quinone
qA	9,10-quinone (circle type)

The argument SUBSLIST is employed to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is seledted to be an arabic numeral between 1 and 10.

Example:

```
\phenanthrenev[q]{9D==O;{{10}D}==O;2==COOH}\hskip1.5cm
\phenanthrenev[qA]{9D==O;{{10}D}==O;2==COOH}
```

These commands produce:

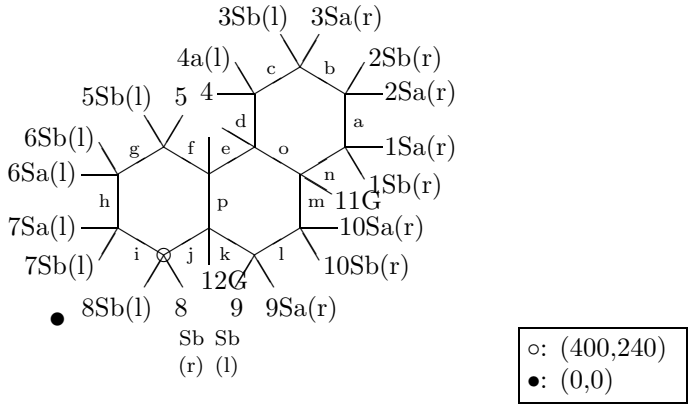


5.2.2 Command for General Use

The macro `\hphenantherev` (carom.sty) is a more general macro than `phenanthrenev`, where the latter is a short-cut command based on the former. The format of the `\hphenantherev` command is as follows:

```
\hphenanthrenev[BONDLIST]{SUBSLIST}
```

Locant numbers (1–12) for designating substitution positions and bond descriptors (a–p) are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses, where the designation of overcrowded positions is abbreviated.

The option argument BONDLIST is based on the assignment of characters (a–p) to respective bonds as shown in the above diagram and Table 5.5. A bond modifier in the argument SUBSLIST for $n = 1$–10 can be one of bond modifiers shown in Table 2.2. The substitution at the bridgehead positions is similar to that designated in Table 5.3 for `\hanthracenev`.

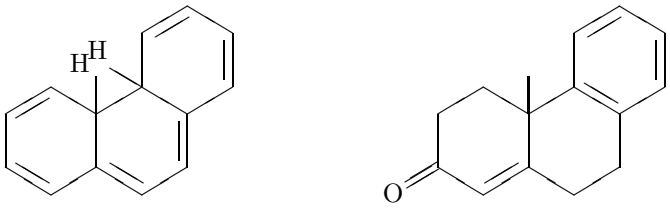
Example:

```
\hphenanthrenev[acgikm]{{{11}F}=={\kern-3em\raise1ex\hbox{H}};%
{{12}F}==\lmoiety{H~~}}\hskip1.5cm
\hphenanthrenev[acoj]{7D==O;{{12}FB}==}
```

These commands produce:

Table 5.5: Argument BONDLIST for commands \hphenanthrenev

Character	Printed structure
none	perhydro-phenanthrene
a	1,2-double bond
b	2,3-double bond
c	3,4-double bond
d	4,4a-double bond
e	4a,4b-double bond
f	4b,5-double bond
g	5,6-double bond
h	6,7-double bond
i	7,8-double bond
j	8,8a-double bond
k	8a,9-double bond
l	9,10-double bond
m	10,10a-double bond
n	1,10a-double bond
o	4a,10a-double bond
p	4b,8a-double bond
A	right aromatic circle
B	central aromatic circle
C	left aromatic circle

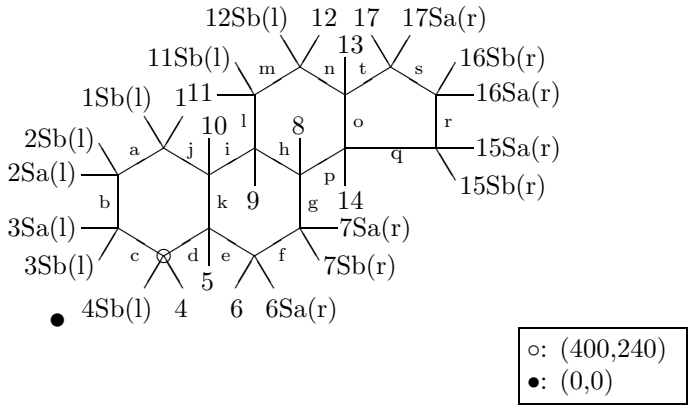


5.3 Drawing Steroid Derivatives

The macro \steroid (carom.sty) typsets a steroid derivative without the side chain. The format of this command is as follows:

```
\steroid[BONDLIST]{SUBSLIST}
```

Locant numbers (1–17) for designating substitution positions and bond descriptors (a–t) are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses, where the designation of overcrowded positions is abbreviated.

The option argument BONDLIST is based on the assignment of characters (a–t) to respective bonds as shown in the above diagram and Table 5.6.

Table 5.6: Argument BONDLIST for commands \steroid

Character	Printed structure	Character	Printed structure
none	steroid skeleton		
a	1,2-double bond	b	2,3-double bond
c	3,4-double bond	d	4,5-double bond
e	6,5-double bond	f	6,7-double bond
g	7,8-double bond	h	9,8-double bond
i	9,10-double bond	j	1,10-double bond
k	5,10-double bond	l	9,11-double bond
m	12,11-double bond	n	12,13-double bond
o	14,13-double bond	p	8,14-double bond
q	14,15-double bond	r	15,16-double bond
s	17,16-double bond	t	17,13-double bond
A	aromatic A ring	B	aromatic B ring
C	aromatic C ring		

A bond modifier in the argument SUBSLIST for n = 1–17 (except fused positions) is selected from the list of bond modifiers (Table 2.2). The substitution at the fused positions (n = 5,8,9,10,13 and 14) is similarly designated as for fused bicylic or tricyclic compounds (Table 5.7).

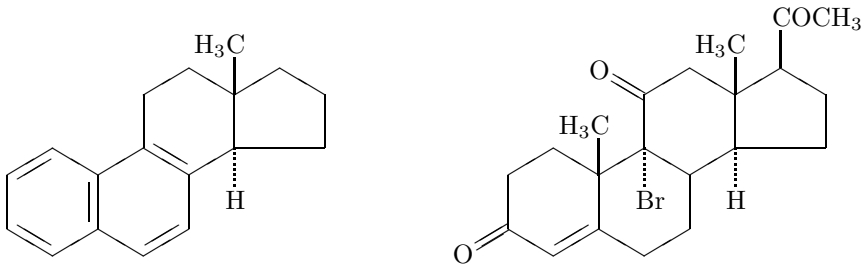
Table 5.7: SUBSLIST for fused positions in \steroid

Character	Printed structure
n or nS	exocyclic single bond at n-atom
nA	alpha single bond at n-atom (boldface)
nB	beta single bond at n-atom (dotted line)
nU	unspecified single bond at n-atom

Example:

```
\steroid[ackhf]{{{13}B}==\lmoiety{H$_{3}$C};{{14}A}==H}\hskip1cm
\steroid[d]{3D==O;9A==Br;{{11}D}==O;%
{{17}B}==COCH$_3$;{{14}A}==H;%
{{13}B}==\lmoiety{H$_3$C};{{10}B}==\lmoiety{H$_3$C}}
```

These commands produce:

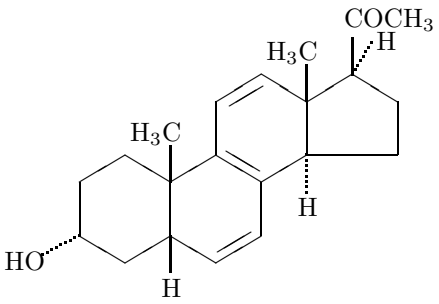


In order to avoid the overcrowding of substitution, you can use TEX primitive commands such as `\raise` and `\kern`.

Example:

```
\steroid[fhm]{3A==HO;5B==H;{{10}B}==\lmoiety{H$_{3}$C};%
{{13}B}==\lmoiety{H$_{3}$C};%
{{14}A}==H;{{17}B}==\raise.5ex\hbox{COCH$_{3}$};%
{{17}SA}=={\kern.5em\lower1.5ex\hbox{H}}}
```

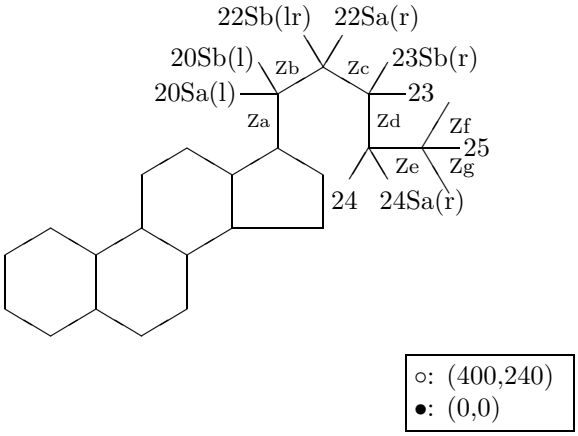
These commands produce:



The macro `\steroidchain` (carom.sty) is to draw a steroid derivative with the side chain. The format of this command is as follows:

```
\steroidchain[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions and bond descriptors for the side chain are represented by the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses, where the designation of overcrowded positions is abbreviated.

The option argument BONDLIST is based on the assignment of characters (a–t) to respective bonds as shown in the above diagram and Table 5.6. The locant-numbering of chain carbons is also designated with the BONDLIST in the form of two-character indicators (Za–Zg) as collected in Table 5.8. A bond

Table 5.8: Argument BONDLIST for chain carbons (\steroidchain)

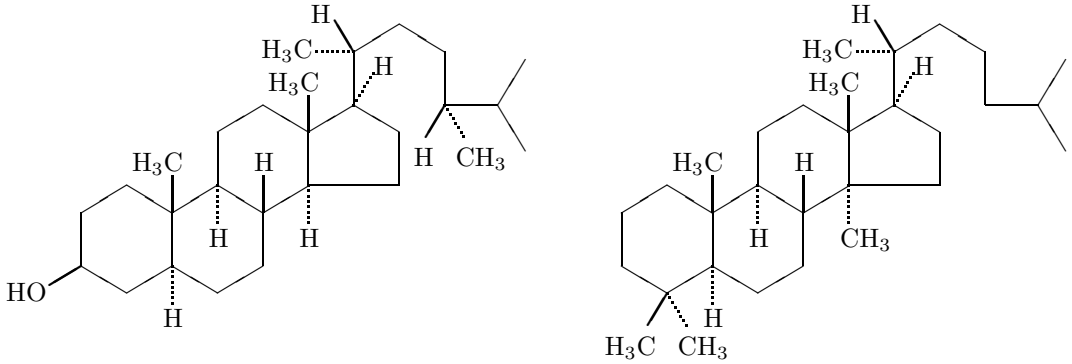
Character	Printed structure	Character	Printed structure
Z	no action		
Za	17,20-double bond	Zb	20,22-double bond
Zc	22,23-double bond	Zd	23,24-double bond
Ze	24,25-double bond	Zf	25,26-double bond
Zg	25,27-double bond		

modifier in the argument SUBSLIST for n = 1–25 (except fused positions and terminal positions not to be specified, *e.g.*, 18) can be one of bond modifiers shown in Table 2.2. On the other hand, a bond modifier in the argument SUBSLIST for n = 5, 8, 9, 10, 13, 14, or 25 (fused positions *etc.*) can be selected from bond modifiers shown in Table 5.7.

For example, the \steroidchain macro prints (24*R*)-24-methyl-5α-cholestan-3β-ol (campestanol) and 5α-lanostane only by replacing substituents in argument SUBSLIST. Thus, the statements

```
\steroidchain{3B==HO;5A==H;{{10}B}==\lmoiety{H$_3$C};9A==H;8B==H;%
{{17}SA}==\lower1ex\hbox{ H};{{13}B}==\lmoiety{H$_3$C};{{14}A}==H;%
{{20}SA}==H$_3$C;{{20}SB}==H;{{24}SA}==CH$_3$;{{24}SB}==H}
\steroidchain{4SB==\lmoiety{H$_3$C};4SA==CH$_3$;5A==H;%
{{17}SA}==\lower1ex\hbox{ H};%
{{10}B}==\lmoiety{H$_3$C};9A==H;8B==H;{{13}B}==\lmoiety{H$_3$C};%
{{14}A}==CH$_3$;{{20}SA}==\lmoiety{H$_3$C};{{20}SB}==H}
```

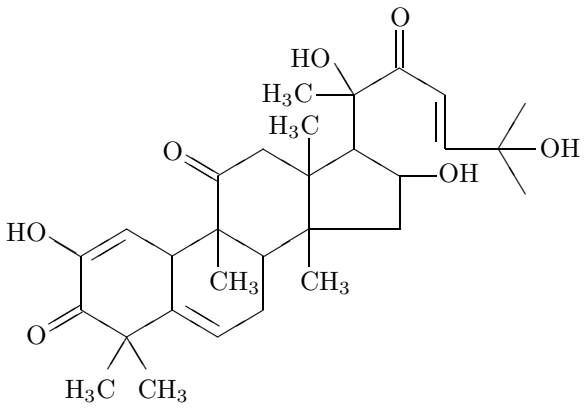
typeset the following structural diagrams:



The following example of drawing cucurbitacin I illustrates the designation of double bonds in the side chain. Thus, a single macro is capable of covering a wide variety of derivatives by altering the description in arguments BONDLIST and SUBSLIST.

```
\steroidchain[ae{Zd}]{2==HO;3D==O;4Sb==\lmoiety{H$_3$C};4Sa==CH$_3$;%
9Sa==CH$_{3}$;{{11}D}==O;{{13}}==\lmoiety{H$_3$C};%
{{14}}==CH$_3$;{{20}Sa}==\lmoiety{H$_3$C};{{20}Sb}==HO;%
{{16}Sa}==OH;{{22}D}==O;{{25}}==OH}
```

produces



Chapter 6

Five- or Lower-Membered Carbocycles

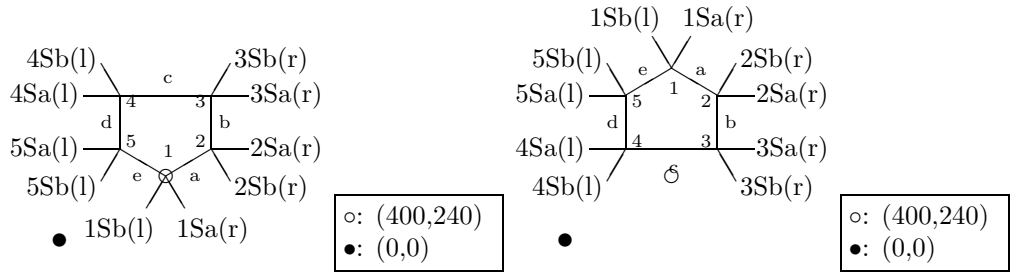
6.1 Drawing Five-Membered Carbocycles

6.1.1 Vertical Forms

The macro `\cyclopentanev` and the corresponding inverse macro are used to draw five-membered carbocyclic compounds of vertical type (lowcycle.sty). The formats of these commands are as follows:

```
\cyclopentanev[BONDLIST]{SUBSLIST}
\cyclopentanevi[BONDLIST]{SUBSLIST}
```

The following diagrams show the numbering of the commands for designating substitution positions (1–5) and bond descriptors (a–e):



In drawing five-membered rings, only commands for general use are ready to use so that they can be employed to typeset both saturated and unsaturated derivatives. Commands for specified use have not been developed since they are not so desirable as compared with the counterparts of six-membered rings.

The optional argument BONDLIST shows bonds to be doubled as shown in Table 6.1. The default structure is a fully saturated form.

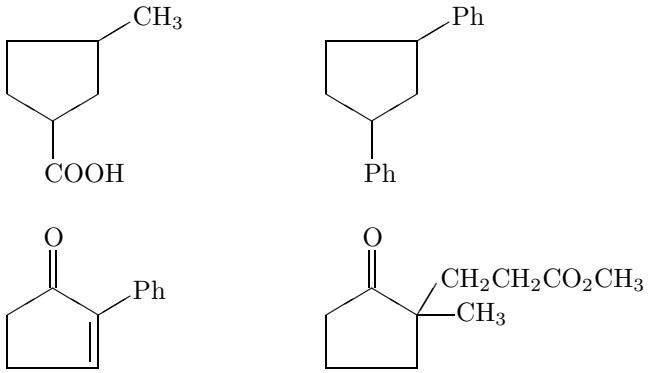
The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 5. For example, the statements,

```
\cyclopentanev{1==COOH;3==CH$_{3}$}\qquad\qquad
\cyclopentanev{1==Ph;3==Ph} \par
\cyclopentanevi[b]{1D==O;2==Ph}\qquad\qquad
\cyclopentanevi{1D==O;2Sa==CH$_{3}$;%
2Sb==CH$_{2}$CH$_{2}$CO$_{2}$CH$_{3}$}
```

produce the following structures:

Table 6.1: BONDLIST for commands \cyclopentanev and \cyclopentaneh

Character	Printed structure
none	mother nucleus
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,1-double bond
A	aromatic circle
{n+}	plus at the n-nitrogen atom (n = 1 to 5)
{0+}	plus (or minus) at the center



The command is capable of typesetting a delocalized and a localized form of cyclopentadienyl anion as follows:

```
\cyclopentanev[A{0{$-$}}]{} \qquad
\cyclopentanev[bd{1{\lower1.2ex\hbox{$-$}}}]{}
```

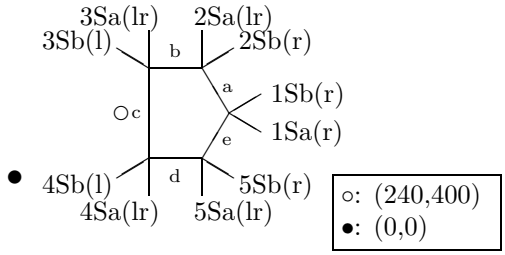
where the charges are designated in terms of the BONDLIST (Table 6.1). These statements produce

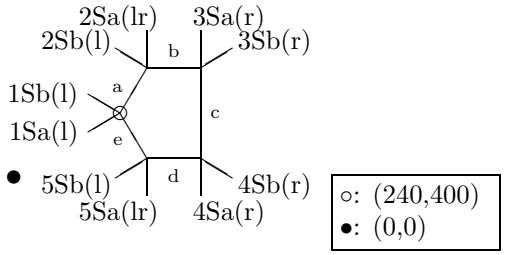
6.1.2 Horizontal Forms

The macro \cyclopentaneh and the corresponding inverse macro are used to draw five-membered carbocyclic compounds of horizontal type (lowcycle.sty). The formats of these commands are as follows:

```
\cyclopentaneh[BONDLIST]{SUBSLIST}
\cyclopentanehi[BONDLIST]{SUBSLIST}
```

The following diagrams show locant numbers for designating substitution positions as well as bond descriptors for showing double bonds:



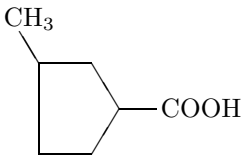


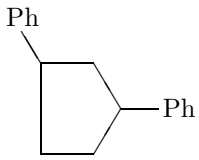
in which the same macro is used to typeset both saturated and unsaturated derivatives. For BONDLIST, see Table 6.1.

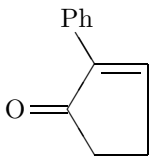
Example:

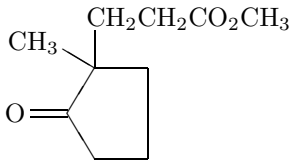
```
\cyclopentaneh{1==COOH;3==CH$_{3}$}\qquad\qquad
\cyclopentaneh{1==Ph;3==Ph} \par
\cyclopentanehi[b]{1D==O;2==Ph}\qquad\qquad
\cyclopentanehi{1D==O;2Sb==CH$_{3}$;%
2Sa==CH$_{2}$CH$_{2}$CO$_{2}$CH$_{3}$}
```

produce the following structures:







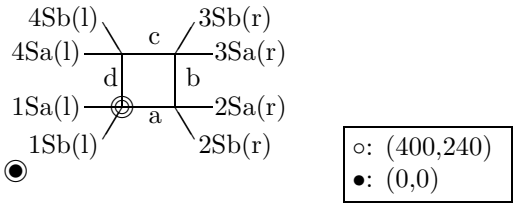


6.2 Drawing Four-Membered Carbocycles

The macro `\cyclobutane` is a command for drawing four-membered carbocycles by using the following format (lowcycle.sty).

```
\cyclobutane[BONDLIST]{SUBSLIST}
```

Locant numbers (1–4) and bond descriptors (a–d) are shown in the following diagram:



The handedness for each oriented position is shown with a character set in parentheses. The optional argument BONDLIST specifies double bonds as shown in Table 6.2.

Table 6.2: BONDLIST for commands \cyclobutane

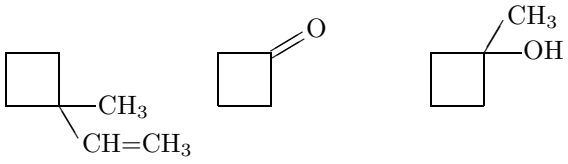
Character	Printed structure	Character	Printed structure
none	mother skeleton (fully saturated)		
a	1,2-double bond	b	2,3-double bond
c	3,4-double bond	d	4,1-double bond
{n+}	plus at the n-nitrogen atom (n = 1 to 4)		

The argument SUBSLIST is filled in to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 4.

Example:

```
\cyclobutane{2Sa==CH$_{3}$;2Sb==CH=CH$_{3}$}
\cyclobutane{3D==O}
\cyclobutane{3Sa==OH;3Sb==CH$_{3}$}
```

produce the following structures:

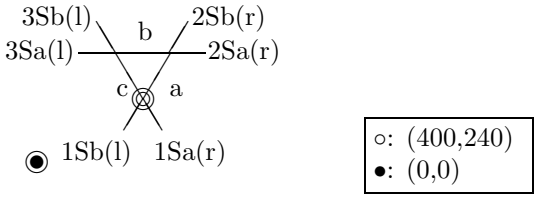


6.3 Drawing Three-Membered Heterocycles

The macro \cyclopropane, which is defined in lowcyclo.sty for drawing three-membered carbocycles, has the following format.

```
\cyclopropane[BONDLIST]{SUBSLIST}
```

Thus, the locant numbering (1–3) and the bond description (a–c) are common as shown in the following diagram:



The handedness for each oriented position is shown with a character set in parentheses. The optional argument BONDLIST is written down to specify double bonds as shown in Table 6.3.

Table 6.3: Argument BONDLIST for commands `\cyclopropane`

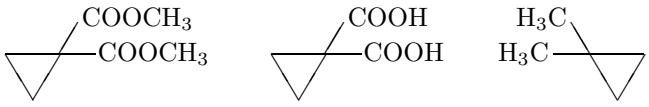
Character	Printed structure
none	saturated
a	1,2-double bond
b	2,3-double bond
c	3,1-double bond
A	aromatic circle
{n+}	plus at the n-hetero atom (n = 1 to 3)
	$n = 4$ – outer plus at 1 position
	$n = 5$ – outer plus at 2 position
	$n = 6$ – outer plus at 3 position
{0+}	plus at the center of a cyclopropane ring

The argument SUBSLIST is entered to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 3.

Example:

```
\cyclopropane{2Sa==COOCH$_{3}$;2Sb==COOCH$_{3}$}\qquad
\cyclopropane{2Sa==COOH;2Sb==COOH}\qquad\qquad
\cyclopropane{3Sa==H$_{3}$C;3Sb==H$_{3}$C}
```

produce the following structures:

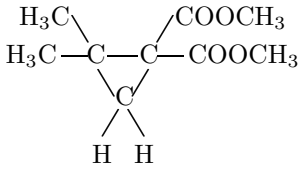


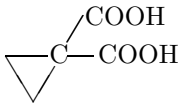
This macro is based on the macro `\threehetero` in which the ATOMLIST of the latter command is set beforehand. In order to draw a carbon atom on a cyclopropane ring, you can use the command `\threehetero` instead of `\cyclopropane`.

Example:

```
\threehetero[H]{1==C;2==C;3==C}%
 {1Sa==H;1Sb==H;2Sa==COOCH$_{3}$;2Sb==COOCH$_{3}$;%
  3Sa==H$_{3}$C;3Sb==H$_{3}$C}\qquad\qquad\qquad
\threehetero[H]{2==C}{2Sa==COOH;2Sb==COOH}
```

produce the following structures:





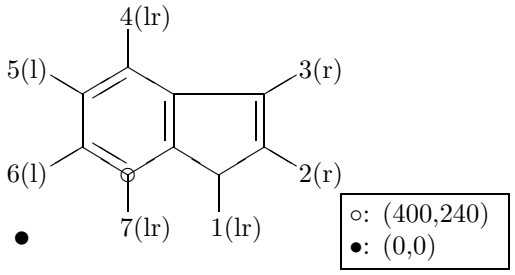
6.4 Drawing Indane Derivatives

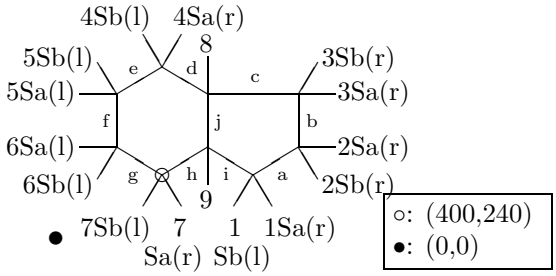
6.4.1 Vertical Forms of Indanes

Since the macros `\indanev` and `\indanevi` are included in the package file 'lowcycle.sty', this package should be introduced by using `\usepackage` in the preamble of your article. The format of `\indanev` is:

```
\indanev[BONDLIST]{SUBSLIST}
```

The locant numbering (1–9) and the bond description (a–j) have a common format as shown in the following diagrams:





The handedness for each oriented or double-sided position is shown with a character set in parentheses. Each character in the optional argument BONDLIST indicates a specific double bond as shown in Table 6.4. The default setting of BONDLIST produces a fully unsaturated structure, when the option BONDLIST is omitted. If you want to draw a fully saturated structure, you should write down a null option ([]) or [H].

Table 6.4: Argument BONDLIST for commands `\indolev` and others

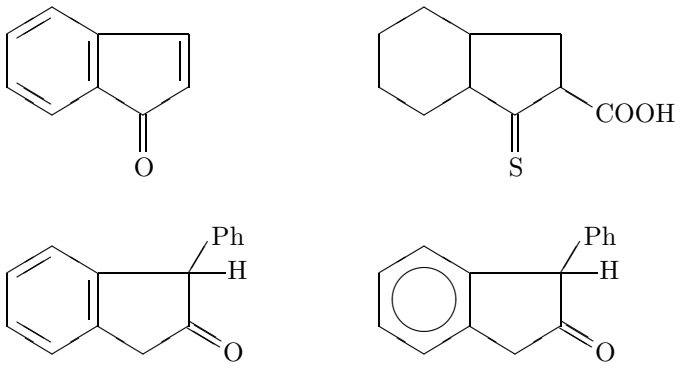
Character	Printed structure	Character	Printed structure
none or r	aromatic six-membered ring	H or []	fully saturated form
a	1,2-double bond	b	2,3-double bond
c	3,3a-double bond	d	4,3a-double bond
e	4,5-double bond	f	5,6-double bond
g	6,7-double bond	h	7,7a-double bond
i	1,7a-double bond	j	3a,4a-double bond
A	aromatic circle (six-membered ring)		
B	aromatic circle (five-membered ring)		
{n+}	plus at the n-nitrogen atom (n = 1 to 9)		

The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 7. Substitution on 8 (3a position) or 9 (7a position) can be assigned in the same way.

Example:

```
\indanev{1D==O} \qquad
\indanev[H]{1D==S;2==COOH}\par
\indanev[egj]{2D==O;3Sa==H;3Sb==Ph}\qquad
\indanev[A]{2D==O;3Sa==H;3Sb==Ph}
```

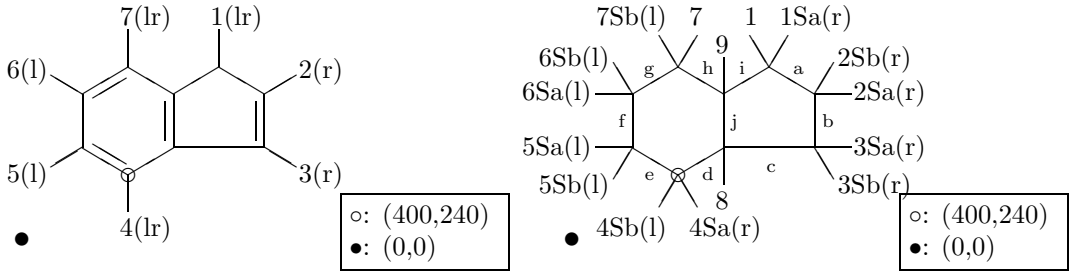
produce the following structures:



The macro `\indanevi` for drawing indane derivatives of inverse vertical type has the following format (lowcycle.sty).

```
\indanevi[BONDLIST]{SUBSLIST}
```

The locant numbering and the bond description are common with the vertical counterpart as shown in the following diagrams:

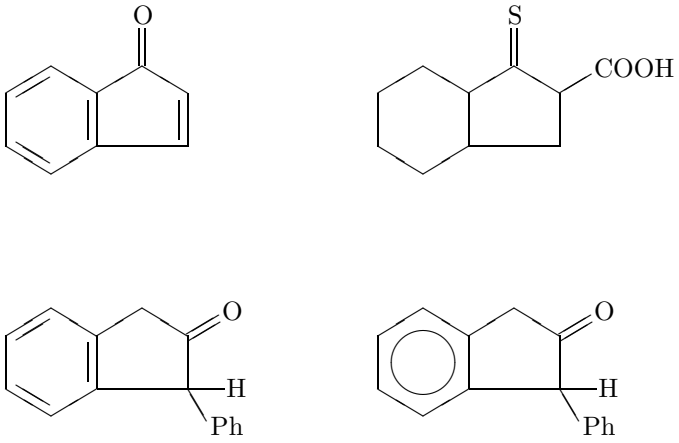


The handedness for each oriented or double-sided position is shown with a character set in parentheses; this is however omitted in the overcrowded position (between 7 and 1). The optional argument BONDLIST specifies bonds to be doubled as shown in Table 6.4.

Example:

```
\indanevi{1D==O} \qquad
\indanevi[H]{1D==S;2==COOH}\par
\indanevi[egj]{2D==O;3Sa==H;3Sb==Ph}\qquad
\indanevi[A]{2D==O;3Sa==H;3Sb==Ph}
```

produce the following structures:

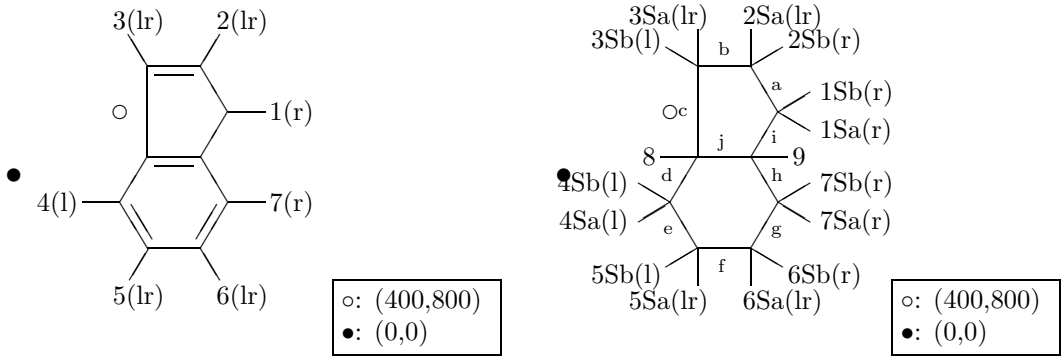


6.4.2 Horizontal Forms of Indanes

The macro `\indaneh` for drawing indane derivatives of horizontal type is defined in lowcycle.sty to have the following format.

```
\indaneh[BONDLIST]{SUBSLIST}
```

Locant numbers (1–9) for designating substitution positions and bond descriptors (a–j) are represented by the following diagram:

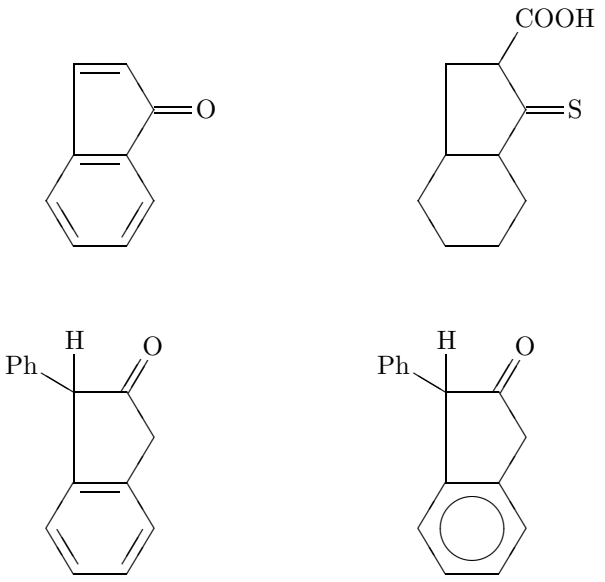


The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument BONDLIST gives the description of double bonds as shown in Table 6.4.

Example:

```
\indaneh{1D==O} \qquad
\indaneh[H]{1D==S;2==COOH}\par
\indaneh[egj]{2D==O;3Sa==H;3Sb==Ph}\qquad
\indaneh[A]{2D==O;3Sa==H;3Sb==Ph}
```

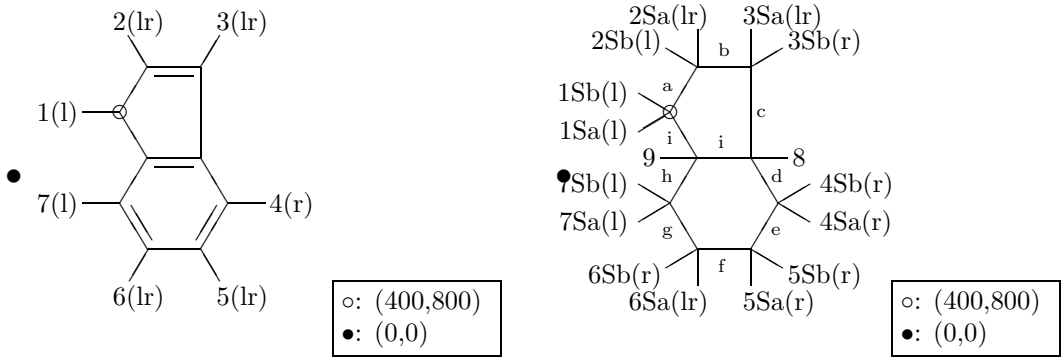
produce the following structures:



The macro `\indanehi` is the inverse counterpart of `\indaneh`, which aims at drawing indane derivatives of inverse horizontal type (lowcycle.sty).

```
\indanehi[BONDLIST]{SUBSLIST}
```

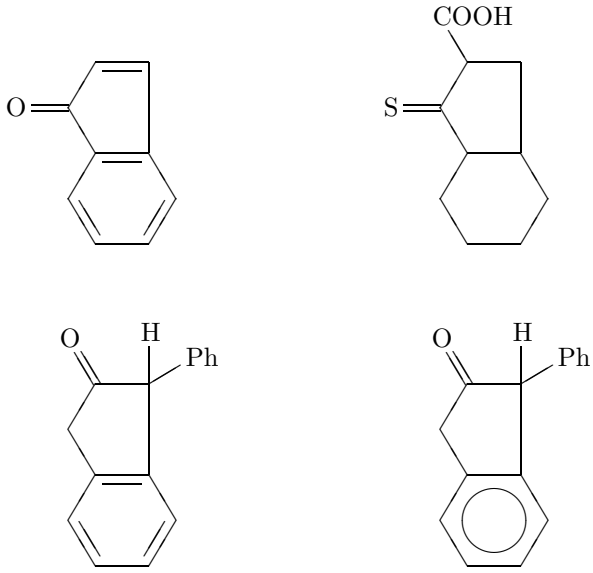
Locant numbers (1–9) for designating substitution positions and characters (a–j) for describing double bonds are shown in the following diagram:



Example:

```
\indanehi{1D==O} \qquad
\indanehi[H]{1D==S;2==COOH}\par
\indanehi[egj]{2D==O;3Sa==H;3Sb==Ph}\qquad
\indanehi[A]{2D==O;3Sa==H;3Sb==Ph}
```

produce the following structures:


O
COOH
S
O
H
Ph
O
H
Ph

Chapter 7

Six-Membered Heterocycles

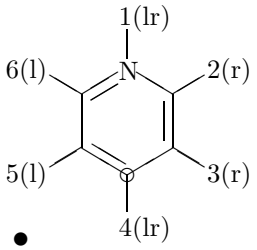
7.1 Drawing Vertical Forms of Six-Membered Heterocycles

7.1.1 Commands for Specified Use

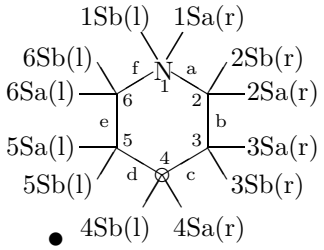
The macro `\pyridinev` and related macros are used to draw six-membered heterocyclic compounds of vertical type (hetarom.sty). Each of these commands typesets heterocycles with the specific arrangement of heteroatoms on its skeleton. The formats of these commands are as follows:

```
\pyridinev[BONDLIST]{SUBSLIST}
\pyrazinev[BONDLIST]{SUBSLIST}
\pyrimidinev[BONDLIST]{SUBSLIST}
\pyridazinev[BONDLIST]{SUBSLIST}
\triazinev[BONDLIST]{SUBSLIST}
```

By using the command `\pyridinev` as an example, the mode of locant numbering for designating substitution positions is shown as follows along with the bond descriptors for assigning inner double bonds:



○: (400,240)
●: (0,0)



○: (400,240)
●: (0,0)

The optional argument BONDLIST specifies bonds to be doubled as shown in Table 7.1. Since a specific character is assigned to a specific bond of each heterocycle, the concrete meaning of the character is different from one heterocycle to another. However, the methodology is common in drawing all heterocycles so that the commands of XeLaTeX are easy to use.

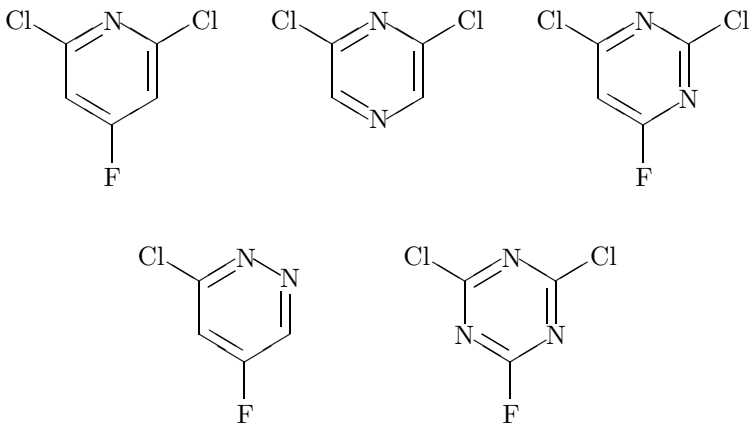
The argument SUBSLIST shows each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 6. For example, the statements,

```
\pyridinev{2==Cl;6==Cl;4==F}
\pyrazinev{2==Cl;6==Cl}
\pyrimidinev{2==Cl;6==Cl;4==F}
\pyridazinev{6==Cl;4==F}
\triazinev{2==Cl;6==Cl;4==F}
```

produce the following structures:

Table 7.1: Argument BONDLIST for commands \pyridinev, *etc.*

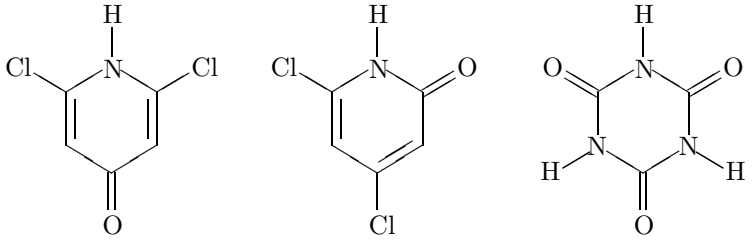
Character	Printed structure
none or r	pyridine (right-handed)
l	pyridine (left-handed)
H or [ ]	fully saturated ring
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,6-double bond
f	6,1-double bond
A	aromatic circle
{n+}	plus at the n-nitrogen atom (n = 1 to 6)



It should be noted that the default setting of the BONDLIST is to depict a fully unsaturated ring (usually an aromatic ring). By setting an appropriate character string, a sigle macro is used to typeset both partially saturated and unsaturated derivatives. Moreover, a fully saturated ring can be obtained by setting a null argument or H in BONDLIST. This specification can be illustrated with the following examples.

```
\pyridinev[be]{1==H;2==Cl;6==Cl;4D==O}\qquad
\pyridinev[ce]{1==H;4==Cl;6==Cl;2D==O}\qquad
\triazinev[H]{2D==O;4D==O;6D==O;1==H;3==H;5==H}
```

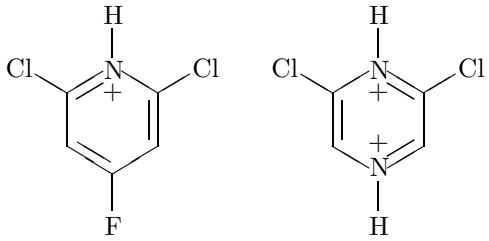
produce the following structures:



In order to depict a charge on a nitrogen, you write the statements, for example:

```
\pyridinev[r{1+}]{1==H;2==Cl;6==Cl;4==F}\qquad
\pyrazinev[l{1+}{4+}]{1==H;4==H;2==Cl;6==Cl}
```

Then you obtain the following structures:



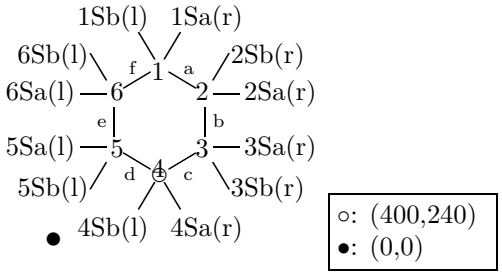
In these cases, a character 'r' or 'l' should be added to the argument BONDLIST, since the defaults are hidden by writing other characters in the BONDLIST.

7.1.2 Commands for General Use

The command `\sixheterov` is a general macro used to draw six-memered heterocyclic derivatives of vertical type (hetarom.sty). It is especially useful to draw heterocyclic compounds having other skeletal atoms than nitrogen atoms. The format of this command is as follows:

```
\sixheterov[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbers for designating substitution positions and characters for bond-description are shown in the following diagram:



Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros because of the lack of spaces to print otherwise.

The option argument BONDLIST is an character string in a pair of brackets, where each character indicates the presence of a double bond at the edge specified by the character. The bond-specification is rather arbitrary in some cases but conforms to chemical conventions as faithfully as possible if such conventions are presence (Table 7.2). Since the default prints a fully unsaturated form, an option argument [H] should be written to typeset a saturated form.

The augument ATOMLIST is a list of heteroatoms, *e.g.*, 1==N for a nitrogen atom at 1-position. It should be emphasized that, in order to typeset a heteroatom at a given position, the edges incident to the heteroatom are automatically truncated to put space for printing the heteroatrom. Compare the following examples.

The argument SUBSLIST for this macro takes a general format, in which the modifiers listed in Table 2.2 are used.

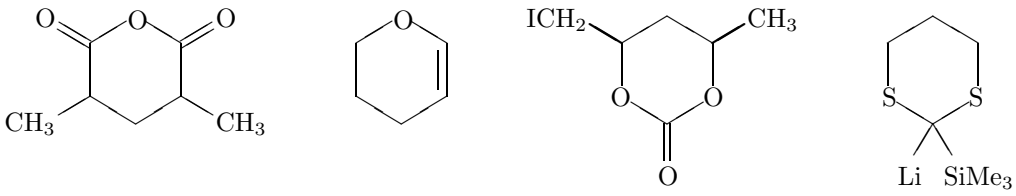
Example:

```
\sixheterov[H]{1==O}{2D==O;6D==O;3==CH$_{3}$;5==CH$_{3}$}\qquad
\sixheterov[b]{1==O}{}\qquad
\sixheterov[H]{3==O;5==O}{4D==O;6B==ICH$_{2}$;2B==CH$_{3}$}\qquad
\sixheterov[H]{3==S;5==S}{4Sa==SiMe$_{3}$;4Sb==Li}
```

Table 7.2: Argument BONDLIST for commands \sixheterov and \sixheterovi

Character	Printed structure
none or r	sixhetero (right-handed)
l	sixhetero (left-handed)
H or []	fully saturated form
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,6-double bond
f	6,1-double bond
A	aromatic circle
{n+}	plus at the n-nitrogen atom (n = 1 to 6)

produce

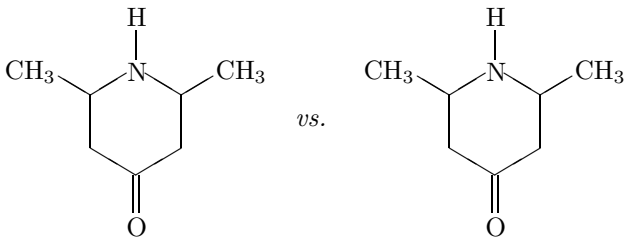


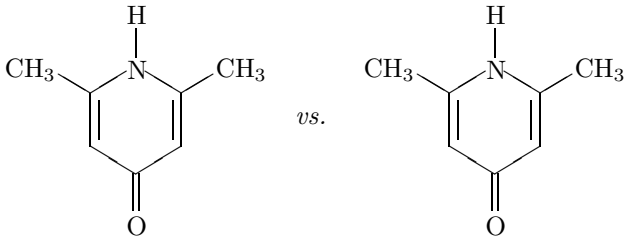
It should be noted that the same compound can be drawn in different ways. This fact is obvious because all the commands, \pyridinev, \pyrazinev, \pyrimidinev, \pyridazinev, and \triazinev, are based on the macro \sixheterov.

Example:

```
\pyridinev[H]{1==H;4D==O;2==CH$_{3}$;6==CH$_{3}$}\qquad
\raisebox{1.5cm}{\em vs.} \qquad
\sixheterov[H]{1==N}{1==H;4D==O;2==CH$_{3}$;6==CH$_{3}$} \par
\bigskip
\pyridinev[be]{1==H;4D==O;2==CH$_{3}$;6==CH$_{3}$}\qquad
\raisebox{1.5cm}{\em vs.} \qquad
\sixheterov[be]{1==N}{1==H;4D==O;2==CH$_{3}$;6==CH$_{3}$}
```

produce

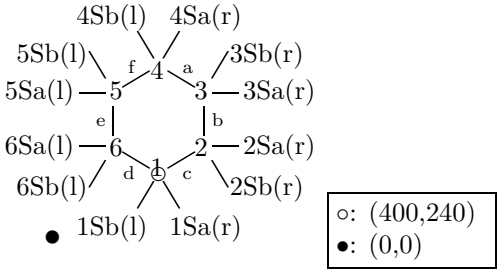




The command `\sixheterovi` is a general macro used to draw six-memered heterocyclic derivatives of inverse vertical type (hetarom.sty). The format of this command is as follows:

```
\sixheterovi[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbers (1–6) for designating substitution positions and bond descriptors (a–f) are shown in the following diagram:

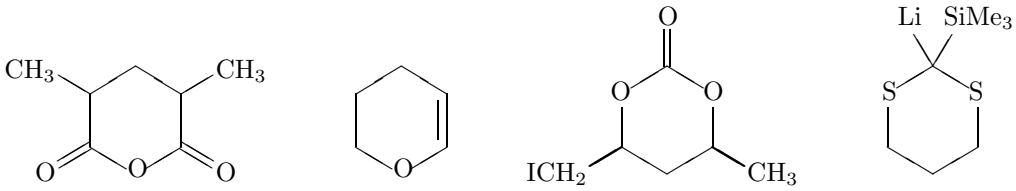


The following structures are obtained as a result of replacing the commands `\sixheterov` in the above examples by the `\sixheterovi` command.

Example:

```
\sixheterovi[H]{1==O}{2D==O;6D==O;3==CH$_{3}$;5==CH$_{3}$}\qquad
\sixheterovi[b]{1==O}{}\qquad
\sixheterovi[H]{3==O;5==O}{4D==O;6B==ICH$_{2}$;2B==CH$_{3}$}\qquad
\sixheterovi[H]{3==S;5==S}{4Sa==SiMe$_{3}$;4Sb==Li}
```

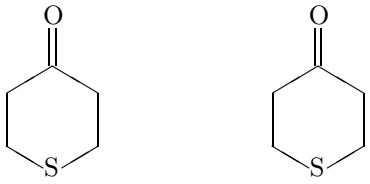
produce



The commands `\sixheterov` and `\sixheterovi` can yield the equivalent results if the modes of numbering are altered in ATOMLIST and SUBSLIST. For example, the following two statements

```
\sixheterov[H]{4==S}{1D==O} \qquad
\sixheterovi[H]{1==S}{4D==O}
```

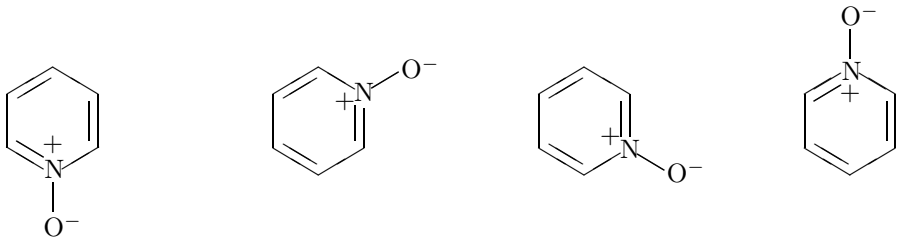
produce the same structure as follows.



However, the latter is preferred to the former because the numbering of the ring atoms comforms to the chemical nomenclature. This is the reason why we have made such macros of inverse type.

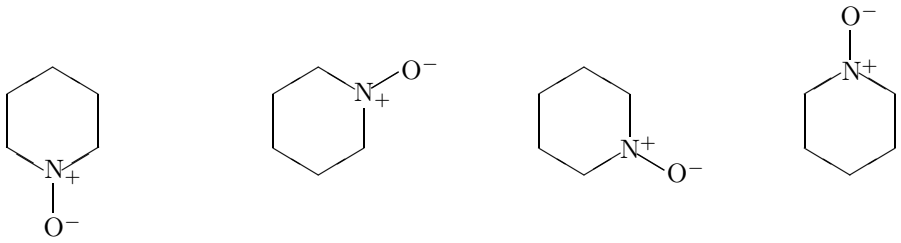
The command `\pyridinev` (or `\pyridinevi`) places a nitrogen atom on a fixed position of a pyridine ring. For printing a nitrogen atom on another position, the command `\sixheterov` should be used as shown in the following examples of drawing different formulas of pyridine N-oxide.

```
\pyridinevi[r{1+}]{1==O$^{-}$} \qquad
\sixheterov[r{2+}]{2==N}{2==O$^{-}$} \qquad
\sixheterov[r{3+}]{3==N}{3==O$^{-}$} \qquad
\pyridinev[r{1+}]{1==O$^{-}$}
```



A charge on an inner nitrogen can be alternatively typset by putting a charged atom in the ATOMLIST of the `\sixhetrov` command.

```
\sixheterov{4==N$_{+}$}{4==O$^{-}$} \qquad
\sixheterov{2==N$_{+}$}{2==O$^{-}$} \qquad
\sixheterov{3==N$^{+}$}{3==O$^{-}$} \qquad
\sixheterov{1==N$^{+}$}{1==O$^{-}$}
```



7.2 Drawing Horizontal Forms of Six-Membered Heterocycles

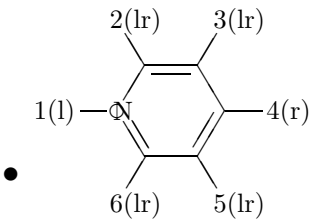
7.2.1 Commands for Specified Use

The macro `\pyridineh` and related macros are used to draw six-membered heterocyclic compounds of horizontal type (hetaromh.sty). The formats of these commands are as follows:

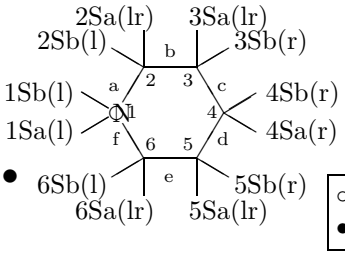
```
\pyridineh[BONDLIST]{SUBSLIST}
\pyrazineh[BONDLIST]{SUBSLIST}
```

```
\pyrimidineh[BONDLIST]{SUBSLIST}
\pyridazineh[BONDLIST]{SUBSLIST}
\triazineh[BONDLIST]{SUBSLIST}
```

The following diagrams show the numbering for designating substitution positions as well as the bond specification for placing double bonds:



○: (240,400)
●: (0,0)

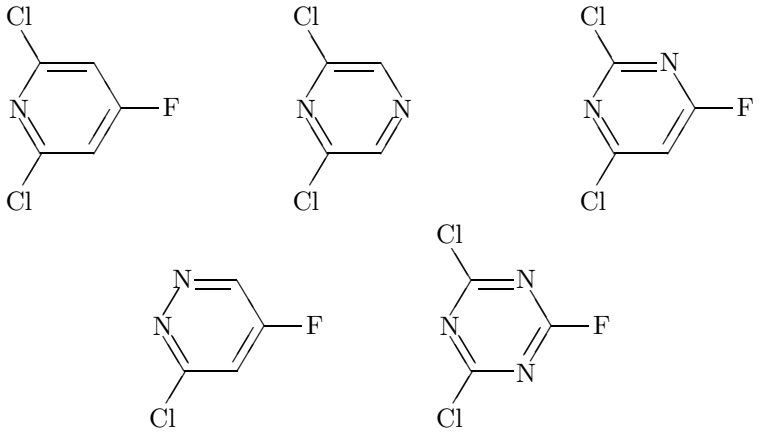


○: (240,400)
●: (0,0)

Each macro can be used to typeset both saturated and unsaturated derivatives. For example, the statements,

```
\pyridineh{2==Cl;6==Cl;4==F}
\pyrazineh{2==Cl;6==Cl}
\pyrimidineh{2==Cl;6==Cl;4==F}
\pyridazineh{6==Cl;4==F}
\triazineh{2==Cl;6==Cl;4==F}
```

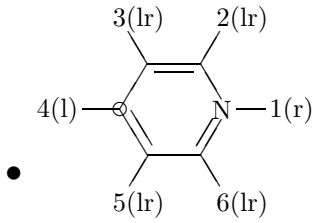
produce the following structures:



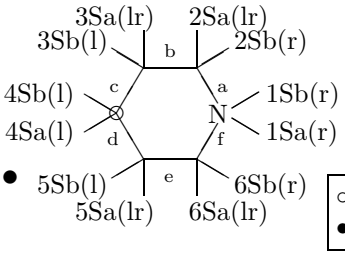
The macro `\pyridinehi` and related macros are used to draw six-membered heterocyclic compounds of inverse horizontal type (hetarom.sty). The formats of these commands are as follows:

```
\pyridinehi[BONDLIST]{SUBSLIST}
\pyrazinehi[BONDLIST]{SUBSLIST}
\pyrimidinehi[BONDLIST]{SUBSLIST}
\pyridazinehi[BONDLIST]{SUBSLIST}
\triazinehi[BONDLIST]{SUBSLIST}
```

The numbering for designating substitution positions and the bond specification are shown in the following diagrams:



○: (240,400)
●: (0,0)

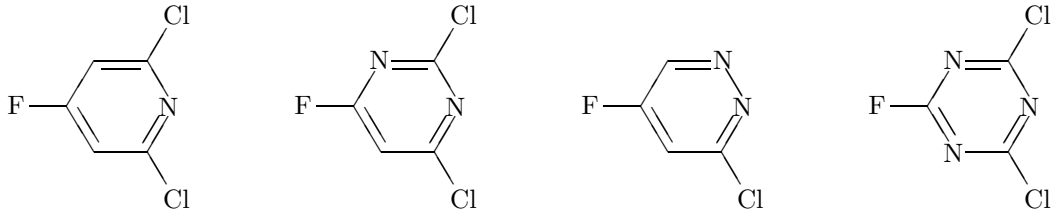


○: (240,400)
●: (0,0)

Each macro can typeset both saturated and unsaturated derivatives, where the default produces a fully unsaturated (aromatic) one. For example, the statements,

```
\pyridinehi{2==Cl;6==Cl;4==F}
\pyrimidinehi{2==Cl;6==Cl;4==F}
\pyridazinehi{6==Cl;4==F}
\triazinehi{2==Cl;6==Cl;4==F}
```

produce the following structures:

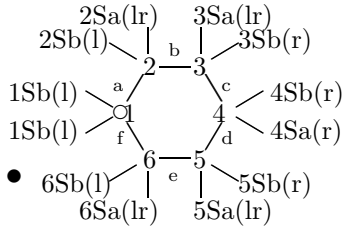


7.2.2 Commands for General Use

The macro `\sixheteroh` is a general user command for drawing six-membered heterocycles of horizontal type (hetaromh.sty). The format of this command is as follows:

```
\sixheteroh[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbers for designating substitution positions as well as bond descriptors are shown in the following diagram:



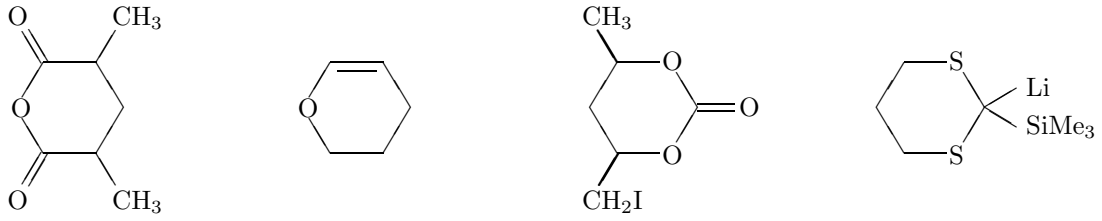
○: (240,400)
●: (0,0)

Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros. The SUBSLIST and the BONDLIST format are shown in Table 2.2 and 7.2, respectivey.

Example:

```
\sixheteroh[H]{1==O}{2D==O;6D==O;3==CH$_{3}$;5==CH$_{3}$}\qquad
\sixheteroh[b]{1==O}{}\qquad
\sixheteroh[H]{3==O;5==O}{4D==O;6B==CH$_{2}$I;2B==CH$_{3}$}\qquad
\sixheteroh[H]{3==S;5==S}{4Sa==SiMe$_{3}$;4Sb==Li}
```

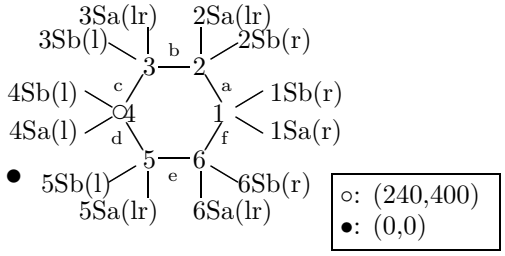
produce



The macro `\sixheterohi` defined in hetaromh.sty is used to draw six-membered hetrocycles of inverse horizontal type. The format of this command is as follows:

```
\sixheterohi[BONDLIST]{ATOMLIST}{SUBSLIST}
```

The following diagram shows locant numbers (1–6) for designating substitution positions along with bond descriptors (a–f) for setting double bonds.

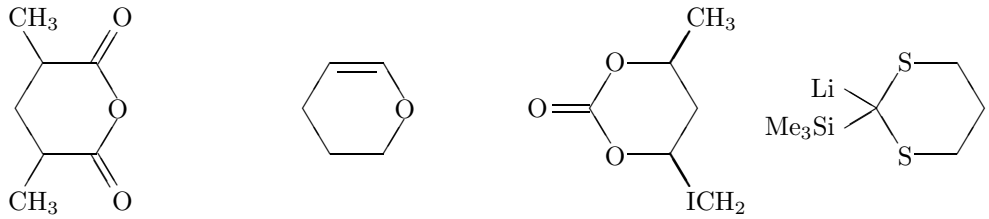


Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros. The SUBSLIST and the BONDLIST format are shown in Table 2.2 and Table 7.2, respectively.

Example:

```
\sixheterohi[H]{1==O}{2D==O;6D==O;3==CH$_{3}$;5==CH$_{3}$}\qquad
\sixheterohi[b]{1==O}{}\qquad
\sixheterohi[H]{3==O;5==O}{4D==O;6B==ICH$_{2}$;2B==CH$_{3}$}\qquad
\sixheterohi[H]{3==S;5==S}{4Sa==Me$_{3}$Si;4Sb==Li}
```

produce



Compare these structures by `\sixheterohi` with the above counterparts obtained by `\sixheteroh`.

Chapter 8

Five- or Lower-Membered Heterocycles

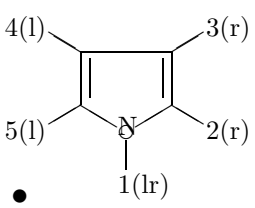
8.1 Drawing Vertical Forms of Five-Membered Heterocycles

8.1.1 Commands for Specified Use

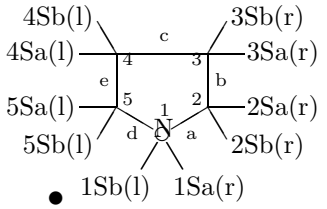
The macro `\pyrrolev` and related macros typeset five-membered heterocyclic compounds of vertical type (hetarom.sty). The formats of these commands are as follows:

```
\pyrrolev[BONDLIST]{SUBSLIST}
\pyrazolev[BONDLIST]{SUBSLIST}
\imidazolev[BONDLIST]{SUBSLIST}
\isoxazolev[BONDLIST]{SUBSLIST}
\oxazolev[BONDLIST]{SUBSLIST}
```

The following diagrams show the numbering for designating substitution positions as well as the bond specification for writing double bonds:



○: (400,240)
●: (0,0)



○: (400,240)
●: (0,0)

Each of the macros is capable of typesetting both saturated and unsaturated derivatives. The optional argument BONDLIST specifies bonds to be doubled as shown in Table 8.1. The default setting is to produce a fully unsaturated ring; on the other hand a null augument or H in BONDLIST produces a fully saturated ring.

The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 5. For example, the statements,

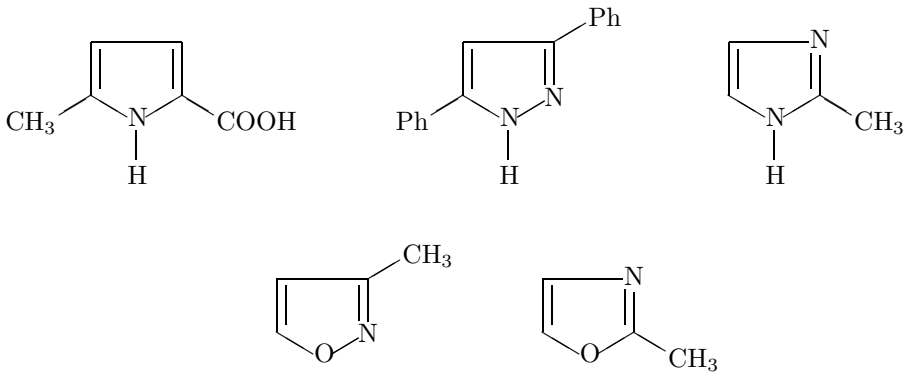
```
\pyrrolev{1==H;2==COOH;5==CH$_{3}$}\qquad\qquad\qquad
\pyrazolev{1==H;3==Ph;5==Ph}\qquad
\imidazolev{1==H;2==CH$_{3}$}
```

Table 8.1: Argument BONDLIST for commands \pyrrolev, *etc.*

Character	Printed structure
none	mother nucleus
H or [ ]	fully saturated form
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,1-double bond
A	aromatic circle
{n+}	plus at the n-nitrogen atom (n = 1 to 5)

```
\isoxazolev{3==CH$_{3}$}\qquad
\oxazolev{2==CH$_{3}$}
```

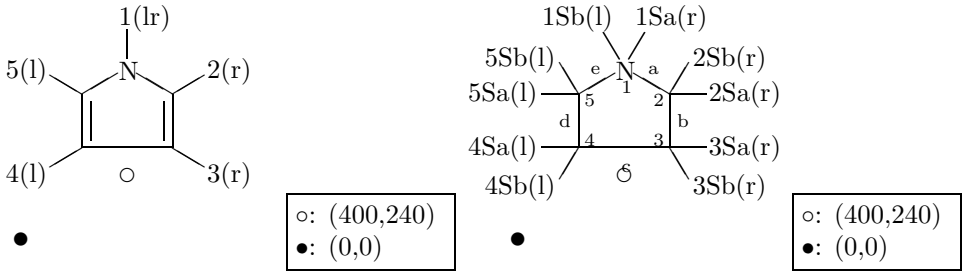
produce the following structures:



The macro \pyrrolevi and related macros are used to draw five-membered heterocyclic compounds of inverse vertical type (hetarom.sty). The formats of these commands are as follows:

```
\pyrrolevi[BONDLIST]{SUBSLIST}
\pyrazolevi[BONDLIST]{SUBSLIST}
\imidazolevi[BONDLIST]{SUBSLIST}
\isoxazolevi[BONDLIST]{SUBSLIST}
\oxazolevi[BONDLIST]{SUBSLIST}
```

The locant numbering and the bond specification are shown in the following diagrams.

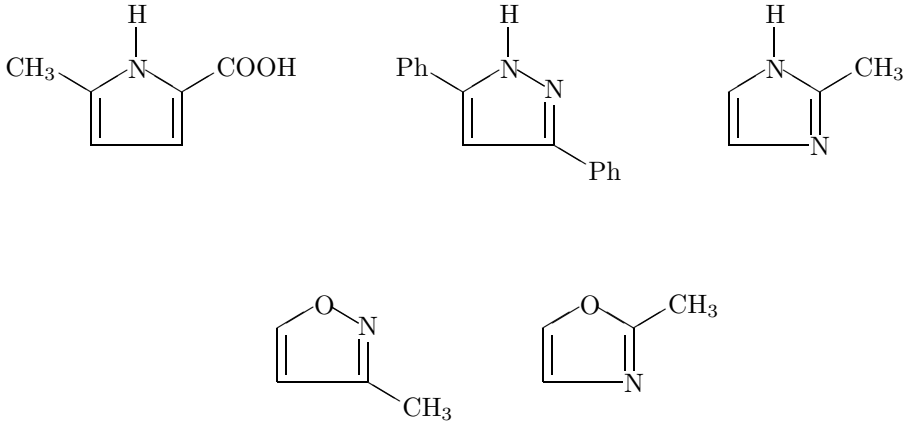


The arguments BONDLIST and SUBSLIST have the same formats as above (Tables 8.1 and 2.2).

Example:

```
\pyrrolevi{1==H;2==COOH;5==CH$_{3}$}\qquad\qquad\qquad
\pyrazolevi{1==H;3==Ph;5==Ph}\qquad
\imidazolevi{1==H;2==CH$_{3}$}\par
\isoxazolevi{3==CH$_{3}$}\qquad
\oxazolevi{2==CH$_{3}$}
```

produce the following structures:

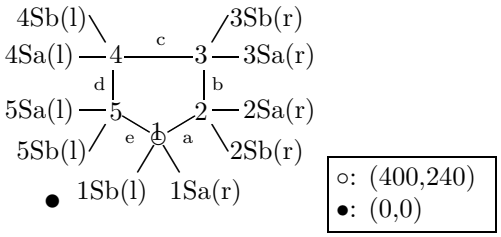


8.1.2 Commands for General Use

The command `\fiveheterov` is a general macro used to draw five-memered heterocyclic derivatives of vertical type (hetarom.sty). The format of this command is as follows:

```
\fiveheterov[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbers for designating substitution positions and bond descriptors for setting double bonds are represented by the following diagram:



Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros.

The option argument BONDLIST is a character string in a pair of brackets, where each character indicates the presence of a double bond at the edge specified by the character (Table 8.2).

Since the default of BONDLIST prints a fully saturated form, the `\fiveheterov` requires no option argument [H] in contrast to `\sixheterov`.

The argument SUBSLIST for this macro takes a general format, in which the modifiers listed in Table 2.2 are used.

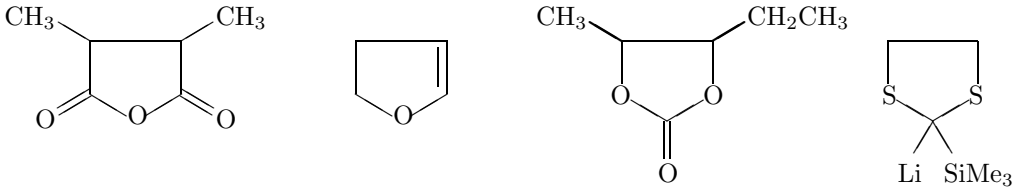
Example:

Table 8.2: Argument BONDLIST for commands \fiveheterov and \fiveheterovi

Character	Printed structure
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,1-double bond
A	aromatic circle
{n+}	plus at the n-nitrogen atom (n = 1 to 6)
{0+}	plus (or minus) at the center

```
\fiveheterov{1==O}{2D==O;5D==O;3==CH$_{3}$;4==CH$_{3}$}\qquad
\fiveheterov[b]{1==O}{}\qquad
\fiveheterov{2==O;5==O}{1D==O;3B==CH$_{2}$CH$_{3}$;4B==CH$_{3}$}\qquad
\fiveheterov{2==S;5==S}{1Sa==SiMe$_{3}$;1Sb==Li}
```

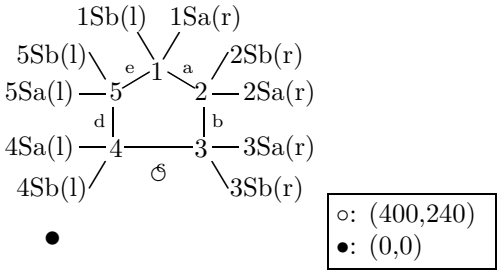
produce



The macro \fiveheterovi is employed to draw five-membered heterocyclic compounds of inverse vertical type (hetarom.sty). The format of this command is as follows:

```
\fiveheterovi[BONDLIST]{ATOMLIST}{SUBSLIST}
```

The following diagram shows the numbering for designating substitution positions:



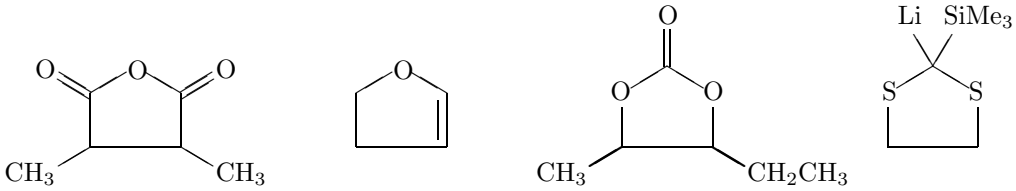
The optional argument BONDLIST specifies double bonds as shown in Table 8.2, where any combinations of characters (a–d) enable us to draw both saturated and unsaturated derivatives.

The argument SUBSLIST specifies each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 5.

Example:

```
\fiveheterovi{1==O}{2D==O;5D==O;3==CH$_{3}$;4==CH$_{3}$}\qquad
\fiveheterovi[b]{1==O}{}\qquad
\fiveheterovi{2==O;5==O}{1D==O;3B==CH$_{2}$CH$_{3}$;4B==CH$_{3}$}\qquad
\fiveheterovi{2==S;5==S}{1Sa==SiMe$_{3}$;1Sb==Li}
```

produce



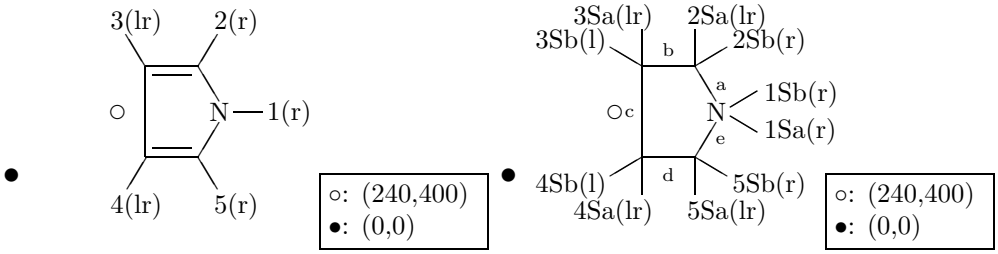
8.2 Drawing Horizontal Forms of Five-Membered Heterocycles

8.2.1 Commands for Specified Use

The macro `\pyrroleh` and related macros are used to draw five-membered heterocyclic compounds of horizontal type (hetaromh.sty). The formats of these commands are as follows:

```
\pyrroleh[BONDLIST]{SUBSLIST}
\pyrazoleh[BONDLIST]{SUBSLIST}
\imidazoleh[BONDLIST]{SUBSLIST}
\isoxazoleh[BONDLIST]{SUBSLIST}
\oxazoleh[BONDLIST]{SUBSLIST}
```

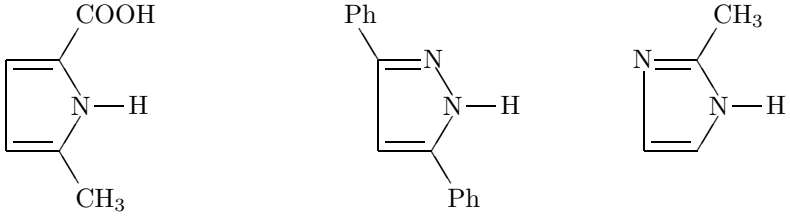
The following diagrams show the numbering for designating substitution positions:



For BONDLIST, see Table 8.2. For example, the statements,

```
\pyrroleh{1==H;2==COOH;5==CH$_{3}$}\qquad\qquad\qquad
\pyrazoleh{1==H;3==Ph;5==Ph}\qquad
\imidazoleh{1==H;2==CH$_{3}$} \par
\isoxazoleh{3==CH$_{3}$}\qquad
\oxazoleh{2==CH$_{3}$}
```

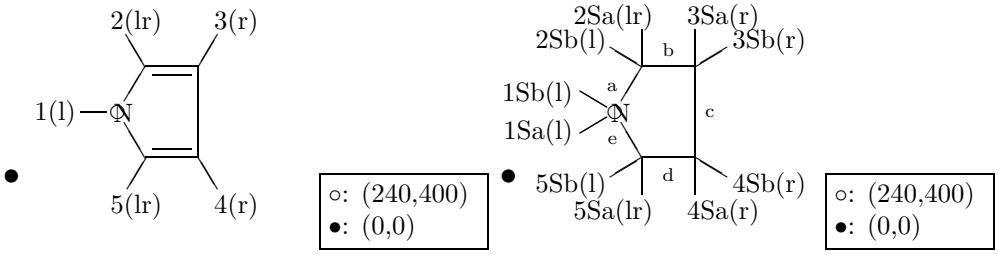
produce the following structures:



The macro \pyrrolehi and related macros are used to draw five-membered heterocyclic compounds of inverse horizontal type (hetaromh.sty). The formats of these commands are as follows:

```
\pyrrolehi[BONDLIST]{SUBSLIST}
\pyrazolehi[BONDLIST]{SUBSLIST}
\imidazolehi[BONDLIST]{SUBSLIST}
\isoxazolehi[BONDLIST]{SUBSLIST}
\oxazolehi[BONDLIST]{SUBSLIST}
```

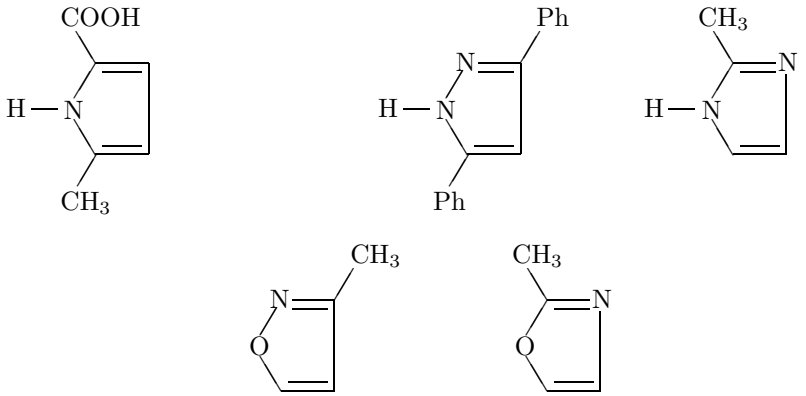
The locant numbering for designating substitution positions and the bond specification for setting double bonds are shown in the following diagrams:



For example, the statements,

```
\pyrrolehi{1==H;2==COOH;5==CH$_{3}$}\qquad\qquad\qquad
\pyrazolehi{1==H;3==Ph;5==Ph}\qquad
\imidazolehi{1==H;2==CH$_{3}$}\par
\isoxazolehi{3==CH$_{3}$}\qquad
\oxazolehi{2==CH$_{3}$}
```

produce the following structures:

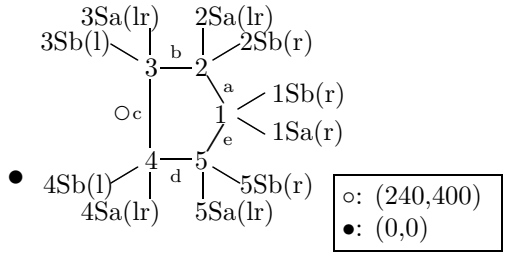


8.2.2 Commands for General Use

The command `\fiveheteroh` is a general macro used to draw five-memered heterocyclic derivatives of horizontal type (hetaromh.sty). The format of this command is as follows:

```
\fiveheteroh[BONDLIST]{ATOMLIST}{SUBSLIST}
```

The locant numbers for designating substitution positions are represented by the following diagram:



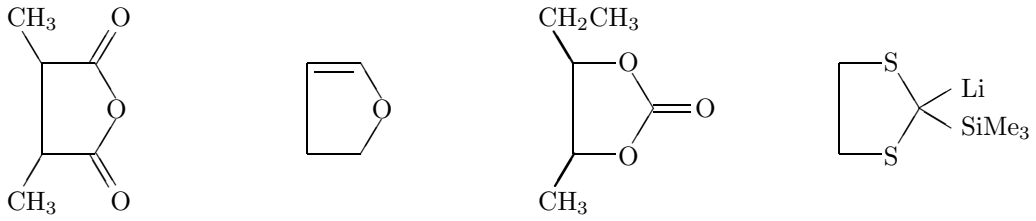
Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros.

The option argument BONDLIST is an character string in a pair of brackets, where each character indicates the presence of a double bond at the edge corresponding to the character (Table 8.2). The argument ATOMLIST lists a set of atoms placed on the edges assigned. The argument SUBSLIST for this macro takes a general format, in which the modifiers listed in Table 2.2 are used.

Example:

```
\fiveheteroh{1==O}{2D==O;5D==O;3==CH$_{3}$;4==CH$_{3}$}\qquad
\fiveheteroh[b]{1==O}{}\qquad
\fiveheteroh{2==O;5==O}{1D==O;3B==CH$_{2}$CH$_{3}$;4B==CH$_{3}$}\qquad
\fiveheteroh{2==S;5==S}{1Sa==SiMe$_{3}$;1Sb==Li}
```

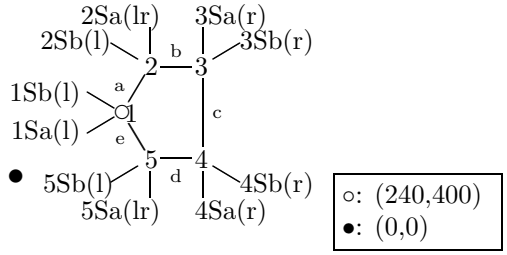
produce



The command `\fiveheterohi` is a general macro for drawing five-memered heterocyclic derivatives of inverse horizontal type (hetaromh.sty). The format of this command is as follows:

```
\fiveheterohi[BONDLIST]{ATOMLIST}{SUBSLIST}
```

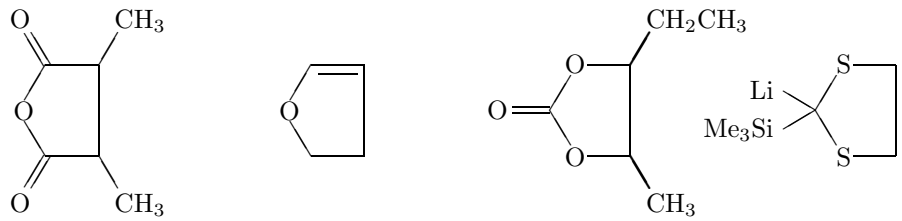
The following diagram shows the numbering for designating substitution positions as well as the bond specification for writing double bonds:



Example:

```
\fiveheterohi{1==O}{2D==O;5D==O;3==CH$_{3}$;4==CH$_{3}$}\qquad
\fiveheterohi[b]{1==O}{}\qquad
\fiveheterohi{2==O;5==O}{1D==O;3B==CH$_{2}$CH$_{3}$;4B==CH$_{3}$}\qquad
\fiveheterohi{2==S;5==S}{1Sa==Me$_{3}$Si;1Sb==Li}
```

produce the following structures:



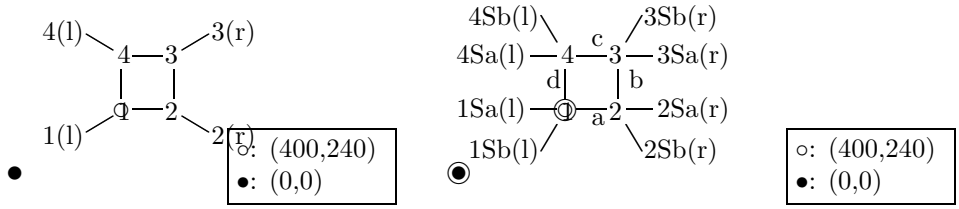
Thus, the use of `\fiveheterohi` and `\fiveheteroh` with the same arguments produces the same structural formulas with opposite directions.

8.3 Drawing Four-Membered Heterocycles

The macro `\fourhetero` is a command for general use, which is capable of giving skeletal atoms as an ATOMLIST. This macro is designed for drawing four-membered heterocycles by using the following format (hetarom.sty).

```
\fourhetero[BONDLIST]{ATOMLIST}{SUBSLIST}
```

The locant numbering is common in these commands as shown in the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument BONDLIST is used for the bond specification shown in Table 8.3.

Table 8.3: Argument BONDLIST for commands `\fourhetero` and others

Character	Printed structure	Character	Printed structure
none	mother compound (fully saturated)		
a	1,2-double bond	b	2,3-double bond
c	3,4-double bond	d	4,1-double bond
{n+}	plus at the n-nitrogen atom (n = 1 to 4)		

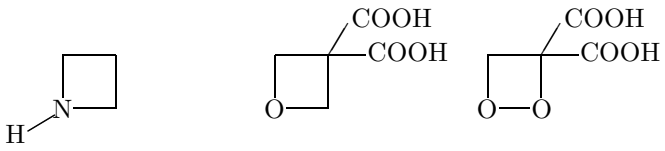
The argument ATOMLIST takes a usual format with respect to heteroatoms attached to $n = 1$ to 4, *e.g.*, 1==N for a nitrogen atom at 1-position.

The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 4.

Example:

```
\fourhetero{1==N}{1==H}
\fourhetero{1==O}{3Sa==COOH;3Sb==COOH}
\fourhetero{1==O;2==O}{3Sa==COOH;3Sb==COOH}
```

produce the following structures:

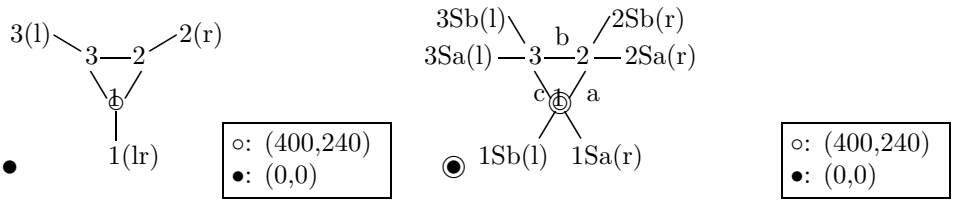


8.4 Drawing Three-Membered Heterocycles

The macro `\threehetero`, which is defined in hetarom.sty for drawing three-membered heterocycles, has the following format:

```
\threehetero[BONDLIST]{ATOMLIST}{SUBSLIST}
```

The locant numbering is common as shown in the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument BONDLIST specifies double bonds as shown in Table 8.4.

The argument ATOMLIST takes a usual format with respect to heteroatoms attached to $n = 1$ to 3, *e.g.*, 1==N for a nitrogen atom at 1-position.

The argument SUBSLIST describes each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 3.

Example:

```
\threehetero{1==N}{1==H;2Sa==COOCH$_{3}$;2Sb==COOCH$_{3}$}\qquad
\threehetero{1==O}{2Sa==COOH;2Sb==COOH}\qquad\qquad
\threehetero{1==S}{3Sa==H$_{3}$C;3Sb==H$_{3}$C}
```

produce the following structures:

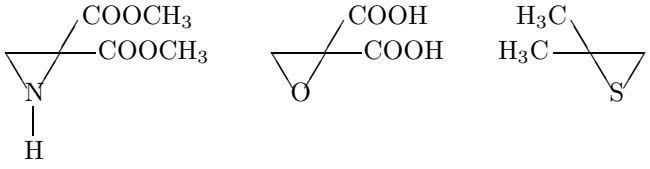


Table 8.4: Argument BONDLIST for commands \threehetero and others

Character	Printed structure
none	saturated
a	1,2-double bond
b	2,3-double bond
c	3,1-double bond
A	aromatic circle
{n+}	plus at the n-hetero atom (n = 1 to 3)
	$n = 4$ – outer plus at 1 position
	$n = 5$ – outer plus at 2 position
	$n = 6$ – outer plus at 3 position
{0+}	plus at the center of a cyclopropane ring

Chapter 9

Heterocycles with Fused Six-to-Six-Membered Rings

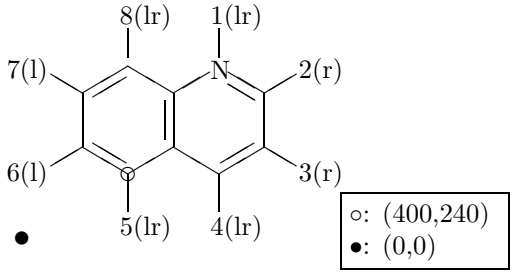
9.1 Drawing Vertical Forms

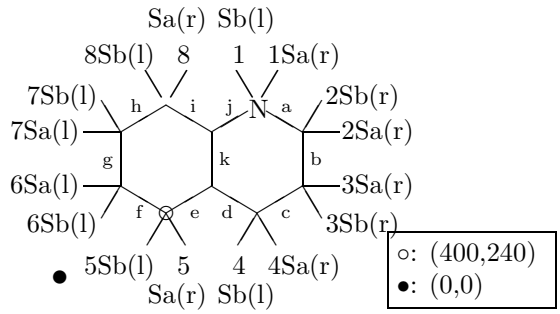
9.1.1 Commands for Specified Use

The macro `\quinolinev` is used to draw quinoline derivatives of vertical type (hetarom.sty). Macros for drawing other fused heterocycles are also defined. The formats of these commands are as follows:

```
\quinolinev[BONDLIST]{SUBSLIST}
\isoquinolinev[BONDLIST]{SUBSLIST}
\quinoxalinev[BONDLIST]{SUBSLIST}
\quinazolinev[BONDLIST]{SUBSLIST}
\cinnolinev[BONDLIST]{SUBSLIST}
\pteridinev[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions as well as bond descriptors for setting double bonds are shown in the following diagrams:





The handedness for each oriented or double-sided position is shown with a character set (r, l, or lr) in parentheses. Each character in the optional argument BONDLIST specifies an inner (endocyclic) double bond as shown in Table 9.1.

The argument SUBSLIST is employed to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 8. Sibstitution on 9 (4a position) or 10 (8a position) can be assigned in the usual way.

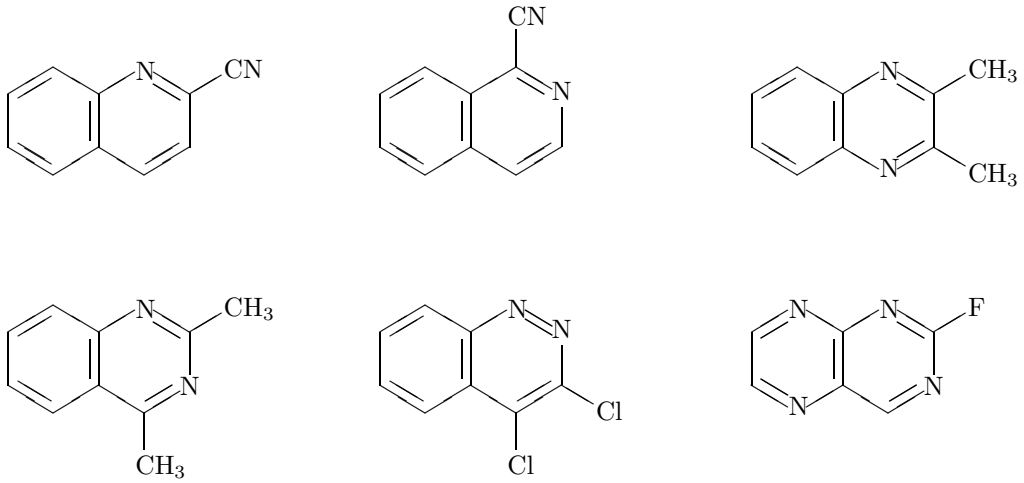
Example:

Table 9.1: Argument BONDLIST for commands \quinolinev and others

Character	Printed structure	Character	Printed structure
none or r	decahetero (right-handed)	l	decahetero (left-handed)
H or [ ]	fully saturated form		
a	1,2-double bond	b	2,3-double bond
c	4,3-double bond	d	4,4a-double bond
e	4a,5-double bond	f	5,6-double bond
g	6,7-double bond	h	7,8-double bond
i	8,8a-double bond	j	1,8a-double bond
k	4a,8a-double bond		
A	aromatic circle	B	aromatic circle
{n+}	plus at the n-nitrogen atom (n = 1 to 10)		

```
\quinolinev{2==CN} \qquad
\isoquinolinev{1==CN}\qquad
\quinoxalinev{2==CH$_{3}$;3==CH$_{3}$} \par
\quinazolinev{2==CH$_{3}$;4==CH$_{3}$}\qquad
\cinnolinev{4==Cl;3==Cl} \qquad
\pteridinev{2==F}
```

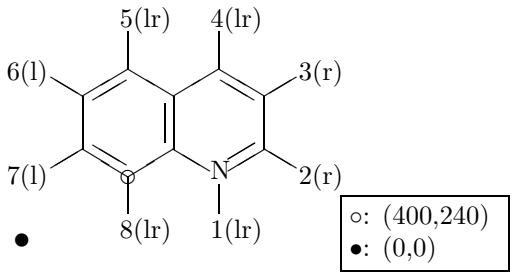
produce the following structures:

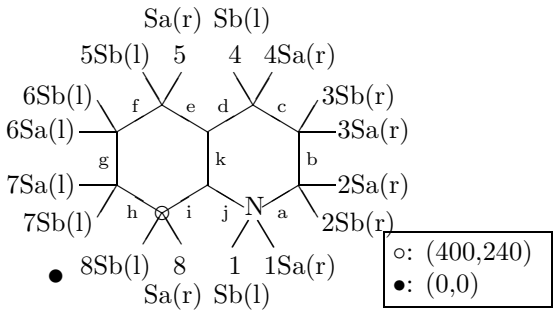


The macro \quinolinevi is used to draw quinoline derivatives of inverse vertical type (hetarom.sty). Macros for depicting other fused heterocycles are also defined. The formats of these commands are as follows:

```
\quinolinevi[BONDLIST]{SUBSLIST}
\isoquinolinevi[BONDLIST]{SUBSLIST}
\quinazolinevi[BONDLIST]{SUBSLIST}
\cinnolinevi[BONDLIST]{SUBSLIST}
\pteridinevi[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions along with bond descriptors are represented by the following diagrams:

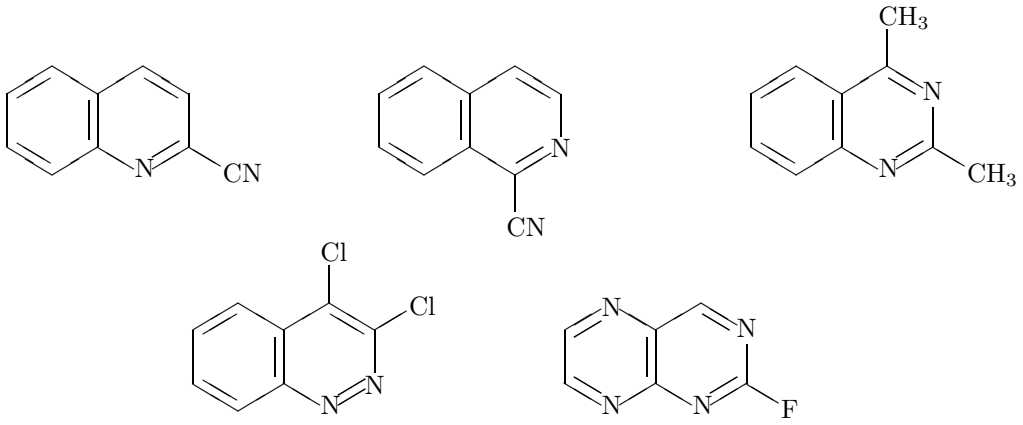




Example:

```
\quinolinevi{2==CN} \qquad
\isoquinolinevi{1==CN}\qquad
\quinazolinevi{2==CH$_{3}$;4==CH$_{3}$}\par
\cinnolinevi{4==Cl;3==Cl} \qquad
\pteridinevi{2==F}
```

produce the following structures:

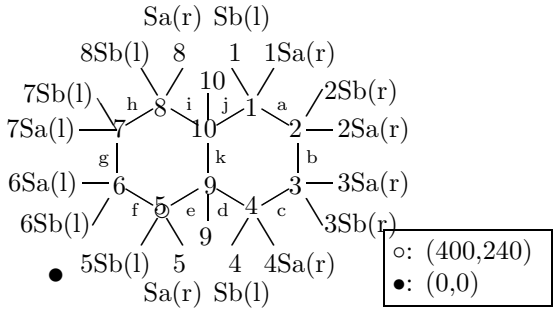


9.1.2 Commands for General Use

Such commands as `\quinolinev` (for specified use) are based on the macro `\decaheterov`, which is generally used to draw six-six-fused heterocycles of vertical type (hetarom.sty).

```
\decaheterov[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbers for designating substitution positions as well as characters for setting double bonds are shown in the following diagram:



The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument BONDLIST is used to specify a bond pattern as shown in Table 9.1. Note that the default structure is the fully unsaturated one.

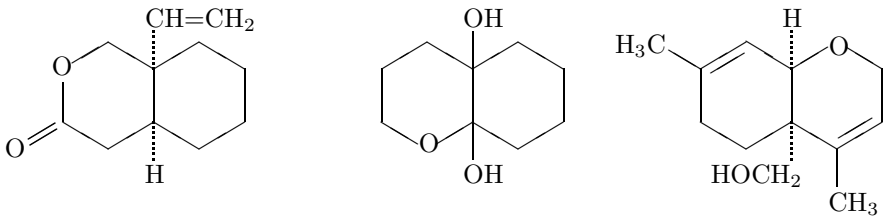
The argument ATOMLIST has a similar format concerning the positions of $n = 1$ to 8. A hetero-atom on the 4a-position is designated to be 4a==N or 9==N; and a hetero-atom on the 8a-position is given as to be 8a==N or {{10}}==N.

The argument SUBSLIST for this macro takes a general format, in which the modifiers listed in Table 2.2 are used. Note that 9 and 10 should be used for designating 4a and 8a positions.

Example:

```
\decaheterov[H]{7==O}{6D==O;9A==H;{{10}A}==CH=CH$_{2}$}
\decaheterov[H]{5==O}{9==OH;{{10}}==OH}
\decaheterov[ch]{1==O}{9A==\lmoiety{HOCH$_{2}$};{{10}A}==H;%
4==CH$_{3}$;7==H$_{3}$C}
```

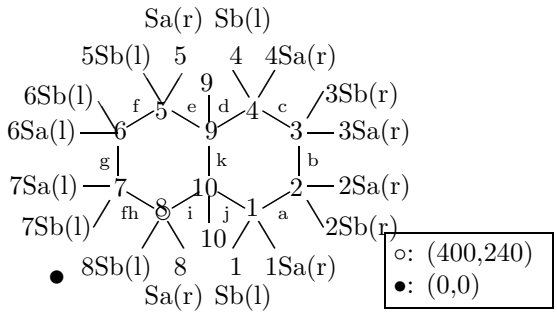
produce the following structures:



The command `\decaheterovi` defined in hetarom.sty is the inverse-type macro of `\decaheterov` described above.

```
\decaheterovi[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbers (1–10) for designating substitution positions and bond descriptors (a–f) for setting double bonds are represented by the following diagram:

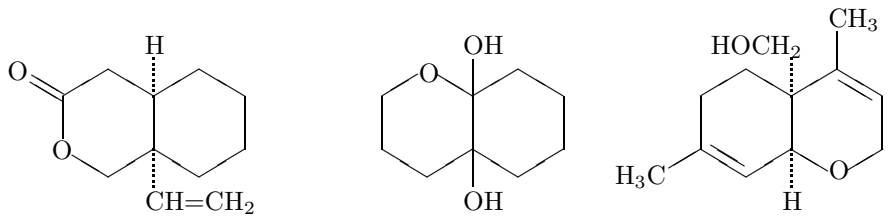


The handedness for each oriented or double-sided position is shown with a character set in parentheses. The arguments are given in the same way as in `\decaheterov`.

Example:

```
\decaheterovi[H]{7==O}{6D==O;9A==H;{{10}A}==CH=CH$_{2}$}
\decaheterovi[H]{5==O}{9==OH;{{10}}==OH}
\decaheterovi[ch]{1==O}{9A==\lmoiety{HOCH$_{2}$};{{10}A}==H;%
4==CH$_{3}$;7==H$_{3}$C}
```

produce the following structures:



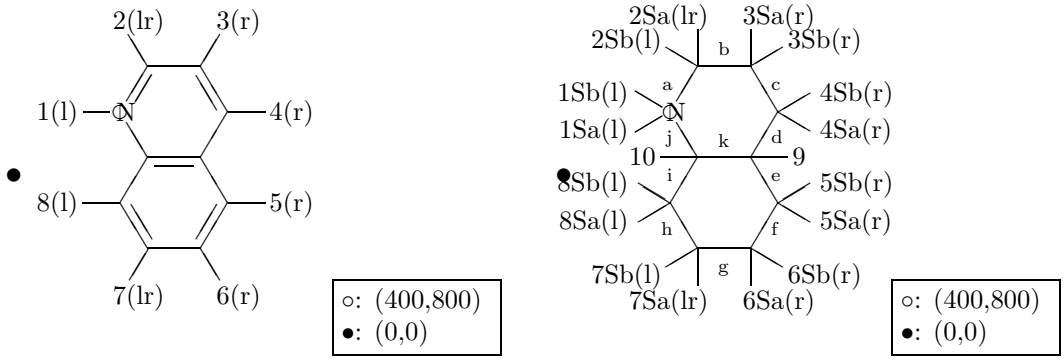
9.2 Drawing Horizontal Forms

9.2.1 Commands for Specified Use

The horizontal counterparts of the commands of vertical type (`\quinolinev`, *etc*) are defined similarly in the package file, hetaromh.sty.

```
\quinolineh[BONDLIST]{SUBSLIST}
\isoquinolineh[BONDLIST]{SUBSLIST}
\quinoxalineh[BONDLIST]{SUBSLIST}
\quinazolineh[BONDLIST]{SUBSLIST}
\cinnolineh[BONDLIST]{SUBSLIST}
\pteridineh[BONDLIST]{SUBSLIST}
```

Locant numbers (1–10) for designating substitution positions and bond descriptors (a–k) are found in the following diagram:

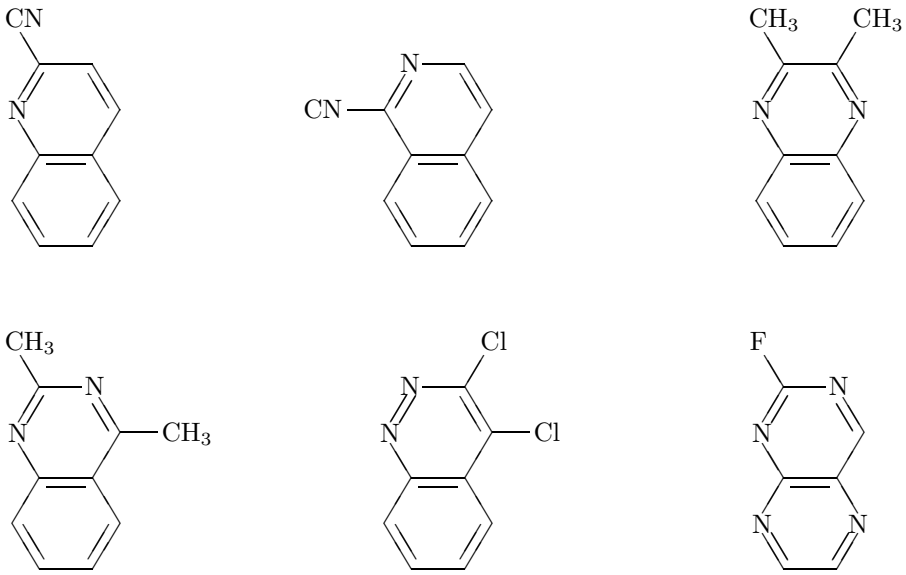


The handedness for each oriented or double-sided position is shown with a character set (r, l, or lr) in parentheses. The optional argument BONDLIST specifies double bonds to be typeset, as shown in Table 9.1.

Example:

```
\quinolineh{2==CN} \qquad
\isoquinolineh{1==CN}\qquad
\quinoxalineh{2==CH$_{3}$;3==CH$_{3}$} \par
\quinazolineh{2==CH$_{3}$;4==CH$_{3}$}\qquad
\cinnolineh{4==Cl;3==Cl} \qquad
\pteridineh{2==F}
```

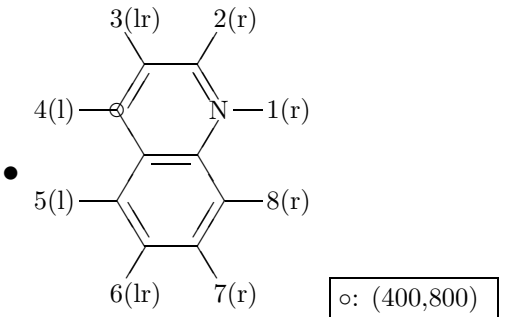
produce the following structures:

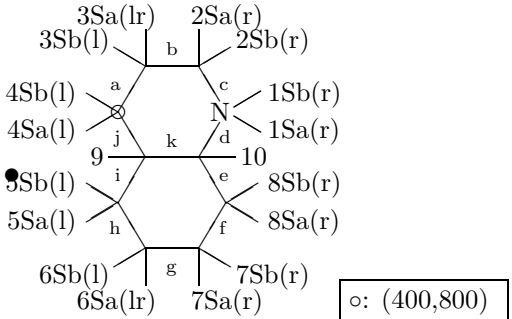


The macro `\quinolinehi` defined in the package hetaromh.sty is used to draw quinoline derivatives of inverse horizontal type. Macros for drawing other fused heterocycles are also defined in the package hetaromh.sty. The formats of these commands are as follows:

```
\quinolinehi[BONDLIST]{SUBSLIST}
\isoquinolinehi[BONDLIST]{SUBSLIST}
\quinazolinehi[BONDLIST]{SUBSLIST}
\cinnolinehi[BONDLIST]{SUBSLIST}
\pteridinehi[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions and characters (a–f) for designating bouble bonds are shown in the following diagram:

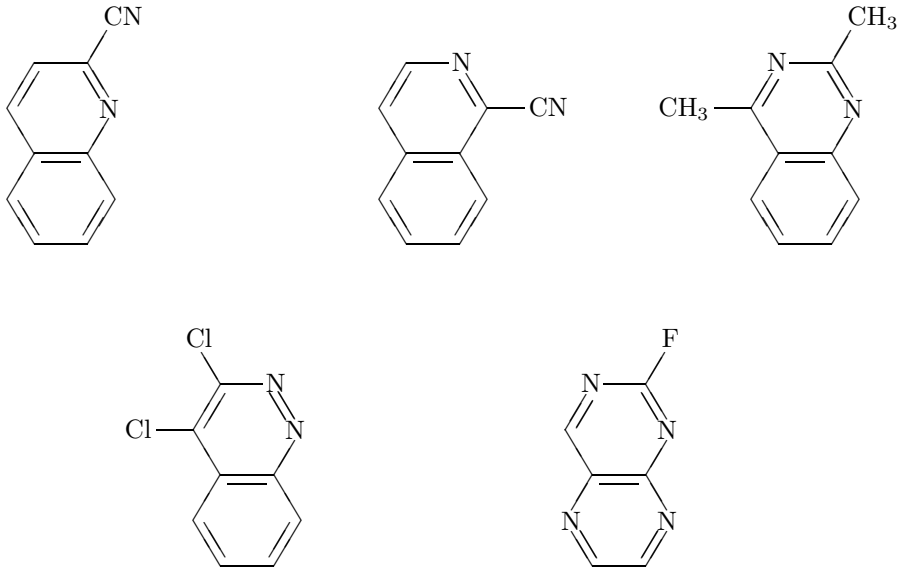




Example:

```
\quinolinehi{2==CN} \qquad
\isoquinolinehi{1==CN}\qquad
\quinazolinehi{2==CH$_{3}$;4==CH$_{3}$}\par
\cinnolinehi{4==Cl;3==Cl} \qquad
\pteridinehi{2==F}
```

produce the following structures:

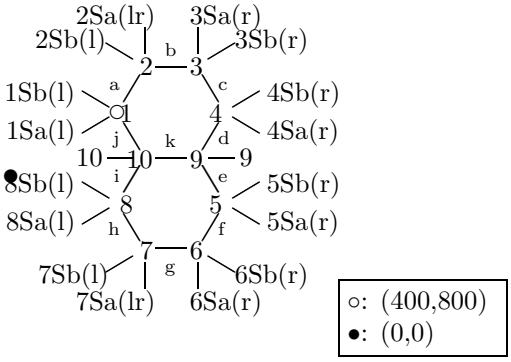


9.2.2 Commands for General Use

The macro `\decaheteroh` (carom.sty) is the horizotal counterpart of `\decaheterov`. The format and the assignment of BONDLIST and SUBSLIST of the former macro are the same as those of the latter described above (see Tables 2.2 and 9.1).

```
\decaheteroh[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbers for designating substitution positions and bond descriptors for typesetting double bonds are represented by the following diagram:

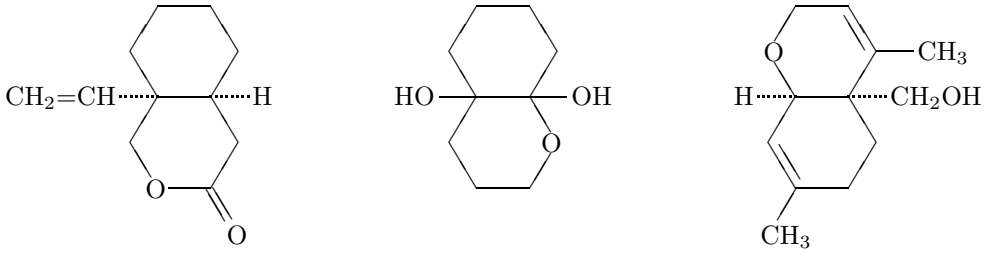


The handedness for each oriented or double-sided position is shown with a character set (r, l, or lr) in parentheses. The optional argument BONDLIST contains a characher string selected from Table 9.1.

Example:

```
\decaheteroh[H]{7==O}{6D==O;9A==H;{{10}A}==CH$_{2}$=CH}
\decaheteroh[H]{5==O}{9==OH;{{10}}==HO}
\decaheteroh[ch]{1==O}{9A==CH$_{2}$OH;{{10}A}==H;%
4==CH$_{3}$;7==CH$_{3}$}
```

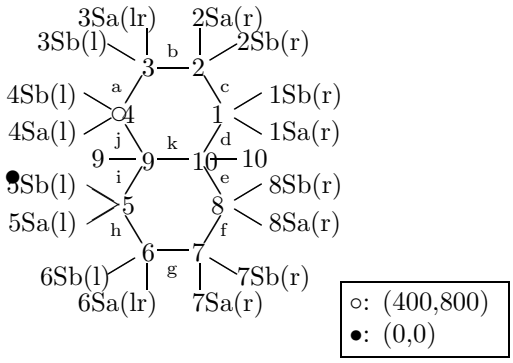
produce the following structures:



The macro `\decaheterohi` (carom.sty) is the inverse counterpart of `\decaheteroh`. The format and the assignment of BONDLIST and SUBSLIST of the former macro are the same as the latter (see Tables 2.2 and 9.1).

```
\decaheterohi[BONDLIST]{ATOMLIST}{SUBSLIST}
```

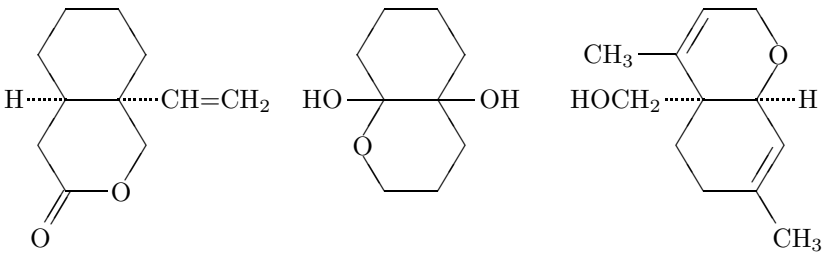
Locant numbers for designating substitution positions and bond descriptors for setting double bonds are represented by the following diagram:



Example:

```
\decaheterohi[H]{7==O}{6D==O;9A==H;{{10}A}==CH=CH$_{2}$}
\decaheterohi[H]{5==O}{9==HO;{{10}}==OH}
\decaheterohi[ch]{1==O}{9A==\lmoiety{HOCH$_{2}$};{{10}A}==H;%
4==CH$_{3}$;7==CH$_{3}$}
```

produce the following structures:



Chapter 10

Heterocycles with Fused Six-to-Five-Membered Rings

10.1 Drawing Vertical Forms

10.1.1 Commands for Specified Use

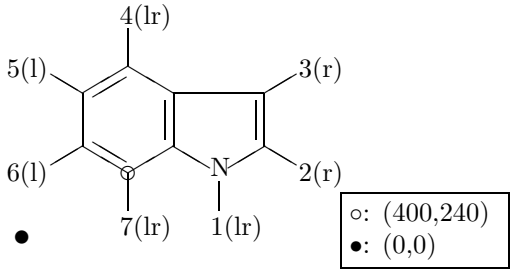
XyMTEX involves macros for drawing representative fused N-heterocycles that consist of a six- and a five-membered ring. These macros have the following formats (hetarom.sty).

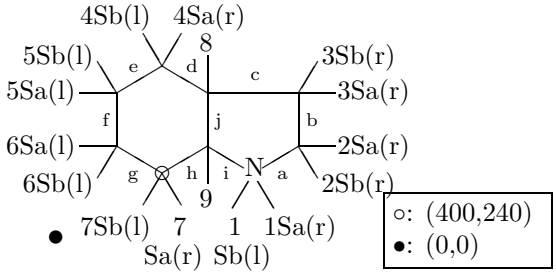
```
\indolev[BONDLIST]{SUBSLIST}
\isoindolev[BONDLIST]{SUBSLIST}
\purinev[BONDLIST]{SUBSLIST}
```

Macros for drawing fused N,O-heterocycles are also available (hetarom.sty). They are the same formats of arguments.

```
\benzofuranev[BONDLIST]{SUBSLIST}
\isobenzofuranev[BONDLIST]{SUBSLIST}
\benzoxazolev[BONDLIST]{SUBSLIST}
```

The locant numbering is common in these commands as shown in the following diagrams:





The handedness for each oriented or double-sided position is shown with a character set (r or l) in parentheses. The optional argument BONDLIST specifies edges with a double bond (Table 10.1).

The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 2.2, in which n is an arabic numeral between 1 and 7. Sibstitution on 8 (3a position) or 9 (7a position) can be assigned in the usual way of specifying bridgehead positions.

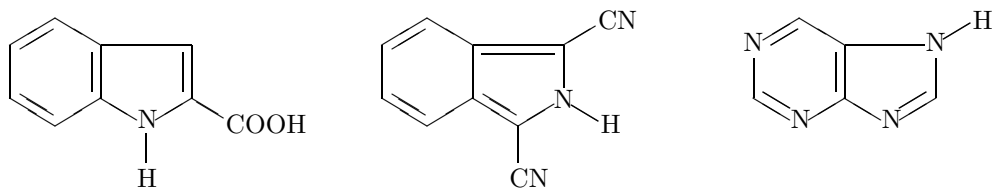
Example:

Table 10.1: Argument BONDLIST for commands \indolev and others

Character	Printed structure	Character	Printed structure
none or r	aromatic six-membered ring	H or []	fully saturated form
a	1,2-double bond	b	2,3-double bond
c	3,3a-double bond	d	4,3a-double bond
e	4,5-double bond	f	5,6-double bond
g	6,7-double bond	h	7,7a-double bond
i	1,7a-double bond	j	3a,4a-double bond
A	aromatic circle (six-membered ring)		
B	aromatic circle (five-membered ring)		
{n+}	plus at the n-nitrogen atom (n = 1 to 9)		

```
\indolev{1==H;2==COOH}\qquad
\isoindolev{2==H;1==CN;3==CN}\qquad
\purinev{3==H}
```

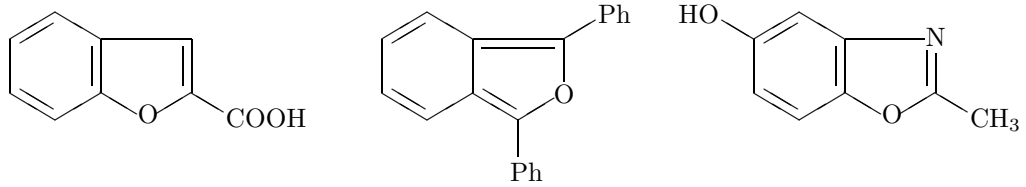
produce the following structures:



Example:

```
\benzofuranev{2==COOH}\qquad
\isobenzofuranev{1==Ph;3==Ph}\qquad
\benzoxazolev{2==CH$_{3}$;5==HO}
```

produce the following structures:



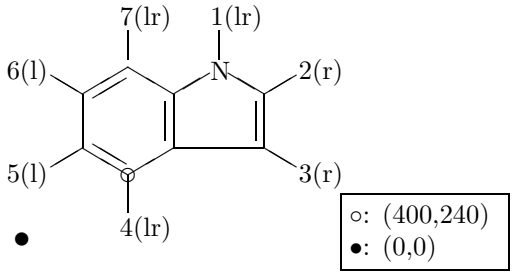
The package hetarom.sty also involves macros for drawing fused N-heterocycles of inverse vertical type. They have the following formats:

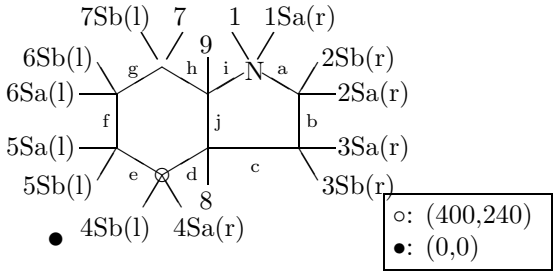
```
\indolevi[BONDLIST]{SUBSLIST}
\isoindolevi[BONDLIST]{SUBSLIST}
\purinevi[BONDLIST]{SUBSLIST}
```

Macros for drawing fused N,O-heterocycles of inverse vertical type have the following formats. They are also contained in the package file hetarom.sty.

```
\benzofuranevi[BONDLIST]{SUBSLIST}
\isobenzofuranevi[BONDLIST]{SUBSLIST}
\benzoxazolevi[BONDLIST]{SUBSLIST}
```

They are the counterparts of the commands without suffix 'i' described above. The locant numbering is common in these commands as shown in the following diagrams:



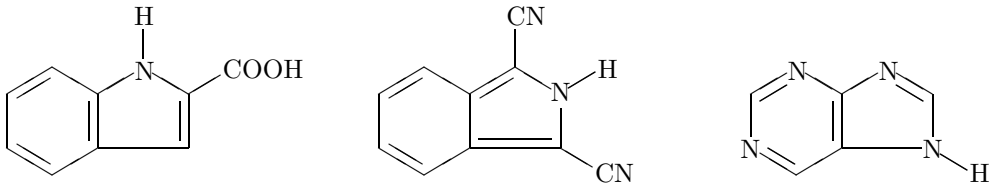


The handedness for each oriented or double-sided position is shown with a character set (l, r, or lr) in parentheses. The optional argument BONDLIST is used in a usual way (Table 10.1).

Example:

```
\indolevi{1==H;2==COOH}\qquad
\isoindolevi{2==H;1==CN;3==CN}\qquad
\purinevi{3==H}
```

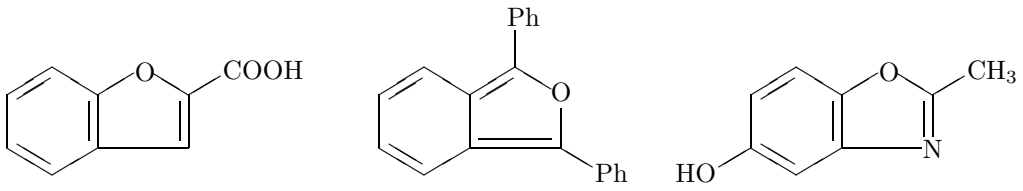
produce the following structures:



Example:

```
\benzofuranevi{2==COOH}\qquad
\isobenzofuranevi{1==Ph;3==Ph}\qquad
\benzoxazolevi{2==CH$_{3}$;5==HO}
```

produce the following structures:

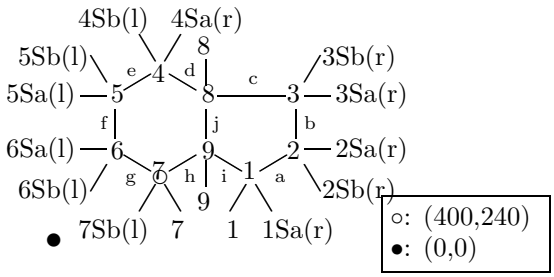


10.1.2 Commands for General Use

Macros for specified use such as `\indolev` are short-cut commands defined on the basis of a general command `\nonaheterov`. Such a general command enables us to draw a further variety of heterocyclic compounds. It has the following format, the definition of which is contained also in hetarom.sty.

```
\nonaheterov[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbers (1–9) and bond descriptors (a–j) are defined as shown in the following diagram:



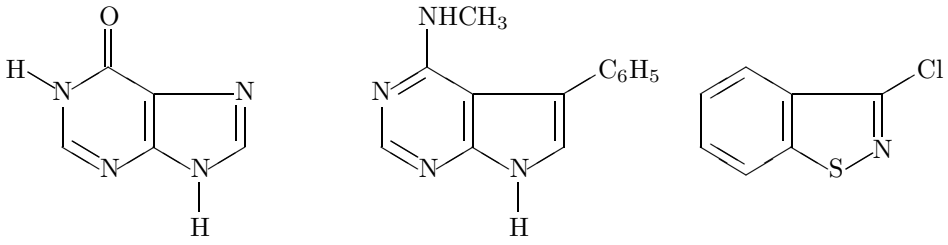
The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument BONDLIST is used to specify double bonds as shown in Table 10.1.

The argument ATOMLIST takes a usual format with respect to heteroatoms attached to $n = 1$ to 7, *e.g.*, 1==N for a nitrogen atom at 1-position. Hetero atoms at 3a- and 7a-positions are represented as to be 3a==N (or 8==N) for a nitrogen at 3a-position, 7a==N (or 9==N) for at a nitrogen at 7a-position, and so on. The augument SUBSLIST takes a usual format except that the locant numbers 3a and 7a are replaced by 8 and 9.

Example:

```
\nonaheterov[bjg]{1==N;3==N;5==N;7==N}{1==H;5==H;4D==O}
\nonaheterov[bjge]{1==N;5==N;7==N}%
 {1==H;3==C$_{6}$H$_{5}$;4==NHCH$_{3}$}
\nonaheterov[bjge]{1==S;2==N}{3==Cl}
```

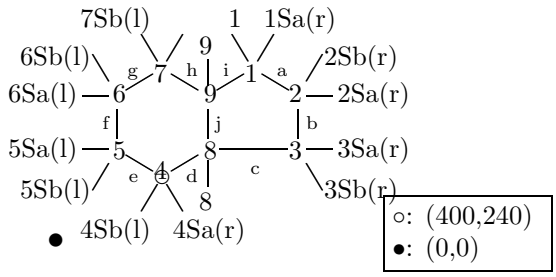
produce the following structures:



The inverse conterpart `\nonaheterovi` has the following format:

```
\nonaheterovi[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Locant numbering is usually defined as shown in the following diagram:

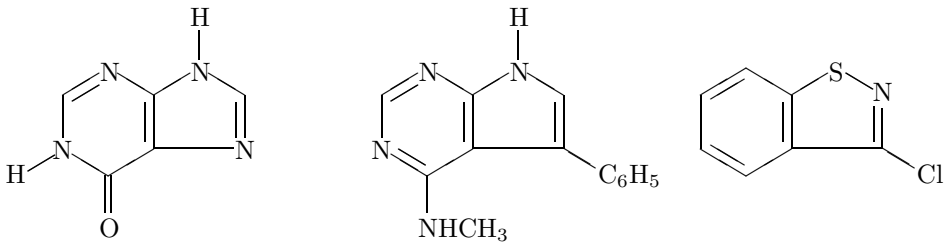


The handedness for each oriented or double-sided position is shown with a character set in parentheses. The same format of the optional argument BONDLIST is used as shown in Table 10.1.

Example:

```
\nonaheterovi[bjg]{1==N;3==N;5==N;7==N}{1==H;5==H;4D==O}
\nonaheterovi[bjge]{1==N;5==N;7==N}%
 {1==H;3==C$_{6}$H$_{5}$;4==NHCH$_{3}$}
\nonaheterovi[bjge]{1==S;2==N}{3==Cl}
```

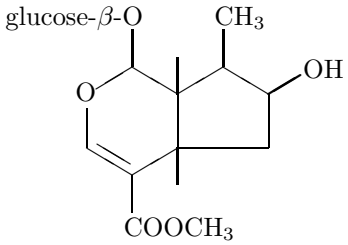
produce the following structures:



A subsituent on a bridgehead position can be designated in the SUBSLIST of the `\nonahetrovi` command (or `\nonahetrov`). The code

```
\nonaheterovi[e]{6==O}{1B==CH$_{3}$;2B==OH;4==COOCH$_{3}$;%
 7B==\lmoiety{glucose-$\beta$-O};8B==;9B==}
```

typesets the structural formula of loganin:



10.2 Drawing Horizontal Forms

10.2.1 Commands for Specified Use

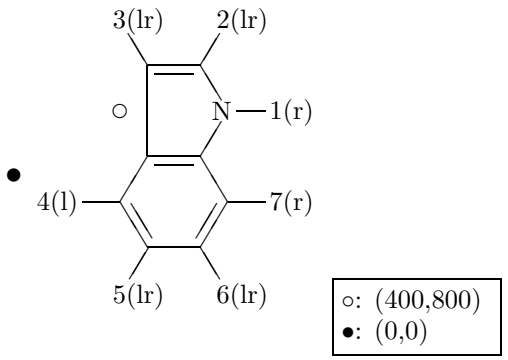
Macros for drawing N-heterocycles of horizontal type have the following formats (hetaromh.sty).

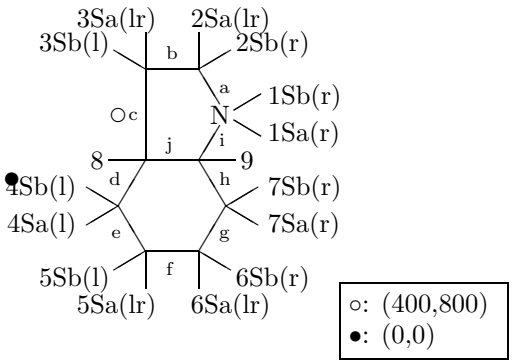
```
\indoleh[BONDLIST]{SUBSLIST}
\isoindoleh[BONDLIST]{SUBSLIST}
\purineh[BONDLIST]{SUBSLIST}
```

Macros for drawing N,O-heterocycles are available by setting the package file hetaromh.sty. They have the following formats.

```
\benzofuraneh[BONDLIST]{SUBSLIST}
\isobenzofuraneh[BONDLIST]{SUBSLIST}
\benzoxazoleh[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



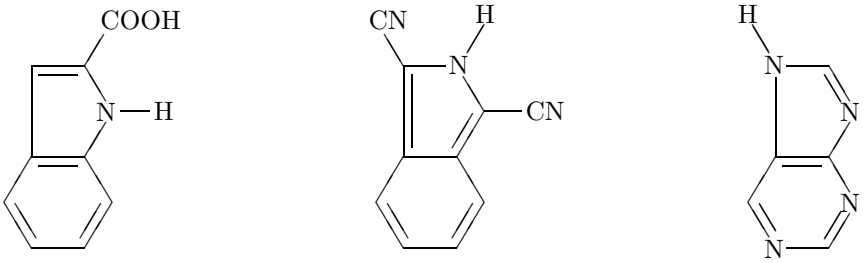


The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument BONDLIST specifies double bonds by using chracters selected from Table 9.1.

Example:

```
\indoleh{1==H;2==COOH}\qquad
\isoindoleh{2==H;1==CN;3==CN}\qquad
\purineh{3==H}
```

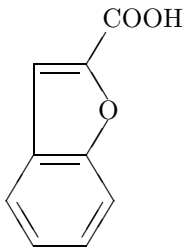
produce the following structures:

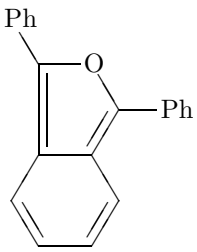


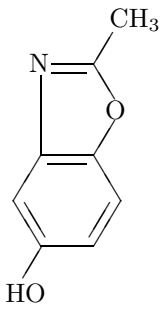
Example:

```
\benzofuraneh{2==COOH}\qquad
\isobenzofuraneh{1==Ph;3==Ph}\qquad
\benzoxazoleh{2==CH$_{3}$;5==HO}
```

produce the following structures:



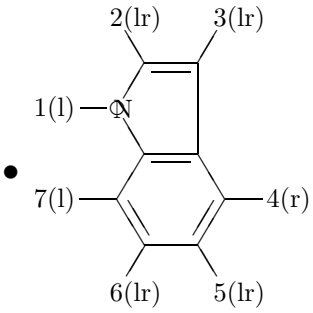




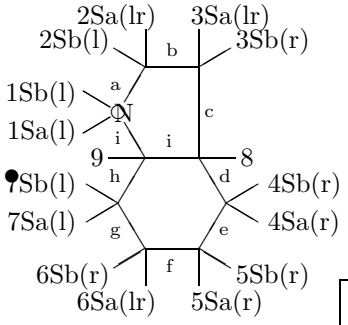
The macro `\indolehi` is used to draw inodele derivatives of inverse horizontal type (hetaromh.sty). Macros for drawing other fused heterocycles are also defined in the package hetaromh.sty. The format of these commands is as follows:

```
\indolehi[BONDLIST]{SUBSLIST}
\isoindolehi[BONDLIST]{SUBSLIST}
\purinehi[BONDLIST]{SUBSLIST}
\benzofuranehi[BONDLIST]{SUBSLIST}
\isobenzofuranehi[BONDLIST]{SUBSLIST}
\benzoxazolehi[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions and characters for describing bonds to be doubled are shown in the following diagram:



○: (400,800)
●: (0,0)

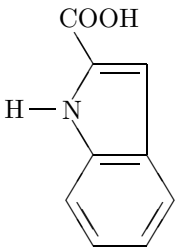


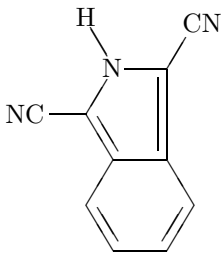
○: (400,800)
●: (0,0)

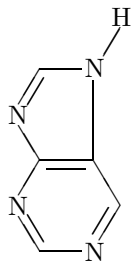
Example:

```
\indolehi{1==H;2==COOH}\qquad
\isoindolehi{2==H;1==NC;3==CN}\qquad
\purinehi{3==H}
```

produce the following structures:



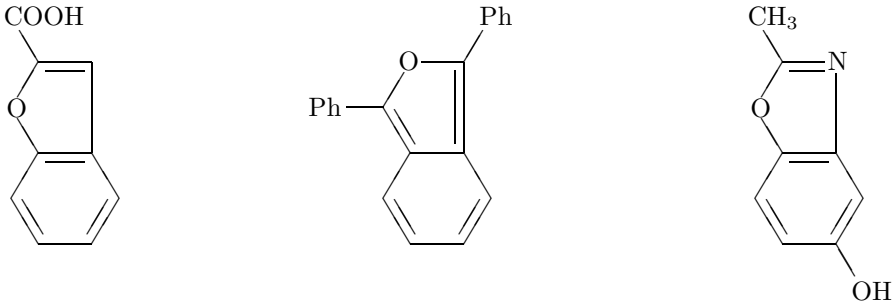




Example:

```
\benzofuranehi{2==COOH}\qquad
\isobenzofuranehi{1==Ph;3==Ph}\qquad
\benzoxazolehi{2==CH$_{3}$;5==OH}
```

produce the following structures:

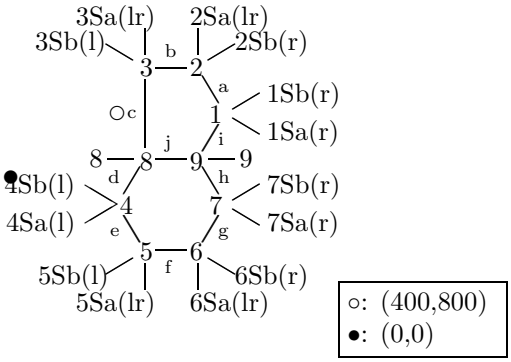


10.2.2 Commands for General Use

Macros for specified use such as `\indoleh` are based on a general command `\nonaheteroh`. This command enable us to draw a further variety of heterocyclic compounds. This macro has the following format which is contained also in hetaromh.sty.

```
\nonaheteroh[BONDLIST]{ATOMLIST}{SUBSLIST}
```

The locant numbering is common in these commands except their directions, as shown in the following diagram:



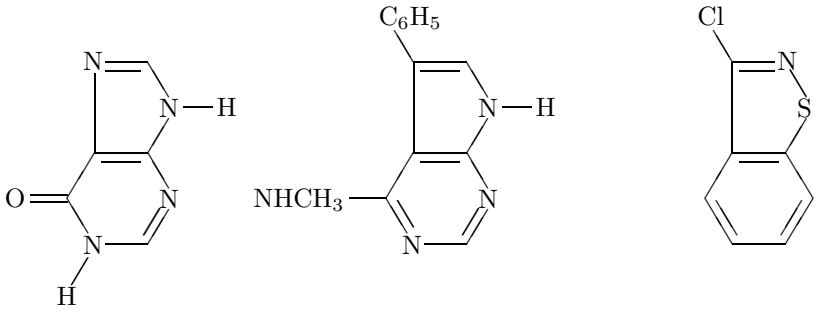
The handedness for each oriented or double-sided position is shown with a character set (l, r, or lr) in parentheses. Characters listed in Table 10.1 are also used to specify double bonds by setting them into the optional argument BONDLIST.

The argument ATOMLIST takes a usual format with respect to heteroatoms attached to $n = 1$ to 7, *e.g.*, 1==N for a nitrogen atom at 1-position. Hetero-atoms at 3a- and 7b-positions are represented as to be 3a==N (or 8==N) for a nitrogen at 3a-position, 7a==N (or 9==N) for at a nitrogen at 7a-position, and so on. The augument SUBSLIST takes a usual format except that the locant numbers 3a and 7a are replaced by 8 and 9.

Example:

```
\nonaheteroh[bjg]{1==N;3==N;5==N;7==N}{1==H;5==H;4D==O}
\nonaheteroh[bjge]{1==N;5==N;7==N}%
 {1==H;3==C$_{6}$H$_{5}$;4==NHCH$_{3}$}
\nonaheteroh[bjge]{1==S;2==N}{3==Cl}
```

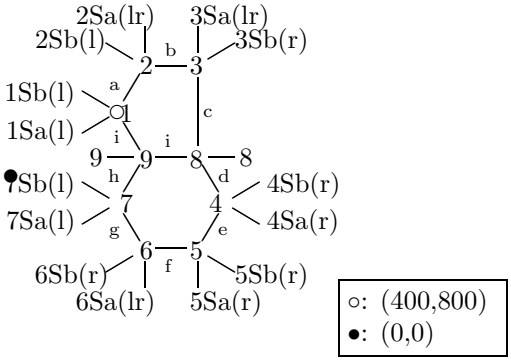
produce the following structures:



The inverse conterpart `\nonaheterohi` are also available by setting '`hetaromh.sty`'. Its format is:

```
\nonaheterohi[BONDLIST]{ATOMLIST}{SUBSLIST}
```

Its locant numbering is common with the normal counterpart as shown in the following diagram:

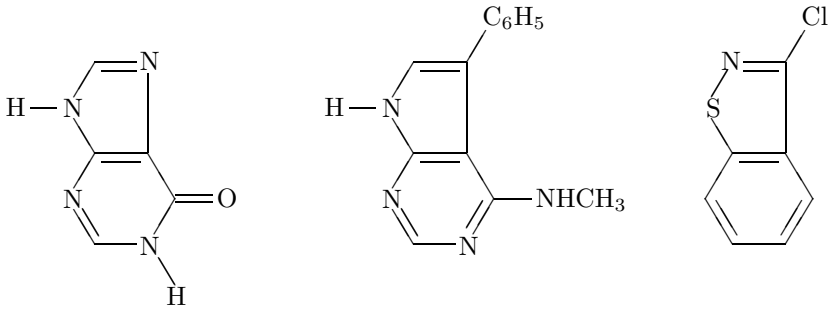


The handedness for each oriented or double-sided position is shown with a character set in parentheses. The optional argument BONDLIST uses characters listed in Table 10.1 so as to specify double bonds.

Example:

```
\nonaheterohi[bjg]{1==N;3==N;5==N;7==N}{1==H;5==H;4D==O}
\nonaheterohi[bjge]{1==N;5==N;7==N}%
 {1==H;3==C$_{6}$H$_{5}$;4==NHCH$_{3}$}
\nonaheterohi[bjge]{1==S;2==N}{3==Cl}
```

produce the following structures:



Chapter 11

Building Blocks

11.1 Six-Membered Blocks

The macro `\sixunitv` provides a six-membered fragment that can be fused to another ring structure, producing a new ring system.

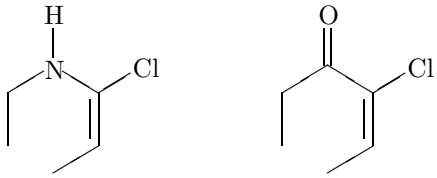
```
\sixunitv[BONDLIST]{ATOMLIST}{SUBSLIST}{OMIT}
```

The modes of numbering for positions and for edges are the same as those of `\sixheterov`. Hence, the arguments BONDLIST, ATOMLIST, and SUBSLIST are written in the same way. The argument OMIT indicates bonds to be deleted, where characters are selected from the table prepared for the BONDLIST of `\sixheterov`. It should be noted that the assignment of a null argument to ATOMLIST produces a carbocyclic building block.

Example:

```
\sixunitv[b]{1==N}{1==H;2==Cl}{d} \qquad
\sixunitv[b]{}{1D==O;2==Cl}{d}
```

produce the following structures:



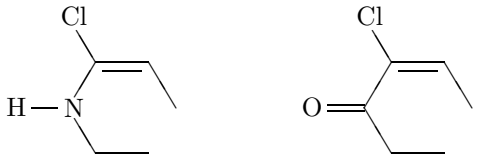
The macro `\sixunith` is the horizontal counterpart of `\sixunitv`. It produces a six-membered fragment of horizontal type,

```
\sixunith[BONDLIST]{ATOMLIST}{SUBSLIST}{OMIT}
```

Example:

```
\sixunith[b]{1==N}{1==H;2==Cl}{d} \qquad
\sixunith[b]{}{1D==O;2==Cl}{d}
```

produce the following structures:



11.2 Five-Membered Blocks

The macro `\fiveunitv` produces a five-membered fragment that can be fused to another ring structure to produce a new ring system.

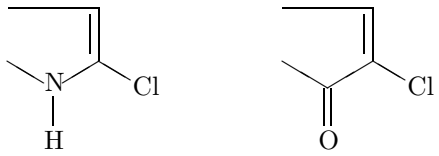
```
\fiveunitv[BONDLIST]{ATOMLIST}{SUBSLIST}{OMIT}
```

The numbering of positions and the designation of edges are the same as those of `\fiveheterov`. Hence, the arguments BONDLIST, ATOMLIST, and SUBSLIST are written in the same way. The argument OMIT indicates bonds to be deleted. It is a list of the same format as the BONDLIST of `\fiveheterov`.

Example:

```
\fiveunitv[b]{1==N}{1==H;2==Cl}{d} \qquad
\fiveunitv[b]{}{1D==O;2==Cl}{d}
```

produce the following structures:



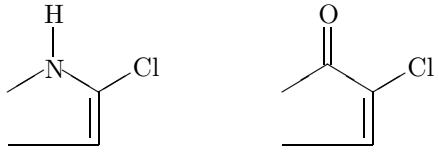
The macro `\fiveunitvi` produces a five-membered fragment of inverse type that can be fused to another ring structure to produce a new ring system.

```
\fiveunitvi[BONDLIST]{ATOMLIST}{SUBSLIST}{OMIT}
```

Example:

```
\fiveunitvi[b]{1==N}{1==H;2==Cl}{d} \qquad
\fiveunitvi[b]{}{1D==O;2==Cl}{d}
```

produce the following structures:



The macro `\fiveunith` is a five-membered fragment of horizontal type that can be fused to another ring structure to produce a new ring system.

```
\fiveunith[BONDLIST]{ATOMLIST}{SUBSLIST}{OMIT}
```

Example:

```
\fiveunith[b]{1==N}{1==H;2==Cl}{d} \qquad
\fiveunith[b]{}{1D==O;2==Cl}{d}
```

produce the following structures:

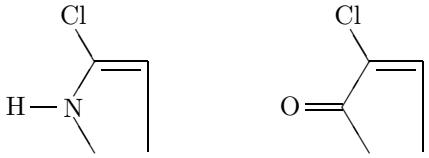
The macro `\fiveunithi` is the inverse counterpart of `\fiveunith`. It produces a five-membered fragment of another horizontal type so that it can be fused to another ring structure to produce a new ring system.

```
\fiveunithi[BONDLIST]{ATOMLIST}{SUBSLIST}{OMIT}
```

Example:

```
\fiveunithi[b]{1==N}{1==H;2==Cl}{d} \qquad
\fiveunithi[b]{}{1D==O;2==Cl}{d}
```

produce the following structures:



11.3 Setting Locant Numbers

The `\bdloocant` prints six characters on the edges of a six-membered ring of vertical type (locant.sty).

```
\bdloocant{#1}{#2}{#3}{#4}{#5}{#6}
```

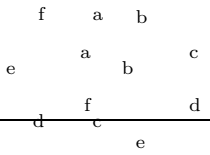
On the other hand, the `\bdloocnth` prints six characters on the edges of a six-membered ring of horizontal type.

```
\bdloocnth{#1}{#2}{#3}{#4}{#5}{#6}
```

Example:

```
\bdloocant{a}{b}{c}{d}{e}{f} \qquad
\bdloocnth{a}{b}{c}{d}{e}{f}
```

produce the following structures:



The `\sxloocant` prints six characters on the vertices of a six-membered ring of vertical type.

```
\sxloocant{#1}{#2}{#3}{#4}{#5}{#6}
```

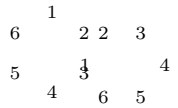
On the other hand, the `\sxloocnth` prints six characters on the vertices of a six-membered ring of horizontal type.

```
\sxloocnth{#1}{#2}{#3}{#4}{#5}{#6}
```

Example:

```
\sxloocant{1}{2}{3}{4}{5}{6} \qquad
\sxloocnth{1}{2}{3}{4}{5}{6}
```

produce the following structures:

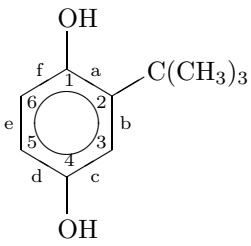


These commands can be used by combining with a structure-drawing macro in a picture environment to show the locant numbering of the structure.

Example:

```
\begin{picture}(1000,1000)(0,0)
\put(0,0){\sxloocant{1}{2}{3}{4}{5}{6}}
\put(0,0){\bdloocant{a}{b}{c}{d}{e}{f}}
\put(0,0){\bzdrv[c]{1==OH;2==C(CH$_{3}$)$_{3}$;4==OH}}
\end{picture}
```

produces the following structure:



Chapter 12

Further Cyclic Compounds

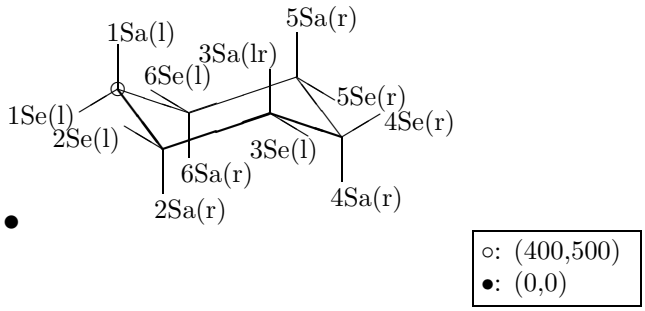
12.1 Drawing Chair Form of Cylohexane

12.1.1 Starndard formula

The macro `\chair` is used to draw cyclohexane derivatives of chair-form (ccycle.sty). The format of this command is as follows:

```
\chair[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



Each character set in parentheses represents the handedness of the corresponding postion, which is fixed in this type of macros.

The option argument BONDLIST is an character string in a pair of brackets, where each character indicates the presence of a double bond at the edge corresponding to the character. The bond-correspondence is rather arbitrary in some cases but conforms to chemical conventions as faithfully as possible if such conventions are presence (Table 12.1).

The argument SUBSLIST for this macro takes a general format, except that modifiers are selected from 'Sa' for an axial substitutent, 'Se' for an equatorial substitutent, and 'D' for a substitutent through an double bond.

Example:

```
\chair{1D==O;2Se==H$_{3}$C;2Sa==CH$_{3}$;%
6Se==CH$_{3}$;6Sa==CH$_{3}$} \qquad\qquad
\chair[b]{1D==O;4Se==CH$_{3}$;4Sa==CH$_{3}$}
```

Table 12.1: Argument BONDLIST for the commands \chair and \chairi

Character	Printed structure
none	cyclohexane
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,6-double bond
f	6,1-double bond

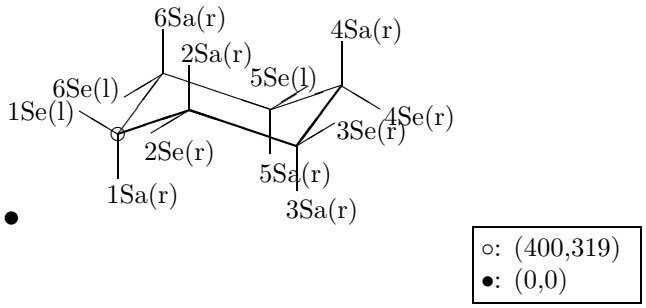
produce the following diagrams:

12.1.2 Inverse formula

The macro \chairi is used to draw cyclohexane derivatives of inverse chair-form (ccycle.sty). The format of this command is as follows:

```
\chairi[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



Each character set in parentheses represents the handedness of the corresponding postion, which is fixed in this type of macros.

The option argument BONDLIST has the same meaning as the counterpart of the \chair command (Table 12.1). The argument SUBSLIST for this macro takes the same format as described in the \chair macro, *e.g.*, 'Sa', 'Se', and D.

Example:

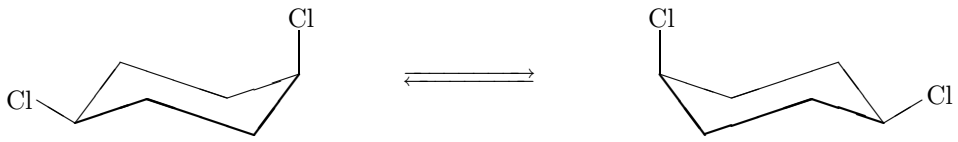
```
\chairi{1D==O;2Se==CH$_{3}$;2Sa==CH$_{3}$;%
6Se==CH$_{3}$;6Sa==CH$_{3}$} \qquad\qquad
\chairi[b]{1D==O;4Se==CH$_{3}$;4Sa==CH$_{3}$}
```

produce the following diagrams:

The following example shows the ring flipping of two chair forms, which are drawn with the macros `\chair` and `\chairi`. In addition, we use the counters *compd* and *deriv* which are available from the commands `\nocompd` and `\derivlabel` of the package chemist.sty.

```
\begin{quotation}
Let us consider {\em cis\/}-1,4-dichlorocyclohexane as an example.
This derivative is generated by putting chlorine atoms on
the 1- and 4-positions.
Thus, we take account of the following pair
of conformers (\cref{cf:06a} and of \cref{cf:06b}):
\end{quotation}
\begin{center}
\nocompd{}
\cdtwocell{0pt}{160pt}{\chairi{1Se==Cl;4Sa==Cl}\vskip-10pt}%
 {\derivlabel{cf:06a}}
\reacteqarrow{10pt}{50pt}{}{}
\cdtwocell{0pt}{160pt}{\chair{1Sa==Cl;4Se==Cl}\vskip-10pt}%
 {\derivlabel{cf:06b}}
\end{center}
```

> Let us consider *cis*-1,4-dichlorocyclohexane as an example. This derivative is generated by putting chlorine atoms on the 1- and 4-positions. Thus, we take account of the following pair of conformers (**1a** and of **1b**):



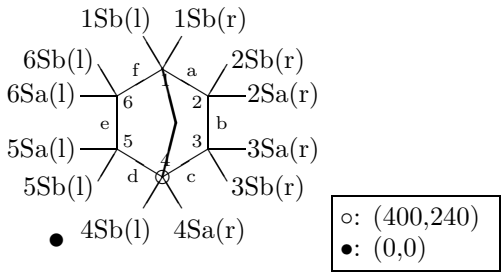
1a **1b**

12.2 Drawing Bicyclo[2.2.1]heptane

The macro `\bicychepv` is used to draw bicyclo[2.2.1]heptane derivatives of vertical type in a flat fashion. The format of this command is as follows:

```
\bicychepv[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions and bond descriptors for assigning double bonds are shown in the following diagram:



Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros.

The option argument BONDLIST is an character string in a pair of brackets, where each character indicates the presence of a double bond at the edge corresponding to the character (Table 12.2).

Table 12.2: Argument BONDLIST for commands `\bicychepv` and `\bicycheph`

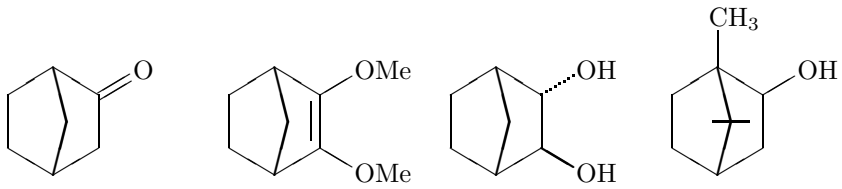
Character	Printed structure
none	bicyclo[2.2.1]heptane
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,6-double bond
f	6,1-double bond
A	aromatic circle
7	7,7-dimethyl

The argument SUBSLIST for this macro takes a general format, in which the modifiers listed in Table 2.2 are used.

Example:

```
\bicychepv{2D==O}
\bicychepv[b]{2==OMe;3==OMe}
\bicychepv{3B==OH;2A==OH}
\bicychepv[7]{1==CH$_{3}$;2==OH}
```

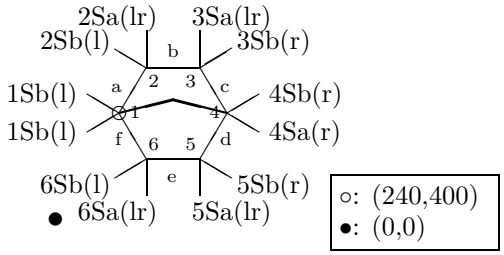
produce the following diagrams:



The macro `\bicycheph` is used to draw bicyclo[2.2.1]heptane derivatives of vertical type in a flat fashion. The format of this command is as follows:

```
\bicycheph[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions along with bond desciptors are found in the following diagram:

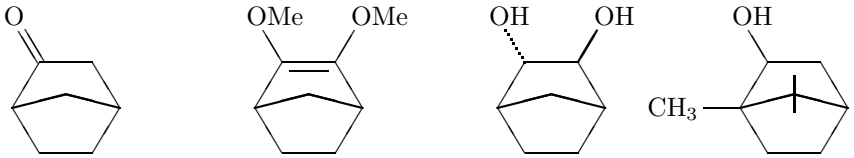


Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros. For the SUBSLIST and the BONDLIST, see Table 2.2 and 12.2. .

Example:

```
\bicycheph{2D==O}
\bicycheph[b]{2==OMe;3==OMe}
\bicycheph{3B==OH;2A==OH}
\bicycheph[7]{1==CH$_{3}$;2==OH}
```

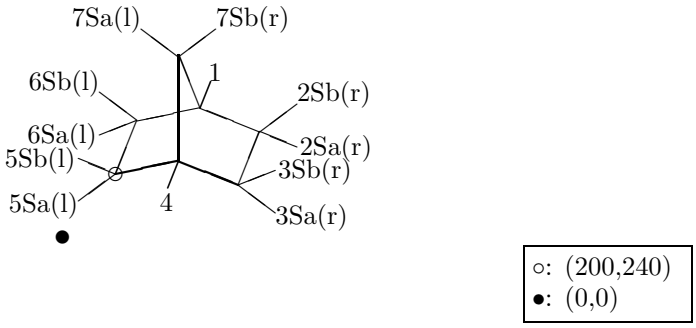
produce the following diagrams:



For the purpose of depicting the stereochemisty of bicyclo[2.2.1]heptane derivatives, you can use the command `\bornane` instead of the commands `\bicychepv` and `\bicycheph`. The format of this command is as follows:

```
\bornane[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros.

The option argument BONDLIST is an character string in a pair of brackets, where each character indicates the presence of a double bond at the edge corresponding to the character (Table 12.3). The SUBSLIST format is shown in Table 2.2.

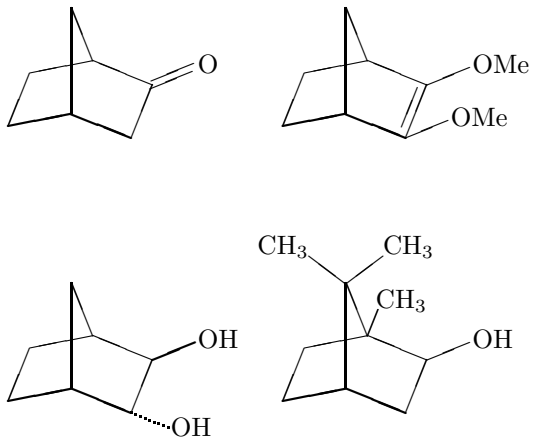
Example:

Table 12.3: Argument BONDLIST for commands \bornane

Character	Printed structure
none	bornane
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,6-double bond
f	6,1-double bond
g	1,7-double bond
h	4,7-double bond

```
\bornane{2D==O}
\bornane[b]{2==OMe;3==OMe}\par
\bornane{3B==OH;2A==OH}
\bornane{7Sa==CH$_{3}$;7Sb==CH$_{3}$;1==CH$_{3}$;2==OH}
```

produce the following diagrams:

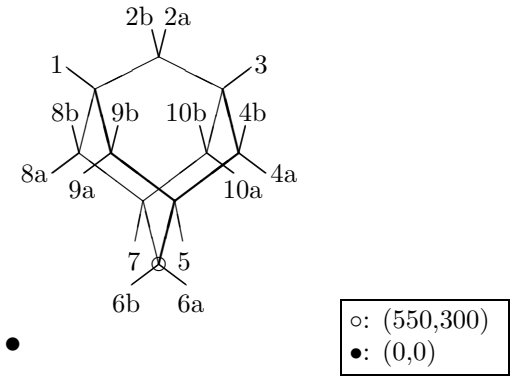


12.3 Drawing Adamantane Derivatives

The macro \adamantane prints adamantane derivatives (vertical formulas) by means of the following format:

```
\adamantane[BONDLIST]{SUBSLIST}
```

where BONDLIST is not effective in the present specification.



The argument SUBSLIST is slightly different from general conventions, as shown in Table 12.4.

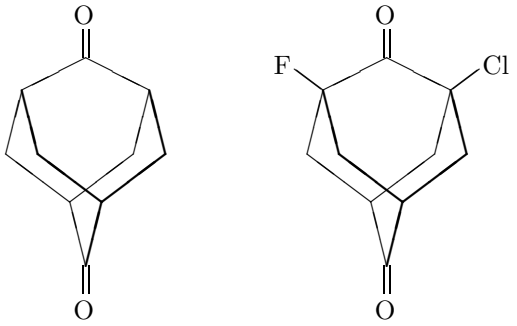
Table 12.4: Argument SUBSLIST for commands `\adamantane` and `\hadamantane`

Character	Printed structure
for $n = 1, 3, 5$, and 7 (bridgeheads)	
n or na	exocyclic single bond at n-atom
for $n = 2, 4, 6, 8, 9$, and 10 (bridges)	
na	exocyclic single bond at n-atom (axial)
nb	exocyclic single bond at n-atom (equatorial)
nD	exocyclic double bond at n-atom (2 and 6)

Example:

```
\adamantane{2D==O;6D==O}
\adamantane{2D==O;6D==O;1==F;3==Cl}
```

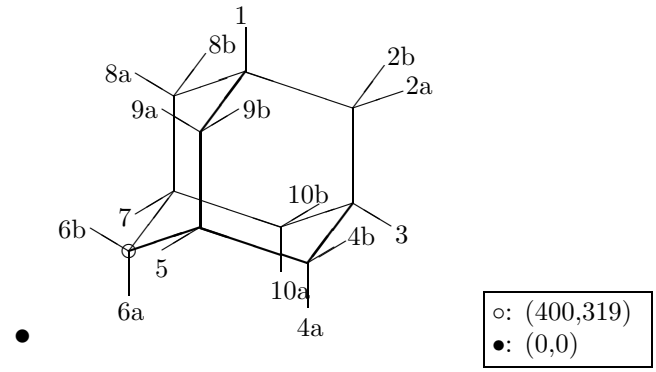
produce the following diagrams:



The macro `\hadamantane` prints adamantane derivatives (horizontal formulas) by means of the following format:

```
\hadamantane[BONDLIST]{SUBSLIST}
```

where BONDLIST is not effective in the present specification.

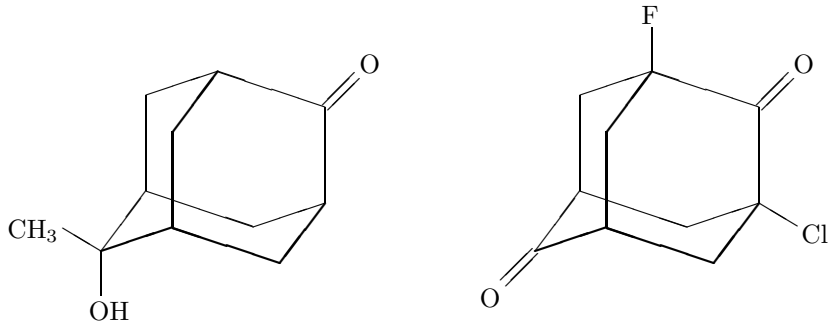


The argument SUBSLIST has the same meanings as that of `\adamantane`, as shown in Table 12.4.

Example:

```
\hadamantane{2D==O;6a==OH;6b==CH$_{3}$}
\hadamantane{2D==O;6D==O;1==F;3==Cl}
```

produce the following diagrams:

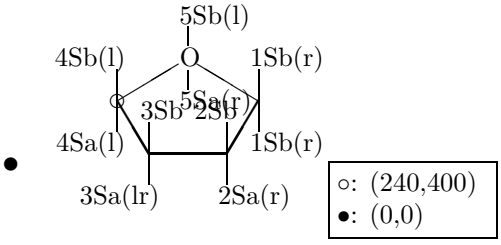


12.4 Drawing Furanoses and Pyranoses

For drawing furanoses, the command `\furanose` is used. The format of this command is as follows:

```
\furanose[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions and bond descriptors are represented by the following diagram:



Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros.

Table 12.5: Argument BONDLIST for commands \bicychepv and \bicycheph

Character	Printed structure
none	mother skeleton
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,1-double bond

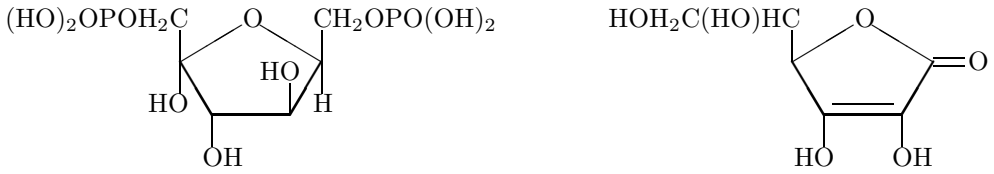
The option argument BONDLIST is an character string in a pair of brackets, where each character indicates the presence of a double bond at the edge corresponding to the character (Table 12.5).

The argument SUBSLIST for this macro takes a general format, in which the modifiers listed in Table 2.2 are used.

Example:

```
\furanose{1Sa==H;1Sb==CH$_{2}$OPO(OH)$_{2}$;2Sb==\lmoiety{HO};3Sa==OH;%
4Sb==(HO)$_{2}$OPOH$_{2}$C;4Sa==HO}\hskip5cm
\furanose[b]{1D==O;2Sa==OH;3Sa==\lmoiety{HO};4Sb==HOH$_{2}$C(HO)HC}
```

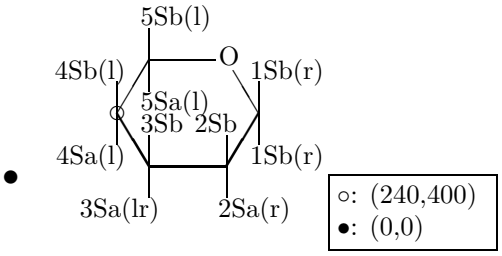
produce the following diagrams:



For drawing furanoses, the command \pyranose is used. The format of this command is as follows:

```
\pyranose[BONDLIST]{SUBSLIST}
```

Locant numbers for designating substitution positions are represented by the following diagram:



Each character set in parentheses represents the handedness of the corresponding position, which is fixed in this type of macros.

The option argument BONDLIST is an character string in a pair of brackets, where each character indicates the presence of a double bond at the corresponding edge (Table 12.6).

The argument SUBSLIST for this macro takes a general format, in which the modifiers listed in Table 2.2 are used.

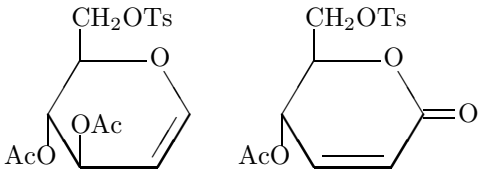
Example:

Table 12.6: Argument BONDLIST for commands \bicychepv and \bicycheph

Character	Printed structure
none	mother skeleton
a	1,2-double bond
b	2,3-double bond
c	4,3-double bond
d	4,5-double bond
e	5,6-double bond
f	6,1-double bond

```
\pyranose[a]{3Sb==OAc;4Sa==AcO;5Sb==CH$_{2}$OTs}
\pyranose[b]{2D==O;5Sb==CH$_{2}$OTs}
```

produce the following diagrams:



Chapter 13

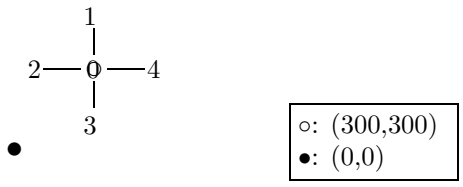
Aliphatic Compounds

13.1 Drawing Tetrahedral Units

The macro `\tetrahedral` is used to draw a tetrahedral unit (aliphat.sty). The format of this command is as follows:

```
\tetrahedral[AUXLIST]{SUBSLIST}
```

The following diagram shows the numbering for designating substitution positions:



in which the same macro is used to typeset both saturated and unsaturated derivatives.

The optional argument AUXLIST is used to specify a charge on the central atom: *i.e.*, {0+} represents a + charge (or another one chararacter) on the center.

The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 13.1, in which n is an arabic numeral between 1 and 4.

Table 13.1: SUBSLIST for `\tetrahedral`

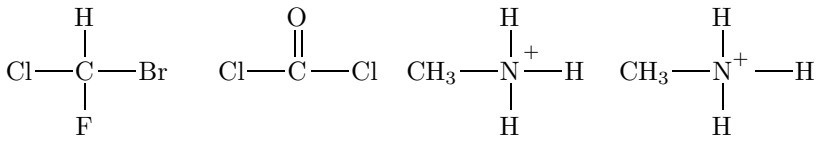
Character	Structures printed
nT	triple bond at n-atom
nD	double bond at n-atom
n or nS	single bond at n-atom
nA	alpha single bond at n-atom
nB	beta single bond at n-atom

The central carbon atom is assigned by writing 0==C in the SUBSLIST. The structural formula of an ammonium ion can also be obtained with this command.

Example:

```
\tetrahedral{0==C;1==H;2==Cl;3==F;4==Br}\qquad
\tetrahedral{0==C;1D==O;2==Cl;4==Cl}\qquad
\tetrahedral[{}{0+}]{0==N;1==H;2==CH$_{3}$;3==H;4==H}\qquad
\tetrahedral{0==N$^{\raise.5ex\hbox{\scriptsize +}}$;%
 1==H;2==CH$_{3}$;3==H;4==H}
```

produce the following structures:



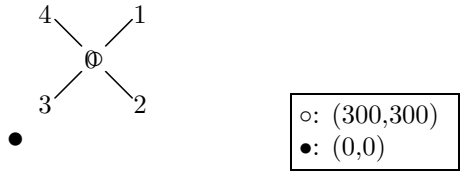
Note that the AUXLIST of the third formula contains a dummy pair of braces {}, which is necessary for corrrect processing. For such a dummy argument, see also the minus charge of a 1,3-dithiane anion on page 145.

On the other hand, the fourth formula shows another way of typesetting the same ammonium ion, where the plus charge is designated as a superscript of the central nitrogen atom in the SUBSLIST after vertical adjustment with the command `\raise` and fontsize adjustment with the command `\scriptsize`.

The macro `\square` is used to draw a tetrahedral unit of another type (aliphat.sty). The format of this command is as follows:

```
\square[AUXLIST]{SUBSLIST}
```

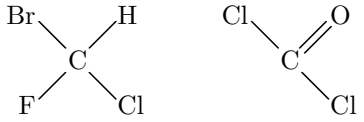
The following diagram shows the numbering for designating substitution positions:



Example:

```
\square{0==C;1==H;2==Cl;3==F;4==Br}\qquad
\square{0==C;1D==O;2==Cl;4==Cl}
```

produce the following structures:



13.2 Drawing Trigonal Units

The macro `\rtrigonal` and `\ltrigonal` are used to draw right-handed and left-handed trigonal units (aliphat.sty). The formats of these commands are as follows:

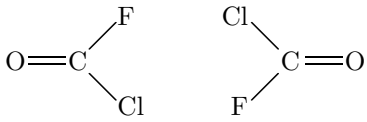
```
\rtrigonal[AUXLIST]{SUBSLIST}
\ltrigonal[AUXLIST]{SUBSLIST}
```

The bond angles of 2–0–3 are 90° in the trigonal units printed with these commands. The arguments AUXLIST and SUBSLIST are the same as those of `\tetrahedral`.

Example:

```
\rtrigonal{0==C;1D==O;2==Cl;3==F}\qquad
\ltrigonal{0==C;1D==O;2==Cl;3==F}
```

produce the following structures:



The macros `\utrigonal` and `\Utrigonal` are used to draw upward trigonal units with angles 90° and 120° (aliphat.sty). The formats of these commands are as follows:

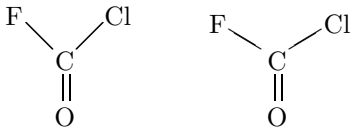
```
\utrigonal[AUXLIST]{SUBSLIST}
\Utrigonal[AUXLIST]{SUBSLIST}
```

The arguments AUXLIST and SUBSLIST are the same as those of `\tetrahedral`.

Example:

```
\utrigonal{0==C;1D==O;2==Cl;3==F}\qquad
\Utrigonal{0==C;1D==O;2==Cl;3==F}
```

produce the following structures:



On the other hand, the macros `\dtrigonal` and `\Dtrigonal` are used to draw downward trigonal units with angles 90° and 120° (aliphat.sty). The formats of these commands are as follows:

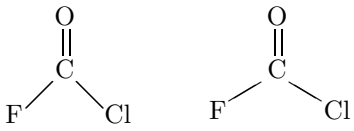
```
\dtrigonal[AUXLIST]{SUBSLIST}
\Dtrigonal[AUXLIST]{SUBSLIST}
```

The arguments AUXLIST and SUBSLIST are the same as those of `\tetrahedral`.

Example:

```
\dtrigonal{0==C;1D==O;2==Cl;3==F}\qquad
\Dtrigonal{0==C;1D==O;2==Cl;3==F}
```

produce the following structures:



13.3 Drawing Ethylene Derivatives

The macro `\ethylene` is used to draw ethylene derivatives with angles 90° (aliphat.sty). The format of this command is as follows:

```
\ethylene[BONDLIST]{ATOMLIST}{SUBSLIST}
```

The following diagram shows the numbering for designating substitution positions:

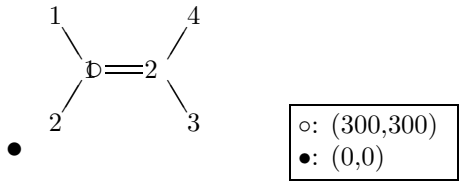


Table 13.2: BONDLIST for `\ethylene`

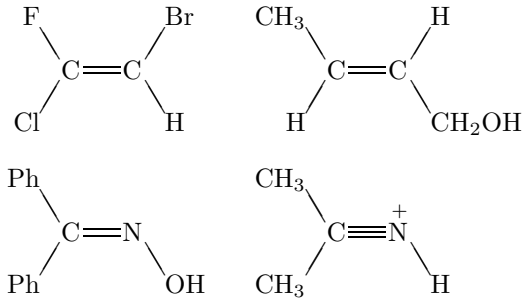
Character	Structures printed
{n+}	+ charge (or another one chararacter) on n-atom
d	inner double bond (between centers 1 and 2)
t	inner triple bond (between centers 1 and 2)

The argument ATOMLIST is used for giving central atoms. The argument SUBSLIST is the same as that of `\tetrahedral`.

Example:

```
\ethylene{1==C;2==C}{1==F;2==Cl;3==H;4==Br}\qquad
\ethylene{1==C;2==C}{1==CH$_{3}$;2==H;3==CH$_{2}$OH;4==H}\par
\ethylene{1==C;2==N}{1==Ph;2==Ph;3==OH}\qquad
\ethylene[t{2+}]{1==C;2==N}{1==CH$_{3}$;2==CH$_{3}$;3==H}
```

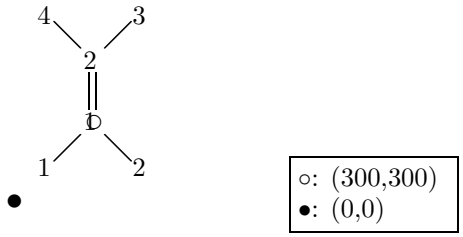
produce the following structures:



The macro `\ethylenev`, which is the vertical counterpart of `\ethylene`, is used to draw ethylene derivatives with angles 90° (aliphat.sty). The format of this command is as follows:

```
\ethylenev[BONDLIST]{ATOMLIST}{SUBSLIST}
```

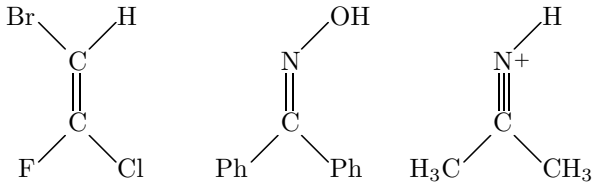
The following diagram shows the numbering for designating substitution positions:



Example:

```
\ethylenev{1==C;2==C}{1==F;2==Cl;3==H;4==Br}\qquad
\ethylenev{1==C;2==N}{1==Ph;2==Ph;3==OH}\qquad
\ethylenev[t{2+}]{1==C;2==N}{1==H$_{3}$C;2==CH$_{3}$;3==H}
```

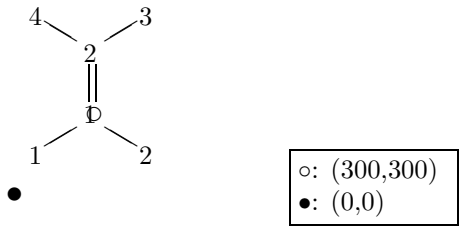
produce the following structures:



The macro `\Ethylenev` is used to draw ethylene derivatives with angles 120° (aliphat.sty). It is the vertical counterpart of `\ethylene`. The format of the command is as follows:

```
\Ethylenev[BONDLIST]{ATOMLIST}{SUBSLIST}
```

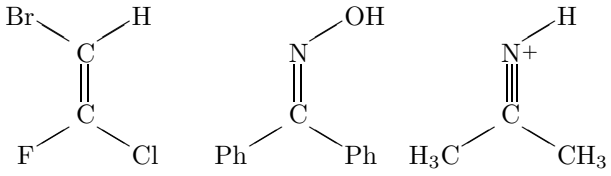
The following diagram shows the numbering for designating substitution positions:



Example:

```
\Ethylenev{1==C;2==C}{1==F;2==Cl;3==H;4==Br}\qquad
\Ethylenev{1==C;2==N}{1==Ph;2==Ph;3==OH}\qquad
\Ethylenev[t{2+}]{1==C;2==N}{1==H$_{3}$C;2==CH$_{3}$;3==H}
```

produce the following structures:



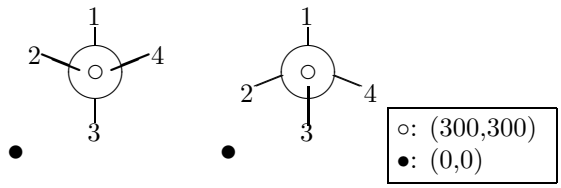
13.4 Drawing Configurations

The macros `tetrastereo` and `dtetrastereo` typset fragments which show actural configuration of a tetrahedral carbon in different modes of projections. The formats of these commands are as follows:

```
\tetrastereo[AUXLIST]{SUBSLIST}
\dtetrastereo[AUXLIST]{SUBSLIST}
```

The arguments AUXLIST and SUBSLIST are the same as those of `\tetrahedral`.

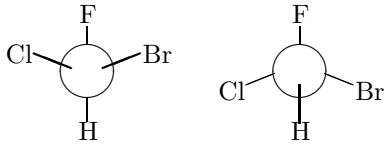
The following diagram shows the numbering for designating substitution positions:



Example:

```
\tetrastereo{1==F;2==Cl;3==H;4==Br}\qquad
\dtetrastereo{1==F;2==Cl;3==H;4==Br}
```

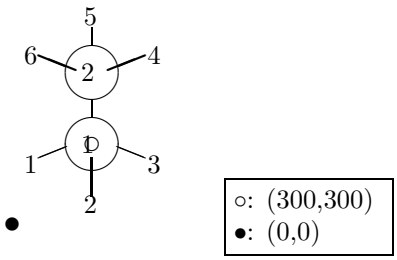
produce the following structures:



The configuration of ethane is typeset by the macro `\ethanestereo`. The format of the command is as follows:

```
\ethanestereo[AUXLIST]{ATOMLIST}{SUBSLIST}
```

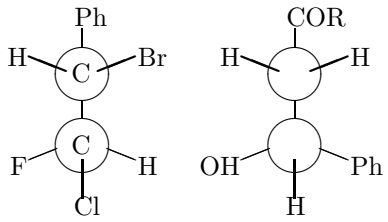
The argument ATOMLIST is used for giving central atoms. The arguments AUXLIST and SUBSLIST are the same as those of `\tetrahedral`.



Example:

```
\ethanestereo{1==C;2==C}{1==F;2==Cl;3==H;4==Br;6==H;5==Ph}\qquad
\ethanestereo{}{1==OH;2==H;3==Ph;4==H;5==COR;6==H}
```

produce the following structures:



Further examples of typesetting the configurations of ethane derivatives have been described in an article concerning sterochemistry [14].

Chapter 14

Combining structures

14.1 General Remarks

14.1.1 Coordinates of the Picture Environment

The macros described in the other chapters of this manual can be combined to construct a more complicated structural formula. This treatment is based on the fact that two or more picture environments of LaTeX can be nested, recognizing each inner picture environment as a LaTeX picture box.

A picture environment of LaTeX is set up with the following statement:

\begin{picture}$(L_x, L_y)(S_x, S_y)$,
⋮
\end{picture}

This command produces an $L_x \times L_y$ area for drawing a structural formula, where the origin (0, 0) can be shifted by giving differences (S_x, S_y).

The \put$(P_x, P_y)$ command places an inner picture box (*e.g.*, a fragment created by a macro of the present paper) so that the reference point of the inner picture is located on the (P_x, P_y) point of the outer picture environment.

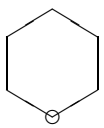
14.1.2 Reference Points and Inner Origins

A macro of XeLaTeX is based on an inner picture environment, which has an original point for drawing a structure and a set of sifted values. Suppose the definition of the macro contains an inner picture environment represented by

$$\texttt{\textbackslash begin\{picture\}}(\ell_x, \ell_y)(s_x, s_y).$$

The point (s_x, s_y) of the inner environment becomes the (0,0) point of the structure generated by the macro. This point is called the reference point of the macro in the present manual. On the other hand, the origin of the inner environment becames the $(-s_x, -s_y)$ point of the generated structure. It is called the inner origin of the macro.

For example, the command \cyclohexanev generates a skeleton,



•

in which the symbol ∘ represents the inner origin and the symbol • represents a reference point. The macro `\cyclohexanev` is defined on the basis of an inner picture environment:

```
\begin{picture}(800, 800)(-400, -240).
```

As a result, the inner origin is referred to as the (400, 200) point of the resulting macro; and the inner $(-400, -240)$ point is regarded as the new origin (0, 0), which is the reference point of the macro.

Each macro is characterized by the reference point and the inner origin, which are shown in a framed box beside the specification of the macro. The reference point and the inner origin of each macro are revealed by stating `\origpttrue` in the preamble of a manuscript. Then, they are printed with the symbols ∘ and •; and the values of them appear on the display. For example, the above cyclohexane structure is typeset by the following statement:

```
{\origpttrue
\begin{center}
\cyclohexanev{}
\end{center}}
```

or by an equivalent statement:

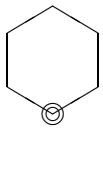
```
\begin{xymspec}
\cyclohexanev{}
\end{xymspec}
```

14.1.3 Setting Coordinates

The command `\put` typesets an object, which may be a character string, a structure generated with a macro, or others. When a macro is written as an argument of the command `\put` in an outer picture environment, a structure due to the macro is typeset so that the reference point of the macro is placed on the point designated by the `\put` command. For example,

```
{\origpttrue
\begin{picture}(1000,700)(0,0)
\put(0,0){\cyclohexanev{}}
\put(0,0){\circle{80}}
\put(400,240){\circle{80}}
\end{picture}
}
```

produces

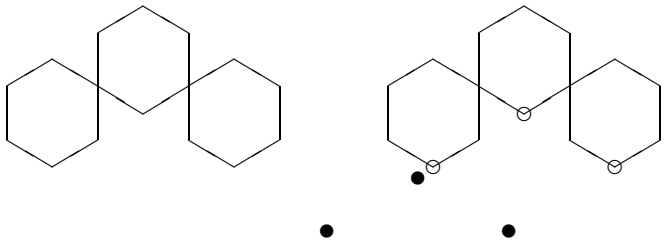


The reference point with • is encircled by an outer circle representing the origin of the outer picture environment. The inner origin represented with an open circle is encircled by an outer circle centered at the $(400, 240)$ point of the outer environment, resulting in a double circle.

Since we adopt a bond length equal to 200 and a bond slope of (5, 3) or (3, 5), such values as 200, 171, and 103 (and any combinations of these values) appear frequently in typsetting combined structures. Note that 171 is approximately equal to $200 \times (5/\sqrt{34})$ and 103 is aproximately equal to $200 \times (3/\sqrt{34})$, where we have $\sqrt{3^2 + 5^2} = \sqrt{34}$ for both the slopes (5, 3) and (3, 5). For example, a spiro compound can be typeset by the statement:

```
\begin{picture}(1200,900)(0,0)
\put(0,0){\cyclohexanev{}}
\put(342,200){\cyclohexanev{}}
\put(684,0){\cyclohexanev{}}
\end{picture}
```

The resulting structures are



The right-hand structure is to show the reference points and the inner origins of the fragements used. In this case, the shifted values $(-400, -240)$ of each fragment are equal to those of another fragment, since each fragment is generated by the same macro. Hence, the argument coordinates of the `\put` can be calculated without considering such shifted values. Thus, the value 342 is equal to 171×2, and 684 is equal to 171×4.

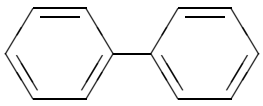
14.2 Combination of Macros Through a Bond

Since each of the macros described in the other chapters of the present manual is based on a picture environment, one of the simplest ways of combining structures is to place individual structures into an outer picture environment. Such combination of macros increases the variety of structural diagrams.

In order to illustrate the method of calculating coordinates, we take the drawing of biphenyl as the first example:

```
\begin{picture}(1200,600)(0,0)
\put(0,0){\bzdrh{4==}}
\put(546,0){\bzdrh{}}
\end{picture}
```

This statement produces the following structure:



The first argument of each `\put` command represents the coordinates of the point on which the structure is printed.

The inner origin of the macro `\bzdrh` is the leftmost position of the benzene ring. The structure typeset by the command `\put(0,0){\bzdrh{4==}}` has a rightmost terminal point at $(546, 0)$ with respect to the inner picture environment.[1] The value 546 is calculated by $406 + 140$, where 406 is the length of the horizontal hexagon $(= 103 + 200 + 103)$ and 140 is the bond length produced by the argument `{4==}`.

The command `\put(546,0){\bzdrh{}}` prints another benzene ring so that the inner origin of this benzene is placed on the terminal position of the former benzene ring. Note again that the argument

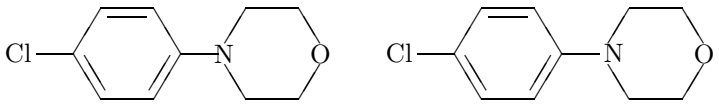
[1] The absolute cooridnates with respect to the outer picture environment is $(546 + 400, 0 + 240)$ in this case.

coordinates of \put can be calculated without considering such shifted values, (400, 240), since each of the fragments is generated by the same macro.

Let us draw 1-chloro-4-morphorinobenzene by means of two different ways.

```
\begin{picture}(1200,600)(0,0)
\put(0,0){\bzdrh{1==Cl;4==}}
\put(546,0){\sixheteroh[H]{1==N;4==O}{}}
\end{picture}
\qquad
\begin{picture}(1200,600)(0,0)
\put(0,0){\bzdrh{1==Cl;4==N}}
\put(566,0){\sixheteroh[H]{1=={ };4==O}{}}
\end{picture}
```

These statements produce essentially the same structure:

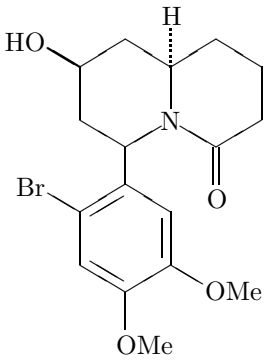


It should be noted that {4==} in SUBSLIST produces a bond without a substituent at the 4-position of the benzene ring, while 1=={␣} in ATOMLIST creates a vacancy to accomodate the nitrogen atom. In other words, the nitrogen atom of the first formula is regarded as a ring nitrogen of the morphorine ring, while the nitrogen of the second formula is considered to be a substituent of the benzene ring.

The following example illustrates a more complicated structure with a vertical bond linking two fragment structures. Thus, the statement:

```
\begin{picture}(1200,1500)(-200,-600)
\put(0,0){\decaheterov[H]{4a==N}{4D==O;7B==HO;{{10}A}==H}}
\put(0,-546){\bzdrv{1==;3==OMe;4==OMe;6==Br}}
\end{picture}
```

prints the following diagram:

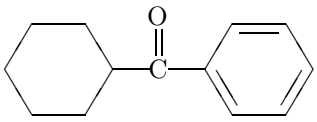


The first argument of each \put command represents the coordinates of the point on which the structure is printed. The value −546 is calculated by 140 (bond length) + 406 (the height of the hexagon = 103 + 200 + 103), because the inner origin of the structure printed by \decaheterov is position 5 (the left carbon atom adjacent to the nitrogen atom) and that of the latter structure is position 4 (the bottom carbon attached by the methoxy group).

The following example illustrates a combined structure in which two cyclic substructures are linked through an aliphatic unit

```
\begin{picture}(800,1000)(0,0)
\put(0,0){\cyclohexaneh[H]{4==}}
\put(754,0){\bzdrh{1==}}
\put(520,100){\tetrahedral{0==C;1D==O}}
\end{picture}
```

This statement produces



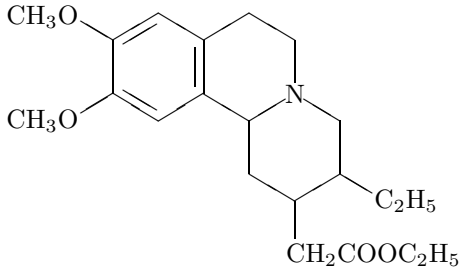
The calculation of the values for `\tetrahedral` is slightly complicated, since its inner origin is different from those of the other commands. The right terminal position due to the `\cyclohexaneh` is the point $(546, 0)$, the x-coordinate of which is equal to the length of a benzene ring (406) plus a bond length (140). The left terminal position due to the `\bzdrh` is the point $(614, 0)$, because $754 - 140 = 614$. Then, the aliphatic unit (`\tetrahedral`) should be placed at the average position of x-coordinate $(546 + 614)/2 = 580$. Since the inner origin of the the `\cyclohexaneh` is $(400, 240)$ and that of the `\tetrahedral` is $(300, 300)$, the x-coordinate is calculated to be $580-(300-240) = 520$ while the y-coordinate is calculated to be $400 - 300 = 100$.

14.3 Using Building Blocks

Macros `\sixunitv` and `\fiveunitv` are designed for building a new fused ring system:

```
\sixunitv[BONDLIST]{ATOMLIST}{SUBSLIST}{OMIT}
\fiveunitv[BONDLIST]{ATOMLIST}{SUBSLIST}{OMIT}
```

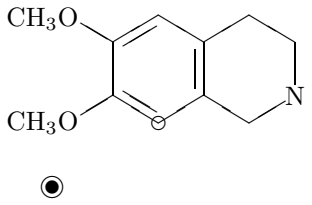
where argument OMIT (a–f for `\sixunitv` and a–e for `\fiveunitv`) is a list of characters, each of which represents one bond to be omitted. The other arguments have the same formats as described in the preceding sections. These macros produce five- and six-membered fragments respectively, in which the bond assigned by argument OMIT is deleted to be a fused vacant bond. They include the mechanism of the hetroatom truncation in order to be applied to typesetting both carbocycles and heterocycles. For example, a tricyclic system



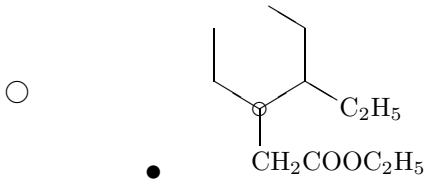
is typeset with the statement:

```
\begin{picture}(1200,1500)(-200,-600)
\put(0,0){\decaheterov[fhk]{3==N}{6==CH$_3$O;7==CH$_3$O}}
\put(513,-303){\sixunitv[H]{1=={ }}%
  {3==C$_2$H$_5$;4==CH$_2$COOC$_2$H$_5$}{f}}
\end{picture}
```

In order to examine the details of the combination, let us typeset the structure with the first command:



and separately the structure with the secdond command:



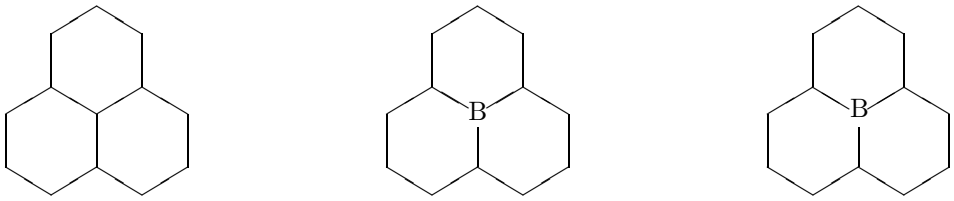
It should be noted that argument ATOMLIST in the `\sixunitv` macro contains the assignment "`1=={␣}`" which assures the vacant bridgehead position. This vacancy is occupied by the bridgehead nitrogen printed by the `\decaheterov` macro.

The following examples illustrate combinations of `decaheterov` and `\sixunitv` to produce a borane and the related carbocycle.

```
\begin{picture}(1200,1000)(0,0)
\put(0,0){\decaheterov[H]{}{}}
\put(171,303){\sixunitv[H]{4=={ }}{}{cd}}
\end{picture}
\qquad
\begin{picture}(1200,1000)(0,0)
\put(0,0){\decaheterov[H]{8a==B}{}}
\put(171,303){\sixunitv[H]{4=={ }}{}{cd}}
\end{picture}
```

```
\qquad
\begin{picture}(1200,1000)(0,0)
\put(0,0){\sixunitv[H]{2=={ }}{}{a}}
\put(342,0){\sixunitv[H]{}{}{ef}}
\put(171,303){\sixheterovi[H]{1==N}{}}
\end{picture}
```

These statements produce the following structures.

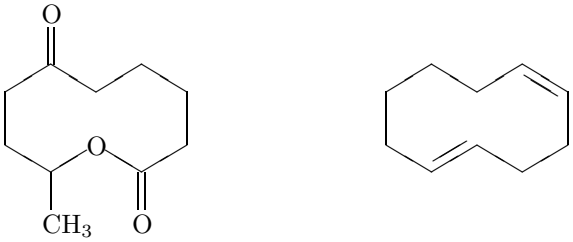


The second and third examples above show alternative ways to depict the carborane. Note that the argument `{2=={␣}}` in the third example is necessary to print the desired structures.

For illustrating the wide applicability of the building-block technique, we show several examples for the combination of two or more building blocks. The following two exmaples involve a combination of two `\sixunitv` commands.

```
\begin{picture}(1200,800)(0,0)
\put(0,0){\sixunitv[H]{3==O}{1D==O;4==CH$_{3}$}{b}}
\put(342,0){\sixunitv[H]{5=={ }}{4D==O}{e}}
\end{picture}
\qquad
\begin{picture}(1200,800)(0,0)
\put(0,0){\sixunitv[c]{}{}{b}}
\put(342,0){\sixunitv[a]{}{}{e}}
\end{picture}
```

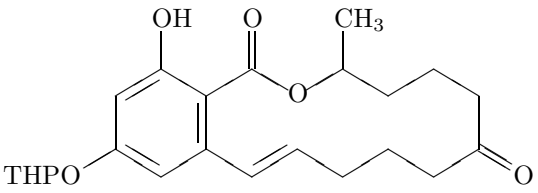
These statements provide



The following exmaple involves a combination of three `\sixunitv` commands and one `\bzdrv` command.

```
\begin{picture}(1600,800)(0,0)
\put(0,0){\bzdrv[r]{1==OH;5==THPO}}
\put(342,0){\sixunitv[c]{2==O}{1D==O}{be}}
\put(684,0){\sixunitv[H]{6=={ }}{1==CH$_{3}$}{be}}
\put(1026,0){\sixunitv[H]{}{3D==O}{e}}
\end{picture}
```

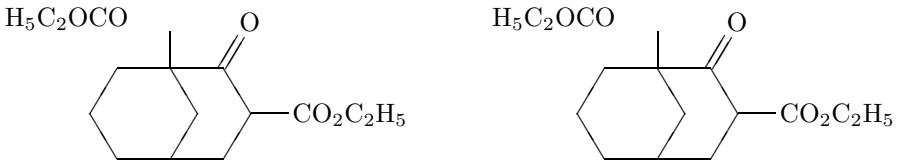
This statement provides



A bicyclo[3.3.1]nonane is typeset by this technique.

```
\begin{picture}(1600,800)(0,0)
\put(0,0){\cyclohexaneh[H]{3Sa==\lmoiety{H$_{5}$C$_{2}$OCO\kern2em}}}
\put(200,0){\sixunith[H]{}{3D==O;4==CO$_{2}$C$_{2}$H$_{5}$}{af}}
\end{picture}
\qquad
\begin{picture}(1600,800)(0,0)
\put(0,0){\cyclohexaneh[H]{3Sa==\lmoiety{H$_{5}$C$_{2}$OCO\kern2em}}}
\put(200,0){\fiveunith[H]{}{2D==O;1==CO$_{2}$C$_{2}$H$_{5}$}{c}}
\end{picture}
```

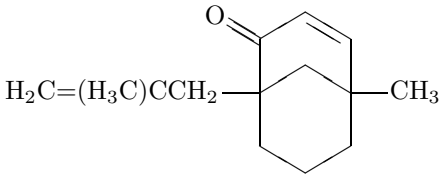
These statements provide equivalent results as follows.



A vertical form of bicyclo[3.3.1]nonane is also typeset by this technique.

```
\begin{picture}(800,1000)(0,0)
\put(0,0){\cyclohexanev[H]{6Sa==\lmoiety{H$_{2}$C=(H$_{3}$C)CCH$_{2}$};
2Sa==CH$_{3}$}}
\put(0,200){\sixunitv[a]{}{6D==O}{cd}}
\end{picture}
```

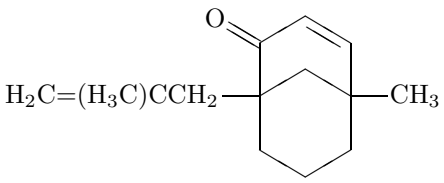
This statement provides



If you want to draw the one-carbon bridge of the bicyclononane in a thick line, you can combine three `\sixunitv` commands, where you use `\thicklines` command to draw the middle building block for the bridge.

```
\begin{picture}(800,1000)(0,0)
\put(0,0){\sixunitv[H]{}{6Sa==\lmoiety{H$_{2}$C=(H$_{3}$C)CCH$_{2}$};%
2Sa==CH$_{3}$}{af}}
\thicklines
\put(0,0){\sixunitv[H]{}{}{bcde}}
\thinlines
\put(0,200){\sixunitv[a]{}{6D==O}{cd}}
\end{picture}
```

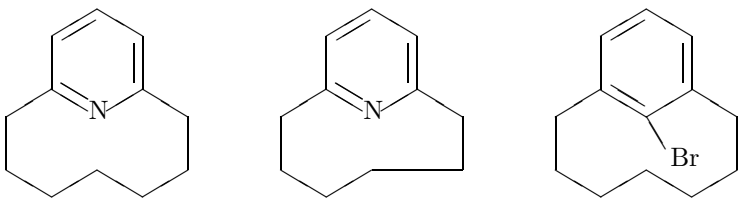
This statement provides



[7](2,6)- and [6](2,6)Pyridinophanes are other examples typeset by this technique. In a similar way, 13-Bromo-(2,6)metacyclophane can be printed easily.

```
\begin{picture}(1200,1000)(0,0)
\put(0,0){\sixunitv[H]{2==\{ \}}{}{ab}}
\put(342,0){\sixunitv[H]{6==\{ \}}{}{ef}}
\put(171,303){\pyridinevi{}}
\end{picture}
\qquad
\begin{picture}(1200,1000)(0,0)
\put(0,0){\sixunitv[H]{2==\{ \}}{}{ab}}
\put(342,0){\fiveunitvi[H]{5==\{ \}}{}{de}}
\put(171,303){\pyridinevi{}}
\end{picture}
\qquad
\begin{picture}(1200,1000)(0,0)
\put(0,0){\sixunitv[H]{2==\{ \}}{}{ab}}
\put(342,0){\sixunitv[H]{6==\{ \}}{}{ef}}
\put(171,303){\bzdrv{4Sa==\kern.5em\raise1ex\hbox{Br}}}
\end{picture}
```

These statements provide

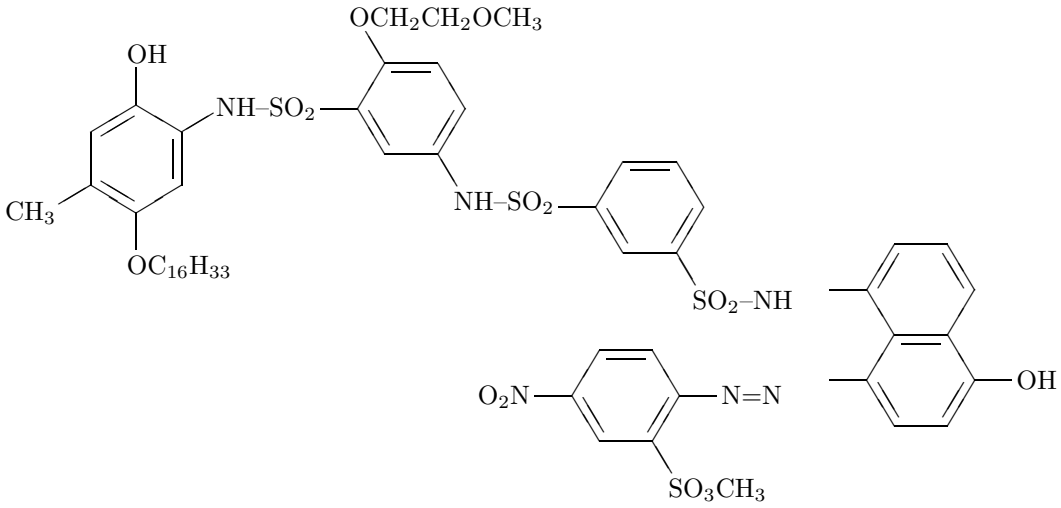


We have reported an article on dye releasers for instant color photography, in which a variety of structural formulas have been typeset by means of original utilities of the LaTeX picture environment

[15]. The present XyMTEX provides us with a more versatile tool of drawing such complex molecules. Thus, the formula of a cyan dye releaser for instant color photography is typeset by the statement:

```
\begin{picture}(4000,2000)(0,-1000)
\put(0,0){\bzdrv{1==OH;2==NH--SO$_{2}$;4==OC$_{16}$H$_{33}$;5==CH$_{3}$$}}
\put(993,230){\bzdrh{1==;2==OCH$_{2}$CH$_{2}$OCH$_{3}$;%
5==NH--SO$_{2}$$}}
\put(1890,-140){\bzdrh{1==;5==SO$_{2}$--NH}}
\put(2750,-850){\naphdrh{1==;5==OH;8==}}
\put(1800,-850){\bzdrh{1==O$_{2}$N;5==SO$_{3}$CH$_{3}$$;4==N=N}}
\end{picture}
%}
```

These commands produce the following structure:



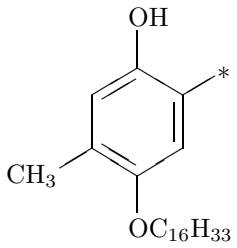
Chapter 15

Large Substituents

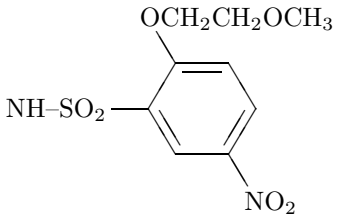
15.1 Basic Ideas

In all of the preceding chapters, any substituents described in SUBSLIST are rather simple ones, which at most vary from an atom of one- or two-character to a group of several characters. How about such a complex substituent as produced by a macro?

Let us consider the substitution of



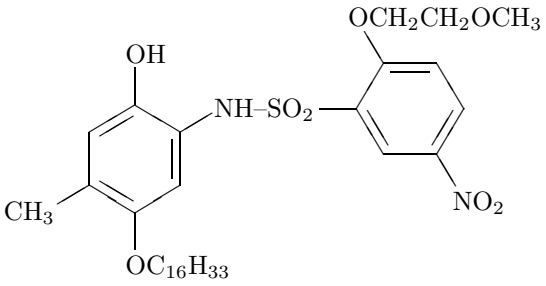
at the 2-position (∗)with the substituent represented by



This task can be accomplished in the light of the technique introduced in the preceding chapter. Thus, the statement

```
\begin{picture}(2000,1000)(0,0)
\put(0,0){\bzdrv{1==OH;2==;4==OC$_{16}$H$_{33}$;5==CH$_{3}$}}
\put(993,230){\bzdrh{1==NH--SO$_{2}$;%
2==OCH$_{2}$CH$_{2}$OCH$_{3}$;5==NO$_{2}$}}
\end{picture}
```

provides



This methodology implies that both of the parts are regarded as fragments to be combined together.

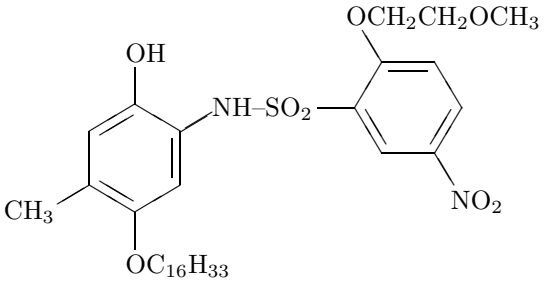
On the other hand, another useful technique is available, if you use the TEX command `\setbox` and the related commands. In the light of this technique, either one is regarded as a substituent of the other. First, a structure regarded as a substituent is constructed in a `\hbox` and stored in `\box4` by means of the command `\setbox` as follows:

```
\setbox4=\hbox{%
 \begin{picture}(0,0)(-285,370)%
%  \put(-285,370){\circle{50}}%change reference point
  \put(0,0){\bzdrh{1==NH--SO$_{2}$;2==OCH$_{2}$CH$_{2}$OCH$_{3}$;%
   5==NO$_{2}$}}%
 \end{picture}}%
```

The inner picture environment has the width of 0pt and the height of 0pt, where the reference point is shifted into the $(-285, 370)$ point which is the rightmost point of the NH–SO_2 group. This reference point is regared as the $(0, 0)$ point of the substituent stored in `\box4`. Then, the substituent `\box4` is written in SUBSLIST of the command `\bzdrv`, *i.e.*,

```
\bzdrv{1==OH;2==\box4;4==OC$_{16}$H$_{33}$;5==CH$_{3}$}
```

This statement produces



It should be noted that the token `2==\box4` creates such a complex fragment that is impossible to be directly assigned to an argument list. This technique is also useful to avoid the overcrowding of substituents, since the reference point of the substituent can be changed appropriately.

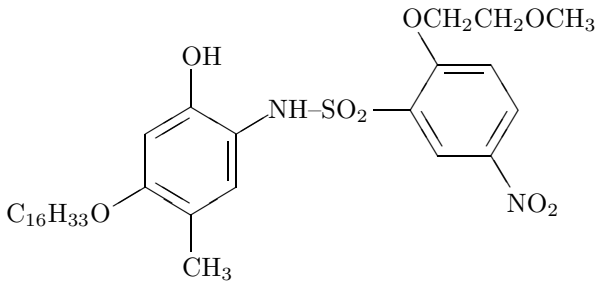
When you multiple times use the stored substituent, you can use the command `\copy` instead of `\box`:

```
\bzdrv{1==OH;2==\copy4;4==OC$_{16}$H$_{33}$;5==CH$_{3}$}
```

Then, you are able to use the stored substituent in another context.

```
\setbox5=\hbox{%
\bzdrv{1==OH;2==\box4;5==C$_{16}$H$_{33}$O;4==CH$_{3}$}}%
\mbox{\box5}
```

This statement provides another derivative having the same substituents.

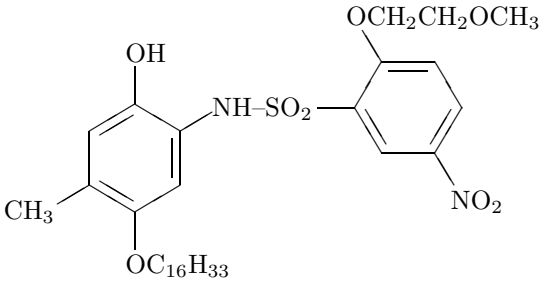


An alternative way of treating a large substituent is to use the definition of a tentative macro such as `\phsulphonyl`:

```
\def\phsulphonyl#1{%
 \begin{picture}(0,0)(-285,370)%
%  \put(-285,370){\circle{50}}%change reference point
  \put(0,0){\bzdrh{1==NH--SO$_{2}$;2==OCH$_{2}$CH$_{2}$OCH$_{3}$;%
   5==#1}}%
 \end{picture}}%
```

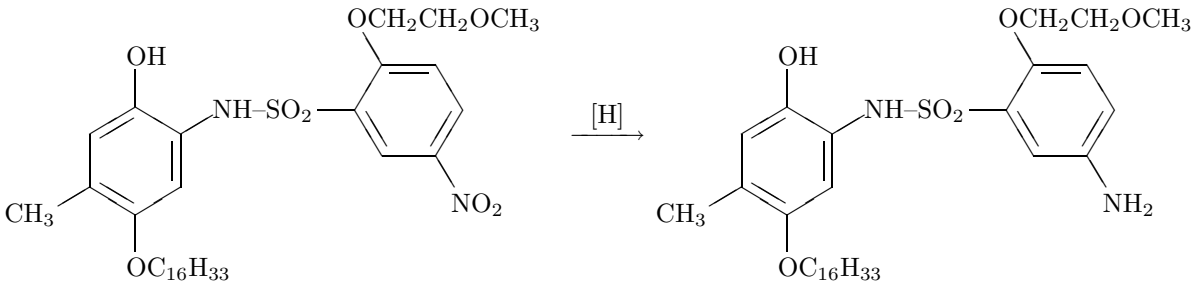
which has an argument to select a phenyl subsituent. Then, the macro can be used in the SUBSLIST of a mother structure command. For example, the compound described above can be typeset as follows:

```
\bzdrv{1==OH;2=={\phsulphonyl{NO$_{2}$}};%
 4==OC$_{16}$H$_{33}$;5==CH$_{3}$}
```



Since the substituent printed by `\phsulphonyl` is regarded to have no width and no height, the size of the resulting formula should be reset to have an appropriate width and height for further use. The following example shows a resetting method to make a box of an appropriate size by means of a picture environment.

```
\begin{trivlist}\item[]
\begin{picture}(2000,1000)(-100,0)
\put(0,0){\bzdrv{1==OH;2=={\phsulphonyl{NO$_{2}$}};%
 4==OC$_{16}$H$_{33}$;5==CH$_{3}$}}
\end{picture}
\hfill \reactrarrow{50pt}{1cm}{[H]}{\strut} \hfill
\begin{picture}(2000,1000)(0,0)
\put(0,0){\bzdrv{1==OH;2=={\phsulphonyl{NH$_{2}$}};%
 4==OC$_{16}$H$_{33}$;5==CH$_{3}$}}
\end{picture}
\end{trivlist}
```



15.2 Nested Substituents

The latest sample reveals that a structure constructed by the present techique can be further nested to be a substituent of another macro. The following example illustrates multiple nesting for drawing the same dye releaser as depicted in the preceding chapter.

First, the formula of 2-methanesulfonyl-4-nitro-phenyl-1-azo group (A) is constructed in the box `\box4` by means of following statement:

```
\setbox4=\hbox{%
 \begin{picture}(0,0)(996,370)%
%  \put(996,370){\circle{50}}%
  \put(0,0){\bzdrh{1==O$_{2}$N;5==SO$_{3}$CH$_{3}$;4==N=N}}%
 \end{picture}}%
```

Note that the value (996,370) results in the shift of the reference point into the rightmost terminal of the azo group, which is a linking point in the next step. The formula (A) strored in `\box4` is placed at the 8-position of a naphalene ring. The resulting formula (B) is, in turn, stored into `\box5`.

```
\setbox5=\hbox{%
 \begin{picture}(0,0)(-250,712)%
%  \put(-250,712){\circle{50}}%
  \put(0,0){\naphdrh{1==SO$_{2}$NH;5==OH;8==\box4}}%
 \end{picture}%
 }%
```

The value $(-250, 712)$ shifts the reference point into the leftmost terminal of the sulfonamido group at the 1-position of the naphthalene ring. The formula B an a sulfamoyl group are placed at the meta position of a benzen ring to produce formula C.

```
\setbox4=\hbox{%
 \begin{picture}(0,0)(-285,370)%
%  \put(-285,370){\circle{50}}%
  \put(0,0){\bzdrh{1==NH--SO$_{2}$;5==\box5}}%
 \end{picture}%
}%
```

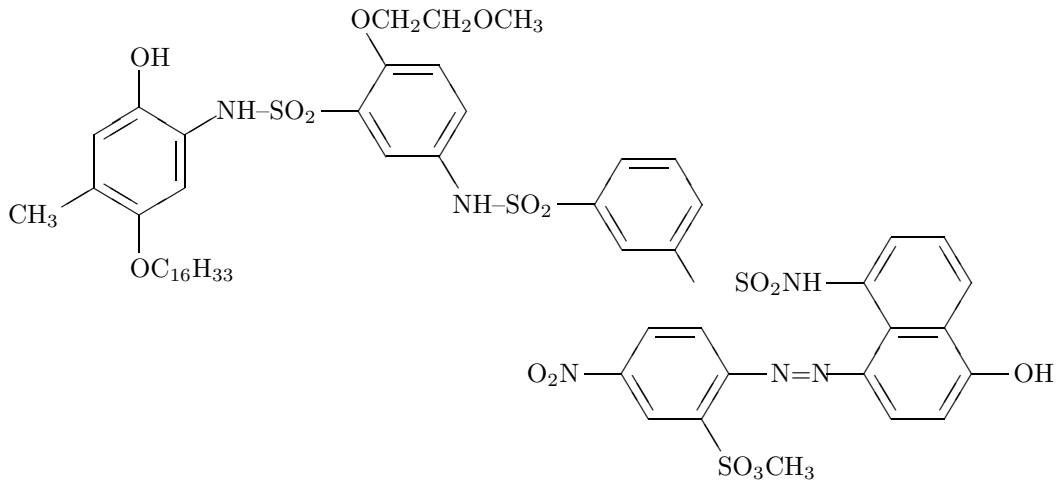
The resulting formula C is further placed on another benzene ring to generate the formula (D) of a complex substituent.

```
\setbox5=\hbox{%
 \begin{picture}(0,0)(-285,370)%
%  \put(-285,370){\circle{50}}%
  \put(0,0){\bzdrh{1==NH--SO$_{2}$;2==OCH$_{2}$CH$_{2}$OCH$_{3}$;%
   5==\box4}}%
 \end{picture}%
}%
```

Finally, the substituent D is placed at the ortho position to a hydroxyl group on a benzene ring.

```
\setbox4=\hbox{%
\bzdrv{1==OH;2==\box5;4==OC$_{16}$H$_{33}$;5==CH$_{3}$}}%
\mbox{\box4}
```

The formula stored in \box4 is printed by means of the command \box4, giving the following structure.



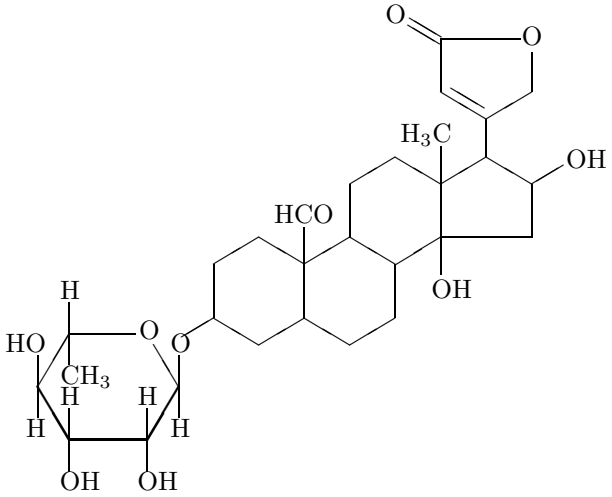
One of the merits of the present methodology is that we can use relative coordinates in each step of combining two structures. Hence, the calculation of coordinates is simpler than that based on the method of the preceding chapter.

The structural formula of adonitoxin can be written in a similar way, where two complex substituents stored in \box0 and \box1 are placed on a steroid skeleton.

```
\setbox0=\hbox{%
 \begin{picture}(0,0)(369,257)%
%  \put(369,257){\circle{50}}%
  \put(0,0){\fiveheterov[e]{3==O}{4D==O}}
 \end{picture}}%
\setbox1=\hbox{%
 \begin{picture}(0,0)(772,530)%
%  \put(772,530){\circle{50}}%
  \put(0,0){\pyranose{1Sb==O;1Sa==H;2Sb==H;2Sa==OH;3Sb==H;3Sa==OH;%
  4Sb==HO;4Sa==H;5Sb==H;5Sa==CH$_{3}$}}%
 \end{picture}}%
\setbox2=\hbox{%
   \steroid{{{10}}==\lmoiety{HCO\kern-.7em};{{14}}==OH;%
   {{13}}==\lmoiety{H$_{3}$C};%
   {{16}}==OH;{{17}}==\box0;3==\box1}}%
\medskip
\begin{center}
%\fbox{
\begin{picture}(2500,1800)(-600,-300)
\put(0,0){\mbox{\box2}}
\end{picture}
```

```
%}
\end{center}
```

These commands produce



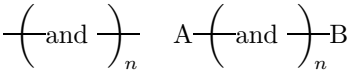
Chapter 16

Polymers

16.1 Polymer Delimeters as Substituents

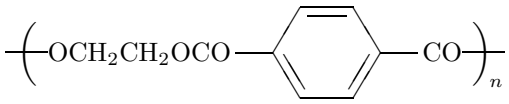
The commands `\leftpolymer` and `\rightpolymer` draw parentheses when used in the SUBSLIST of a XyMTEX command. For example, they are combined the `\sbond` command (for drawing a single bond) to typeset a sigle bond crossed with a parenthesis:

```
\leftpolymer{}\sbond and \sbond\rightpolymer{}{n} \qquad
\leftpolymer{A}\sbond and \sbond\rightpolymer{B}{n}
```



They are used pairwise to indicate a polymer unit, as shown in the following code for drawing polyethylene terephthalate.

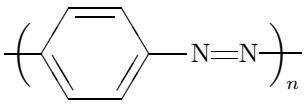
```
\hskip6cm
\bzdrh{1=={\leftpolymer{}\sbond OCH$_{2}$CH$_{2}$OCO};%
4=={CO\sbond\rightpolymer{}{n}}}
```



Note that the designation of the polymer is treated as the description of a substituent by the command `\bzdrh`. A long subsituent appearing in the left-hand side of this example is regarded as having no length; hence the adjustment with `\hskip6cm` is necessary.

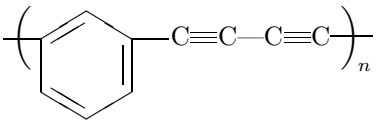
A poly-azophenylene can be drawn in the light of the same methodology.

```
\bzdrh{1=={\leftpolymer{}};%
4=={N\dbond N\sbond \rightpolymer{}{n}}}
```



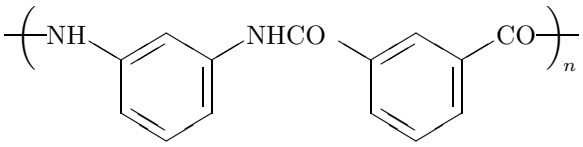
The following example uses the \bzdrv command, where horizontal valences at the 2- and 6- positions are typeset by the bond modifier Sa.

```
\bzdrv{6Sa=={\leftpolymer{}};%
2Sa=={C\tbond C---C\tbond C\sbond \rightpolymer{}{n}}}
```



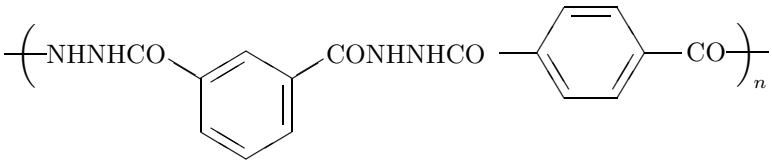
The following code for drawing poly-*m*-phenylene isophthalamide contains two successive \bzdrv commands.

```
\bzdrv{6=={\leftpolymer{}\sbond NH};2==NHCO}\kern14pt
\bzdrv{2=={CO\sbond\rightpolymer{}{n}};6==}
```



The following code uses the commands \bzdrv and \bzdrh for drawing an aromatic polyhydrazide. Since their original points for drawing are different to each other, the vertical adjustment with the \raisebox command is necessary to assure the horizontal alignment of bonds.

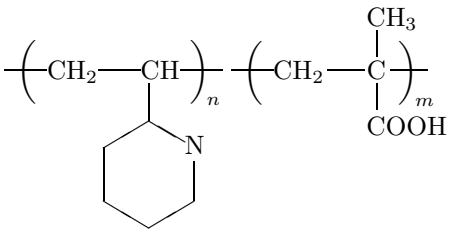
```
\bzdrv{6=={\leftpolymer{}\sbond NHNHCO};2==CONHNHCO}
\kern420\unitlength
\raisebox{24pt}{\bzdrh{4=={CO\sbond\rightpolymer{}{n}};1==}}
```



The combination of \sixheterov and \tetrahedral in an outer picture environment enables us to draw 2-vinylpyridine-methacrylic acid block polymer, where \leftpolymer and \rightpolymer are used to show a polymer unit.

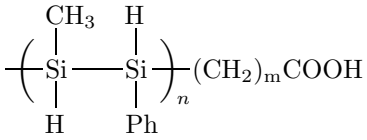
```
\begin{picture}(1700,1100)(-200,-500)
\put(0,-535){\sixheterov{2==N}{1=={%
\hbox to0pt{\hss\leftpolymer{}\sbond CH$_{2}$\sbond}%
CH\sbond\rightpolymer{}{n}}}}
```

```
\put(960,0){\tetrahedral{0==C;1==CH$_{3}$;%
2=={\leftpolymer{}\sbond CH$_{2}$};%
3==COOH;4=={\rightpolymer{}{m}}}}
\end{picture}
```



The following example shows the use of two `\tetrahedral` commands in an outer picture environment.

```
\begin{picture}(1000,600)(0,0)
\put(0,0){\tetrahedral{%
 0==Si;1==CH$_3$;%
 2=={\leftpolymer{}};% no teminal atoms
 3==H;4==}}
\put(300,0){\tetrahedral{%
 0==Si;1==H;%
 4=={\rightpolymer{(CH$_{2}$)$_{\mbox{\scriptsize\rm m}}$COOH}{n}};%
 2==;3==Ph}}
\end{picture}
```

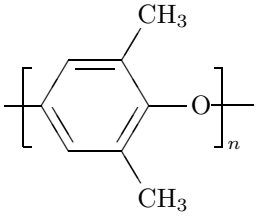


The height of parentheses can be changed by using `\leftPolymer` and `\rightPolymer` in which a desired delimeter is designated as an argument. Thus, an ethyenimine-succinimide copolymer is typeset by this technique.

```
\leftPolymer{(}{}\sbond CH$_{2}$CH$_{2}$CONHCH$_{2}$CH$_{2}$NH%
\sbond\rightPolymer{)}{}{n}
```

The parentheses can be changed into barackets by using `\leftsqrpolymer` and `\rightsqrpolymer`.

```
\bzdrh{%
1=={\leftsqrpolymer{}};%
3==CH$_{3}$;5==CH$_{3}$;%
4=={O\sbond\rightsqrpolymer{}{n}}}
```

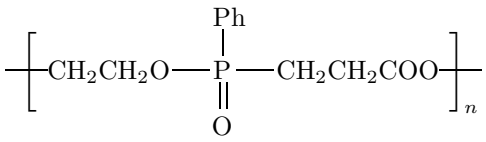


In a similar way, poly-*p*-xylylene is drawn as follows.

```
\bzdrh{%
1=={\leftsqrpolymer{}\sbond CH$_{2}$};%
4=={CH$_{2}$\sbond\rightsqrpolymer{}{n}}}
```

The following example uses the `\tetrahedral` command for drawing phenyldioxaphosphorane-acrylic acid copolymer, where thick-line brackets produced by `\leftSqrpolymer` and `\rightSqrpolymer` are used as polymer delimiters.

```
\tetrahedral{%
0==P;1==Ph;%
2=={\leftSqrpolymer{}\sbond CH$_{2}$CH$_{2}$O};%
3D==O;%
4=={CH$_{2}$CH$_{2}$COO\sbond\rightSqrpolymer{}{n}}}
```

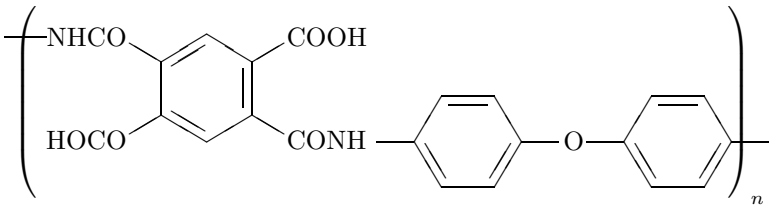


16.2 Polymer delimeters as whole enclosures

The command `\mpolymer` takes two arguments; the first argument is a polymer unit and the second is a repeating number. It measures the height of the polymer unit and surrounds the unit with parentheses. Thus, the code

```
\mpolymer{%
\begin{picture}(2600,700)(-240,200)
\put(0,158){\bzdrv{2==COOH;3==CONH;5==HOCO;6=={\sbond NHCO}}}
\put(940,0){\bzdrh{1==;4==O}}
\put(1730,0){\bzdrh{1==;4==}}
\end{picture}}{n}
```

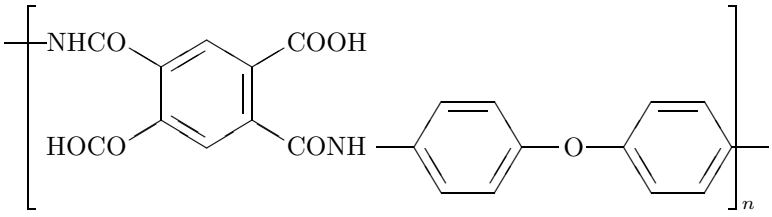
produces



The command `\sqrpolymer` has the same function as `\mpolymer` except that it surrounds a polymer unit with thin-line brackets. The short declaration `\sqrpolymer{TEXT}{}` produces $\left[\mathrm{TEXT}\right]$ and the statement:

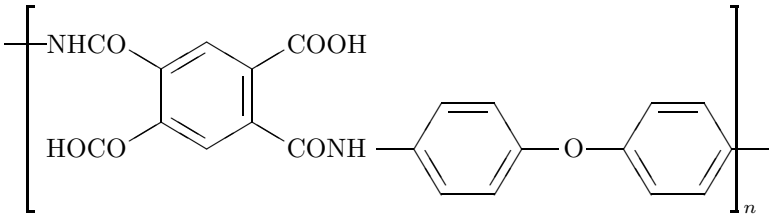
```
\sqrpolymer{%
\begin{picture}(2600,700)(-240,200)
\put(0,158){\bzdrv{2==COOH;3==CONH;5==HOCO;6=={\sbond NHCO}}}
\put(940,0){\bzdrh{1==;4==O}}
\put(1730,0){\bzdrh{1==;4==}}
\end{picture}}{n}
```

produces a polymer:



On the other hand, the command `\Sqrpolymer` surrounds a polymer unit with thick-line brackets.

```
\Sqrpolymer{%
\begin{picture}(2600,700)(-240,200)
\put(0,158){\bzdrv{2==COOH;3==CONH;5==HOCO;6=={\sbond NHCO}}}
\put(940,0){\bzdrh{1==;4==O}}
\put(1730,0){\bzdrh{1==;4==}}
\end{picture}}{n}
```



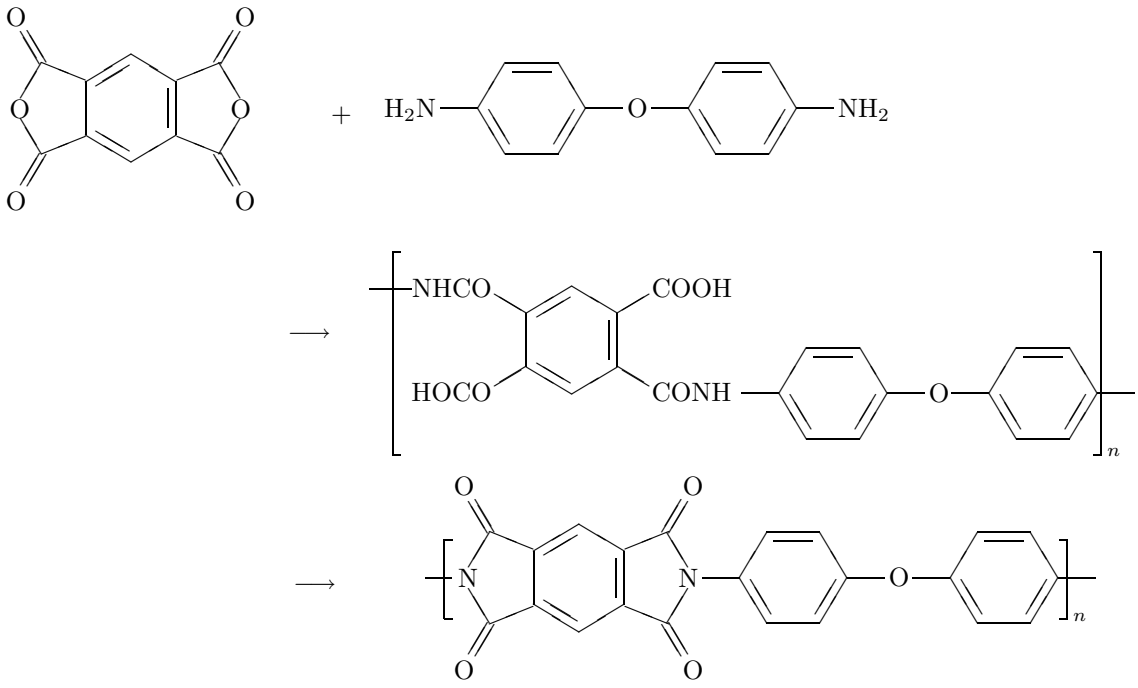
A scheme for preparing a polymer is typeset by combining the commands described above. Let us first prepre a macro `\pyromellitic` for drawing pyromellitic acid derivatives.

```
\def\pyromellitic#1#2#3#4{%
\begin{picture}(1200,0)(0,400)
\def\kktmp{#1}
\ifx\kktmp\empty
\put(0,0){\fiveunithi{1==#3}{2D==O;5D==O}{bcd}}
\else
\put(0,0){\fiveunithi{1==#3}{1==#1;2D==O;5D==O}{bcd}}
\fi
\put(343,573){\line(5,-2){165}}
\put(343,227){\line(5,2){165}}
\put(280,-43){\bzdrv{}}
\put(1019,573){\line(-5,-2){165}}
\put(1019,227){\line(-5,2){165}}
\def\kktmp{#2}
\ifx\kktmp\empty
\put(468,0){\fiveunith{1==#4}{2D==O;5D==O}{bcd}}
\else
\put(468,0){\fiveunith{1==#4}{1==#2;2D==O;5D==O}{bcd}}
\fi
\end{picture}}
```

The first and second arguments of the `\pyromellitic` show the presence of exocyclic valences for polymerization. The third and fourth arguments show the hetero atoms on the five-membered rings.

The preparation of a poly-pyromellitimide from pyromellitic anhydride and an diamine is illustrated as follows.

```
% the first line
\raisebox{400\unitlength}{\pyromellitic{}{}{O}{O}}
\qquad \raisebox{350\unitlength}{+} \qquad
\bzdrh{1==H$_{2}$N;4==O}
\hskip-120\unitlength
\bzdrh{1==;4==NH$_{2}$}
% the 2nd line
\begin{flushright}
\raisebox{400\unitlength}{$\longrightarrow$} \qquad
\sqrpolymer{%
\begin{picture}(2600,700)(-240,200)
\put(0,158){\bzdrv{2==COOH;3==CONH;5==HOCO;6=={\sbond NHCO}}}
\put(940,0){\bzdrh{1==;4==O}}
\put(1730,0){\bzdrh{1==;4==}}
\end{picture}}{n} \\[5pt]
% the 3rd line
\raisebox{350\unitlength}{$\longrightarrow$} \qquad
\raisebox{400\unitlength}{%
\pyromellitic{{\leftsqrpolymer{}}}{}{N}{N}}
\hskip-190\unitlength
\bzdrh{1==;4==O}
\hskip-120\unitlength
\bzdrh{1==;4=={\rightsqrpolymer{}{n}}}
\end{flushright}
```

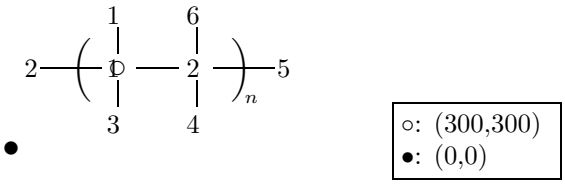


16.3 Polyethylene unit

The command `\polyethylene` is used to draw polyethylene derivatives, in which each substituent is designated by the SUBSLIST. The format of this command is as follows:

```
\polyethylene[AUXLIST]{CALIST}{SUBSLIST}
```

The following diagram shows the numbering for designating substituents and center-atom positions.



in which the same macro is used to typeset both saturated and unsaturated derivatives.

The optional argument AUXLIST is used to specify a charge on the central atoms: *i.e.*, {n+} represents a + charge (or another one chararacter) on the n-center.

The argument CALIST indicates central atoms 1 and 2, *e.g.*, `1==C` and `2==Si`. A double bond and a triple bond between the centeral atoms can be designated by writing `0D==` an `0T==`, respectively.

The argument SUBSLIST is used to specify each substituent with a locant number and a bond modifier shown in Table 16.1, in which n is an arabic numeral between 1 and 4.

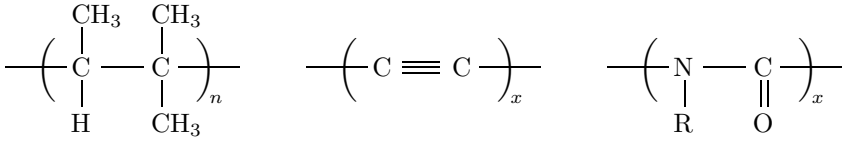
Table 16.1: SUBSLIST for `\polyethylene`

Character	Structures printed
nT	triple bond at n-atom
nD	double bond at n-atom
n or nS	single bond at n-atom
nA	alpha single bond at n-atom
nB	beta single bond at n-atom

Example:

```
\polyethylene{1==C;2==C}%
 {1==CH$_{3}$;2==;3==H;4==CH$_{3}$;5==;6==CH$_{3}$;0==n}
\polyethylene{1==C;2==C;0T==}{2==;5==}
\polyethylene{1==N;2==C}{2==;5==;3==R;4D==O}
```

produce the following structures:



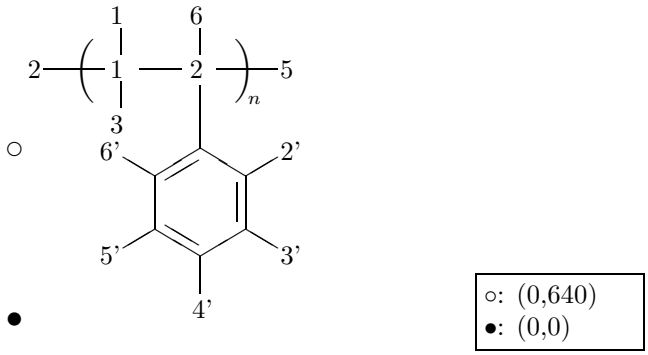
16.4 Polystyrene unit

The command `\polystyrene` is used to draw polystyrene derivatives, in which substituents on both the polymer chain and the phenyl group are designated by the SUBSLIST and PHSUBSLIST, respectively.

The format of this command is as follows:

```
\polystyrene[AUXLIST]{CALIST}{SUBSLIST}{PHSUBSLIST}
```

The following diagram shows the numbering for designating substituents and center-atom positions.



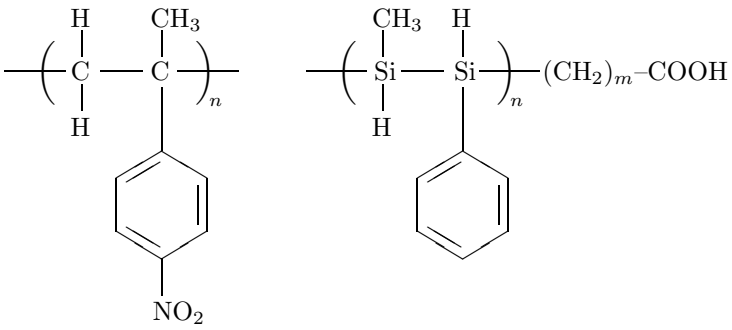
in which the same macro is used to typeset both saturated and unsaturated derivatives.

The arguments AUXLIST, CALIST and SUBSLIST have the same meanings as described for the command `\polyethylene` (see Table 16.1). The argument PHSUBSLIST is used to indicate substituents on the phenyl group. For example, `n` or `nS` shows the presence of a single bond at the n-atom of the phenyl.

Example:

```
\polystyrene{}%
{1==H;2==;3==H;5==;6==CH$_{3}$;0==n}{4==NO$_{2}$}
\polystyrene{1==Si;2==Si}{6==H;2==;3==H;%
5=={(CH$_{2}$)$_{m}$--COOH};%
1==CH$_{3}$;0==n}{}
```

produce the following structures:



Chapter 17

Chemical Environments

This chapter deals with several useful tools provided by chemist.sty.

17.1 Chemical Equations

17.1.1 The 'chem' Version

If you write a chemical equation by using the equation environment of LaTeX, *e.g.*,

```
\begin{equation}
2H_{2} + O_{2} \rightarrow 2H_{2}O
\end{equation}
```

you obtain an insufficient result:

$$2H_2 + O_2 \rightarrow 2H_2O \tag{17.1}$$

where alphabetical characters for atomic symbols are typeset in an italic typeface, though they should be printed in a roman typeface. For remedying this situation, you may write a chemical equation in the equation environment by using the `\textrm` command as follows.

```
\begin{equation}
2\textrm{H}_{2} + \textrm{O}_{2} \rightarrow 2\textrm{H}_{2}\textrm{O}
\end{equation}
```

$$2\mathrm{H}_2 + \mathrm{O}_2 \rightarrow 2\mathrm{H}_2\mathrm{O} \tag{17.2}$$

However, this way of writing chemical equations is somewhat tedious because *every* roman character is designated with the `\textrm` command.

If you have to typeset an equation containing subscripts and/or superscripts, you can use the `\mathrm` command as shown in the follwing example.

```
\begin{equation}
\mathrm{2KMn^{VII}O_{4} + 16HCl =
2Mn^{II}Cl_{2} + 5Cl_{2} + 2KCl + 8H_{2}O}
\end{equation}
```

$$\mathrm{2KMn^{VII}O_4 + 16HCl = 2Mn^{II}Cl_2 + 5Cl_2 + 2KCl + 8H_2O} \tag{17.3}$$

This method is simpler than the `\textrm` method described above. However, it is necessary to declare two or more `\mathrm` commands when an eqnarray environment is used, *e.g.*,

```
\begin{eqnarray}
\mathrm{2KMn^{VII}O_{4} + 16HCl} & = &
\mathrm{2Mn^{II}Cl_{2} + 5Cl_{2} + 2KCl + 8H_{2}O} \\
\mathrm{2In^{0}(s) + 3Hg^{I}{}_{2}SO_{4}(s)} & = &
\mathrm{In^{III}{}_{2}(SO_{4})_{3}(aq) + 6Hg^{0}(l)}
\end{eqnarray}
```

$$\begin{aligned} \mathrm{2KMn^{VII}O_4 + 16HCl} &= \mathrm{2Mn^{II}Cl_2 + 5Cl_2 + 2KCl + 8H_2O} && (17.4)\\ \mathrm{2In^0(s) + 3Hg^I{}_2SO_4(s)} &= \mathrm{In^{III}{}_2(SO_4)_3(aq) + 6Hg^0(l)} && (17.5) \end{aligned}$$

The chemist package of XƎMTEX provides the 'chem' version to settle this problem. Note that the chem version is declared as an argument of the `\mathversion`, since it is an additional mathematical version added to the original 'normal' and 'bold' vesrions of LaTeX 2_ε.

```
\mathversion{chem}
\begin{equation}
2H_{2} + O_{2} \rightarrow 2H_{2}O \label{chemenv:a1}
\end{equation}
\begin{equation}
2KMn^{VII}O_{4} + 16HCl =
2Mn^{II}Cl_{2} + 5Cl_{2} + 2KCl + 8H_{2}O
\end{equation}
```

This statement provides the following equation.

$$\mathrm{2H_2 + O_2 \rightarrow 2H_2O} \tag{17.6}$$

$$\mathrm{2KMn^{VII}O_4 + 16HCl = 2Mn^{II}Cl_2 + 5Cl_2 + 2KCl + 8H_2O} \tag{17.7}$$

Since the chem version is a declaration-type command, the above declation is effective afterward. Thus, the following eqnarray environment is typeset under such effect without using any further `mathversion` command.

```
\begin{eqnarray}
Me_{3}B + LiMe & \rightarrow & Me_{4}B^{-}Li^{+} \\
Ph_{5}B + LiPh & \rightarrow & Ph_{6}B^{-}Li^{+}
\end{eqnarray}
```

$$\begin{aligned} \mathrm{Me_3B + LiMe} &\rightarrow \mathrm{Me_4B^-Li^+} && (17.8)\\ \mathrm{Ph_5B + LiPh} &\rightarrow \mathrm{Ph_6B^-Li^+} && (17.9) \end{aligned}$$

If you write a mathematical equation, the mathematical formula version of your manuscript should be returned to the 'normal' version as follows.

```
\mathversion{normal}
\begin{eqnarray}
x^{3}+y^{3} & = & (x+y)(x^{2}-xy+y^{2}) \\
x^{5}+y^{5} & = & (x+y)(x^{4}-x^{3}y+x^{2}y^{2}-xy^{3}+y^{4})
\end{eqnarray}
```

$$\begin{aligned} x^3 + y^3 &= (x+y)(x^2 - xy + y^2) && (17.10)\\ x^5 + y^5 &= (x+y)(x^4 - x^3y + x^2y^2 - xy^3 + y^4) && (17.11) \end{aligned}$$

17.1.2 Chemical Corrections

A chemical symbol with both a subscript and a superscript has the subscript at a different vertical level from the counterpart of a symbol with a subscript only. For example, the subscript of M_2^{II} is different in the bottom level from that of FeO_4 in the formula $M_2^{II}FeO_4$ (see [1] Chapter 18). For aligning the bottoms of such subscripts, we introduce 'chemical corrections' and define a macro `\chemform`. Compare the following examples with special attention to the printing places of the subscripts.

```
$\mathrm{M}_{2}^{\mathrm{II}}\mathrm{FeO}_{4}$ (uncorrected)
{\it vs.}
\chemform{M_{2}^{II}FeO_{4}} (corrected)
```

$$\mathrm{M}_{2}^{\mathrm{II}}\mathrm{FeO}_{4}\ \text{(uncorrected)}\ \textit{vs.}\ \mathrm{M}_{2}^{\mathrm{II}}\mathrm{FeO}_{4}\ \text{(corrected)}$$

Since the `\chemform` command is defined to take account of the chem version automatically, it is unnecessary to use the command `\mathrm` or `\textrm`.

17.1.3 The 'chemmath' and 'chemeqn' environments

For typesetting chemical equations, we use the chemmath environment of the chemist package, which corresponds to the displaymath environment of LaTeX (see [8] Chapter 9). Note that it requires no declaration of the chem version by `\mathversion`.

```
\begin{chemmath}
H_{2} + Cl_{2} \rightarrow 2HCl
\end{chemmath}
```

$$\mathrm{H}_{2} + \mathrm{Cl}_{2} \rightarrow 2\mathrm{HCl}$$

For numbering chemical equations, we use the chemeqn environment of the chemist package. The environment corresponds to the equation environment of LaTeX; but it requires no declaration of the chem version by `\mathversion`. Compare eq. 17.12 with eq. 17.6.

```
\begin{chemeqn}
2In^{0}(s) + 3Hg^{I}{}_{2}SO_{4}(s) =
In^{III}{}_{2}(SO_{4})_{3}(aq) + 6Hg^{0}(l) \label{chemenv:a2}
\end{chemeqn}
```

$$2\mathrm{In}^{0}(\mathrm{s}) + 3\mathrm{Hg}^{\mathrm{I}}{}_{2}\mathrm{SO}_{4}(\mathrm{s}) = \mathrm{In}^{\mathrm{III}}{}_{2}(\mathrm{SO}_{4})_{3}(\mathrm{aq}) + 6\mathrm{Hg}^{0}(\mathrm{l}) \tag{17.12}$$

17.1.4 The 'chemeqnarray' and like environments

Two or more chemical equations related to each other are typeset by using the chemeqnarray environment of the chemist package, which corresponds to the eqnarray environment of LaTeX.

```
\begin{chemeqnarray}
Me_{3}B + LiMe & \rightarrow & Me_{4}B^{-}Li^{+} \\
Ph_{5}B + LiPh & \rightarrow & Ph_{6}B^{-}Li^{+}
\end{chemeqnarray}
```

$$\begin{aligned} \mathrm{Me}_{3}\mathrm{B} + \mathrm{LiMe} &\rightarrow \mathrm{Me}_{4}\mathrm{B}^{-}\mathrm{Li}^{+} && (17.13)\\ \mathrm{Ph}_{5}\mathrm{B} + \mathrm{LiPh} &\rightarrow \mathrm{Ph}_{6}\mathrm{B}^{-}\mathrm{Li}^{+} && (17.14) \end{aligned}$$

They are also typeset by using the chemeqnarray$*$ environment of the chemist package without numbering. This corresponds to the eqnarray$*$ environment of LaTeX.

```
\begin{chemeqnarray*}
Me_{3}B + LiMe & \rightarrow & Me_{4}B^{-}Li^{+} \\
Ph_{5}B + LiPh & \rightarrow & Ph_{6}B^{-}Li^{+}
\end{chemeqnarray*}
```

$$\begin{aligned} \mathrm{Me_3B + LiMe} &\rightarrow \mathrm{Me_4B^-Li^+} \\ \mathrm{Ph_5B + LiPh} &\rightarrow \mathrm{Ph_6B^-Li^+} \end{aligned}$$

The mode of numbering can be changed by using the chemeqnarraya environment of of the chemist package, which corresponds to the eqnarraya environment of the mathchem package [8].

```
\begin{chemeqnarraya}
Me_{3}B + LiMe & \rightarrow & Me_{4}B^{-}Li^{+}
 \label{chemenv:a3} \\
Ph_{5}B + LiPh & \rightarrow & Ph_{6}B^{-}Li^{+}
\end{chemeqnarraya}
```

$$\begin{aligned} \mathrm{Me_3B + LiMe} &\rightarrow \mathrm{Me_4B^-Li^+} && \text{(17.15a)} \\ \mathrm{Ph_5B + LiPh} &\rightarrow \mathrm{Ph_6B^-Li^+} && \text{(17.15b)} \end{aligned}$$

Sub-numbering references can be referred by using such a command as `\ref{chemenv:a3}`, which results in, *e.g.*, 17.15a.

17.2 Arrows for Chemical Equations

Various arrows have been defined in chemist.sty for typsetting chemical equations (Table 17.1). See Ref. [8] for detailed description.

Table 17.1: Various arrows for chemical equations

		`\leftrihgtarrowfill`
`\Leftarrowfill`	`\Rightarrowfill`	`\Leftrihgtarrowfill`
`\reactrarrow`	`\reactlarrow`	`\reactlrarrow`
		`\reacteqarrow`
`\schemerarrow`	`\schemelarrow`	`\schemelrarrow`

Note that the commands `\leftarrowfill` and `\rightarrowfill` have been already defined in the original TEX/LATEX, though Table 17.1 does not involve them.

17.3 Boxes for Chemical Formulas

Various boxes have been defined in chemist.sty for typsetting chemical formulas (Table 17.2). See Ref. [8] for detailed description.

17.4 Cross-References of Compounds

17.4.1 Counters for Compounds

The macro `\compd` assigns a reference number to a chemical compound. The reference number can be referred to by the LATEX cross-reference techinique with `\label` and `\ref`. In the light of chemist.sty, the

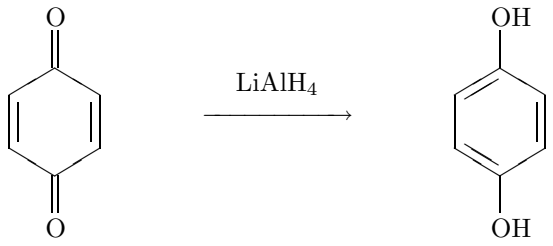
Table 17.2: Various boxes for chemical equations

\compdfbox	\compdmbox	\lbcompdpbox
\derivfbox	\derivmbox	\lbderivpbox
\cdonecell	\cdtwocell	

\cref command can be used for cross-references of chemical compounds, where a compound number is typeset with a boldfaced character.

```
Example: 1,4-Benzoquinone (\cref{cmd:a1}) is reduced into
hydroquinone (\cref{cmd:a2}) by lithium aluminum hydride as
shown in the following scheme.
\begin{center}
\begin{tabular}{c}
\bzdrv[p]{1D==O;4D==O} \\[.3cm] \compd \label{cmd:a1} \\
\end{tabular}
\begin{tabular}{c}
LiAlH$_{4}$ \\
\parbox{2cm}{\rightarrowfill} \\[1cm] \mathstrut \\
\end{tabular}
\begin{tabular}{c}
\bzdrv{1==OH;4==OH} \\[.3cm] \compd \label{cmd:a2} \\
\end{tabular}
\end{center}
```

Example: 1,4-Benzoquinone (**1**) is reduced into hydroquinone (**2**) by lithium aluminum hydride as shown in the following scheme.



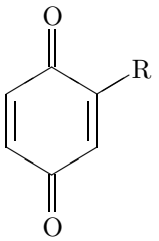
1 **2**

Derivatives of a compound are numbered by using the \deriva command, after the \compd command is declared. An alternative numbering can be obtained by using the \deriv command, after the \nocompd command is used.

```
\begin{center}
\begin{tabular}{c}
\bzdrv[p]{1D==O;4D==O;2==R} \\ \compd \label{cmd:a3} \\
\end{tabular}
\begin{tabular}{ll}
\deriva \label{drv:a3a} & R = H \\
\deriva \label{drv:a3b} & R = CH$_{3}$ \\
\deriva \label{drv:a3c} & R = C(CH$_{3}$)$_{3}$ \\
\deriva \label{drv:a3d} & R = Cl \\
```

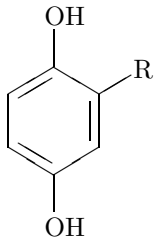
```
\end{tabular}
\qquad
\begin{tabular}{c}
\bzdrv{1==OH;4==OH;2==R} \\ \nocompd \label{cmd:a4} \\
\end{tabular}
\begin{tabular}{ll}
\deriv \label{drv:a4a} & R = H \\
\deriv \label{drv:a4b} & R = CH$_{3}$ \\
\deriv \label{drv:a4c} & R = C(CH$_{3}$)$_{3}$ \\
\deriv \label{drv:a4d} & R = Cl \\
\end{tabular}
\end{center}
```

The left-hand formula is an example of the former and the right-hand one is an example of the latter technique.



3

a R = H
b R = CH_3
c R = $C(CH_3)_3$
d R = Cl



4a R = H
4b R = CH_3
4c R = $C(CH_3)_3$
4d R = Cl

In both of the cases, such cross-references as `\cref{drv:a3a}` and `\cref{drv:a4c}` typeset compound numbers as being **3a** and **4c**. The parent numbers such as **3** and **4** can be printed by using `\cref{cmd:a3}` and `\cref{cmd:a4}`.

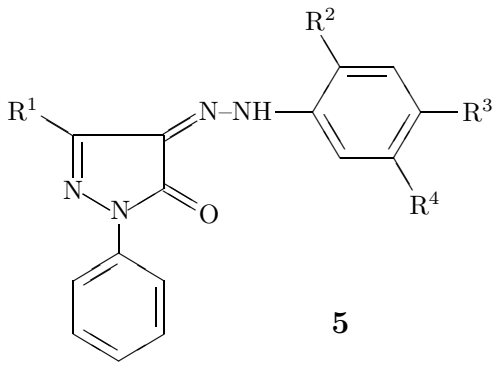
The following tabular matter illustrates a more complicated application of cross-reference.[1] The strucutural formula of this example is typeset in a box `\yellowdyeII` that is prepared by using the `\newsavebox` command of LaTeX. Then, it is printed by means of the `\usebox` command. The output is shown in Table 17.3.

```
\begin{table}[h]
\begin{quotation}
The preparation of
Several dye-sulphonyl chlorides (\cref{cmj01}) were
prepared from the corresponding dye-sulfonic acid
reacted with POCl$_3$/DMA (Table \ref{tabj01}).
The use of POCl$_3$/DMF in place of POCl$_3$/DMA
did not give \cref{cmj01a}; but
resulted in the formation of a sulphonyl chloride (\cref{cmj01g}),
in which the original carbamoyl group is changed into
a cyano group.
\end{quotation}
\newsavebox{\yellowdyeII}
\savebox{\yellowdyeII}{%
\begin{picture}(2000,1500)(0,-400)
\put(0,0){\fiveheterov[d]{1==N;5==N}{1==;2D==O;3D==N--NH;4==R$^1$}}
\put(890,230){\bzdrh{1==;2==R$^2$;4==R$^3$;5==R$^4$}}
\put(0,-546){\bzdrv{}}
```

[1] S. Fujita, K. Koyama, S. Ono, *Nippon Kagaku Kai-Shi*, 1–12 (1991).

The preparation of Several dye-sulphonyl chlorides (**5**) were prepared from the corresponding dye-sulfonic acid reacted with $POCl_3$/DMA (Table 17.3). The use of $POCl_3$/DMF in place of $POCl_3$/DMA did not give **5a**; but resulted in the formation of a sulphonyl chloride (**5g**), in which the original carbamoyl group is changed into a cyano group.

Table 17.3: Dye-sulfonyl chloride prepared by $POCl_3$/DMA



5

Dye	R^1	R^2	R^3	R^4	Yield (%)
5a	$CONH_2$	OCH_3	H	SO_2Cl	93
5b	$CONHCH_3$	OCH_3	SO_2Cl	H	94
5c	$CONHCH_3$	OCH_3	H	SO_2Cl	90
5d	$CONHCH_3$	CH_3	SO_2Cl	H	70
5e	$CONHCH_3$	H	CH_3	SO_2Cl	95
5f	CN	H	$OCH_2CH_2OCH_3$	SO_2Cl	95
5g	CN	OCH_3	H	SO_2Cl	95

```
\put(1200,-200){\compd\label{cmj01}}
\end{picture}}
\begin{center}
\caption{Dye-sulfonyl chloride prepared by POCl$_3$/DMA}
\label{tabj01}
%\fbox{%
\usebox{\yellowdyeII}
%}
\begin{tabular}{cccccc}
\hline
Dye & R$^1$ & R$^2$ & R$^3$ & R$^4$ & Yield (\%) \\
\hline
\deriv\label{cmj01a} & CONH$_2$ & OCH$_3$ & H & SO$_2$Cl & 93 \\
\deriv\label{cmj01b} & CONHCH$_3$ & OCH$_3$ & SO$_2$Cl & H & 94 \\
\deriv\label{cmj01c} & CONHCH$_3$ & OCH$_3$ & H & SO$_2$Cl & 90 \\
\deriv\label{cmj01d} & CONHCH$_3$ &  CH$_3$ & SO$_2$Cl & H  & 70 \\
\deriv\label{cmj01e} & CONHCH$_3$ &  H & CH$_3$ & SO$_2$Cl & 95 \\
\deriv\label{cmj01f} & CN &  H & OCH$_2$CH$_2$OCH$_3$ & SO$_2$Cl & 95 \\
\deriv\label{cmj01g} & CN & OCH$_3$ & H & SO$_2$Cl & 95 \\
\hline
\end{tabular}
\end{center}
```

```
\end{table}
```

17.5 Verbatim Environment

The verbatim environment of LaTeX prints every control sequence with a top backslash symbol. The top symbol of every control sequence is changed into a yen symbol (¥), when you use the chemist package in a default setting. For example, the control sequence for typesetting the LaTeX logo is '`\LaTeX`' in the original LaTeX, while it is '`¥LaTeX`' in the chemist package.

If you want to obtain the backslash-type representation when you use the chemist package, you shoud declare `\verbswitchfalse` in the preamble of your tex file, as shown in the main file of this document (xymtex.tex).

In such a document as having the declaration of `\verbswitchfalse`, you can change the backslash-type representation into the yen-type one by declaring `\verbswichtrue`, as shown in the following example.

When `\verbswitchfalse` is effective, we have:

```
\TeX{} \LaTeX{} \XyMTeX
```

When `\verbswitchtrue` is effective, we have:

```
¥TeX{} ¥LaTeX{} ¥XyMTeX
```

Chapter 18

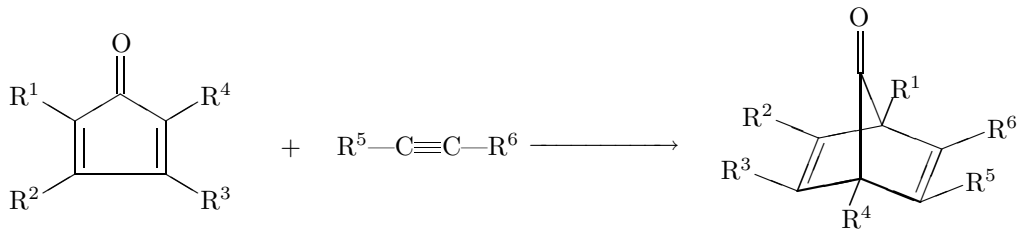
Reaction Schemes

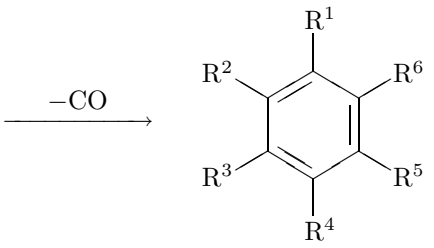
A reaction scheme contains several structural formulas and arrows that are aligned consequtively. This chapter deals with the drawing of such reaction schemes by using XƎMTEX commands combined with the tools of the chemist package.

18.1 XƎMTEX Formulas as TEX Boxes

Structural formulas typeset by XƎMTEX are regarded as boxes; hence, they obey the typesetting rule of TEX/LATEX in the same way as usual TEX boxes. In the following example, XƎMTEX boxes are aligned consequtively to construct a reaction scheme, where vertical adjustment is carried out by using the `\cdonecell` command of the chemist package.

```
\cdonecell{0pt}{4cm}{%
\cyclopentanevi[bd]{1D==O;2==R$^{4}$;3==R$^{3}$;%
4==R$^{2}$;5==R$^{1}$}} + \hskip10pt
\mbox{R$^{5}$---C\tbond C---R$^{6}$}
\reactrarrow{0pt}{2cm}{}{}
\cdonecell{10pt}{5cm}{%
\bornane[be]{7D==O;1==R$^{1}$;4==R$^{4}$;%
2==R$^{6}$;3==R$^{5}$;5==R$^{3}$;6==R$^{2}$}} \par
\hspace*{3cm}
\reactrarrow{0pt}{2cm}{$-$CO}{\strut}
\cdonecell{0pt}{4cm}{%
\bzdrv{1==R$^{1}$;4==R$^{4}$;%
2==R$^{6}$;3==R$^{5}$;5==R$^{3}$;6==R$^{2}$}}
```

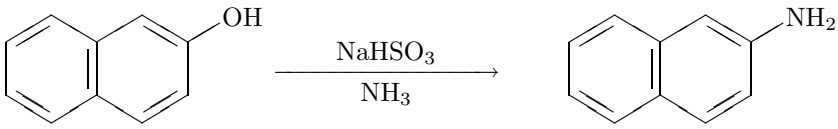




18.2 XyMTEX Commands in the Center Environment

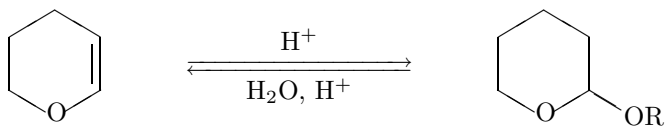
The Bucherer reaction (replacement of a hydroxy group by an amino group) is typset by using the `\naphdrv` command in the center environment. The arrow between the substrate and the product is drawn by means of the `\reactrarrow` command, where reagents used are designated on the upper side and lower side of the arrow.

```
\begin{center}
\naphdrv{2==OH}
\reactrarrow{40pt}{3cm}{NaHSO$_{3}$}{NH$_{3}$}
\naphdrv{2==NH$_{2}$}
\end{center}
```



If you want to use cross-reference, a compound number can be attached to each formula by means of a one-column tabular environment. Since the tabular environment causes the vertical centering of the formula included, no vertical adjustment is necessary to shift the arrow produced by the `\reacteqarrow` command; thus, its first argument is 0pt.

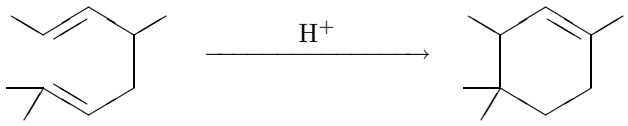
```
\begin{center}
\begin{tabular}{c}
\sixheterovi[b]{1==O}{} \\[-.5cm] \compd \label{reac:c1} \\
\end{tabular}
\reacteqarrow{0pt}{3cm}{H$^{+}$}{H$_{2}$O, H$^{+}$}
\begin{tabular}{c}
\sixheterovi[]{1==O}{2==OR} \\[-.5cm] \compd \label{reac:c2} \\
\end{tabular}
\end{center}
```



1 **2**

The \shortstack command can be used for the same purpose. This method requires the vertical ajustment of placing the arrow produced by \reactrarrow.

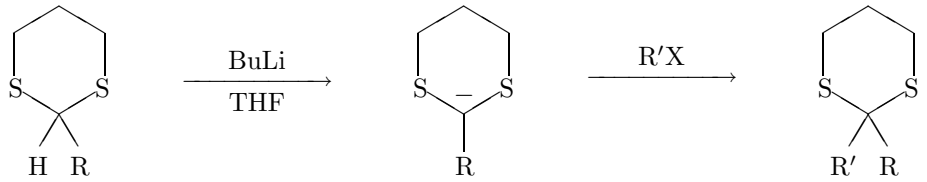
```
\begin{center}
\shortstack{%
\sixunitv[df]{}{2==;6==;5Sa==;5Sb==}{e}
\\[-.5cm] \compd \label{reac:c3}}
\reactrarrow{40pt}{3cm}{H$^{+}$}{\strut}
\shortstack{%
\cyclohexanev[a]{2==;6==;5Sa==;5Sb==}
\\[-.5cm] \compd \label{reac:c4}}
\end{center}
```



3 **4**

An alternative method of numbering compounds is to use the \cdtwocell command. It is applied to draw a scheme of alkylation of 1,3-dithianes. The minus charge on the 2-carbon atom of 1,3-dithiane is printed by using the BONDLIST argument of the \sixheterovi command, where a dummy argument 'H' is necessary to obtain a correct result. For such a dummy argument, see also the plus charge of an ammonium ion on page 104.

```
\begin{center}
\cdtwocell{0pt}{3cm}{%
 \sixheterovi[]{2==S;6==S}{1Sa==R;1Sb==H}}%
 {\compd \label{reac:c5}}
\reactrarrow{0pt}{2cm}{BuLi}{THF}
\cdtwocell{0pt}{3cm}{%
 \sixheterovi[H{1{{\lower1ex\hbox{$-$}}}}]{2==S;6==S}{1==R}}%
 {\compd \label{reac:c6}}
\reactrarrow{0pt}{2cm}{R$^{\prime}$X}{\strut}
\cdtwocell{0pt}{3cm}{%
 \sixheterovi[]{2==S;6==S}{1Sa==R;1Sb==R$^{\prime}$}}%
 {\compd \label{reac:c6a}}
\end{center}
```



5 **6** **7**

Another example of using \cdtwocell and \noderiv is the following table cited from *Yuki Gosei Kagaku Kyokai-Shi*.[1] Note that a set of \noderiv commands along with \label commands are declared

[1] S. Fujita, *Yuki Gosei Kagaku Kyokai-Shi*, **40**, 307–320 (1982).

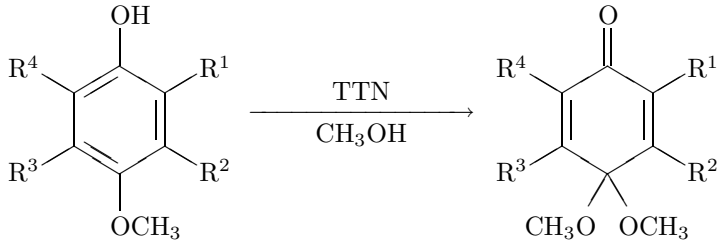
after the \compd command for each of the structural formulas; thereby, only compound numbers with no derivative numbers are typset there. On the other hand, a set of \cref commands are used in the tabular environment to print compond numbers. The output is shown in Table 18.1.

```
\begin{table}
\begin{quotation}
McKillop reported a method of preparing quinone monoacetals (\cref{cmb12})
by the oxidation of $p$-methoxyphenols (\cref{cmb11}) with
thallium(III) nitrate (TTN)/methanol.  The method can be
applied to the preparation of a wide vaiety of quinone monoacetals.
The reactions of
the quinone monoacetal with no subsituents (\cref{cmb12a})
with nucleophilic reagents have been examined in detail.
\end{quotation}

\caption{The preparation of quinone monoacetals}
\label{tt:chen01}
\begin{center}
\cdtwocell{0pt}{90pt}{\bzdrv{1==OH;4==OCH$_3$;%
 2==R$^1$;3==R$^2$;5==R$^3$;6==R$^4$}}{\compd\label{cmb11}%
 \noderiv\label{cmb11a}
 \noderiv\label{cmb11b}\noderiv\label{cmb11c}
 \noderiv\label{cmb11d}\noderiv\label{cmb11e}
 \noderiv\label{cmb11f}\noderiv\label{cmb11g}\noderiv\label{cmb11h}}
 \reactrarrow{10pt}{3cm}{TTN}{CH$_3$OH}
\cdtwocell{0pt}{90pt}{\bzdrv[pa]{1D==O;4Sa==OCH$_3$;4Sb==CH$_3$O;%
  2==R$^1$;3==R$^2$;5==R$^3$;6==R$^4$} }{\compd\label{cmb12}%
  \noderiv\label{cmb12a}\noderiv\label{cmb12b}\noderiv\label{cmb12c}
  \noderiv\label{cmb12d}\noderiv\label{cmb12e}
  \noderiv\label{cmb12f}\noderiv\label{cmb12g}\noderiv\label{cmb12h}}
 \\[10pt]
\begin{tabular}{ccccccccc}
\hline
derivatives & \cref{cmb12a} & \cref{cmb12b} &
\cref{cmb12c} & \cref{cmb12d} & \cref{cmb12e} &
\cref{cmb12f} & \cref{cmb12g} & \cref{cmb12h} \\
\hline
R$^1$ & H & CH$_3$ & CH$_3$ & C(CH$_3$)$_3$ & H       &H      &Cl&Br\\
R$^2$ & H &  H     &  H     &  H           & OCH$_3$ &OCH$_3$ &H &H\\
R$^3$ & H &  H     &  H     &  H           & OCH$_3$ &OCH$_3$ &H &H\\
R$^4$ & H &  H     &CH$_3$  & C(CH$_3$)$_3$ & H   &OCOCH$_3$  &H &H\\
\hline
yields (\%) & 97& 89 & 87     &    96         & 95      &92  & 97 & 91\\
\hline
\end{tabular}
\end{center}
\end{table}
```

McKillop reported a method of preparing quinone monoacetals (**9**) by the oxidation of *p*-methoxyphenols (**8**) with thallium(III) nitrate (TTN)/methanol. The method can be applied to the preparation of a wide vaiety of quinone monoacetals. The reactions of the quinone monoacetal with no subsituents (**9a**) with nucleophilic reagents have been examined in detail.

Table 18.1: The preparation of quinone monoacetals



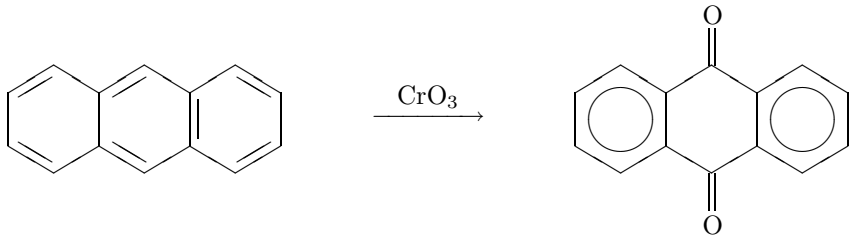
8 **9**

derivatives	**9a**	**9b**	**9c**	**9d**	**9e**	**9f**	**9g**	**9h**
R^1	H	CH_3	CH_3	$C(CH_3)_3$	H	H	Cl	Br
R^2	H	H	H	H	OCH_3	OCH_3	H	H
R^3	H	H	H	H	OCH_3	OCH_3	H	H
R^4	H	H	CH_3	$C(CH_3)_3$	H	$OCOCH_3$	H	H
yields (%)	97	89	87	96	95	92	97	91

18.3 XyMTEX in the Equation Environment

A reaction scheme can be written in the equation environment of LaTeX, where a vertical adjustment is necessary by means of the `\cdonecell` command of the chemist package.

```
\begin{equation}
\cdonecell{0pt}{5cm}{\anthracenev{}}\hskip1cm
\reactrarrow{0pt}{1.5cm}{CrO$_{3}$}{\strut}
\cdonecell{0pt}{5cm}{\anthracenev[pA]{9D==O;{{10}D}==O}}
\end{equation}
```



(18.1)

18.4 XyMTEX in the Picture Environment

A structural formula prepared by XyMTEX is a TEX box containing inner picture environments. It can be placed in an outer picture enviroment.

```
\begin{figure}
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
```

```
\def\bmC{\mbox{\boldmath $C$}}
\def\bmD{\mbox{\boldmath $D$}}
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
Example:
\begin{quotation}
The pair with [6,6,0,0] and $\bmD_{3d}$-symmetry
represents a r-1,t-2,c-3,t-4,c-5,t-6-hexasub\-stituted cyclohexane.
The pattern of substitution of the six Xs and the six hydrogens
in the pair strictly complies with the equation
which predicts the appearance of two six-membered orbits.
Since the pseudo-point group $\bmD_{3d}$ is anisoenergetic,
the starting molecule (\cref{cf:107a}) is
diastereomeric to the product molecule (\cref{cf:107b}).
The diastereomeric nature stems from the fact that
the six Xs are all equatorial in \cref{cf:107a} and
all axial in \cref{cf:107b}.
In the light of the present notation,
such an anisoenergetic pseudo-point group is
easily recognized, since it is represented by a symbol without a hat.
\end{quotation}

\begin{center}
%\fbox{%
%\begin{picture}(4100,3900)(0,0)
\begin{picture}(4100,3500)(0,0)
%
\put(0,2600){\chairi{1Se==X;2Se==X;3Se==X;4Se==X;5Se==X;6Se==X}}
\nocompd
\put(700,2600){\derivlabel{cf:107a}}
\put(1750,3000){\vector(1,0){500}}
\put(2250,2950){\vector(-1,0){500}}
\put(1750,2750){\hbox to50pt{%
 \hss$[6,6,6,0]$, $\bmD_{3d}$\hss}}
\put(2400,2600){\chair{1Sa==X;2Sa==X;3Sa==X;4Sa==X;5Sa==X;6Sa==X}}
\put(3200,2600){\derivlabel{cf:107b}}
%
\put(0,1300){\chairi{1Se==X;1Sa==X;3Se==X;3Sa==X;5Se==X;5Sa==X}}
\nocompd
\put(700,1300){\derivlabel{cf:108a}}
 \put(1750,1700){\vector(1,0){500}}
\put(2250,1650){\vector(-1,0){500}}
\put(1750,1450){\hbox to50pt{\hss$[6,6,6,0]$, $\widehat{\bmD}_{3h}$\hss}}
\put(2400,1300){\chair{1Se==X;1Sa==X;3Se==X;3Sa==X;5Se==X;5Sa==X}}
\put(3100,1300){\derivlabel{cf:108b}}
%
\put(0,0){\chairi{1Se==X;2Sa==X;3Se==X;4Sa==X;5Se==X;6Sa==X}}
\nocompd
\put(700,0){\derivlabel{cf:109a}}
\put(1750,400){\vector(1,0){500}}
\put(2250,350){\vector(-1,0){500}}
\put(1750,150){\hbox to50pt{%
 \hss$[6,6,6,0]$, $\widehat{\bmC}_{6v}^{\prime}$\hss}}
```

```
\put(2400,0){\chair{1Sa==X;2Se==X;3Sa==X;4Se==X;5Sa==X;6Se==X}}
\put(3200,0){\derivlabel{cf:109b}}
\end{picture}%
%}
\end{center}
\caption{[6,6,0,0]-Cyclohexane Derivatives of Higher Symmetries}
\label{ff:105}
\bigskip
\rightline{S. Fujita, {\it Bull. Chem. Soc. Jpn}, {\bf 67}, 2935 (1994)}
\end{figure}
```

Example:

The pair with [6,6,0,0] and $\boldsymbol{D}_{3d}$-symmetry represents a r-1,t-2,c-3,t-4,c-5,t-6-hexasubstituted cyclohexane. The pattern of substitution of the six Xs and the six hydrogens in the pair strictly complies with the equation which predicts the appearance of two six-membered orbits. Since the pseudo-point group $\boldsymbol{D}_{3d}$ is anisoenergetic, the starting molecule (**10a**) is diastereomeric to the product molecule (**10b**). The diastereomeric nature stems from the fact that the six Xs are all equatorial in **10a** and all axial in **10b**. In the light of the present notation, such an anisoenergetic pseudo-point group is easily recognized, since it is represented by a symbol without a hat.

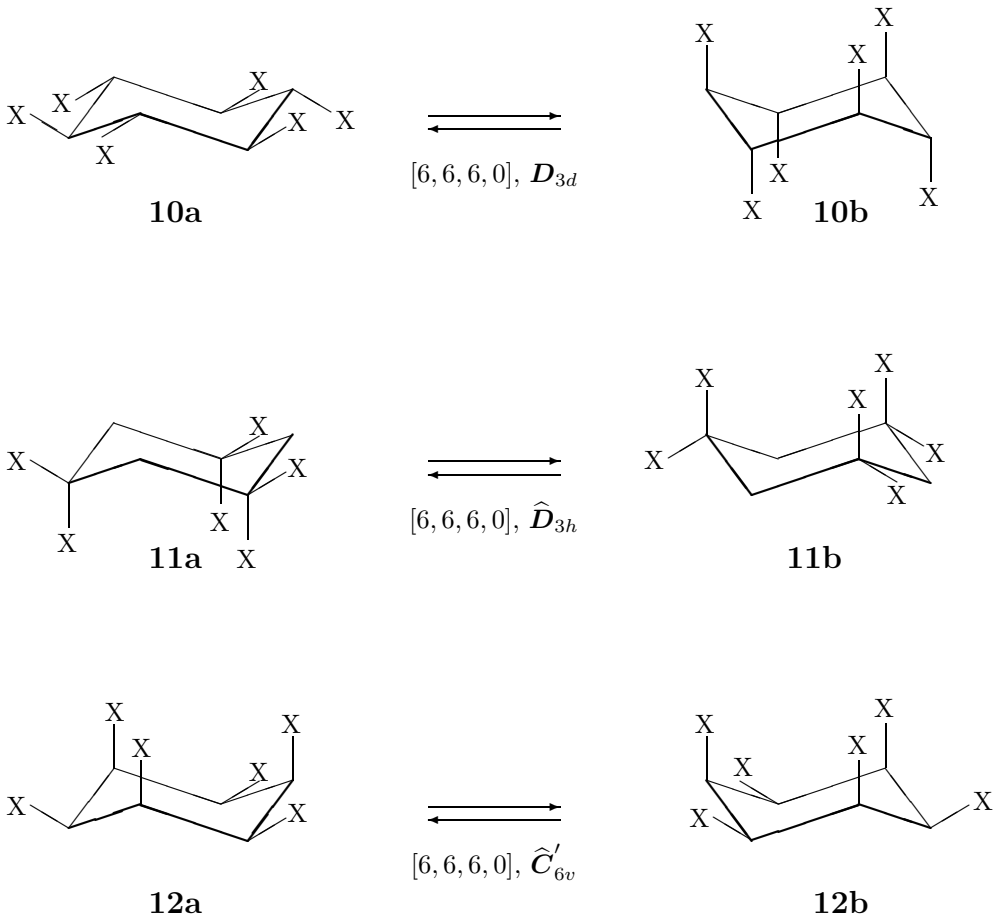


Figure 18.1: [6,6,0,0]-Cyclohexane Derivatives of Higher Symmetries

S. Fujita, *Bull. Chem. Soc. Jpn*, **67**, 2935 (1994)

Chapter 19

Boxes

This chapter deals with useful tools provided by chemist.sty for drawing boxes with a frame.

19.1 Environments for Drawing Framed Boxes

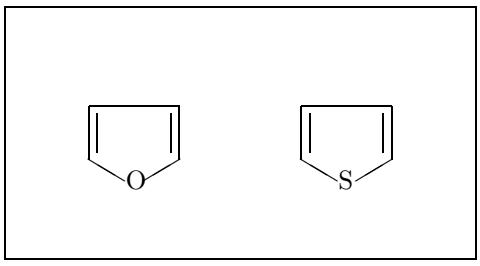
The frameboxit environment of the chemist package has one argument as follows;

```
\begin{frameboxit}{BWIDTH}
(text)
\end{frameboxit}
```

where the argument BWIDTH represents the width of the box produced by this environment. For example, you write a statement such as

```
\begin{frameboxit}{6cm}
\centering
\fiveheterov[bd]{1==O}{}
\fiveheterov[bd]{1==S}{}
\end{frameboxit}
```

Then, you obtain the following result.



Note that each structural formula drawn by XeLaTeX has a space around itself, which will be used for typesetting possible substituents.

Since the frameboxit environment is based on the fr@meboxit environment of the chemist package, the use of the latter inner environment enable us to change the line thickness of the frame (LWIDTH) and the margin (SPACE) around the text included:

```
\begin{fr@meboxit}{LWIDTH}{SPACE}{BWIDTH}
(text)
\end{fr@meboxit}
```

The default values of them are equal to those of the `\fbox` command of LaTeX.

The following example shows changes of such parameters. Note that the comands `\makeatletter` and `\makeatother` should be used for the special treatment of the `@` character.

```
\makeatletter
\begin{frameboxit}{5cm}
Default Parameters are selected to be 0.4pt for the line thickness and
3pt for the margin space.
\end{frameboxit}
\begin{fr@meboxit}{1pt}{10pt}{5cm}
Parameters are changed into 1pt for the line thickness and
10pt for the margin space.
\end{fr@meboxit}
\makeatother
```

This statement produces the following result.

Default Parameters are selected to be 0.4pt for the line thickness and 3pt for the margin space.

Parameters are changed into 1pt for the line thickness and 10pt for the margin space.

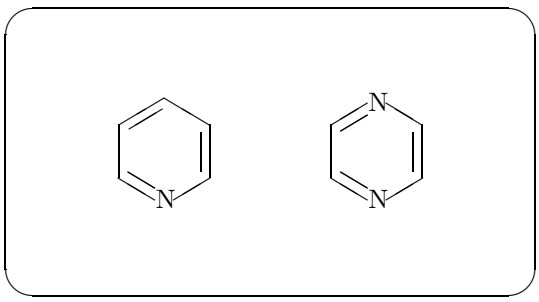
The miniscreen environment of the chemist package has one argument as follows;

```
\begin{miniscreen}{BWIDTH}
(text)
\end{miniscreen}
```

where the argument `BWIDTH` represents the width of the box produced by this environment. For example, by writing a statement such as

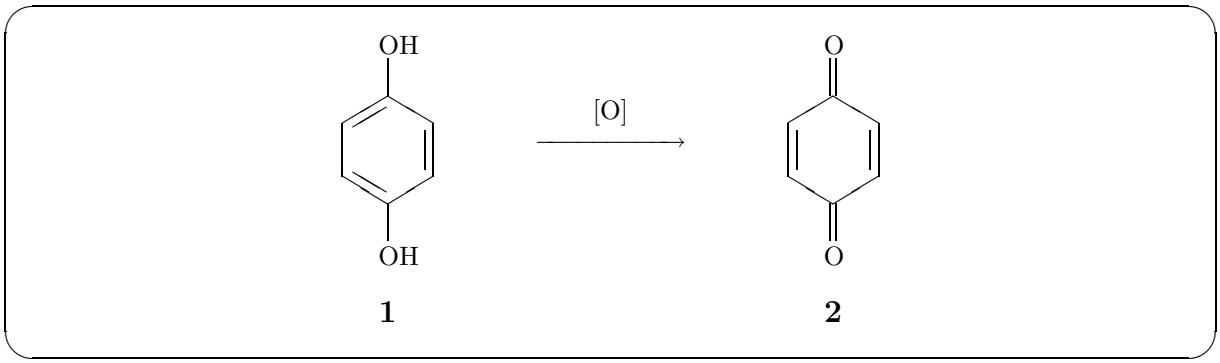
```
\begin{miniscreen}{7cm}
\centering
\pyridinevi{}
\pyrazinev{}
\end{miniscreen}
```

you obtain the following result.



The screen environment, which has originally defined in the package ascmac.sty, is used to generate a framed text of width `\textwidth`. It can be redefined on the basis of the definition of the miniscreen enviroment described above. The redfined screen environment can be used as follows.

```
\begin{screen}
\begin{center}
\begin{tabular}{c}
\bzdrv{1==OH;4==OH} \\[.3cm] \compd \label{box:a2} \\
\end{tabular}
\begin{tabular}{c}
[O] \\ \parbox{2cm}{\rightarrowfill} \\[1cm] \mathstrut \\
\end{tabular}
\begin{tabular}{c}
\bzdrv[p]{1D==O;4D==O} \\[.3cm] \compd \label{box:a3} \\
\end{tabular}
\end{center}
\end{screen}
```



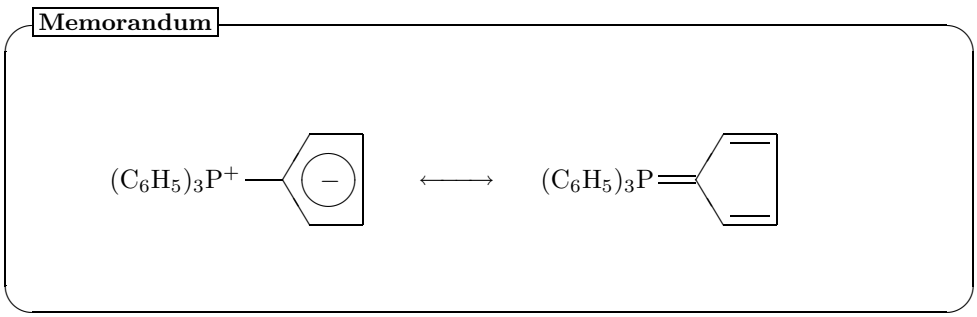
The tboxminiscreen environment of the chemist package is used to generate a miniscreen box with a heading title (the default title is "Memorandum"). It has one argument as follows;

```
\begin{tboxminiscreen}{BWIDTH}
(text)
\end{tboxminiscreen}
```

where the argument BWIDTH represents the width of the box produced by this environment. For example, by writing such a statement as

```
\begin{tboxminiscreen}{0.8\textwidth}
\setbox0=\hbox{$\mathrm{(C_{6}H_{5})_{3}P}$}
\hspace*{50pt}
\cdonecell{0pt}{4cm}{\cyclopentanehi[A{0{$-$}}]{1=={\copy0$^{+}$}}}
\hskip-25pt\reactlrarrow{-5pt}{1cm}{}{}\hskip35pt
\cdonecell{0pt}{4cm}{\cyclopentanehi[bd]{1D==\box0}}
\end{tboxminiscreen}
```

you obtain the following result.



For changing the heading title, you redefine the control sequence `\tboxtitle` by means of the comand `\def` or `\renewcommand`. For example, the statament

```
\def\tboxtitle{\bf Summary Notes}
\begin{tboxminiscreen}{0.8\textwidth}
The Beckmann rearrangement is a transformation of
an oxime into an amide under an acidic condition.
Since a substrate oxime can be easily obtained from
a ketone (or aldehyde) and hydroxylamine,
the Beckmann rearrangement is important as one of
valuable industrial processes.
\end{tboxminiscreen}
```

typesets the following miniscreen box with a changed title.

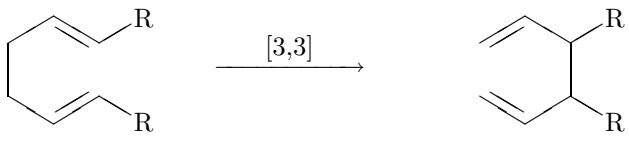
Summary Notes

The Beckmann rearrangement is a transformation of an oxime into an amide under an acidic condition. Since a substrate oxime can be easily obtained from a ketone (or aldehyde) and hydroxylamine, the Beckmann rearrangement is important as one of valuable industrial processes.

The tboxscreen environment provides a frame spreading for `\textwidth`.

```
\def\tboxtitle{\bf [3,3]Sigmatropic Rearrangement}
\begin{tboxscreen}
\centering
\cdonecell{0pt}{4cm}{\sixunitv[ac]{}{2==R;3==R}{b}}
\reactrarrow{0pt}{2cm}{[3,3]}{\strut}
\cdonecell{0pt}{4cm}{\sixunitv[df]{}{2==R;3==R}{e}}
\end{tboxscreen}
```

[3,3]Sigmatropic Rearrangement



19.2 Envitonment for Drawing Shadow Boxes

19.2.1 With a Left-Hand Shadow

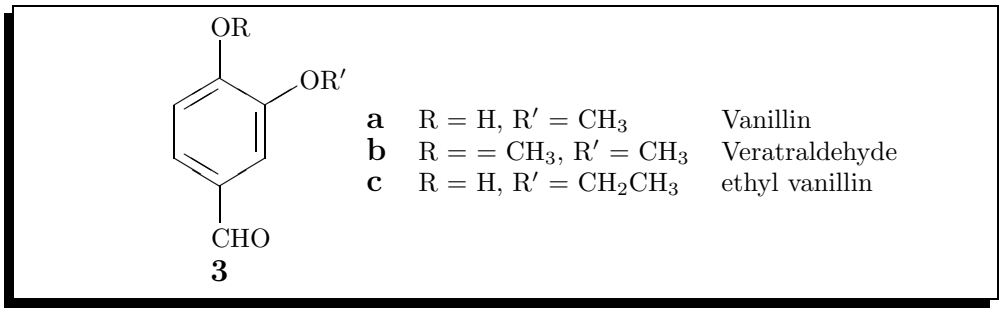
The lshfboxit environment of the chemist package is used to generated a frameed box with a left-hand shadow. It has one argument as follows;

```
\begin{lshfboxit}{BWIDTH}
(text)
\end{lshfboxit}
```

where the argument BWIDTH represents the width of the box produced by this environment. For example, you write such a statement such as

```
\begin{lshfboxit}{0.8\textwidth}
\begin{center}
\begin{tabular}{c}
\bzdrv{1==OR;4==CHO;2==OR$^{\prime}$} \\ \compd \label{va:a3} \\
\end{tabular}
\begin{tabular}{lll}
\deriva \label{va:a3a} &
 R = H, R$^{\prime}$ = CH$_{3}$ & Vanillin \\
\deriva \label{va:a3b} &
 R = = CH$_{3}$, R$^{\prime}$ = CH$_{3}$ & Veratraldehyde \\
\deriva \label{va:a3c} &
 R = H, R$^{\prime}$ = CH$_{2}$CH$_{3}$ & ethyl vanillin \\
\end{tabular}
\end{center}
\end{lshfboxit}
```

Then, you obtain the following result.



The lshfboxit environment is based on the lshfr@meboxit environment of the chemist package. Hence, we can use the latter inner environment to change the thickness of the horizontal shadow (HSWIDTH), the thickness of the vertical shadow (VSWIDTH), and the margin spacing (SPACE) around the text included.

```
\begin{lshfr@meboxit}{HSWIDTH}{VSWIDTH}{SPACE}{BWIDTH}
(text)
\end{lshfr@meboxit}
```

Note that the line thickness of the frame (LWIDTH) is fixed to be 0.4pt.

The following example shows changes of such parameters.

```
\makeatletter
\begin{lshfboxit}{5cm}
Default Parameters are selected
3pt for the thickness of the horizontal shadow,
3pt for the thickness of the vertical shadow, and
3pt for the margin space.
\end{lshfboxit}
\qquad
\begin{lshfr@meboxit}{5pt}{10pt}{10pt}{5cm}
Parameters are changed into
5pt for the thickness of the horizontal shadow,
10pt for the thickness of the vertical shadow, and
```

```
10pt for the margin space.
\end{lshfr@meboxit}
\makeatother
```

This statement produces the following result.

Default Parameters are selected 3pt for the thickness of the horizontal shadow, 3pt for the thickness of the vertical shadow, and 3pt for the margin space.

Parameters are changed into 5pt for the thickness of the horizontal shadow, 10pt for the thickness of the vertical shadow, and 10pt for the margin space.

19.2.2 With a Right-Hand Shadow

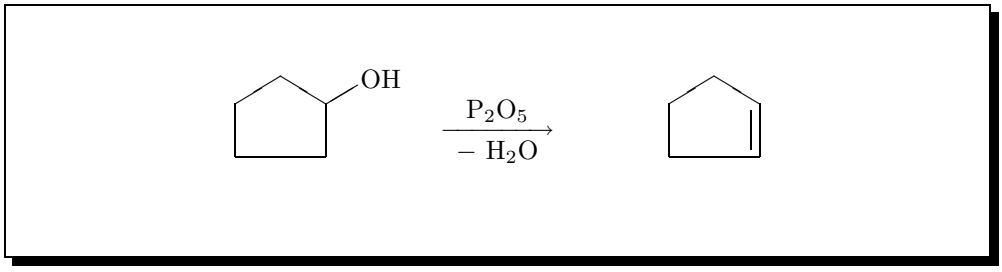
On the other hand, the rshfboxit environment of the chemist package is used to generated a frameed box with a right-hand shadow. It has one argument as follows;

```
\begin{rshfboxit}{BWIDTH}
(text)
\end{rshfboxit}
```

where the argument BWIDTH represents the width of the box produced by this environment. For example, you write such a statement such as

```
\begin{rshfboxit}{0.8\textwidth}
\centering
\cdonecell{0pt}{4cm}{\cyclopentanevi[]{2==OH}}
\reactrarrow{0pt}{1.5cm}{P$_{2}$O$_{5}$}{$-$ H$_{2}$O}
\cdonecell{0pt}{4cm}{\cyclopentanevi[b]{}}
\end{rshfboxit}
```

Then, you have



The rshfboxit environment is based on the rshfr@meboxit environment of the chemist package. Hence, we can use the latter inner environment to change parameters in a similar way to the lshfr@meboxit environment.

```
\begin{rshfr@meboxit}{HSWIDTH}{VSWIDTH}{SPACE}{BWIDTH}
(text)
\end{rshfr@meboxit}
```

Note that the line thickness of the frame (LWIDTH) is fixed to be 0.4pt, while changeable parameters are the thickness of the horizontal shadow (HSWIDTH), the thickness of the vertical shadow (VSWIDTH), and the margin spacing (SPACE) around the text included.

The following example shows changes of such parameters.

```
\makeatletter
\begin{rshfboxit}{5cm}
Default Parameters are selected
3pt for the thickness of the horizontal shadow,
3pt for the thickness of the vertical shadow, and
3pt for the margin space.
\end{rshfboxit}
\qquad
\begin{rshfr@meboxit}{10pt}{5pt}{10pt}{5cm}
Parameters are changed into
10pt for the thickness of the horizontal shadow,
5pt for the thickness of the vertical shadow, and
10pt for the margin space.
\end{rshfr@meboxit}
\makeatother
```

This statement produces the following result.

Default Parameters are selected 3pt for the thickness of the horizontal shadow, 3pt for the thickness of the vertical shadow, and 3pt for the margin space.

Parameters are changed into 10pt for the thickness of the horizontal shadow, 5pt for the thickness of the vertical shadow, and 10pt for the margin space.

19.2.3 With a Left-Hand Gradient Shadow

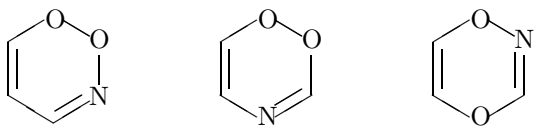
The glshfboxit environment of the chemist package is used to generated a frameed box with a left-hand gradient shadow. It has one argument as follows;

```
\begin{glshfboxit}{BWIDTH}
(text)
\end{glshfboxit}
```

where the argument BWIDTH represents the width of the box produced by this environment. For example, you write such a statement such as

```
\begin{glshfboxit}{0.7\textwidth}
\centering
1,2,3-, 1,2,4- and 1,4,2-dioxazine \\
\sixheterov[ce]{1==O;2==O;3==N}{}
\sixheterov[ce]{1==O;2==O;4==N}{}
\sixheterov[be]{1==O;2==N;4==O}{}
\end{glshfboxit}
```

1,2,3-, 1,2,4- and 1,4,2-dioxazine



The glshfboxit environment is based on the glshfr@meboxit environment of the chemist package. Hence, we can use the latter inner environment to change the thickness of the horizontal shadow (HSWIDTH), the thickness of the vertical shadow (VSWIDTH), and the margin spacing (SPACE) around the text included.

```
\begin{glshfr@meboxit}{HSWIDTH}{VSWIDTH}{SPACE}{BWIDTH}
(text)
\end{glshfr@meboxit}
```

Note that the line thickness of the frame (LWIDTH) is fixed to be 0.4pt.

The following example shows the way of changing such parameters.

```
\makeatletter
\begin{glshfboxit}{5cm}
Default Parameters are selected
3pt for the thickness of the horizontal shadow,
3pt for the thickness of the vertical shadow, and
3pt for the margin space.
\end{lshfboxit}
\qquad
\begin{glshfr@meboxit}{5pt}{10pt}{10pt}{5cm}
Parameters are changed into
5pt for the thickness of the horizontal shadow,
10pt for the thickness of the vertical shadow, and
10pt for the margin space.
\end{lshfr@meboxit}
\makeatother
```

This statement produces the following result.

Default Parameters are selected 3pt for the thickness of the horizontal shadow, 3pt for the thickness of the vertical shadow, and 3pt for the margin space.

Parameters are changed into 5pt for the thickness of the horizontal shadow, 10pt for the thickness of the vertical shadow, and 10pt for the margin space.

It should be noted that two large valules assigned to HSWIDTH/ and VSWIDTH may result in printing jagged edges, since the sloped edges consist of a fixed number of lines drawn repeatedly.

19.2.4 With a Right-Hand Gradient Shadow

On the other hand, the grshfboxit environment of the chemist package is used to generated a frameed box with a right-hand gradient shadow. It has one argument as follows;

```
\begin{grshfboxit}{BWIDTH}
(text)
\end{grshfboxit}
```

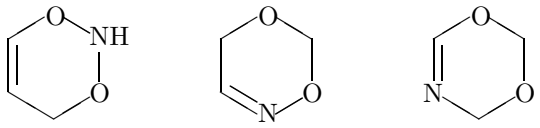
where the argument BWIDTH represents the width of the box produced by this environment. For example, you write such a statement such as

```
\begin{grshfboxit}{0.7\textwidth}
\centering
```

```
2H,4H-1,3,2-, 2H,6H-1,3,4- and 2H,4H-1,3,5-dioxazine \\
\sixheterov[e]{1==O;2==NH;3==O}{}
\sixheterov[d]{1==O;3==O;4==N}{}
\sixheterov[e]{1==O;3==O;5==N}{}
\end{grshfboxit}
```

Then, you have

2H,4H-1,3,2-, 2H,6H-1,3,4- and 2H,4H-1,3,5-dioxazine



The grshfboxit environment is based on the grshfr@meboxit environment of the chemist package. Hence, we can use the latter inner environment to change parameters. See also the glshfr@meboxit environment descrived above.

```
\begin{grshfr@meboxit}{HSWIDTH}{VSWIDTH}{SPACE}{BWIDTH}
(text)
\end{grshfr@meboxit}
```

Note that the line thickness of the frame (LWIDTH) is fixed to be 0.4pt, while changeable parameters are the thickness of the horizontal shadow (HSWIDTH), the thickness of the vertical shadow (VSWIDTH), and the margin spacing (SPACE) around the text included.

The following example shows changes of such parameters.

```
\makeatletter
\begin{grshfboxit}{5cm}
Default Parameters are selected
3pt for the thickness of the horizontal shadow,
3pt for the thickness of the vertical shadow, and
3pt for the margin space.
\end{grshfboxit}
\qquad
\begin{grshfr@meboxit}{10pt}{5pt}{10pt}{5cm}
Parameters are changed into
10pt for the thickness of the horizontal shadow,
5pt for the thickness of the vertical shadow, and
10pt for the margin space.
\end{grshfr@meboxit}
\makeatother
```

This statement produces the following result.

Default Parameters are selected 3pt for the thickness of the horizontal shadow, 3pt for the thickness of the vertical shadow, and 3pt for the margin space.

Parameters are changed into 10pt for the thickness of the horizontal shadow, 5pt for the thickness of the vertical shadow, and 10pt for the margin space.

19.3 Commands for Framed Boxes

The `\fboxit` command is used for surrounding a text of one line with a frame,

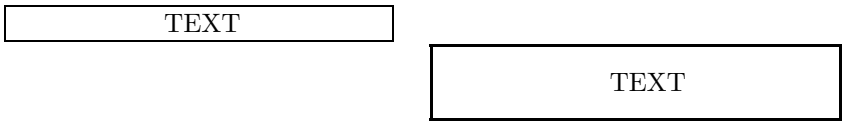
```
\fboxit{BWIDTH}{TEXT}
```

where `BWIDTH` represent the width of the frame box. This command is based on the inner command `\fb@xit` having changeable parameters.

```
\fb@xit{LWIDTH}{SPACE}{BWIDTH}{TEXT}
```

where `LWIDTH` represents line thickness, `SPACE` represents spacing around the text, and `BWIDTH` represents the width of the box.

```
\makeatletter
\fboxit{5cm}{\centering TEXT} \qquad
\fb@xit{1pt}{10pt}{5cm}{\centering TEXT}
\makeatother
```



The `\leftshframe` command is used for surrounding a text of one line with a frame having a left-hand shadow,

```
\leftshframe{BOXWIDTH}{TEXT}
```

where `BWIDTH` represent the width of the frame box. On the other hand, the `\leftshfbox` command generates a frame having a left-hand shadow in accordance with the length of the text included.

```
\leftshfbox{TEXT}
```

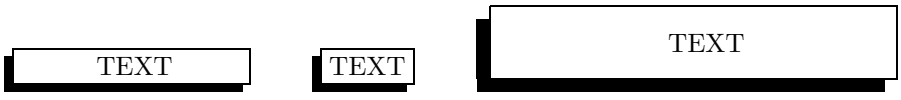
These commands are based on the inner command `\leftshfr@me` having changeable parameters.

```
\leftshfr@me{SWIDTH}{SPACE}{BWIDTH}{TEXT}
```

where `SWIDTH` represents the thickness of a shadow, `SPACE` represents spacing around the text, and `BWIDTH` represents the width of the box.

The following examples show the difference between these box-generating commands.

```
\makeatletter
\leftshframe{3cm}{\centering TEXT} \qquad
\leftshfbox{\centering TEXT} \qquad
\leftshfr@me{5pt}{10pt}{5cm}{\centering TEXT}
\makeatother
```



The `\rightshframe` command is used for surrounding a text of one line with a frame having a right-hand shadow,

```
\rightshframe{BOXWIDTH}{TEXT}
```

where `BWIDTH` represent the width of the frame box. On the other hand, the `\rightshfbox` command generates a frame having a right-hand shadow in accordance with the length of the text included.

```
\rightshfbox{TEXT}
```

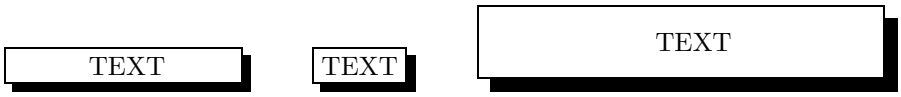
These commands are based on the inner command `\rightshfr@me` having changeable parameters.

```
\rightshfr@me{SWIDTH}{SPACE}{BWIDTH}{TEXT}
```

where `SWIDTH` represents the thickness of a shadow, `SPACE` represents spacing around the text, and `BWIDTH` represents the width of the box.

The following examples show the difference between these box-generating commands.

```
\makeatletter
\rightshframe{3cm}{\centering TEXT} \qquad
\rightshfbox{\centering TEXT} \qquad
\rightshfr@me{5pt}{10pt}{5cm}{\centering TEXT}
\makeatother
```



19.4 Several Symbols

The commands `\Yubin` and `\yubin` are used to print zip codes of Japanese style. For example, the statement

```
\Yubin{250-01} Minami-Ashigara-Shi, Nakanuma, 210 \\
\yubin{250-01} Minami-Ashigara-Shi, Nakanuma, 210
```

prints

〒250-01 Minami-Ashigara-Shi, Nakanuma, 210
〒250-01 Minami-Ashigara-Shi, Nakanuma, 210

Temperatures are designated by `\degC`, `\Cent` or `\degF`, *e.g.*,

```
5\Cent, $-5\degC$, 40\degF
```

5℃, −5℃, 40℉

Several logos can be typeset as follows.

```
\jLaTeX, \pTeX, \jTeX, \jBibTeX,
\tpic, \PiCTeX, \PostScript,
```

JLaTeX, pTeX, JTeX, JBibTeX, Tpic, PiCTeX, PostScript,

Bibliography

[1] Knuth D. E., *The TEXbook*, Addison-Wesley, Reading (1984).

[2] Ito T., *Kagaku to Kogyo*, **46**, 66 (1993). This article described that the Chemical Society of Japan changed the publishing process of its journal (*Bull. Chem. Soc. Jpn.*) on January 1993 into the one that is based on the combination of TEX and SGML (Standard Generalized Mark-up Language).

[3] Lamport L., *LATEX. A document Preparation System*, Addison-Wesley, Reading (1986).

[4] Haas R. T. & O'Kane K. C., *Comput. Chem.*, **11**, 251 (1987).

[5] Ramek, M., in Clark, M. (ed), TEX: Applications, Uses, Methods, Ellis Horwood, London (1990), p. 277.

[6] Fujita S., "Typesetting structural formulas with the text formatter TEX/LATEX", *Comput. Chem.*, **18**, 109 (1994).

[7] Fujita S., "XƎMTEX for Drawing Chemical Structural Formulas", *TUGboat*, **16** (1), 80 (1995).

[8] Fujita S., *Kagakusha-Seikagakusha no tame no LATEX (LATEX for Chemists and Biochemists)*, Tokyo Kagaku Dozin, Tokyo (1993).

[9] NIFTY-Serve achives, FPRINT libray No. 7, Item Nos. 201, 202, 204.

[10] CTAN, tex-archive/macros/latex209/contrib/xymtex/.

[11] Lamport L., *LATEX. A document Preparation System*, 2nd ed. for LATEX 2_ε, Addison-Wesley, Reading (1994).

[12] Goossens, M., Mittelbach, F., & Samarin, A., *The LATEX Companion*, Addison-Wesley, Reading (1994).

[13] Podar S., "Enhancements to the picture environment of LATEX", Manual for Version 1.2 dated July 14, 1986.

[14] Fujita S., "Promolecules with subsymmetry of $\mathbf{D}_{\infty h}$. Combinatorial enumeration and stereochemical properties", *J. Chem. Inf. Comput. Sci.*, **32**, 354–363 (1992).

[15] Fujita S., Koyama K., & Ono S., "Dye releasers for instant color photography", *Rev. Heteroatom Chem.*, **7**, 229–267 (1992).

Appendix A

List of Commands

A.1 aliphat.sty

The style file 'aliphat.sty' contains the following commands

```
%
%  <Macros for tetravalent atoms>
%
%     \tetrahedral                        \@tetrahedral
%     \square                             \@square
%
%  <Macros for trivalent atoms>
%
%     \rtrigonal                          \@rtrigonal
%     \ltrigonal                          \@ltrigonal
%     \utrigonal                          \@utrigonal
%     \Utrigonal                          \@Utrigonal
%     \dtrigonal                          \@dtrigonal
%     \Dtrigonal                          \@Dtrigonal
%
%  <Macros for two-carbon compounds>
%
%     \ethylene                           \@ethylene
%     \ethylenev                          \@ethylenev
%     \Ethylenev                          \@Ethylenev
%
%  <Macros for stereo-projection>
%
%     \tetrastereo                        \@tetrastereo
%     \dtetrastereo                       \@dtetrastereo
%     \ethanestereo                       \@ethanestereo
%
```

A.2 carom.sty

The style file 'carom.sty' contains the following commands.

```
%
```

```
%   <VERTICAL TYPE>
%
%     \cyclohexanev               \@cyclohexanev
%     \bzdrv                      \@bzdrv
%     \decalinev                  \@decalinev
%     \naphdrv                    \@naphdrv
%     \tetralinev                 \@tetralinev
%     \hanthracenev               \@hanthracenev
%     \anthracenev                \@anthracenev
%     \hphenanthrenev             \@hphenanthrenev
%     \phenanthrenev              \@phenanthrenev
%     \steroid                    \@steroid
%     \steroidchain               \@steroidchain
%
%   <HORIZONTAL TYPE>
%
%     \cyclohexaneh               \@cyclohexaneh
%     \bzdrh                      \@bzdrh
%     \decalineh                  \@decalineh
%     \naphdrh                    \@naphdrh
%     \tetralineh                 \@tetralineh
%
```

A.3 lowcycle.sty

The style file 'lowcycle.sty' contains the following commands.

```
%
%     \cyclopentanev                    \@cyclopentanev
%     \cyclopentanevi                   \@cyclopentanevi
%     \cyclopentaneh                    \@cyclopentaneh
%     \cyclopentanehi                   \@cyclopentanehi
%     \indanev                          \@indanev
%     \indanevi                         \@indanevi
%     \indaneh                          \@indaneh
%     \indanehi                         \@indanehi
%     \cyclobutane                      \@cyclobutane
%     \cyclopropane                     \@cyclopropane
%
%
```

A.4 ccycle.sty

The style file 'ccycle.sty' contains the following commands.

```
%
%  <Basic Macros>
%
%     \chair                    \@chair
%     \chairi                   \@chairi
%     \bicychepv                \@bicychepv
```

```
%     \bicycheph                \@bicycheph
%     \bornane                  \@bornane
%     \adamantane               \@damantane
%     \hadamantane              \h@damantane
%
```

A.5 hetarom.sty

The style file 'hetarom.sty' contains the following commands.

```
%
%  <Six-Membered Heterocycles>

%    Basic Macros
%
%     \sixheterov              \@sixheterov
%     \sixheterovi             \@sixheterovi
%
%    Application Macros
%
%     \pyridinev               \@pyridinev
%     \pyridinevi              \@pyridinevi
%     \pyrazinev               \@pyrazinev
%     \pyrimidinev             \@pyrimidinev
%     \pyrimidinevi            \@pyrimidinevi
%     \pyridazinev             \@pyridazinev
%     \pyridazinevi            \@pyridazinevi
%     \triazinev               \@triazinev
%     \triazinevi              \@triazinevi
%
%  <Five-Membered Heterocycles>
%    Basic Macros
%
%     \fiveheterov             \@fiveheterov
%     \fiveheterovi            \@fiveheterovi
%
%    Application Macros
%
%     \pyrrolev                \@pyrrolev
%     \pyrazolev               \@pyrazolev
%     \imidazolev              \@imidazolev
%     \isoxazolev              \@isoxazolev
%     \oxazolev                \@oxazolev
%     \pyrrolevi               \@pyrrolevi
%     \pyrazolevi              \@pyrazolevi
%     \imidazolevi             \@imidazolevi
%     \isoxazolevi             \@isoxazolevi
%     \oxazolevi               \@oxazolevi
%
%  <Six-Six-Fused Heterocycles>
%    Basic Macros
%
```

```
%     \decaheterov              \@decaheterov
%     \decaheterovi             \@decaheterovi
%
%    Application Macros
%
%     \quinolinev               \@quinolinev
%     \quinolinevi              \@quinolinevi
%     \isoquinolinev            \@isoquinolinev
%     \isoquinolinevi           \@isoquinolinevi
%     \quinoxalinev             \@quinoxalinev
%     \quinazolinev             \@quinazolinev
%     \quinazolinevi            \@quinazolinevi
%     \cinnolinev               \@cinnolinev
%     \cinnolinevi              \@cinnolinevi
%     \pteridinev               \@pteridinev
%     \pteridinevi              \@pteridinevi
%
%  <Six-Five-Fused Heterocycles>
%    Basic Macros
%
%     \nonaheterov              \@nonaheterov
%     \nonaheterovi             \@nonaheterovi
%
%    Application Macros
%
%     \purinev                  \@purinev
%     \purinevi                 \@purinevi
%     \indolev                  \@indolev
%     \indolevi                 \@indolevi
%     \indolizinev              \@indolizinev
%     \indolizinevi             \@indolizinevi
%     \isoindolev               \@isoindolev
%     \isoindolevi              \@isoindolevi
%     \benzofuranev             \@benzofuranev
%     \benzofuranevi            \@benzofuranevi
%     \isobenzofuranev          \@isobenzofuranev
%     \isobenzofuranevi         \@isobenzofuranevi
%     \benzoxazolev             \@benzoxazolev
%     \benzoxazolevi            \@benzoxazolevi
%
%  <Four-Membered Heterocycles>
%
%     \fourhetero               \@fourhetero
%
%  <Four-Membered Heterocycles>
%
%     \threehetero              \@threehetero
%
%  <Building Units>
%
%     \sixunitv                 \@sixunitv
%     \fiveunitv                \@fiveunitv
```

```
%     \fiveunitvi               \@fiveunitvi
%
```

A.6 hetaromh.sty

The style file 'hetaromh.sty' contains the following commands.

```
%
%  <Six-membered heterocycles (horizontal)>
%
%    Basic Macros
%
%     \sixheteroh               \@sixheteroh
%     \sixheterohi              \@sixheterohi
%
%    Application Macros
%
%     \pyridineh                \@pyridineh
%     \pyridinehi               \@pyridinehi
%     \pyrazineh                \@pyrazineh
%     \pyrimidineh              \@pyrimidineh
%     \pyrimidinehi             \@pyrimidinehi
%     \pyridazineh              \@pyridazineh
%     \pyridazinehi             \@pyridazinehi
%     \triazineh                \@triazineh
%     \triazinehi               \@triazinehi
%
%  <Five-membered heterocycles (horizontal)>
%
%    Basic Macros
%
%     \fiveheteroh              \@fiveheteroh
%     \fiveheterohi             \@fiveheterohi
%
%    Application Macros
%
%     \pyrroleh                 \@pyrroleh
%     \pyrazoleh                \@pyrazoleh
%     \imidazoleh               \@imidazoleh
%     \isoxazoleh               \@isoxazoleh
%     \oxazoleh                 \@oxazoleh
%     \pyrrolehi                \@pyrrolehi
%     \pyrazolehi               \@pyrazolehi
%     \imidazolehi              \@imidazolehi
%     \isoxazolehi              \@isoxazolehi
%     \oxazolehi                \@oxazolehi
%
%  <Six-Six fused heterocycles (horizontal)>
%
%    Basic Macros
%
%     \decaheteroh              \@decaheteroh
```

```
%     \decaheterohi              \@decaheterohi
%
%    Application Macros
%
%     \quinolineh                \@quinolineh
%     \quinolinehi               \@quinolinehi
%     \isoquinolineh             \@isoquinolineh
%     \isoquinolinehi            \@isoquinolinehi
%     \quinoxalineh              \@quinoxalineh
%     \quinazolineh              \@quinazolineh
%     \quinazolinehi             \@quinazolinehi
%     \cinnolineh                \@cinnolineh
%     \cinnolinehi               \@cinnolinehi
%     \pteridineh                \@pteridineh
%     \pteridinehi               \@pteridinehi
%
%  <Six-Five fused heterocycles (horizontal)>
%
%    Basic Macros
%
%     \nonaheteroh               \@nonaheteroh
%     \nonaheterohi              \@nonaheterohi
%
%    Application Macros
%
%     \purineh                   \@purineh
%     \purinehi                  \@purinehi
%     \indoleh                   \@indoleh
%     \indolehi                  \@indolehi
%     \indolizineh               \@indolizineh
%     \indolizinehi              \@indolizinehi
%     \isoindoleh                \@isoindoleh
%     \isoindolehi               \@isoindolehi
%     \benzofuraneh              \@benzofuraneh
%     \benzofuranehi             \@benzofuranehi
%     \isobenzofuraneh           \@isobenzofuraneh
%     \isobenzofuranehi          \@isobenzofuranehi
%     \benzoxazoleh              \@benzoxazoleh
%     \benzoxazolehi             \@benzoxazolehi
%
%   <Building units>
%
%     \sixunith                  \@sixunith
%     \fiveunith                 \@fiveunith
%     \fiveunithi                \@fiveunithi
%
```

A.7 hcycle.sty

The style file 'hcycle.sty' contains the following commands.

```
%
```

```
% <Pyranose and Furanose>
%
%     \pyranose                  \@pyranose
%     \furanose                  \@furanose
%
```

A.8 chemstr.sty

The style file 'chemstr.sty' contains the following commands.

```
%
%  <Convention>
%
%     \rmoiety
%     \lmoiety
%     \putlatom
%     \putratom
%     \putlratom
%     \Putlratom

%   <Setting of atoms and bonds>
%
%     \setsixringv      (on a vertical six-membered ring)
%     \setdecaringv     (on a decaline (six-six) ring)
%     \setfusedbond     (at fused bond positions)
%     \setatombond      (at an appropiate position)
%     \setsixringh       (on a horizontal six-membered ring)
%
```

A.9 locant.sty

The style file 'locant.sty' contains the following commands.

```
%
%     \sxlocant      \sxloocant
%     \bdlocant      \bdloocant
%     \sxlocnth      \sxloocnth
%     \bdlocnth      \bdloocnth
%
```

A.10 polymers.sty

The style file 'polymers.sty' contains the following commands.

```
%
%  <Delimeters for polymer units>
%
%    \leftPMdelim      \rightPMdelim
%    \leftpmdelim      \rightpmdelim
%
%    \leftPolymer       \leftpolymer
```

```
%    \rightPolymer      \rightpolymer
%
%    \leftsqrPolymer    \leftSqrpolymer
%    \lsqrdelimiter     \leftsqrpolymer
%
%    \rightsqrPolymer   \rightSqrpolymer
%    \rsqrdelimiter     \rightsqrpolymer
%
%    \@sqrpolymer
%    \sqrpolymer        \Sqrpolymer
%    \mpolymer
%
%  <Bonds for polymers>
%
%    \WestPbond         \EastPbond
%    \sbond
%
%  <Polymers>
%
%    \polyethylene      \@polyethylene
%    \polystyrene       \@polystyrene
%
```

A.11 chemist.sty

The style file 'chemist.sty' contains the following commands.

```
%
%  <counters for compounds>
%
%     \thecompd           \compd
%     \nocompd
%     \compdlabel         \nocompdlabel
%     \cref
%
%     \compdfbox          \compdmbox
%     \lbcompdpbox
%
%  <counters for derivative>
%
%     \thderiv
%     \deriv              \noderiv
%     \derivlabel         \noderivlabel
%     \derivfbox          \derivmbox
%     \lbderivpbox
%     \deriva             \noderiva
%     \derivalabel        \noderivalabel
%
%  <Boxes for compounds>
%
%     \cdonecell
%     \cdtwocell
```

```
%
%  <Arrows with variable length>
%
%     \leftrightarrowfill     \Leftarrowfill
%     \Rightarrowfill         \Leftrightarrowfill
%
%  <Reaction arrows with variable length>
%
%     \reactrarrow            \reactlarrow
%     \reactlrarrow           \reacteqarrow
%     \schemelarrow           \schemerarrow
%     \schemelrarrow
%
%  <changin line skip>
%
%     \changespace
%
%  <Caption of floats>
%
%     \tbcaption              \fgcaption
%     \smcaption
%
%  <Redefinition of verbatim>
%
%     \yen
%     \ifverbswitch
%     \verb
%     verbatim          @verbatim
%
%  <Shadowed Boxes>
%
%     \fboxit           \f@boxit
%     \leftshframe      \leftshfr@me
%     \leftshfbox
%     \rightshframe     \rightshfr@me
%     \rightshfbox
%     ffboxit           \ffparbox
%
%  <Framed box environment without shadows>
%
%     frameboxit        fr@meboxit
%
% <Framed box environment with shadows>
%
%     lshfboxit         lshfr@meboxit
%     rshfboxit         rshfr@meboxit
%
%     \hsgr@d           \vsgr@d
%     \vwgr@d           \hwgr@d
%     \vsepgr@d         \hsepgr@d
%     \nrep             \nrepmax
%
```

```
%     grshfboxit        grshfr@meboxit
%     glshfboxit        glshfr@meboxit
%
%  <(Mini)screen environment>
%
%     \@oval
%     \h@h@x            \@vrf@
%     miniscreen        screen
%
%  <(Mini)screen environment with a title>
%
%     \tboxscreentoprule
%     \tboxtitle
%     tboxminiscreen    tboxscreen
%     \sboxit
%
%  <Boxes with no height or no width>
%
%     \bury             \pushtowall
%
%  <Symbols for instant photography>
%
%     \cyan       \magenta     \yellow
%     \cyandv     \magentadv   \yellowdv
%     \agxlatent  \agxdv       \agx
%
%  <Various arrows with fixed length>
%
%     \llongrightarrow        \llongleftarrow
%     \llongleftrightarrow    \Equilibarrow
%     \Llongrightarrow        \Llongleftarrow
%     \Llongleftrightarrow
%     \llongleftharpoondown   \llongrightharpoondown
%     \equilibarrow           \eqproton
%
%     \lllongrightarrow       \lllongleftarrow
%     \lllongleftrightarrow   \Equiliblongarrow
%     \Lllongrightarrow       \Lllongleftarrow
%     \Lllongleftrightarrow
%
%     \lllongleftharpoondown  \lllongrightharpoondown
%     \equiliblongarrow       \deHBr
%
%  <Correction commands>
%
%     \chemcorr               \leavechemcorr
%
%
%  <Chem Version>
%
%     \ifnewl@tex
%     \chemform
```

```
%     chemmath       chemeqn
%
%     chemeqnarray   chemeqnarray*
%     chemeqnarraya
%
%  <Useful commands>
%
%     \endash
%     \Cent          \degC          \degF
%     \yubin         \Yubin
%     \xlethead      \horizon
%
%  <Logos>
%
%     \resetfontsize
%     \Bib           \BibTeX         \jBibTeX
%     \jTeX          \pTeX
%     \Sub           \Subbib
%     \La            \LaTeX          \jLaTeX
%     \PiC           \PiCTeX
%     \tpic
%     \Post          \Script         \PostScript
%
%  <Double and triple bonds>
%
%     \triplebond   \doublebond
%     \tbond        \dbond
```