

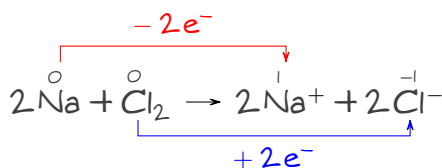
the chemmacros bundle

v3.0c 2012/02/05

packages **chemmacros**, **chemformula** and **ghsystem**

Clemens NIEDERBERGER

<https://bitbucket.org/cgnieder/chemmacros/>
contact@mychemistry.eu



Contents		
	7.3. Latin Phrases	10
	8. Units for the Usage With siunitx	11
I. Preliminaries	3	
1. Licence, Requirements and README	3	
2. Motivation	3	
3. Installation, Loading the Bundle	4	
4. Package Options	4	
5. Setup	6	
II. chemmacros	7	
6. Particles, Ions and Symbols	7	
6.1. Predefined	7	
6.2. Own Particles	8	
7. Stereo Descriptors, Nomenclature, Latin Phrases	8	
7.1. IUPAC Names	8	
7.2. Stereo Descriptors and Nomenclature	9	
	10. Oxidation Numbers, Real and Formal Charges	12
	10.1. Ion Charges	12
	10.2. Oxidation Numbers	13
	10.3. Partial Charges and Similar Stuff	14
	11. Reaction Mechanisms	15
	12. Redox Reactions	16
	13. (Standard) State, Thermodynamics	18
	13.1. Thermodynamic Variables . . .	18
	13.1.1. Create New Variables . .	19
	13.1.2. Redefine Variables . . .	20
	13.2. State	20
	14. Spectroscopy	21
	15. Commands for mhchem	22
	16. Reaction Environments	23
	16.1. Defined by CHEMMACROS	23

16.2. Own Reactions	24	28.3. Customization	43
16.3. List of Reactions	25		
17. Phases	26	29. Names	44
		29.1. Syntax	44
18. Newman Projections	27	29.2. Customization	45
19. s, p, and Hybrid Orbitals	28	30. Format and Font	45
20. CHEMFORMULA support	30	31. Usage In Math Equations	48
21. Key Overview	31	32. Further Examples	48
III. chemformula	33	IV. ghsystem	50
22. Setup	33	33. Setup	50
23. The Basic Principle	33	34. Get Hazard and Precautionary State- ments	50
24. Stoichiometric Factors	34	34.1. Simple Statements	50
25. Compounds	36	34.2. Statements with Placeholders	51
25.1. Adducts	36	34.3. Statements with Gaps	52
25.2. Subscripts	36	34.4. Combined Statements	52
25.3. Commands	37	35. Pictograms	53
25.4. Charges and Other Superscripts	37	35.1. The Pictures	53
25.5. Bonds	38	35.2. Picture Type Depending on Engine	55
25.6. Customization	38		
26. Special Input Types	40	36. Available Languages	55
27. Escaped Input	40	37. List of All Statements	56
27.1. Text	40		
27.2. Math	41	V. Appendix	65
28. Arrows	41	Suggestions and Bug Reports	65
28.1. Arrow types	41	Index	66
28.2. Labels	42		

Part I.

Preliminaries

1. Licence, Requirements and README

The CHEMMACROS bundle underlies the L^AT_EX project public license (lppl) version 1.3 or later. (<http://www.latex-project.org/lppl.txt>)

CHEMMACROS uses the packages expl3, xparse¹, l3keys2e² and xfrac³, which are part of the bundles l3kernel⁴ and l3packages⁵. expl3 is part of l3kernel, and xparse, l3keys2e and xfrac are part of l3packages.

Obviously, the packages CHEMFORMULA and GHSYSTEM are loaded as they are part of this bundle.

CHEMMACROS also uses the packages siunitx⁶, mathtools⁷, bm⁸ and environ⁹ as well as tikz¹⁰ and the TikZ libraries calc and arrows.

Package option bpchem (section 4) needs the package bpchem¹¹, package option xspace needs the package xspace¹² and package option method = mhchem needs the package mhchem¹³.

CHEMMACROS has been bundled with the new CHEMFORMULA and GHSYSTEM packages. CHEMFORMULA provides an alternative to mhchem. This leads to some internal changes in CHEMMACROS. On the same time the documentation has been redesigned.

You might remember that CHEMMACROS' options all belong to different modules, see section 5 for further information. These are typeset in the left margin when the option is first mentioned. In section 21 all options and the module they belong to are listed. Throughout the document options are typeset green and modules red.

CHEMFORMULA uses the packages CHEMMACROS and nicefrac¹⁴. CHEMFORMULA has no own package options but passes all options on to CHEMMACROS.

GHSYSTEM uses the packages CHEMMACROS, tabu¹⁵, longtable¹⁶, ifpdf¹⁷ and graphicx¹⁸. GHSYSTEM has no own package options but passes all options on to CHEMMACROS.

2. Motivation

CHEMMACROS started some years ago as a growing list of custom macros that I frequently used. I cannot completely recall when and why I decided to release them as a package. Well – here we go and you might find it useful, too, I hope.

I would guess that nearly every chemist using L^AT_EX 2_ε is aware of the great mhchem package. CHEMFORMULA is *not* a replacement for mhchem but an alternative. I thought about writing it for a while. Some minor points in mhchem always bothered me, but they hardly seemed enough for a new package. They weren't even enough for a feature request to the mhchem author. What convinced me at last was the fun and the challenge.

CHEMFORMULA works very similar to mhchem but is more strict as to how compounds, stoichiometric factors and arrows are input. In the same time CHEMFORMULA offers some possibilities to customize the output that mhchem does not. If you are content with the way mhchem works there is probably no need to use CHEMFORMULA. Maybe you are curious anyway.

¹ CTAN: xparse ² CTAN: l3keys2e ³ CTAN: xfrac ⁴ CTAN: l3kernel ⁵ CTAN: l3packages ⁶ CTAN: siunitx
⁷ CTAN: mathtools ⁸ CTAN: bm ⁹ CTAN: environ ¹⁰ CTAN: pgf ¹¹ CTAN: bpchem ¹² CTAN: xspace ¹³ CTAN: mhchem
¹⁴ CTAN: nicefrac ¹⁵ CTAN: tabu ¹⁶ CTAN: longtable ¹⁷ CTAN: ifpdf ¹⁸ CTAN: graphicx

As a chemist you are probably aware of the fact that the UNITED NATIONS have developed the “Globally Harmonized System of Classification and Labelling of Chemicals” (GHS) as a global replacement for the various different systems in different countries. While it has not been implemented by all countries yet¹⁹, it is only a matter of time. The package **GHSYSTEM** now enables you to typeset all the hazard and precautionary statements and pictograms in a very easy way.

3. Installation, Loading the Bundle

The bundle comes with three style files²⁰, a directory called `language/` containing the language-definition files for GHS (ending `def`), and a directory `pictures/` containing `eps`, `jpg` and `png` files (the GHS pictograms). If you install the bundle manually *please make sure to place the directories `language/` and `pictures/` in the same directory as the style files.*

Loading **CHEMMACROS** with

```
1 \usepackage{chemmacros} % 'chemmacros', 'formula' and 'ghs' are loaded
```

will also load **CHEMFORMULA** and **GHSYSTEM**. However, you can prevent **CHEMMACROS** from loading **GHSYSTEM**:

```
1 \usepackage[ghs=false]{chemmacros} % 'chemmacros' and 'formula' are loaded
```

Loading **CHEMFORMULA** or **GHSYSTEM** explicitly is possible and will also load **CHEMMACROS** if it hasn't been loaded yet, and will therefore implicitly load the other package, too.

```
1 \usepackage{chemformula} % 'chemmacros', 'formula' and 'ghs' are loaded
2 or
3 \usepackage[ghs=false]{chemformula} % 'chemmacros' and 'formula' are loaded
```

However, it is recommended to simply use `\usepackage{chemmacros}` and setup the required options with `\chemsetup` (also see section 5).

4. Package Options

CHEMMACROS has several package options. They all are used as key/value pairs like

```
1 \usepackage[option1 = <value1>, option2 = <value2>]{chemmacros}
```

¹⁹ http://www.unec.org/trans/danger/publi/ghs/implementation_e.html ²⁰ Those ending `sty`.

Some also can be used without value (`\usepackage[option1]{chemmacros}`), which means that the underlined value is used.

Both `CHEMFORMULA` and `GHSYSTEM` don't have package options of their own. If you load them explicitly you can give them `CHEMMACROS`' options instead and they will pass them on to `CHEMMACROS`.

- option** `bpchem` = `true/false` This option loads the package `bpchem` and adjusts the layout of the `\NMR` command to the `bpchem` commands `\HNMR` and `\CNMR`. (default = `false`)
- option** `circled` = `formal/all/none` `CHEMMACROS` uses two different kinds of charges²¹, which indicate the usage of real (+/−) and formal (\oplus/\ominus) charges. The option `formal` distinguishes between them, option `none` displays them all without circle, option `all` circles all (default = `formal`)
- option** `circletype` = `chem/math` This option switches between two kinds of circled charge symbols: `\fplus` \oplus and `\oplus` \oplus . (default = `chem`)
- option** `detect-bold` = `true/false` This option determines whether or not macros like `\pKa` recognize a bold font series. **`pKA` text** or **`pKA` text**. (default = `false`)
- option** `EZ` = `chemmacros/cool` The command `\E` is defined by the package `cool`²² as well as by `CHEMMACROS`. With this option you can choose, which definition is used, see page 9. (default = `chemmacros`). This option can only be chosen in the preamble.
- option** `german` = `true/false` This option changes the commands `\pKa`, `\sld` and `\lqd` (default = `false`).
option This option has the alias `ngerman`.
- option** `ghs` = `true/false` Disable the use of the `GHSYSTEM` package. Setting `ghs` = `false` will prevent `CHEMMACROS` from loading `GHSYSTEM`. (default = `true`)
- option** `method` = `chemformula/mhchem` You can choose the method which `CHEMMACROS` will use for the reaction environments (see section 16) and the typesetting of the particles (see section 6). (default = `chemformula`). This option can only be chosen in the preamble.
- option** `Nu` = `chemmacros/mathspec` The package `mathspec`²³ also defines a macro `\Nu`. This option chooses which definition holds, see page 7. (default = `chemmacros`). This option can only be chosen in the preamble.
- option** `strict` = `true/false` Setting `strict` = `true` will turn all warning messages into errors messages. (default = `false`)
- option** `synchronize` = `true/false` The setting `true` will tell `CHEMMACROS` to adapt the font settings of `CHEMFORMULA` if that method has been chosen (default = `false`). In order to demonstrate this feature this document is set with `synchronize` = `true` and the `CHEMFORMULA` setting `\chemsetup[chemformula]{font-spec={[Color=darkgray]Latin Modern Sans}}`.
- option** `version` = `1/2/bundle` This option restores the old definitions of some commands, so documents set with `v1.*` will still compile correctly. (default = `bundle`). Actually `2` and `bundle` are only aliases. This option can only be chosen in the preamble.

option `xspace` = true/false With this option most commands are defined with a `\xspace`. (default = true)

5. Setup

Various of `CHEMMACROS`’, `CHEMFORMULA`’s and `GHSYSTEM`’s commands have key/value pairs with which they can be customized. Most times they can be used as (optional) argument of the commands themselves. They also can most times be used with the `\chemsetup` command.

`\chemsetup[<module>]{<key> = <value>}` or

`\chemsetup{<module>/<key> = <value>}`

The keys each belong to a module, which defines for which commands they are intended for. If a key is presented, you’ll see the module to which it belongs in the left margin. You have two ways to use keys with the `\chemsetup`, as you can see above.

The package options can also be seen as keys belonging to the module **option**. This means they can also be used with the `\chemsetup` command (except for the option `version` = 1/2/3).

```
1 \chemsetup[option]{circled=none}\mch\ \pch\ \fmch\ \fpch\ \el\ \prt \\  
2 \chemsetup[option]{circled=formal}\mch\ \pch\ \fmch\ \fpch\ \el\ \prt \\  
3 \chemsetup[option]{circletype=math}\mch\ \pch\ \fmch\ \fpch\ \el\ \prt \\  
4 \chemsetup{option/circletype=chem,option/circled=all}\mch\ \pch\ \fmch\ \fpch\ \el\  
   \prt \\  
5 \chemsetup{option/circletype=math}\mch\ \pch\ \fmch\ \fpch\ \el\ \prt  
  
- + - + e- p+  
- + ⊖ ⊕ e- p+  
- + ⊖ ⊕ e- p+  
⊖ ⊕ ⊖ ⊕ e⊖ p⊕  
⊖ ⊕ ⊖ ⊕ e⊖ p⊕
```

Keys *not* belonging to a module *cannot* be used with `\chemsetup`!

All options of `CHEMFORMULA` belong to the module **chemformula** and all of `GHSYSTEM`’s options belong to the module **ghs**.

²¹ Thanks to Christoph Schäfer, who pointed out to me, that v1.1 handled the charges too sloppy! ²² CTAN: cool

²³ CTAN: mathspec

Part II.

chemmacros

6. Particles, Ions and Symbols

6.1. Predefined

Some simple macros for displaying often needed particles and a symbol. Please note, that they're displayed differently depending on the package options used, see section 4. These commands can be used in text as well as in math mode.

`\Hp1` H^+ (proton)

`\Hyd` OH^- (hydroxide)

`\HtO` H_3O^+ (oxonium ion) (H three O)

`\water` H_2O

`\el` e^- (electron)

`\prt` p^+ (proton)

`\ntr` n^0 (neutron)

`\Nu` Nu^- (nucleophile). The package `mathspec` also defines a macro `\Nu`. If you chose package option `Nu = mathspec` `CHEMMACROS` defines `\Nuc` instead.

`\El` E^+ (electrophile)

`\ba` ba^- (base)

`\fplus` \oplus

`\fminus` \ominus

`\transitionstatesymbol` \neq (uses `TikZ`)

`\standardstate` \ominus . This symbol is only provided by `CHEMMACROS`, if the package `chemstyle`²⁴ is not loaded; the idea is borrowed from there²⁵.

There is another command which allows to typeset radicals with charges and subscripts.

`\R[<sign>]{<subscript>}`

```
1 \R[+]{tert} \R[-]{sek} \R{prim}  $\text{R}_{\text{tert}}^+ \text{R}_{\text{sek}}^- \text{R}_{\text{prim}}$ 
```

The two particles `\Nu` and `\ba` can be modified. To do that you use the option

`particle` `elpair = false/dots/dash.`

It only has any effect, if the package `chemfig`²⁶ is loaded, since it uses its command `\Lewis`.

```
1 % needs package 'chemfig'
2 \ba[elpair] \Nu[elpair=dash]          ba:~ Nu|~
3
4 \chemsetup{particle}{elpair}          ba:~ Nu:~
5 \ba \Nu
```

6.2. Own Particles

If you like you can define your own particles using

```
\DeclareParticle{<cmd>}{<definition>}
```

Depending on the `method` you chose as option the `<definition>` will either be formula defined with `mhchem` or with `CHEMFORMULA`. The particle defined this way behaves like the predefined ones with one exception: if you chose `method = mhchem` the particle *will not* obey the option `circled`. If you want formal charges with this method you need to use `CHEMMACROS`' commands (see section 10) explicitly. If you chose `method = chemformula` the particle *will* obey the `circled` option.

```
1 \DeclareParticle{\Sod}{Na+}
2 \DeclareParticle{\Pot}{K+}
3 \Sod \Pot \chemsetup{option}{circled=all} \Sod \Pot

Na+ K+ Na⊕ K⊕
```

The particles are only defined if the `<cmd>` you chose is not already used by any other command. If it is already used `CHEMMACROS` will either give a warning or an error, depending on the option `strict`.

7. Stereo Descriptors, Nomenclature, Latin Phrases

7.1. IUPAC Names

Similar to the `bpchem` package `CHEMMACROS` provides a command²⁷ to typeset IUPAC names. Why is that useful? IUPAC names can get very long. So long indeed that they span over more than two lines, especially in two-column documents. This means they must be allowed to be broken more than one time. This is what the following command does.

```
\iupac{<IUPAC name>} Inside this command use \| and \- to indicate a breaking point or a
breaking dash. Use \^ as a shortcut for \textsuperscript28.
```

²⁴ CTAN: `chemstyle` ²⁵ many thanks to the package author [Joseph Wright](#).
²⁶ CTAN: `chemfig` ²⁷ The idea and the implementation is shamelessly borrowed from `bpchem` by Bjørn Pedersen.
²⁸ Actually `\^` uses a `CHEMFORMULA` command instead.


```

1 \begin{minipage}{.4\linewidth}
2 \iupac{Tetra\|cyclo[2.2.2.1^{1,4}]\|-un\|decane-2\|-dodecyl\|-5\|-(hepta\|decyl\|iso\|
   dodecyl\|thio\|ester)}
3 \end{minipage}

Tetracyclo[2.2.2.11,4]-undecane-2-do-
decyl-5-(heptadecylisododecylthioes-
ter)

```

The `\iupac` command is more of a semantic command. Most times you can achieve (nearly) the same thing by using `\-` instead of `\|`, `-` instead of `\-` and `\textsuperscript` instead of `\^`.

7.2. Stereo Descriptors and Nomenclature

The macros in this section are intended to make the writing of IUPAC names more convenient.

Cahn-Ingold-Prelog

`\Rcip` (*R*)

`\Scip` (*S*)

`\cip{<conf>}` e. g.: `\cip{R,S}` (*R,S*)

Fischer

`\Dfi` *D*

`\Lfi` *L*

cis/trans, zusammen/entgegen, syn/anti & tert Please notice, that the commands `\cis` and `\trans` are defined by the `bpchem` package as well. If you load that package, they are redefined by `CHEMMACROS`. With `bpchem` they *always* get a `\xspace`, with `CHEMMACROS` *never*.

`\cis` *cis*

`\trans` *trans*

`\Z` (*Z*)

`\E` (*E*) (`\E` is also defined by the package `cool`. By using the package option `EZ = cool` instead of `\E` and `\Z` `CHEMMACROS` defines `\Ent` and `\Zus`.)

`\syn` *syn*

`\anti` *anti*

`\tert` *tert*

ortho/meta/para

`\ortho` *o*

`\meta` *m*

`\para` *p*

absolute configuration (uses *TikZ*)

`\Rconf[<letter>]` `\Rconf:`  `\Rconf[]:` 

`\Sconf[<letter>]` `\Sconf:`  `\Sconf[]:` 

Examples:

```
1 \iupac{\Dfi\Wein\|s\"aure} = \\
2 \iupac{\cip{2S,3S}\Wein\|s\"aure} \\
3 \iupac{\Dfi\-(\$-\$)\-Threose} = \\
4 \iupac{\cip{2S,3R}\-(\$-\$)\-2,3,4-Tri\|hydroxy\|butanal} \\
5 \iupac{\cis\2\Butene} = \\
6 \iupac{\Z\2\Butene}, \\
7 \iupac{\cip{2E,4Z}\Hexa\|diene} \\
8 \iupac{\meta\Xylol} = \\
9 \iupac{1,3-Di\|methyl\|benzene}
```

D-Weinsäure =
(2*S*,3*S*)-Weinsäure
D-(–)-Threose =
(2*S*,3*R*)-(–)-2,3,4-Trihydroxybutanal
cis-2-Butene =
(*Z*)-2-Butene,
(2*E*,4*Z*)-Hexadiene
m-Xylol =
1,3-Dimethylbenzene

7.3. Latin Phrases

At last there are two commands for common latin phrases.

`\insitu` *in situ*

`\abinitio` *ab initio*

If the package `chemstyle` has been loaded, too²⁹, they are defined using `chemstyle`'s `\latin` command. This means that then the appearance depends on `chemstyle`'s option `abbremp`:

²⁹ `chemstyle` defines other similar commands like *et al.*, *in vacuo*.

1	<code>\insitu, \abinitio\</code>	<i>in situ, ab initio</i>
2	<code>\cstsetup{abbremph=false}</code>	in situ, ab initio
3	<code>\insitu, \abinitio</code>	

If chemstyle hasn't been loaded, they're always in *italics*.

8. Units for the Usage With siunitx

In chemistry some non-SI units are very common. siunitx provides the command `\DeclareSIUnit{<command>}{<unit>}` to add arbitrary units. CHEMMACROS uses that command to provide some units. Like all siunitx units they're only valid inside `\SI{<num>}{<unit>}` and `\si{<unit>}`.

`\atmosphere` atm

`\atm` atm

`\calory` cal

`\cal` cal

`\cmc` cm³ The units `\cmc`, `\molar`, and `\Molar` are defined by the package chemstyle as well. CHEMMACROS only defines them, if chemstyle is not loaded.

`\molar` mol dm⁻³

`\moLar` mol L⁻¹

`\Molar` M

`\MolMass` g mol⁻¹

`\normal` N

`\torr` torr

By the way: `\mmHg` mmHg already is defined by siunitx and chemstyle

9. Acid/Base

Easy representation of pH , pK_A ... (the command `\pKa` depends on the package option `german`, see section 4.)

`\pH` pH

`\pOH` pOH

`\Ka` K_A

`\Kb` K_B

`\Kw` K_W

`\pKa[<num>]` `\pKa:` pK_A , `\pKa[1]:` pK_{A1}

`\pKb[<num>]` `\pKb:` pK_B , `\pKb[1]:` pK_{B1}

`\p{<anything>}` e. g. `\p{\Kw}` pK_W

```
1 \Ka \Kb \pKa \pKa[1] \pKb \pKb[1]\  K_A K_B pK_A pK_{A1} pK_B pK_{B1}
2 \chemsetup[option]{german=true}      K_S K_B pK_S pK_{S1} pK_B pK_{B1}
3 \Ka \Kb \pKa \pKa[1] \pKb \pKb[1]
```

10. Oxidation Numbers, Real and Formal Charges

CHEMMACROS distinguishes between real (+/−) and formal (\oplus/\ominus) charge symbols, also see section 4. All commands using formal charge symbols start with a f.

10.1. Ion Charges

Simple displaying of (real) charges:

`\pch[<number>]` positive charge (plus + charge)

`\mch[<number>]` negative charge (minus + charge)

```
1 \pch, Na\pch, Ca\pch[2]\  +, Na+, Ca2+
2 \mch, F\mch, S\mch[2]    −, F−, S2−
```

The same for formal charges:

`\fpch[<number>]` positive charge

`\fmch[<number>]` negative charge

```
1 \fpch\ \fmch\ \fpch[3] \fmch[3]   $\oplus \ominus 3\oplus 3\ominus$ 
```

There is a key which influences the behaviour of the charges.

charges `append` = true/false if set true, the charge is appended together with an empty group.

This is how the key influences the behaviour:

```
1 \chemsetup[charges]{append=false}
2 \ce{H\pch\aq} \ce{H\aq\pch}      H(aq)+ H(aq)+
3
4 \chemsetup[charges]{append=true}
5 \ce{H\pch\aq} \ce{H\aq\pch}      H(aq)+ H(aq)+
```

In most cases this behaviour will be unwanted. However, in some cases it might be useful, for example together with the `\ox` command (see next section):

```
1 \chemsetup[charges]{append=false}
2 \ce{\ox{1,H}\pch\aq}
3
4 \chemsetup[charges]{append=true}
5 \ce{\ox{1,H}\pch\aq}
```

$$\begin{array}{c} \text{I}^+ \\ \text{H}_{(\text{aq})} \\ \text{I}^+ \\ \text{H}_{(\text{aq})}^+ \end{array}$$

10.2. Oxidation Numbers

Typesetting oxidation numbers:

`\ox[<keyval>]{<number>,<atom>}` places <number> above <atom>; <number> has to be a (rational) number!

```
1 \ox{+1,Na}, \ox{2,Ca}, \ox{-2,S}, \ox{-1,F}
```

$$\begin{array}{c} \text{I} \quad \text{II} \quad -\text{II} \quad -\text{I} \\ \text{Na, Ca, S, F} \end{array}$$

There are a number of keys, that can be used to modify the `\ox` command.

- ox** `parse` = `true`/false when false an arbitrary entry can be used for <number>.
- ox** `roman` = `true`/false switches from roman to arabic numbers.
- ox** `pos` = top/super/side; top places <number> above <atom>, super to the upper right as superscript and side to the right and inside brackets.
- ox** `explicit-sign` = `true`/false shows the + for positiv numbers and the ± for 0.
- ox** `decimal-marker` = comma/point choice for the decimal marker for formal oxidation numbers like $\overset{1.2}{\text{X}}$.

```
1 \ox[roman=false]{2,Ca} \ox{2,Ca} \\\
2 \ox[pos=super]{3,Fe}-Oxide \\\
3 \ox[pos=side]{3,Fe}-Oxide \\\
4 \ox[parse=false]{?,Mn}
```

$$\begin{array}{c} \text{2} \quad \text{II} \\ \text{Ca Ca} \\ \text{Fe}^{\text{III}}\text{-Oxide} \\ \text{Fe(III)-Oxide} \\ \text{?} \\ \text{Mn} \end{array}$$

The `pos` = super variant also can be set with the shortcut `\ox*`:

```
1 \ox{3,Fe} \ox*{3,Fe}
```

$$\overset{\text{III}}{\text{Fe}} \text{ Fe}^{\text{III}}$$

Using the `explicit-sign` key will always show the sign of the oxidation number:

```

1 \chemsetup[ox]{explicit-sign = true}
2 \ox{+1,Na}, \ox{2,Ca}, \ox{-2,S}, \ch{"\ox{0,F}" {}2}

+INa, +IICa, -II±0S, F2

```

```

1 Compare \ox{-1,\ch{O2^2-}} to \ch{"\ox{-1,O}" ^2^2-}

Compare O2-I to O2-I

```

Sometimes one might want to use formal oxidation numbers like 0.5 or $\frac{1}{3}$:

```

1 \ox{.5,\ch{Br2}} \ch{"\ox{1/3,I}" ^3+}

Br20.5 I31/3+

```

The fraction uses the `\sfrac` command of the `xfrac` package. For this purpose the instance `chemmacros-ox-fraction` is defined.

```

1 \DeclareInstance{xfrac}{chemmacros-ox-fraction}{text}
2 {
3   scale-factor          = 1.2 ,
4   denominator-bot-sep  = -.5ex ,
5   numerator-top-sep    = -.3ex ,
6   slash-left-kern      = -.2em ,
7   slash-right-kern     = -.2em ,
8   slash-symbol-font    = lmr
9 }

```

Of course you can redefine it so that it suits your needs as the output often strongly depends on the used font.

10.3. Partial Charges and Similar Stuff

The next ones probably are seldomly needed but nevertheless useful:

`\delp` δ^+ (delta + plus)

`\delm` δ^- (delta + minus)

`\fdelp` δ^\oplus

`\fdelm` δ^\ominus

These macros for example can be used with the `\ox` command or with the `chemfig` package:

```

1 \chemsetup{
2   option/circled = all,
3   ox/parse      = false
4 }
5 \ce{\ox{\del p,H}-\ox{\del m,Cl}} \hspace*{1cm}
6 \chemfig{\chemabove[3pt]{\lewis{246,Br}}{\del m}-\chemabove[3pt]{H}{\del p}}

```



The following macros are useful together with chemfig, too.

`\scrip` +(scriptstyle + plus)

`\scrm` -(scriptstyle + minus)

`\fscrp` \oplus

`\fscrm` \ominus

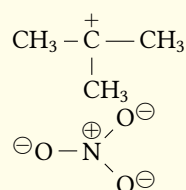
`\fsscrp` \oplus (using `\scriptscriptstyle`)

`\fsscrm` \ominus

```

1 \setatomsep{1.8em}\chemfig{CH_3-\chemabove{C}{\scrip}{-[6]C|H_3}-\vphantom{H_3}CH_3}
2
3 \chemfig{\fmch{}}|O-\chemabove{N}{\fscrp}{-[1]O|\fmch}{-[7]O|\fmch}

```



11. Reaction Mechanisms

With the command

`\mech`[<type>]

one can specify the most common reaction mechanisms. <type> can have one of the following values:

- `\mech` (empty, no opt. argument) nucleophilic substitution S_N
- `\mech[1]` unimolecular nucleophilic substitution S_N1
- `\mech[2]` bimolecular nucleophilic substitution S_N2
- `\mech[se]` electrophilic substitution S_E

- `\mech[1e]` unimolecular electrophilic substitution S_E1
- `\mech[2e]` bimolecular electrophilic substitution S_E2
- `\mech[ar]` electrophilic aromatic substitution $Ar-S_E$
- `\mech[e]` elimination E
- `\mech[e1]` unimolecular elimination $E1$
- `\mech[e2]` bimolecular elimination $E2$
- `\mech[cb]` unimolecular elimination “conjugated base”, i. e. via carbanion $E1_{cb}$

12. Redox Reactions

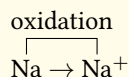
`CHEMMACROS` provides two commands³⁰, to visualize the transfer of electrons in redox reactions. Both commands are using *TikZ*.

`\OX{<name>,<atom>}`

`\redox(<name1>,<name2>)[<tikz>][<num>]{<text>}`

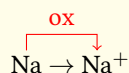
`\OX` places `<atom>` into a node, which is named with `<name>`. If you have set two `\OX`, they can be connected with a line using `\redox`. To do so the names of the two nodes that are to be connected are written in the round braces. Since `\redox` draws a *tikzpicture* with options `remember picture, overlay`, the document needs to be *compiled at least two times*.

```
1 \OX{a,Na} $\rightarrow$ \OX{b,Na}\pch\redox(a,b){oxidation}
```



This line can be customized using *TikZ* keys in `[<tikz>]`:

```
1 \OX{a,Na} $\rightarrow$ \OX{b,Na}\pch\redox(a,b)[->,red]{ox}
```



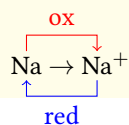
With the argument `[<num>]` the length of the vertical parts of the line can be adjusted. The default length is `.6em`. This length is multiplied with `<num>`. If you use a negative value the line is placed *below* the text.

³⁰ Thanks to [Peter Cao](#) who suggested this feature.


```

1 \OX{a,Na} $\rightarrow$ \OX{b,Na}\pch
2 \redox(a,b)[->,red]{ox}
3 \redox(a,b)[<-,blue][-1]{red}

```



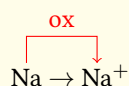
The default length of the vertical lines can be customized with the option

redox **dist** = <dim> A T_EX dimension. Default = .6em

```

1 \chemsetup{redox/dist=1em}
2 \OX{a,Na} $\rightarrow$ \OX{b,Na}\pch\redox(a,b)[->,red]{ox}

```



Additionally the option

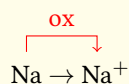
redox **sep** = <dim> Default = .2em

can be used to change the distance between the atom and the beginning of the line.

```

1 \chemsetup{redox/sep=.5em}
2 \OX{a,Na} $\rightarrow$ \OX{b,Na}\pch\redox(a,b)[->,red]{ox}

```

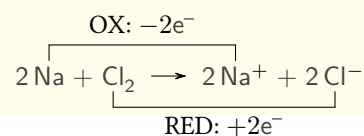


Examples:

```

1 \ch{ 2 "\OX{o1,Na}" + "\OX{r1,Cl}" {}2 -> 2 "\OX{o2,Na}" \pch{} + 2 "\OX{r2,Cl}" \
  mch }
2 \redox(o1,o2){\small OX: $- 2\text{e}^-$}
3 \redox(r1,r2)[][-1]{\small RED: $+ 2\text{e}^-$}

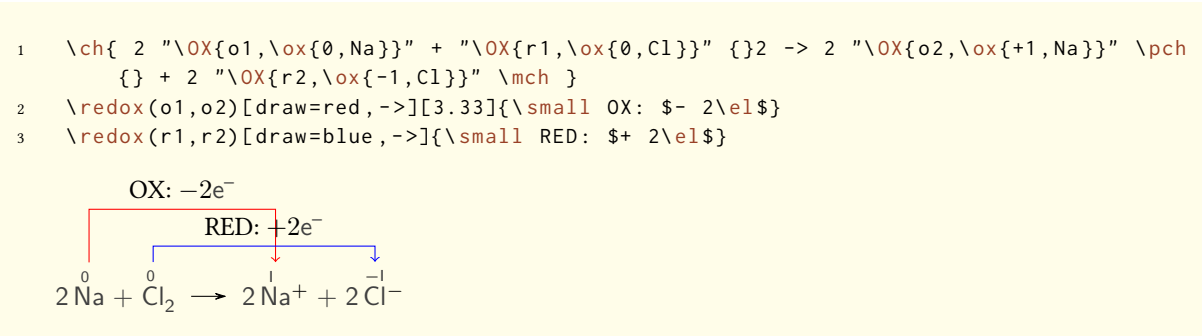
```



```

1 \ch{ 2 "\OX{o1,\ox{0,Na}}"} + "\OX{r1,\ox{0,Cl}}"} {}2 -> 2 "\OX{o2,\ox{+1,Na}}"} \pch
    {} + 2 "\OX{r2,\ox{-1,Cl}}"} \mch }
2 \redox(o1,o2){\small OX: $- 2\el$}
3 \redox(r1,r2)[][{-1}]{\small RED: $+ 2\el$}

```

$$\begin{array}{c}
 \text{OX: } -2e^- \\
 \begin{array}{c}
 \overbrace{2\overset{0}{\text{Na}} + \overset{0}{\text{Cl}}_2} \longrightarrow 2\overset{+1}{\text{Na}} + 2\overset{-1}{\text{Cl}}^- \\
 \underbrace{\hspace{10em}}
 \end{array}
 \end{array}$$


```

1 \ch{ 2 "\OX{o1,\ox{0,Na}}" + "\OX{r1,\ox{0,Cl}}" {}2 -> 2 "\OX{o2,\ox{+1,Na}}" \pch
    {} + 2 "\OX{r2,\ox{-1,Cl}}" \mch }
2 \redox{o1,o2}[green,-stealth]{\small OX}
3 \redox{r1,r2}[purple,-stealth][-1]{\small RED}

```

$$\begin{array}{c}
 \text{OX} \\
 \xrightarrow{\hspace{1.5cm}} \\
 2\overset{0}{\text{Na}} + \overset{0}{\text{Cl}}_2 \rightarrow 2\overset{+1}{\text{Na}} + 2\overset{-1}{\text{Cl}}^- \\
 \xleftarrow{\hspace{1.5cm}}
 \end{array}$$


13.1. Thermodynamic Variables

The following commands use siunitx:

$$\backslash \text{Enthalpy}[\langle \text{keyval} \rangle](\langle \text{subscript} \rangle)\{\langle \text{value} \rangle\}$$

`\Entropy[<keyval>](<subscript>){<value>}`

`\Gibbs[<keyval>](<subscript>){<value>}`

Their usage is pretty much self-explaining:

1	<code>\Enthalpy{123} \\\</code>	$\Delta H^\ominus = 123 \text{ kJ mol}^{-1}$
2	<code>\Entropy{123} \\\</code>	$S^\ominus = 123 \text{ J K}^{-1} \text{ mol}^{-1}$
3	<code>\Gibbs{123}</code>	$\Delta G^\ominus = 123 \text{ kJ mol}^{-1}$

The argument (<subscript>) adds a subscript for specification: `\Enthalpy(r){123}` $\Delta_r H^\ominus = 123 \text{ kJ mol}^{-1}$.

There are several keys to customize the commands.

-none- `exponent` = <anything>
-none- `delta` = <anything>/false
-none- `subscript` = left/right
-none- `unit` = <unit>

The default values depend on the command.

1	<code>\Enthalpy[unit=\kilo\joule]{-285} \\\</code>	$\Delta H^\ominus = -285 \text{ kJ}$
2	<code>\Gibbs[delta=false]{0} \\\</code>	$G^\ominus = 0 \text{ kJ mol}^{-1}$
3	<code>\Entropy[delta=\Delta,exponent=]{56.7}</code>	$\Delta S = 56.7 \text{ J K}^{-1} \text{ mol}^{-1}$

The unit is set corresponding to the rules of siunitx and depends on its settings:

1	<code>\Enthalpy{-1234.56e3} \\\</code>	
2	<code>\sisetup{per-mode=symbol,exponent-product=\cdot,output-decimal-marker={,},group-four-digits=true}</code>	
3	<code>\Enthalpy{-1234.56e3}</code>	

$\Delta H^\ominus = -1234.56 \times 10^3 \text{ kJ mol}^{-1}$
 $\Delta H^\ominus = -1\,234,56 \cdot 10^3 \text{ kJ/mol}$

13.1.1. Create New Variables

You can use the command

`\setnewstate[<keyval>]{<name>}<symbol>{<unit>}`

to create new corresponding commands:

1	<code>\setnewstate{Helmholtz}{A}{\kilo\joule\per\mole}</code>	
2	<code>\setnewstate[subscript-left=false,exponent=]{ElPot}{E}{\volt}</code>	
3	<code>\Helmholtz{123.4} \\\</code>	
4	<code>\ElPot{-1.1} \\\</code>	
5	<code>\ElPot[exponent=0]{\$\ch{Sn} \ch{Sn^2+} \ch{Pb^2+} \ch{Pb}}{0.01}</code>	

$\Delta A^\ominus = 123.4 \text{ kJ mol}^{-1}$
 $\Delta E = -1.1 \text{ V}$
 $\Delta E_{\text{Sn}|\text{Sn}^{2+}||\text{Pb}^{2+}|\text{Pb}}^0 = 0.01 \text{ V}$

The command has some keys with which the default behaviour of the new command can be set.

`exponent` = <anything>

`delta` = <anything>/false

`-none-` `subscript-left` = true/false

`subscript` = <anything>

13.1.2. Redefine Variables

With

`\renewstate`[<keyval>]{<name>}<symbol>{<unit>}

you can redefine the already existing commands:

```
1 \renewstate{Enthalpy}{h}{\joule}
2 \Enthalpy{f}{12.5}
```

$$\Delta_f h^\ominus = 12.5 \text{ J}$$

The command is analogous to `\setnewstate`, i. e. it has the same keys.

So – for following thermodynamic conventions – one could define a molar and an absolute variable:

```
1 \setnewstate[exponent=]{enthalpy}{h}{\kilo\joule\per\mole}% molar
2 \renewstate[exponent=]{Enthalpy}{H}{\kilo\joule}% absolute
3 \enthalpy{-12.3} \Enthalpy{-12.3}
```

$$\Delta h = -12.3 \text{ kJ mol}^{-1} \quad \Delta H = -12.3 \text{ kJ}$$

13.2. State

The commands presented in section 13.1 internally all use the command³¹

`\State`[<keyval>]{<symbol>}{<subscript>}

It can be used to write the thermodynamic variables without value and unit.

Examples:

```
1 \State{A}, \State{G}{f}, \State[subscript-left=false]{E}{\ch{Na}}, \State[exponent=\SI{1000}{\celsius}]{H}
```

$$\Delta A^\ominus, \Delta_f G^\ominus, \Delta E_{\text{Na}}^\ominus, \Delta H^{1000\text{ }^\circ\text{C}}$$

Again there are some keys to customize the command:

`state` `exponent` = <anything>

`state` `subscript-left` = true/false

`state` `delta` = <anything>/false

³¹ Please note that {<subscript>} is an *optional* argument.

14. Spectroscopy

If substances are examined whether they are what they're supposed to, one often needs NMR spectroscopy. Measured results then are written in a way like: $^1\text{H-NMR}$ (400 MHz, CDCl_3): $\delta = 1.59\dots$ **CHEMMACROS** provides a command which simplifies writing this down (uses siunitx).

```
\NMR{<num>,<elem>}(<num>,<unit>)[<solvent>]
```

```
\NMR*{<num>,<elem>}(<num>,<unit>)[<solvent>]
```

All arguments are optional! Without arguments we get³²:

```
1 \NMR \\\                                 $^1\text{H-NMR: } \delta$ 
2 \NMR* \                                 $^1\text{H-NMR}$ 
```

With the first argument you can specify the kind of NMR:

```
1 \NMR{13,C}                              $^{13}\text{C-NMR: } \delta$ 
```

With the second argument the frequency (in MHz) can be specified:

```
1 \NMR(400)                              $^1\text{H-NMR (400 MHz): } \delta$ 
```

You also can change the unit:

```
1 \NMR(4e8,\hertz)                        $^1\text{H-NMR (4} \times 10^8 \text{ Hz): } \delta$ 
```

Please note that the setup of siunitx also has an impact on this command:

```
1 \sisetup{exponent-product=\cdot}\NMR(4e8,\hertz)
 $^1\text{H-NMR (4} \cdot 10^8 \text{ Hz): } \delta$ 
```

And finally with the third argument the solvent can be specified:

```
1 \NMR[CDCl3]                              $^1\text{H-NMR (CDCl}_3\text{): } \delta$ 
```

With the keys

```
nmr    unit = <unit> Default = \mega\hertz
```

nmr **nucleus** = {<num>,<elem>} Default = {1,H}

the default unit and the default nucleus can be changed.

Examples:

```
1  {\chemsetup[nmr]{nucleus={13,C}}\NMR(100) \NMR*(100) } \\  
2  \NMR*{19,F}[CFCl3] \NMR*{19,F}(285)[CFCl3] \\  
3  \NMR(400)[CDCl3] = \num{1.59} (q, 1H, \textit{J} = \SI{11.6}{\hertz}, H-4)  
  
13C-NMR (100 MHz):  $\delta$  13C-NMR (100 MHz)  
19F-NMR (CFCl3) 19F-NMR (285 MHz, CFCl3)  
1H-NMR (400 MHz, CDCl3):  $\delta$  = 1.59 (q, 1H,  $J$  = 11.6 Hz, H-4)
```

15. Commands for mhchem

mhchem isn't loaded automatically any more but only if you've specified **method** = mhchem in the preamble. In the default settings **CHEMMACROS** uses **CHEMFORMULA** instead.

CHEMMACROS provides only one command specifically for mhchem³³. It is meant to place text below of compounds.

\mhName[<keyval>]{<formula>}{<text>}

For example:

```
1  \ce{4 C2H5Cl + Pb / Na -> \mhName{Pb(C2H5)4}{former antiknock additive} + NaCl}  
  
4 C2H5Cl + Pb/Na → Pb(C2H5)4 + NaCl  
                      former antiknock  
                      additive
```

There are several keys to customize **\mhName**.

mhName **align** = <alignment command> the alignment of the text in the box it is placed in, default = **\centering**

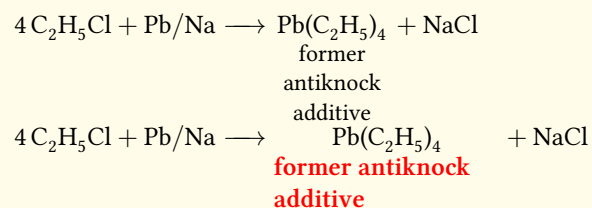
mhName **format** = <anything> the format of the text

mhName **fontsize** = the fontsize of the text, default = **\tiny**

mhName **width** = <dim>/auto the width of the box the text is placed in, default = auto

```
1  \ce{4 C2H5Cl + Pb / Na -> \mhName[fontsize=\footnotesize]{Pb(C2H5)4}{former  
    antiknock additive} + NaCl}\\  
2  \chemsetup[mhName]{align=\raggedright,fontsize=\small,format=\bfseries\color{red},  
    width=3cm}  
3  \ce{4 C2H5Cl + Pb / Na -> \mhName{Pb(C2H5)4}{former antiknock additive} + NaCl}
```

³² All arguments can be combined freely, the command can also be used in math mode. ³³ **CHEMFORMULA** provides its own possibility.



16. Reaction Environments

16.1. Defined by CHEMMACROS

You can use these environments for numbered...

`\begin{reaction} <formula or mhchem code> \end{reaction}`

`\begin{reactions} <formula or mhchem code> \end{reactions}`

...and their starred versions for unnumbered reactions.

`\begin{reaction*} <formula or mhchem code> \end{reaction*}`

`\begin{reactions*} <formula or mhchem code> \end{reactions*}`

With them you can create (un)numbered reaction equations similar to mathematical equations.

The environments `reaction/reaction*` use the `equation/equation*` environments and the environments `reactions/reactions*` use the `align/align*` environments to display the reactions.

<pre> 1 Reaction with counter: 2 \begin{reaction} 3 A -> B 4 \end{reaction} </pre>	<p>Reaction with counter:</p> $A \longrightarrow B \quad \{1\}$
---	---

<pre> 1 Reaction without counter: 2 \begin{reaction*} 3 C -> D 4 \end{reaction*} </pre>	<p>Reaction without counter:</p> $C \longrightarrow D$
--	--

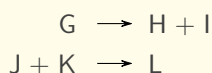
<pre> 1 Several aligned reactions with counter: 2 \begin{reactions} 3 A &-> B + C \\ 4 D + E &-> F 5 \end{reactions} </pre>	<p>Several aligned reactions with counter:</p> $A \longrightarrow B + C \quad \{2\}$ $D + E \longrightarrow F \quad \{3\}$
---	--

```

1 Several aligned reactions without counter:
2 \begin{reactions*}
3   G      &-> H + I \\
4   J + K &-> L
5 \end{reactions*}

```

Several aligned reactions without counter:



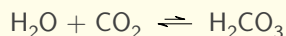
If you want to change the layout of the counter tags, you can use

`\renewtagform{<tagname>}{<format>}{<right delim>}{<left delim>}`³⁴.

```

1 \renewtagform{reaction}[R \textbf]{[]{}
2 \begin{reaction}
3   H2O + CO2 <=> H2CO3
4 \end{reaction}

```



[R 4]

16.2. Own Reactions

You can create new types of reactions with the command:

`\newreaction[<keyval>]{<name>}{<math name>}`

`<name>` will be the name of the new environment. `<math name>` is the used math environment. The command has two keys.

-none- `star` = true/false

-none- `arg` = true/false

There is `star`, which will also define a starred version of the new environment, if the starred math environment exists. If it doesn't exist, this will cause an error.

Then there is `arg`, which is used to define an environment with a mandatory argument. Of course this only works, if the used math environment has a mandatory argument.

The predefined environments are defined via

`\newreaction[star]{reaction}{equation}` and

`\reaction[star]{reactions}{align}`.

Let's suppose, you'd like to have the alignment behaviour of the alignat environment for **CHEM-FORMULA**/mhchem reactions. You could do the following:

`\newreaction[star,arg]{reactionsat}alignat`

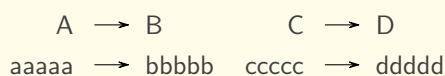
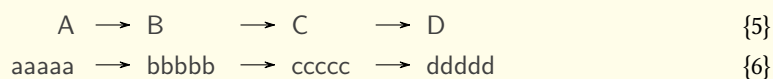
With this the reactionsat environment is defined.

³⁴ Provided by the mathtools package


```

1 \newreaction[star,arg]{reactionsat}{alignat}
2 \begin{reactionsat}{3}
3 A &-> B &&-> C &&-> D \\
4 aaaaa &-> bbbbb &&-> ccccc &&-> ddddd
5 \end{reactionsat}
6 \begin{reactionsat*}{2}
7 A &-> B & C &-> D \\
8 aaaaa &-> bbbbb &\quad{} ccccc &-> ddddd
9 \end{reactionsat*}

```



16.3. List of Reactions

CHEMMACROS also provides a command to display a list of the reactions created with the reaction environment.

`\listofreactions`

```

1 \listofreactions

```

List of reactions

Reaction {1}	23
Reaction {2}	23
Reaction {3}	23
Reaction [4]	24
Reaction {5}	25
Reaction {6}	25
Reaction {7}: Autoprotolyse	26
Reaction {8}: Synthese von Alkanen	48

The Output of this list can be modified by two options:

reaction **list-name** = <name of the list> Let's you set the name of the list manually. Default = List of reactions

reaction **list-entry** = <prefix to each entry> Let's you set a prefix to each list entry. Default = Reaction

Both default option values recognize the package option **german**.

Instead of using the option **list-name** you also could redefine `\reactionlistname`.

The list lists all reactions with a number and disregards reactions without number. All reaction environments without star have an optional argument which let's you add a description (or caption) for the entry in the list.

```
1 \begin{reaction}[Autoprotolyse]
2 2 H2O <=> H3O+ + OH-
3 \end{reaction}
```

$$2 \text{H}_2\text{O} \rightleftharpoons \text{H}_3\text{O}^+ + \text{OH}^- \quad \{7\}$$

17. Phases

These commands are intended to indicate the phase of a compound.

`\sld[<anything>]` (s)

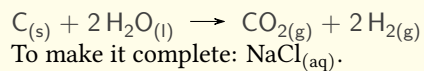
`\lqd[<anything>]` (l)

`\gas` (g)

`\aq` (aq)

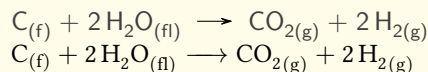
Please notice, that the commands `\solid` and `\liquid` are now called `\sld` and `\lqd`, respectively. Also notice that they *don't* have an optional argument inside `\ch`!

```
1 \ch{C\sld{} + 2 H2O\lqd{} -> CO2\gas{} + 2 H2\gas}\
2 To make it complete: NaCl\aq.
```



With the package option `german` (see section 4) or by using the optional arguments you get the german versions:

```
1 {\chemsetup[option]{german=true}
2 \ch{C\sld{} + 2 H2O\lqd{} -> CO2\gas{} + 2 H2\gas} }\
3 \ce{C \sld[f] + 2 H2O \lqd[f1] -> CO2\gas{} + 2 H2\gas}
```



If you looked closely, you have probably noticed that the german `\lqd` isn't identical to `\lqd[f1]` but to `\lqd[f\1]`. This makes the subscript more readable.

Of course it doesn't matter which command with optional argument you use. Both of them just write a subscript with braces. `\sld[f]` is identical to `\lqd[f]`.

One can think of other uses, too:

```
1 C\sld[graphite]
```

C_(graphite)

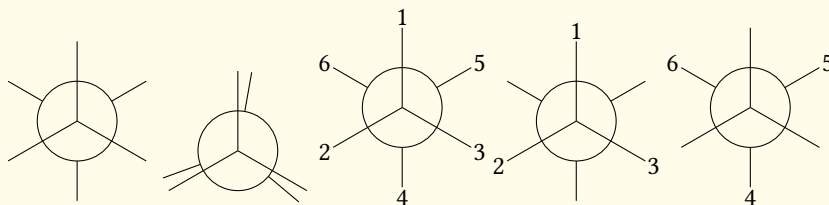
18. Newman Projections

CHEMMACROS provides the command

```
\newman[<keyval>](<angle>){<1>,<2>,<3>,<4>,<5>,<6>}
```

which allows you to create newman projections (uses *TikZ*). With <angle> the back atoms are rotated counter clockwise with respect to the front atoms.

```
1 \newman{} \newman(170){}  
2 \newman{1,2,3,4,5,6} \newman{1,2,3} \newman{,,4,5,6}
```



Several keys allow customization:

newman **angle** = <angle> default angle

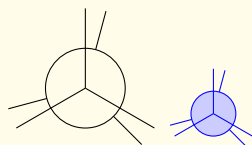
newman **scale** = <factor> scale the whole projection

newman **ring** = <tikz> customize the ring with *TikZ* keys

newman **atoms** = <tikz> customize the nodes within which the atoms are set

newman **back-atoms** = <tikz> explicitly customize the back atoms

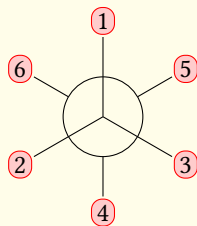
```
1 \chemsetup[newman]{angle=45} \newman{}  
2 \newman[scale=.75,ring={draw=blue,fill=blue!20}]{}
```



```

1 \chemsetup[newman]{atoms={draw=red,fill=red!20,inner sep=2pt,rounded corners}}
2 \newman{1,2,3,4,5,6}

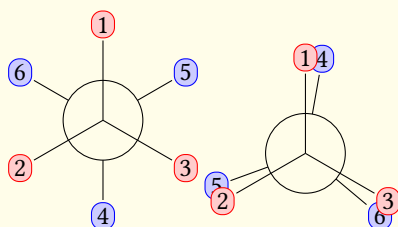
```



```

1 \chemsetup[newman]{
2   atoms = {draw=red,fill=red!20,inner sep=2pt,rounded corners},
3   back-atoms = {draw=blue,fill=blue!20,inner sep=2pt,rounded corners}
4 }
5 \newman{1,2,3,4,5,6} \newman(170){1,2,3,4,5,6}

```



19. s, p, and Hybrid Orbitals

CHEMMACROS provides the following command to create orbitals:

`\orbital[<keyval>]{<type>}`

There are the following types available for <type>:

s

p

sp

sp²

sp³

```

1 \orbital{s} \orbital{p} \orbital{sp} \orbital{sp2} \orbital{sp3}

```



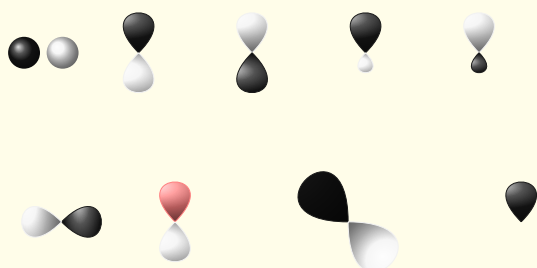
Depending on the type you have different keys to modify the orbitals:

- orbital** **phase** = +/- changes the phase of the orbital (all types)
- orbital** **scale** = <factor> changes the size of the orbital (all types)
- orbital** **color** = <color> changes the color of the orbital (all types)
- orbital** **angle** = <angle> rotates the orbitals with a p contribution counter clockwise (all types except s)
- orbital** **half** = true/false displays only half an orbital (only p)

```

1 \orbital{s} \orbital[phase=-]{s}
2 \orbital{p} \orbital[phase=-]{p}
3 \orbital{sp3} \orbital[phase=-]{sp3}
4
5 \orbital[angle=0]{p} \orbital[color=red!50]{p} \orbital[angle=135,scale=1.5]{p} \
  orbital[half]{p}

```



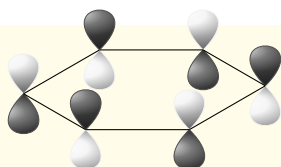
Additionally there are two keys, with which the *TikZ* behaviour can be changed.

- orbital** **overlay** = true/false the orbital “doesn't need space”; it is displayed with the *TikZ* option overlay.
- orbital** **opacity** = <num> the orbital becomes transparent; <value> can have values between 1 (fully opaque) to 0 (invisible).

```

1 \vspace{1cm}\hspace{1cm}
2 \chemsetup[orbital]{
3   overlay,
4   p/color = black!70
5 }
6 \setbondoffset{0pt}
7 \chemfig{?\orbital{p}-[,1.3]{\orbital[phase=-]{p}}-[:30,1.1]\orbital{p}-[:150,.9]{\
  orbital[phase=-]{p}}-[:4,1.3]\orbital{p}-[:150,1.1]{\orbital[phase=-]{p}}?}
8 \vspace{1cm}

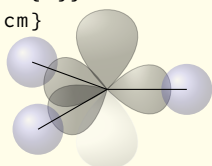
```



```

1  \vspace{2cm}\hspace{2cm}
2  \setbondoffset{0pt}
3  \chemsetup[orbital]{
4    overlay ,
5    opacity = .75 ,
6    p/scale = 1.6 ,
7    s/color = blue!50 ,
8    s/scale = 1.6
9  }
10 \chemfig{\orbital{s}-[: -20]{\orbital[scale=2]{p}}{\orbital[half,angle=0]{p}}{\orbital[angle=170,half]{p}}{\orbital[angle=-150,half]{p}}(-[: -150]\orbital{s})-\orbital{s}}
11 \vspace{2cm}

```



20. CHEMFORMULA support

CHEMFORMULA and **CHEMMACROS** are designed to work together closely. This is not perfect, though. There are even some restrictions. If you want to avoid them simply use `method = mhchem` and forget about this section and the next part.

You don't need to use `\mch` and related commands inside `\ch`. Indeed, you shouldn't use them as they might mess with the subscript and superscript alignment. The **CHEMMACROS** option circled is obeyed by `\ch`.

```

1 \chemsetup[option]{circled=all}      H⊕ + OH⊖ ⇌ H2O
2 \ch{H+ + OH- <=> H2O}

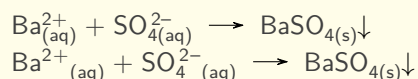
```

The commands `\sld` and `\lqd` don't have an optional argument inside `\ch`.

```

1 \ch{Ba^{2+}\aq{}} + SO4^{2-}\aq{} -> BaSO4\sld{} v} \
2 \ch{Ba^2+ \aq{} + SO4^2- \aq{} -> BaSO4\sld{} v}

```



The option `synchronize` recognizes the format and font selections made for `CHEMFORMULA`.

21. Key Overview

In the table below all keys provided by `CHEMMACROS` for customization are listed. All keys that belong to a module can be set with

`\chemsetup[<module>]{<keyval>}` or

`\chemsetup{<module>/<keyval>}`.

Some keys can be set without value. Then the underlined value is used.

key	module	values	default	
package options:				
<code>bpchem</code>	<code>option</code>	<u>true</u> /false	false	page 5
<code>circled</code>	<code>option</code>	<u>formal</u> /all/none	formal	page 5
<code>circletype</code>	<code>option</code>	<u>chem</u> /math	chem	page 5
<code>detect-bold</code>	<code>option</code>	<u>true</u> /false	false	page 5
<code>EZ</code>	<code>option</code>	<u>chemmacros</u> /cool	chemmacros	page 5
<code>german</code>	<code>option</code>	<u>true</u> /false	false	page 5
<code>ghs</code>	<code>option</code>	<u>true</u> /false	true	page 5
<code>method</code>	<code>option</code>	<u>chemformula</u> /formula	formula	page 5
<code>Nu</code>	<code>option</code>	<u>chemmacros</u> /mathspec	chemmacros	page 5
<code>strict</code>	<code>option</code>	<u>true</u> /false	false	page 5
<code>synchronize</code>	<code>option</code>	<u>true</u> /false	false	page 5
<code>version</code>	<code>option</code>	<u>1/2</u> /bundle	bundle	page 5
<code>xspace</code>	<code>option</code>	<u>true</u> /false	true	page 6
<code>\ba, \Nu:</code>				
<code>elpair</code>	<code>particle</code>	<u>dots</u> /dash/false	false	page 8
<code>\pch, \mch, \fpch, \fmch:</code>				
<code>append</code>	<code>charges</code>	<u>true</u> /false	false	page 12
<code>\ox:</code>				
<code>parse</code>	<code>ox</code>	<u>true</u> /false	true	page 13
<code>roman</code>	<code>ox</code>	<u>true</u> /false	true	page 13

key	module	values	default	
pos	ox	top/super/side	top	page 13
explicit-sign	ox	true/false	false	page 13
decimal-marker	ox	comma/point	point	page 13
\OX, \redox:				
dist	redox	<dim>	.6em	page 17
sep	redox	<dim>	.2em	page 17
\Enthalpy, \Entropy, \Gibbs:				
exponent		<anything>	\standardstate	page 19
delta		<anything>/false		page 19
subscript		left/right		page 19
unit		<unit>		page 19
\setnewstate, \renewstate:				
exponent		<anything>	\standardstate	page 19
delta		<anything>/false		page 19
subscript		<anything>		page 19
subscript-left		true/false		page 20
\State:				
exponent	state	<anything>	\standardstate	page 20
delta	state	<anything>/false		page 20
subscript-left	state	true/false		page 20
\NMR:				
unit	nmr	<unit>	\mega\hertz	page 21
nucleus	nmr	{<num>,<atom symbol>}	{1,H}	page 22
\newreaction:				
star		true/false	false	page 24
arg		true/false	false	page 24
list-name	reaction	<anything>	List of reactions	page 25
list-entry	reaction	<anything>	Reaction	page 25
\mhName:				
align	mhName	<alignment>	\centering	page 22
format	mhName	<commands>		page 22
fontsize	mhName	<fontsize>	\tiny	page 22
width	mhName	<dim>		page 22
\newman:				
angle	newman	<angle>	0	page 27
scale	newman	<factor>	1	page 27
ring	newman	<tikz>		page 27
atoms	newman	<tikz>		page 27
back-atoms	newman	<tikz>		page 27
\orbital <type> = s/p/sp/sp2/sp3:				
phase	orbital/<type>	+/-	+	page 29
scale	orbital/<type>	<factor>	1	page 29
color	orbital/<type>	<color>	black	page 29
angle	orbital/<type>	<angle>	90	page 29
half	orbital/p	true/false	false	page 29
overlay	orbital	true/false	false	page 29
opacity	ornital	<num>	1	page 29

Part III.

chemformula

22. Setup

All of **CHEMFORMULA**'s options belong to the module **chemformula**. This means they can be setup with

```
1 \chemsetup[chemformula]{<options>} or
2 \chemsetup{chemformula/<option1>,chemformula/<option2>}
```

23. The Basic Principle

CHEMFORMULA offers one main command.

\ch[<options>]{<input>}

The usage will seem very familiar to you if you're familiar with mhchem:

1 \ch{H2O} \\	H ₂ O
2 \ch{Sb2O3} \\	Sb ₂ O ₃
3 \ch{H+} \\	H ⁺
4 \ch{CrO4^2-} \\	CrO ₄ ²⁻
5 \ch{AgCl2-} \\	AgCl ₂ ⁻
6 \ch{[AgCl2]-} \\	[AgCl ₂] ⁻
7 \ch{Y^{99}+} \\	Y ⁹⁹⁺
8 \ch{Y^{99+}} \\	Y ⁹⁹⁺
9 \ch{H2_{(aq)}} \\	H _{2(aq)}
10 \ch{NO3-} \\	NO ₃ ⁻
11 \ch{(NH4)2S} \\	(NH ₄) ₂ S
12 \ch{^{227}_{90}Th+} \\	²²⁷ ₉₀ Th ⁺
13 \$V_{\ch{H2O}}\$ \\	V _{H₂O}
14 \ch{Ce^{IV}} \\	Ce ^{IV}
15 \ch{KCr(SO4)2 * 12 H2O}	KCr(SO ₄) ₂ · 12 H ₂ O

However, there are differences. The most notable one: **CHEMFORMULA** distinguishes between different types of input. These different parts *have* to be separated with blanks:

\ch{part1 part2 part3 part4}

A blank in the input *never* is a blank in the output. This role of the blank strictly holds and disregarding it can have unexpected results and even lead to errors.

Another notable difference: **CHEMFORMULA** tries to avoid math mode whenever possible:

<pre>1 \ch{A + B ->[a] C} \\ 2 \ce{A + B ->[a] C}</pre>	$A + B \xrightarrow{a} C$ $A + B \xrightarrow{a} C$
---	---

This means that `\ch{2H2O}` is recognized as a *single* part, which in this case is recognized as a compound.

<pre>1 \ch{2H2O} \\ 2 \ch{2 H2O}</pre>	$_2\text{H}_2\text{O}$ $2\text{H}_2\text{O}$
--	--

This also means, that a part cannot contain a blank since this will automatically divide it into two parts. If you need an extra blank in the output you need to use `~`. However, since commands in most cases gobble a space after them an input like `\ch{\command ABC}` will be treated as a single part. If you want or need to divide them you need to add an empty group: `\ch{\command{} ABC}`. The different input types are described in the following sections.

There are some options to customize the output of the `\ch` command. They can either be applied locally using the optional argument or can be set globally using the setup command. All options of `CHEMFORMULA` belong to the module `chemformula`.

`\chemsetup[chemformula]{<options>}`

24. Stoichiometric Factors

A stoichiometric factor may only contain of numbers and the signs `.`, `_`, `/`

<pre>1 \ch{2} \\ 2 \ch{12} 3 4 % decimals: 5 \ch{3.5} \\ 6 \ch{5,75} 7 8 % fractions: 9 \ch{3/2} \\ 10 \ch{1_1/2}</pre>	2 12 3.5 5.75 $\frac{3}{2}$ $1\frac{1}{2}$
---	--

You have to be a little bit careful with the right syntax but I believe it is rather intuitive.

```
1 this won't work but will result in an error: \ch{1/1_1}
```

There are a few possibilities to customize the output.

`decimal-marker` = `<marker>` the symbol to indicate the decimal. Default = `.`

`frac-style = math/xfrac/nicefrac` determines how fractions are displayed. Default = `math`

`stoich-space = <dim>` The space that is placed after the stoichiometric factor. Default = `.1667em`

```
1 \ch[decimal-marker={,}]{3.5} \ch[decimal-marker={\cdot}]{3,5}
3,5 3·5
```

The option `frac-style = xfrac` uses the `\sfrac` command of the `xfrac` package. The output strongly depends on the font you use.

```
1 \ch[frac-style=xfrac]{3/2} \ch[frac-style=xfrac]{1_1/2}
¾ 1½
```

CHEMFORMULA defines the instance `formula-text-fraction` which you can redefine to your needs. See the `xfrac` documentation for further information. The default definition is this:

```
1 \DeclareInstance{xfrac}{chemformula-text-fraction}{text}
2 {
3   slash-left-kern = -.15em ,
4   slash-right-kern = -.15em
5 }
```

This document uses the font `Linux Libertine` and the following definition:

```
1 \DeclareInstance{xfrac}{chemformula-text-fraction}{text}
2 {
3   scale-factor      = 1 ,
4   denominator-bot-sep = -.2ex ,
5   denominator-format = \scriptsize #1 ,
6   numerator-top-sep = -.2ex ,
7   numerator-format = \scriptsize #1
8 }
```

The option `frac-style = nicefrac` uses the `\nicefrac` command of the `nicefrac` package.

```
1 \ch[frac-style=nicefrac]{3/2} \ch[frac-style=nicefrac]{1_1/2}
¾ 1½
```

The option `stoich-space` allows you to customize the space between stoichiometric factor and the group following after it.

1	<code>\ch{2 H2O} \</code>	$2\text{H}_2\text{O}$
2	<code>\ch[stoich-space=.2em]{2 H2O}</code>	$2\text{H}_2\text{O}$

25. Compounds

CHEMFORMULA determines compounds as the type that “doesn't fit in anywhere else”. This point will become more clear when you know what the other types are.

1	<code>\ch{H2SO4} \</code>	H_2SO_4
2	<code>\ch{[Cu(NH3)4]^2+}</code>	$[\text{Cu}(\text{NH}_3)_4]^{2+}$

25.1. Adducts

CHEMFORMULA has two identifiers which will create adducts.

`\ch{A.B}`

`\ch{A*B}`

1	<code>\ch{CaSO4.H2O} \</code>	$\text{CaSO}_4 \cdot \text{H}_2\text{O}$
2	<code>\ch{CaSO4*H2O}</code>	$\text{CaSO}_4 \cdot \text{H}_2\text{O}$

Since numbers in a compound always are treated as subscripts (see section 25.2) you sometimes need to introduce stoichiometric factors for the right output:

1	<code>\ch{Na3PO4*12H2O} \</code>	$\text{Na}_3\text{PO}_4 \cdot 12\text{H}_2\text{O}$
2	<code>\ch{Na3PO4* 12 H2O} \</code>	$\text{Na}_3\text{PO}_4 \cdot 12\text{H}_2\text{O}$
3	<code>\ch{Na3PO4 * 12 H2O}</code>	$\text{Na}_3\text{PO}_4 \cdot 12\text{H}_2\text{O}$

25.2. Subscripts

All numbers in a compound are treated as subscripts.

1	<code>\ch{H2SO4}</code>	H_2SO_4
---	-------------------------	-------------------------

If you want a letter to be a subscript you can use the math syntax:

1	<code>\ch{A_nB_m}</code>	A_nB_m
---	--------------------------	----------

The subscript recognizes groups. You can also use math inside it.

1	<code>\ch{A_{n\$}B_{m\$}} \\\</code>	A_nB_m
2	<code>\ch{NaCl_{(aq)}}</code>	$\text{NaCl}_{(\text{aq})}$

25.3. Commands

Commands are allowed in a compound:

1	<code>\ch{\textbf{A2}B3} \ch{A2\color{red}B3} A₂B₃ A₂B₃</code>
---	---

However, if the commands demand numbers as argument, e. g. space commands or chemmacros³⁵ `\ox` command the direct use will fail. This is because the numbers are treated as subscripts *before* the command expands.

1	<code>\ch{A\hspace{2mm}B}</code> will raise an error because <code>\hspace</code> sees something like this : <code>\hspace{\$_2\$mm}</code> .
---	--

See section 27.1 for a way around this.

25.4. Charges and Other Superscripts

If a compound *ends* with a plus or minus sign it will be treated as charge sign and typeset as superscript. In other places a plus is treated as a triple bond and a dash will be used as a single bond, see section 25.5.

1	<code>\ch{A+B} \ch{AB+} \\\</code>	$A\equiv B$ AB^+
2	<code>\ch{A-B} \ch{AB-}</code>	$A-B$ AB^-

For longer charge groups or other superscripts you can use the math syntax. It recognizes groups and you can use math inside them. Inside these groups neither + nor - are treated as bonds. If a dot . is inside a superscript it is treated as indicator for a radical.

1	<code>\ch{A^{x-}} \\\</code>	A^{x-}
2	<code>\ch{A^x-} \\\</code>	A^{x-}
3	<code>\ch{A^{x}-} \\\</code>	A^{x-}
4	<code>\ch{A^{x-\$}} \\\</code>	A^{x-}
5	<code>\ch{RNO2^{-.}} \\\</code>	$\text{RNO}_2^{\cdot-}$
6	<code>\ch{^3H} \\\</code>	^3H
7	<code>\ch{^{14}6C} \\\</code>	$^{14}_6\text{C}$
8	<code>\ch{^{58}_{26}Fe}</code>	$^{58}_{26}\text{Fe}$

³⁵ CTAN: chemmacros

Ions and ion composites with more than one charge can be typeset quite as easy:

```
1 \ch{SO4^2-} \ch{Ca^2+ SO4^2-} SO42- Ca2+SO42-
```

25.5. Bonds

There are three kinds of bonds:

```
1 single: \ch{CH3-CH3} \\\           single: CH3-CH3
2 double: \ch{CH2=CH2} \\\          double: CH2=CH2
3 triple: \ch{CH+CH}                triple: CH≡CH
```

25.6. Customization

These options allow you to customize the output of the compounds:

subscript-vshift = <dim> Extra vertical shift of the subscripts. Default = 0pt

subscript-style = text/math Style that is use to typeset the subscripts. Default = text

charge-hshift = <dim> Shift of superscripts when following a subscript. Default = .3ex

charge-style = text/math Style that is use to typeset the superscripts. Default = text

adduct-space = <dim> Space to the left and the right of the adduct point. Default = .1333em

bond-length = <dim> The length of the bonds. Default is the length of an endash as measured by `\settowidth{<len>}{\textendash}`.

Maybe you have noticed that charges of certain ions are a little bit shifted to the right:

```
1 \ch{SO4^2-} \ch{NH4+} \ch{Na+} SO42- NH4+ Na+
```

They are shifted if they *follow* a subscript. The amount of the shift can be set with the option **charge-hshift**.

```
1 \chemsetup[chemformula]{charge-hshift=0pt}
2 \ch{SO4^2-} \ch{NH4+} \ch{Na+} \\\
3 \chemsetup[chemformula]{charge-hshift=1ex}
4 \ch{SO4^2-} \ch{NH4+} \ch{Na+}
```

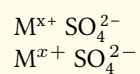
SO₄²⁻ NH₄⁺ Na⁺
SO₄²⁻ NH₄⁺ Na⁺

If you don't want the charges to be typeset in text mode you can switch to math mode:

```

1 \ch{M^x+} \ch{SO4^2-} \\
2 \chemsetup[chemformula]{charge-style = math}
3 \ch{M^x+} \ch{SO4^2-}

```

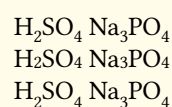


The option `subscript-vshift` can be used to adjust the vertical shift of the subscripts:

```

1 \ch{H2SO4} \ch{Na3PO4} \\
2 \chemsetup[chemformula]{subscript-vshift=.5ex}
3 \ch{H2SO4} \ch{Na3PO4} \\
4 \chemsetup[chemformula]{subscript-vshift=-.2ex}
5 \ch{H2SO4} \ch{Na3PO4}

```

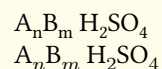


You can choose the mode subscripts are typeset in the same way as it is possible for the charges:

```

1 \ch{A_nB_m} \ch{H2SO4} \\
2 \chemsetup[chemformula]{subscript-style = math}
3 \ch{A_nB_m} \ch{H2SO4}

```

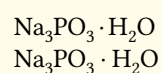


The option `adduct-space` sets the space left and right to the adduct symbol \cdot .

```

1 \ch{Na3PO3*H2O} \\
2 \chemsetup[chemformula]{adduct-space=.2em}
3 \ch{Na3PO3*H2O}

```

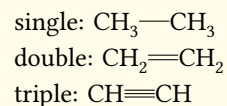


Changing the length of the bonds:

```

1 \chemsetup[chemformula]{bond-length=4mm}%
2 single: \ch{CH3-CH3} \\
3 double: \ch{CH2=CH2} \\
4 triple: \ch{CH+CH}

```



26. Special Input Types

There are some “special type” input groups. These are groups which consist of only one token, namely of the following ones:

`\ch{ + }` + creates the plus sign between compounds with space around it:

`\ch{2 Na + Cl2}` 2Na + Cl₂

`\ch{ v }` ↓ sign for precipitate: `\ch{BaSO4 v}` BaSO₄↓

`\ch{ ^ }` ↑ sign for escaping gas³⁶: `\ch{H2 ^}` H₂↑

The space left and right of the plus sign can be set with this option:

`plus-space` = <dim> Default = .3em

1	<code>\ch{A + B}\</code>	A + B
2	<code>\ch[plus-space=4pt]{A + B}</code>	A + B

27. Escaped Input

In some cases it may be desirable to prevent `CHEMFORMULA` from parsing the input. This can be done in two ways.

27.1. Text

If you put something between " " or ' ' then the input will be treated as normal text, except that spaces are not allowed and have to be input with ~.

`\ch{ "<escaped text>" }`

`\ch{ '<escaped text>' }`

1	<code>\ch{"\ox{2,Ca}" 0} \</code>
2	<code>\ch{"\ldots\," Na + "\ldots\," Cl2 -> "\ldots\," NaCl} \</code>
3	<code>\ch{'A~->~B'}</code>

$$\overset{\text{II}}{\text{CaO}}$$
$$\dots\text{Na} + \dots\text{Cl}_2 \longrightarrow \dots\text{NaCl}$$
$$\text{A} \rightarrow \text{B}$$

In many cases you won't need to escape the input. But when you get into trouble when using a command inside `\ch` try hiding it.

³⁶ Is this the correct English term? Please correct me if it isn't.

27.2. Math

If you especially want to input math you just enclose it with `$ $`. This output is different from the escaped text as it is followed by a space.

`\ch{ $<escaped math>$ }`

1	escaped text: <code>\ch{"\$x\$" H2O} \\</code>	escaped text: $x\text{H}_2\text{O}$
2	escaped math: <code>\ch{\$x\$ H2O} \\</code>	escaped math: $x\text{H}_2\text{O}$
3	<code>\ch{\$2n\$ Na + \$n\$ Cl2 -> \$2n\$ NaCl}</code>	$2n\text{Na} + n\text{Cl}_2 \rightarrow 2n\text{NaCl}$

The space that is inserted after a math group can be edited:

`math-space` = <dim> Default = .1667em

```
1 \ch{$2n$ Na + $n$ Cl2 -> $2n$ NaCl} \\
2 \chemsetup[chemformula]{math-space=.2em}
3 \ch{$2n$ Na + $n$ Cl2 -> $2n$ NaCl} \\
4 \ch{$A->B$}
```

$2n\text{Na} + n\text{Cl}_2 \rightarrow 2n\text{NaCl}$
 $2n\text{Na} + n\text{Cl}_2 \rightarrow 2n\text{NaCl}$
 $A- > B$

28. Arrows

28.1. Arrow types

Arrows are input in the same intuitive way they are with mhchem. There are six different types:

`\ch{ -> }` \rightarrow standard right arrow

`\ch{ <- }` \leftarrow standard left arrow

`\ch{ <-> }` \longleftrightarrow mesomeric arrow

`\ch{ <=> }` \rightleftharpoons equilibrium arrow

`\ch{ <=>> }` \rightleftharpoons unbalanced equilibrium arrow to the right

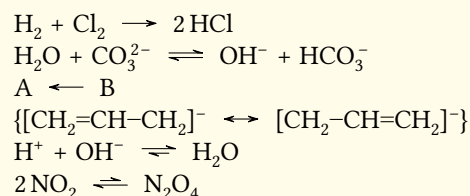
`\ch{ <<=> }` \rightleftharpoons unbalanced equilibrium arrow to the left

All these arrows are drawn with **Ti \boldsymbol{k} Z**.

```

1 \ch{H2 + Cl2 -> 2 HCl} \\
2 \ch{H2O + CO3^2- <=> OH- + HCO3-} \\
3 \ch{A <- B} \\
4 \ch{\{[CH2=CH-CH2]- <-> [CH2-CH=CH2]- \}} \\
5 \ch{H+ + OH- <=> H2O} \\
6 \ch{2 NO2 <=> N2O4}

```



28.2. Labels

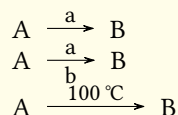
The arrows take two optional arguments to label them.

`\ch{ ->[<above>][<below>] }`

```

1 \ch{A ->[a] B} \\
2 \ch{A ->[a][b] B} \\
3 \ch{A ->[\SI{100}{\celsius}] B}

```

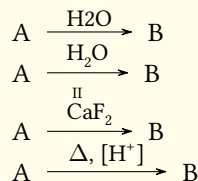


The label text can be parsed separately from the arrow. The recipe is easy: leave blanks.

```

1 \ch{A ->[H2O] B} \\
2 \ch{A ->[ H2O ] B} \\
3 \ch{A ->[ "\ox{2,Ca}" F2 ] B} \\
4 \ch{A ->[\Delta$,~ [H+ ] ] B}

```

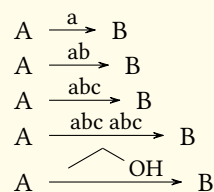


If you leave the blanks `CHEMFORMULA` treats the groups inside the square brackets as separated input types. The arrow reads its arguments *afterwards*. As you can see the arrows “grow” with the length of the labels. What stays constant is the part that protrudes the labels.

```

1 \ch{A ->[a] B} \\
2 \ch{A ->[ab] B} \\
3 \ch{A ->[abc] B} \\
4 \ch{A ->[abc~abc] B} \\
5 % needs the 'chemfig' package:
6 \setatomsep{15pt}
7 \ch{A ->[ "\chemfig{-[:30]-[: -30]OH}" ] B}

```



28.3. Customization

These are the options which enable you to customize the arrows:

arrow-offset = <dim> This is the length that an arrow protrudes a label on both sides. This means an empty arrow's length is two times arrow-offset. Default = 1.5ex

arrow-yshift = <dim> Shifts an arrow up (positive value) or down (negative value). Default = 0pt

arrow-ratio = <factor> The ratio of the arrow lengths of the unbalanced equilibrium. .4 would mean that the length of the shorter arrow is 0.4× the length of the longer arrow. Default = .6

arrow-tips = <type> This specifies the arrow tips. In principle all of *TikZ*' arrows could be used. However, this option requires <type> to be an arrow tip itself *and* to have a left <type> and right <type> to be defined. This means if you don't define any arrow tips yourself you can only choose between cf and to. Default = cf

arrow-style = <tikz> *TikZ* keys for free customization of the arrows.

compound-sep = <dim> The space between compounds and the arrows. Default = 1ex

label-offset = <dim> The space between the labels and the arrows. Default = 2pt

label-style = The relative font size of the labels. Default = \footnotesize

The following code shows the effect of the different options on the <=>> arrow:

```

1 standard: \ch{A <=>>[x][y] B} \\
2 longer: \ch[arrow-offset=12pt]{A <=>>[x][y] B} \\
3 higher: \ch[arrow-yshift=2pt]{A <=>>[x][y] B} \\
4 more balanced: \ch[arrow-ratio=.8]{A <=>>[x][y] B} \\
5 other arrow tips: \ch[arrow-tips=to]{A <=>>[x][y] B} \\
6 labels further away: \ch[label-offset=4pt]{A <=>>[x][y] B} \\
7 larger distance to compounds: \ch[compound-sep=2ex]{A <=>>[x][y] B} \\
8 other TikZ: \ch[arrow-style={thick,red}]{A <=>>[x][y] B} \\
9 smaller labels: \ch[label-style=\tiny]{A <=>>[x][y] B}

```

standard: $A \xrightleftharpoons[x]{x} B$
 longer: $A \xrightleftharpoons[y]{x} B$
 higher: $A \xrightleftharpoons[y]{x} B$
 more balanced: $A \xrightleftharpoons[y]{x} B$
 other arrow tips: $A \xrightleftharpoons[y]{x} B$
 labels further away: $A \xrightleftharpoons[y]{x} B$
 larger distance to compounds: $A \xrightleftharpoons[y]{x} B$
 other TikZ: $A \xrightleftharpoons[y]{x} B$
 smaller labels: $A \xrightleftharpoons[y]{x} B$

If you define own `pgfarrows`³⁷ or the suiting aliases so that you have the arrow tips `<type>`, left `<type>` and right `<type>` available you can use them with the option `arrow-tips`:

```

1 % 'left hook' and 'right hook' are already defined in the tikzlibrary 'arrows'
2 \pgfarrowsdeclarealias{hook}{hook}{hooks}{hooks}%
3 % 'left hook reversed' and 'right hook reversed' are already defined in the
  tikzlibrary 'arrows'
4 \pgfarrowsdeclarealias{hook reversed}{hook reversed}{hooks reversed}{hooks reversed}%
5 \ch[arrow-tips=hook]{A -> B <=> C} \
6 \ch[arrow-tips=hook reversed]{A -> B <=> C}

A \rightarrow B \rightleftharpoons C
A \rightarrow B \rightleftharpoons C

```

29. Names

29.1. Syntax

`CHEMFORMULA` has a built-in syntax to write text under a compound. In a way it works very similar to the arrows.

`\ch{ !(<name>)(<formula>) }`

If an exclamation mark is followed by a pair of parentheses `CHEMFORMULA` will parse it this way:

```

1 \ch{!(ethanol)( CH2CH2OH )}

```

CH₂CH₂OH
ethanol

The same what's true for the arrows arguments holds for these arguments: if you leave blanks the different parts will be treated according to their input type before the text is set below the formula.

³⁷ See the `pgfmanual` for details.

H ₂ O	H ₂ O	H ₂ O	H ₂ O	H ₃ O ⁺
water	water	2n - 1	H ₂ O	oxonium

```
1 \ch{H2O~(!)} \ \ H2O (!)
2 \ch{A!{}}{()} A!()
```

name-width = <dim>/auto The width of the box where the label is put into. auto will detect the width of the name and set the box to this width. Default = auto

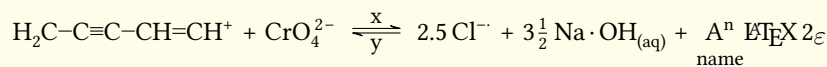
$$\begin{array}{lcl} \text{H}_2\text{SO}_4 & \longrightarrow & \text{B} \\ \text{acid} & & \\ \text{H}_2\text{SO}_4 & \longrightarrow & \text{B} \\ \text{acid} & & \\ \text{H}_2\text{SO}_4 & \longrightarrow & \text{B} \\ \text{N: acid} & & \\ \text{H}_2\text{SO}_4 & \longrightarrow & \text{B} \\ \text{acid} & & \end{array}$$

45

```

1 \newcommand*\sample{\ch{H2C-C≡C-CH=CH+ + CrO4^2- <=>[x][y] 2.5 Cl^- + 3_1/2 Na*OH
  _{(aq)}} + !(name)( A^n ) "\LaTeXe"}}
2 \sample

```

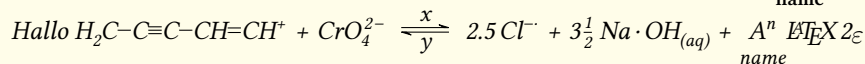
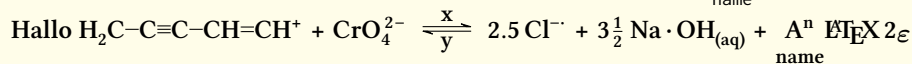
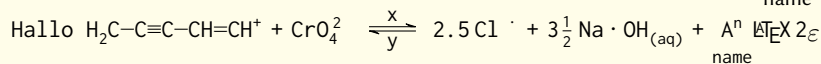
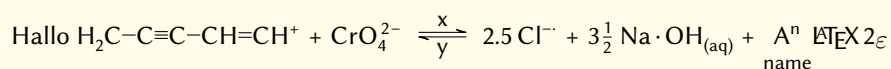


Now we're going to change different aspects of the font a look what happens:

```

1 \sffamily Hallo \sample \
2 \ttfamily Hallo \sample \normalfont \
3 \bfseries Hallo \sample \normalfont \
4 \itshape Hallo \sample

```



As you can see most features adapt to the surrounding font.

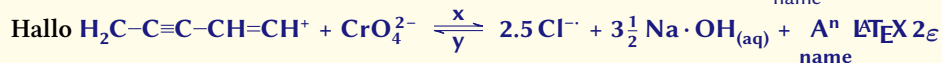
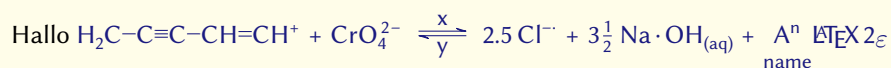
If you want to change the default format you need to use this option:

format = <anything> Adds arbitrary code before the output of **\ch**.

```

1 \definecolor{newblue}{rgb}{.1,.1,.5}\chemsetup[chemformula]{format=\color{newblue}\
  sffamily}
2 \sffamily Hallo \sample \
3 \ttfamily Hallo \sample \normalfont \
4 \bfseries Hallo \sample \normalfont \
5 \itshape Hallo \sample

```



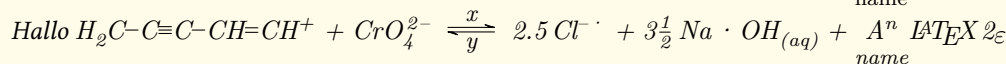
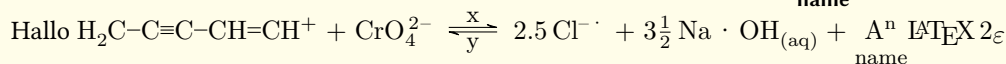
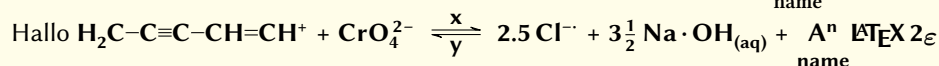
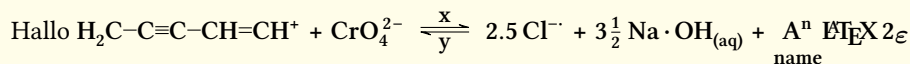
You can also specifically change the fontfamily, fontseries and fontshape of the output.

font-family = <family> Changes the fontfamily of the output with **\fontfamily{<family>}\selectfont**.

font-series = <series> Changes the fontseries of the output with `\fontseries{<series>}\selectfont`.

font-shape = <shape> Changes the fontshape of the output with `\fontshape{<shape>}\selectfont`

```
1 \chemsetup[chemformula]{font-series=bx}
2 Hallo \sample \
3 \sffamily Hallo \sample \normalfont \
4 \chemsetup[chemformula]{font-family=lmr,font-series=m} Hallo \sample \normalfont \
5 \itshape Hallo \sample
```

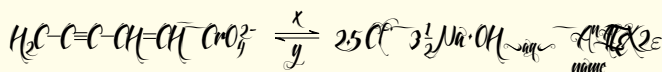
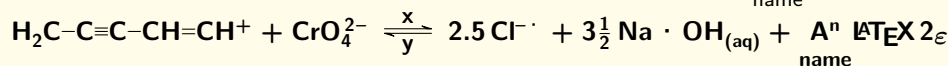
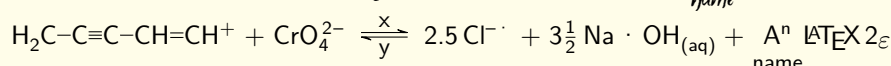
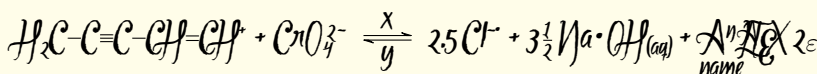
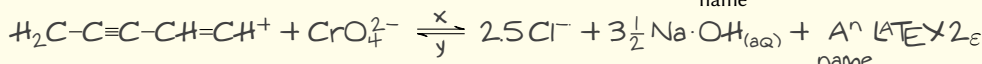
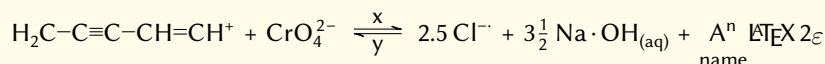


If you're using \LaTeX or Lua\LaTeX and have loaded `fontspec`³⁸ you have the possibility to set the font with it:

font-spec = {} or with options

font-spec = {[<options>]}

```
1 \chemsetup[chemformula]{font-spec={Linux Biolinum O}} \sample \
2 \chemsetup[chemformula]{font-spec={[Color=darkgray]Augie}} \sample \
3 \chemsetup[chemformula]{font-spec={Tipbrush Script}} \sample \
4 \chemsetup[chemformula]{font-spec={Latin Modern Sans}} \sample \
5 \bfseries \sample \normalfont \
6 \chemsetup[chemformula]{font-spec={Feathergraphy Decoration}} \sample
```



³⁸ CTAN: [fontspec](#)

31. Usage In Math Equations

The `\ch` command can be used inside math equations. It recognizes `\\` and `&` and passes them on. However, you can't use the optional arguments of `\\` inside `\ch`.

```

1 \begin{align}
2 \ch[arrow-tips=to]{
3   H2O & \rightarrow[a] H2SO4 \\
4   Cl2 & \rightarrow[x][y] CH4
5 }
6 \end{align}
7 \begin{align*}
8 \ch{
9   RNO2 & \rightleftharpoons[+ e^-] RNO2^{\cdot-} \\
10  RNO2^{\cdot-} & \rightleftharpoons[+ e^-] RNO2^{2-}
11 }
12 \end{align*}

```

$$\begin{array}{rcl}
 \text{H}_2\text{O} & \xrightarrow{\text{a}} & \text{H}_2\text{SO}_4 \\
 \text{Cl}_2 & \xrightarrow[\text{y}]{\text{x}} & \text{CH}_4
 \end{array}
 \begin{array}{l}
 (1) \\
 (2)
 \end{array}$$

$$\begin{array}{rcl}
 \text{RNO}_2 & \rightleftharpoons^{+ e^-} & \text{RNO}_2^{\cdot-} \\
 \text{RNO}_2^{\cdot-} & \rightleftharpoons^{+ e^-} & \text{RNO}_2^{2-}
 \end{array}$$

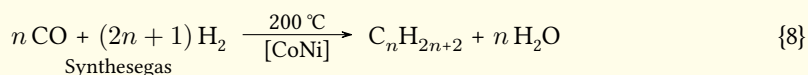
32. Further Examples

This section just presents some examples of a possible usage.

```

1 \begin{reaction}[Synthese von Alkanen]
2   !(Synthesegas)( $n$ CO + $(2n+1)$ H2 ) ->[\SI{200}{\celsius}][ [ CoNi ] ] C_{n$}H
   _{$2n+2$} + $n$ H2O
3 \end{reaction}

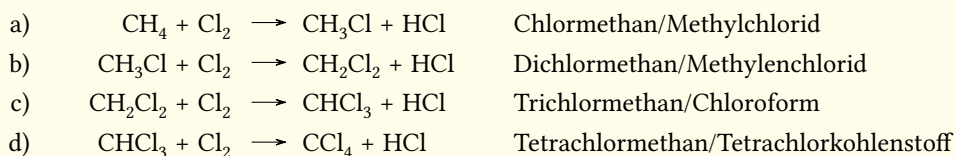
```



```

1 \begin{reactions*}
2 "a)" & \& CH4 + Cl2 & \rightarrow CH3Cl + HCl & \& "{\small Chlormethan/Methylchlorid}" \\
3 "b)" & \& CH3Cl + Cl2 & \rightarrow CH2Cl2 + HCl & \& "{\small Dichlormethan/Methylenchlorid}" \\
4 "c)" & \& CH2Cl2 + Cl2 & \rightarrow CHCl3 + HCl & \& "{\small Trichlormethan/Chloroform}" \\
5 "d)" & \& CHCl3 + Cl2 & \rightarrow CCl4 + HCl & \& "{\small Tetrachlormethan/} \\
   & & & & & Tetrachlorkohlenstoff}"
6 \end{reactions*}

```

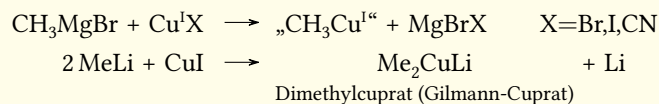


$$\begin{array}{l} \delta^- - \delta^+ \\ \text{C-M} \quad \quad \quad (\leftrightarrow) \quad \delta^+ - \delta^- \\ \text{C-Br} \\ \text{X} \cdot + \cdot \text{Y} \rightleftharpoons \text{X-Y} + \text{Bindungsenergie} \\ \text{CH}_{4(g)} + 2 \text{O}_{2(g)} \longrightarrow \text{CO}_{2(g)} + \text{H}_2\text{O}_{(l)} + \Delta H^\ominus \\ \Delta_f H^\ominus \quad \quad \quad \text{??} \quad \quad \quad 0 \quad \quad \quad -94.3 \quad -57.9 \quad -192.1 \end{array}$$

```

1 \begin{reactions*}
2 CH3MgBr + "\ox*{1,Cu}" X &-> "\glqq" CH3 "\ox*{1,Cu}\grqq" + MgBrX "\quad X\sim=\$~Br
   ,I,CN" \\\
3 2 MeLi + CuI &-> !(Dimethylcuprat~(Gilmann-Cuprat))( Me2CuLi ) + Li
4 \end{reactions*}

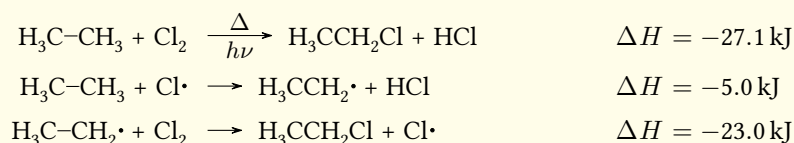
```



```

1 % needs 'chemfig'
2 \begin{reactions*}
3   H3C-CH3 + Cl2 &->[\Delta][h\nu] H3CCH2Cl + HCl
4   && \& "Enthalpy{-27.1}" \\
5   H3C-CH3 + "\Lewis{0.,Cl}" &-> H3CCH2 "\Lewis{0.,\vphantom{H}}" + HCl
6   && \& "Enthalpy{-5.0}" \\
7   H3C-CH2 "\Lewis{0.,\vphantom{H}}" + Cl2 &-> H3CCH2Cl + "\Lewis{0.,Cl}"
8   && \& "Enthalpy{-23.0}"
9 \end{reactions*}

```



Part IV.

ghsystem

33. Setup

All of `GHSYSTEM`'s options belong to the module `ghsystem`. This means they can be setup with

```
1 \chemsetup[ghsystem]{<options>} or
2 \chemsetup{ghsystem/<option1>,ghsystem/<option2>}
```

34. Get Hazard and Precautionary Statements

34.1. Simple Statements

The general usage is simple: you use the command

```
\ghs[<options>]{<type><number>
```

```
\ghs*[<options>]{<type><number>
```

There are three types available: h, euh and p. The {<type>} argument is case insensitive, so just type them in as you like.

1	<code>\ghs{h}{200} \\\</code>	H200: Unstable explosives.
2	<code>\ghs{H}{224} \\\</code>	H224: Extremely flammable liquid and vapour.
3	<code>\ghs{euh}{001} \\\</code>	EUH001: Explosive when dry.
4	<code>\ghs{Euh}{202} \\\</code>	EUH202: Cyanoacrylate. Danger. Bonds skin and eyes in seconds. Keep out of the reach of children.
5	<code>\ghs{p}{201}</code>	P201: Obtain special instructions before use.

The starred version hides the identifier and only gives the statement. If you want to hide the statement itself instead you can use the option:

```
hide = true/false
```

There is an option to customize the output, too.

```
space = <space command> Space between <type> and <number>.
```

1	<code>\ghs{h}{200} \\\</code>	H200: Unstable explosives.
2	<code>\ghs[space=\\]{h}{200} \\\</code>	H 200: Unstable explosives.
3	<code>\ghs*{h}{200} \\\</code>	Unstable explosives.
4	<code>\ghs[hide]{h}{200}</code>	H200

34.2. Statements with Placeholders

Some of the statements contain placeholders. They can be one of the following:

- *<state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard>*
- *<state specific effect if known>*
- *<or state all organs affected, if known>*
- *<name of sensitising substance>*

Except the last one which needs to be filled in, they are hidden per default. They can be made visible with the option

`fill-in = true/false` Show placeholders. Default = false

```
1 \ghs{h}{340} \\
2 \ghs[fill-in]{h}{340} \\
3 \ghs{h}{360} \\
4 \ghs[fill-in]{h}{360} \\
5 \ghs{h}{370} \\
6 \ghs[fill-in]{h}{370} \\
7 \ghs{euh}{208} \\
8 \ghs[fill-in]{euh}{208}
```

H340: May cause genetic defects.

H340: May cause genetic defects. *<state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard>*

H360: May damage fertility or the unborn child.

H360: May damage fertility or the unborn child. *<state specific effect if known> <state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard>*

H370: Causes damage to organs.

H370: Causes damage to organs *<or state all organs affected, if known>. <state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard>*

EUH208: Contains *<name of sensitising substance>*. May produce an allergic reaction.

EUH208: Contains *<name of sensitising substance>*. May produce an allergic reaction.

These placeholders can be replaced with one of these options:

`exposure` = *<text>* exposure placeholder

`effect` = *<text>* effect placeholder

`organs` = *<text>* organ placeholder

`substance` = *<text>* substance placeholder

```

1 \ghs[exposure=This is how you get exposed.]{h}{340} \\
2 \ghs[effect=These are the effects.]{h}{360} \\
3 \ghs[organs=to this organ]{h}{370} \\
4 \ghs[substance=substance]{euh}{208}

```

H340: May cause genetic defects. This is how you get exposed.
H360: May damage fertility or the unborn child. These are the effects.
H370: Causes damage to this organ.
EUH208: Contains substance. May produce an allergic reaction.

34.3. Statements with Gaps

Some of the statements have gaps that can be filled.

```

1 \ghs{p}{301} \\
2 \ghs{p}{401} \\
3 \ghs{p}{411} \\
4 \ghs{p}{413}

```

P301: IF SWALLOWED:
P401: Store ...
P411: Store at temperatures not exceeding °C/°F.
P413: Store bulk masses greater than kg/lbs at temperatures not exceeding °C/°F.

These gaps can be filled using these options:

text = <text>

dots = <text>

C-temperature = <num>

F-temperature = <num>

kg-mass = <num>

lbs-mass = <num>

```

1 \ghs[dots=contact physician!]{p}{301} \\
2 \ghs[text=here]{p}{401} \\
3 \ghs[C-temperature=50, F-temperature=122]{p}{411} \\
4 \ghs[kg-mass=5.0, lbs-mass=11, C-temperature=50, F-temperature=122]{p}{413}

```

P301: IF SWALLOWED:
P401: Store ...
P411: Store at temperatures not exceeding 50 °C/122 °F.
P413: Store bulk masses greater than 5.0 kg/11 lbs at temperatures not exceeding 50 °C/122 °F.

34.4. Combined Statements

There are some combinations of statements. They are input with a + between the numbers:

```
1 \ghs{p}{235+410} \\  
2 \ghs{p}{301+330+331}
```

P235 + P410: Keep cool. Protect from sunlight.

P301 + P330 + P331: IF SWALLOWED: rinse mouth. Do NOT induce vomiting.

Note that you can only get combinations that officially exist. *You can't combine freely.*

35. Pictograms

35.1. The Pictures

The GHS defines a number of pictograms:



The command

```
\ghspic[<options>]{<name>}
```

loads them. Table 2 shows all available pictograms and their names. To be more precise: it shows the names to use with the `\ghspic` command. The file names are `ghsystem_<name>.<filetype>` where `<filetype>` is `eps`, `jpg` or `png`, see also section 35.2.

```
1 \ghspic{skull}
```



If you don't like the default size you can change it using this option:

`scale` = <factor> Scales the pictogram. Default = 1

The pictures are actually quite large. The default setting scales them by a factor of $\frac{1}{20}$.

```
1 \ghspic[scale=2]{skull}
```



If you want to use some specific `\includegraphics` options, e. g. if you want to rotate the pictogram for some reason, use this option:












`includegraphics` = {<includegraphics keyvals>}

```
1 \ghspic[includegraphics={angle=90}]{skull}
```



Table 2: All available GHS pictograms.

name	pictogram	name	pictogram
explos		explos-1	
explos-2		explos-3	
explos-4		explos-5	
explos-6			
flame		flame-2-white	
flame-2-black		flame-3-white	
flame-3-black		flame-4-1	
flame-4-2		flame-4-3-white	
flame-4-3-black		flame-5-2-white	
flame-5-2-black			
flame-0		flame-0-5-1	

name	pictogram	name	pictogram
bottle		bottle-2-black	
bottle-2-white			
acid		acid-8	
skull		skull-2	
skull-6			
exclam			
health			
aqpol			

35.2. Picture Type Depending on Engine

As you probably know you can't use every picture type with every compiler engine. $\text{\texttt{PDF}\TeX}$ in DVI mode *needs* eps pictures while $\text{\texttt{PDF}\TeX}$ in PDF mode, $\text{\texttt{X}\TeX}$ and $\text{\texttt{Lua}\TeX}$ convert eps pictures into pdf files, given they have the rights to write in the directory the pictures are saved in.

However, the latter can include jpg and png without any problems, while $\text{\texttt{PDF}\TeX}$ in DVI mode can't.

To resolve this [GHSYSTEM](#) tests which engine is used and if $\text{\texttt{PDF}\TeX}$ which mode is used and then chooses either eps or png for the pictograms. You are free to choose the picture type yourself with the option

`pic-type = eps/jpg/png`

36. Available Languages

Right now the H and P statements are only available in English or German. The package adapts the package option `german` but does not (yet) recognize language settings by [babel](#)³⁹ or [polyglossia](#)⁴⁰.

You can also choose the language explicitly.

³⁹ CTAN: [babel](#) ⁴⁰ CTAN: [polyglossia](#)

language = english/german

1	<code>\ghs{h}{201}</code>	H201: Explosive; mass explosion hazard.
2		
3	<code>\chemsetup[ghsystem]{language=german}</code>	H201: Explosiv, Gefahr der Massenexplosion.
4	<code>\ghs{h}{201}</code>	

I will add other languages some time in future. This may take a while, though. If you would be willing to contribute and write the statements of another language please feel free to contact me⁴¹. I would provide you with a template file, a pdf containing the official translations, and help to all your questions.

37. List of All Statements

If for some reason you want to list all sentences you can use

`\ghslistall[<options>]`

This command has a number of options to customize the table, which is created with the `longtable` environment of the `longtable` package.

`table-head-number` = <text> Default = Identifier

`table-head-text` = <text> Default = Statement

`table-next-page` = <text> Default = continues on next page

`table-caption` = <text> As <text> in `\caption{<text>}`. Default = All H, EUH, and P Statements.

`table-caption-short` = <text> As <short> in `\caption[<short>]{<text>}`.

`table-label` = <text> The label to refer to the table with `\ref` and similar commands. Default = `tab:ghs-hp-statements`

`table-row-sep` = <dim> The separation of the table rows. A $\text{T}_{\text{E}}\text{X}$ dimension. Default = 3pt

`table-rules` = `default/booktabs/none` The style of the horizontal rules in the table. `default` uses `\hline`, `booktabs` uses `\toprule`, `\midrule` or `\bottomrule`, resp. This option needs the `booktabs`⁴² package which you have to load yourself then. Default = `default`

`table-top-head-rule` = `default/booktabs/none` Change top rule explicitly. Default = `default`

`table-head-rule` = `default/booktabs/none` Change rule below head explicitly. Default = `default`

`table-foot-rule` = `default/booktabs/none` Change foot rule explicitly. Default = `default`

⁴¹ contact@mychemistry.eu ⁴² CTAN: `booktabs`

`table-last-foot-rule = default/booktabs/none` Change last foot rule explicitly. Default = default

The code below shows how Table 3 was created:

```
1 \ghslistall[fill-in,table-rules=booktabs]
```

Table 3: All H, EUH, and P Statements.

Identifier	Statement
H200	Unstable explosives.
H201	Explosive; mass explosion hazard.
H202	Explosive, severe projection hazard.
H203	Explosive; fire, blast or projection hazard.
H204	Fire or projection hazard.
H205	May mass explode in fire.
H220	Extremely flammable gas.
H221	Flammable gas.
H222	Extremely flammable aerosol.
H223	Flammable aerosol.
H224	Extremely flammable liquid and vapour.
H225	Highly flammable liquid and vapour.
H226	Flammable liquid and vapour.
H228	Flammable solid.
H240	Heating may cause an explosion.
H241	Heating may cause a fire or explosion.
H242	Heating may cause a fire.
H250	Catches fire spontaneously if exposed to air.
H251	Self-heating; may catch fire.
H252	Self-heating in large quantities; may catch fire.
H260	In contact with water releases flammable gases which may ignite spontaneously.
H261	In contact with water releases flammable gases.
H270	May cause or intensify fire; oxidiser.
H271	May cause fire or explosion; strong oxidiser.
H272	May intensify fire; oxidiser.
H280	Contains gas under pressure; may explode if heated.

continues on next page

Identifier	Statement
H281	Contains refrigerated gas; may cause cryogenic burns or injury.
H290	May be corrosive to metals.
H300	Fatal if swallowed.
H301	Toxic if swallowed.
H302	Harmful if swallowed.
H304	May be fatal if swallowed and enters airways.
H310	Fatal in contact with skin.
H311	Toxic in contact with skin.
H312	Harmful in contact with skin.
H314	Causes severe skin burns and eye damage.
H315	Causes skin irritation.
H317	May cause an allergic skin reaction.
H318	Causes serious eye damage.
H319	Causes serious eye irritation.
H330	Fatal if inhaled.
H331	Toxic if inhaled.
H332	Harmful if inhaled.
H334	May cause allergy or asthma symptoms or breathing difficulties if inhaled.
H335	May cause respiratory irritation.
H336	May cause drowsiness or dizziness.
H340	May cause genetic defects. <i><state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H341	Suspected of causing genetic defects. <i><state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H350	May cause cancer. <i><state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H351	Suspected of causing cancer. <i><state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H360	May damage fertility or the unborn child. <i><state specific effect if known> <state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H361	Suspected of damaging fertility or the unborn child. <i><state specific effect if known> <state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H362	May cause harm to breast-fed children.

continues on next page

Identifier	Statement
H370	Causes damage to organs <i><or state all organs affected, if known>. <state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H371	May cause damage to organs <i><or state all organs affected, if known>. <state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H372	Causes damage to organs <i><or state all organs affected, if known> through prolonged or repeated exposure. <state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H373	May cause damage to organs <i><or state all organs affected, if known> through prolonged or repeated exposure. <state route of exposure if it is conclusively proven that no other routes of exposure cause the hazard></i>
H400	Very toxic to aquatic life.
H410	Very toxic to aquatic life with long lasting effects.
H411	Toxic to aquatic life with long lasting effects.
H412	Harmful to aquatic life with long lasting effects.
H413	May cause long lasting harmful effects to aquatic life.
H350i	May cause cancer by inhalation.
H360F	May damage fertility.
H360D	May damage the unborn child.
H361f	Suspected of damaging fertility.
H361d	Suspected of damaging the unborn child.
H360FD	May damage fertility. May damage the unborn child.
H361fd	Suspected of damaging fertility. Suspected of damaging the unborn child.
H360Fd	May damage fertility. Suspected of damaging the unborn child.
H360Df	May damage the unborn child. Suspected of damaging fertility.
EUH001	Explosive when dry.
EUH006	Explosive with or without contact with air.
EUH014	Reacts violently with water.
EUH018	In use may form flammable/explosive vapour-air mixture.
EUH019	May form explosive peroxides.
EUH044	Risk of explosion if heated under confinement.
EUH029	Contact with water liberates toxic gas.
EUH031	Contact with acids liberates toxic gas.
EUH032	Contact with acids liberates very toxic gas.

continues on next page

Identifier	Statement
EUH066	Repeated exposure may cause skin dryness or cracking.
EUH070	Toxic by eye contact.
EUH071	Corrosive to the respiratory tract.
EUH059	Hazardous to the ozone layer.
EUH201	Contains lead. Should not be used on surfaces liable to be chewed or sucked by children.
EUH201A	Warning! contains lead.
EUH202	Cyanoacrylate. Danger. Bonds skin and eyes in seconds. Keep out of the reach of children.
EUH203	Contains chromium(VI). May produce an allergic reaction.
EUH204	Contains isocyanates. May produce an allergic reaction.
EUH205	Contains epoxy constituents. May produce an allergic reaction.
EUH206	Warning! Do not use together with other products. May release dangerous gases (chlorine).
EUH207	Warning! Contains cadmium. Dangerous fumes are formed during use. See information supplied by the manufacturer. Comply with the safety instructions.
EUH208	Contains <name of sensitising substance>. May produce an allergic reaction.
EUH209	Can become highly flammable in use.
EUH209A	Can become flammable in use.
EUH210	Safety data sheet available on request.
EUH401	To avoid risks to human health and the environment, comply with the instructions for use.
P101	If medical advice is needed, have product container or label at hand.
P102	Keep out of reach of children.
P103	Read label before use.
P201	Obtain special instructions before use.
P202	Do not handle until all safety precautions have been read and understood.
P210	Keep away from heat/sparks/open flames/hot surfaces. — No smoking.
P211	Do not spray on an open flame or other ignition source.
P220	Keep/Store away from clothing/.../combustible materials.
P221	Take any precaution to avoid mixing with combustibles ...
P222	Do not allow contact with air.
P223	Keep away from any possible contact with water, because of violent reaction and possible flash fire.

continues on next page

Identifier	Statement
P230	Keep wetted with ...
P231	Handle under inert gas.
P232	Protect from moisture.
P233	Keep container tightly closed.
P234	Keep only in original container.
P235	Keep cool.
P240	Ground/bond container and receiving equipment.
P241	Use explosion-proof electrical/ventilating/lighting/... equipment.
P242	Use only non-sparking tools.
P243	Take precautionary measures against static discharge.
P244	Keep reduction valves free from grease and oil.
P250	Do not subject to grinding/shock/.../friction.
P251	Pressurized container: Do not pierce or burn, even after use.
P260	Do not breathe dust/fume/gas/mist/vapours/spray.
P261	Avoid breathing dust/fume/gas/mist/vapours/spray.
P262	Do not get in eyes, on skin, or on clothing.
P263	Avoid contact during pregnancy/while nursing.
P264	Wash ... thoroughly after handling.
P270	Do not eat, drink or smoke when using this product.
P271	Use only outdoors or in a well-ventilated area.
P272	Contaminated work clothing should not be allowed out of the workplace.
P273	Avoid release to the environment.
P280	Wear protective gloves/protective clothing/eye protection/face protection.
P281	Use personal protective equipment as required.
P282	Wear cold insulating gloves/face shield/eye protection.
P283	Wear fire/flammable resistant/retardant clothing.
P284	Wear respiratory protection.
P285	In case of inadequate ventilation wear respiratory protection.
P231 + P231	Handle under inert gas. Protect from moisture.
P235 + P410	Keep cool. Protect from sunlight.
P301	IF SWALLOWED:
P302	IF ON SKIN:
P303	IF ON SKIN (or hair):
P304	IF INHALED:

continues on next page

Identifier	Statement
P305	IF IN EYES:
P306	IF ON CLOTHING:
P307	IF exposed:
P308	IF exposed or concerned:
P309	IF exposed or if you feel unwell:
P310	Immediately call a POISON CENTER or doctor/physician.
P311	Call a POISON CENTER or doctor/physician.
P312	Call a POISON CENTER or doctor/physician if you feel unwell.
P313	Get medical advice/attention.
P314	Get medical advice/attention if you feel unwell.
P315	Get immediate medical advice/attention.
P320	Specific treatment is urgent (see ... on this label).
P321	Specific treatment (see ... on this label).
P322	Specific measures (see ... on this label).
P330	Rinse mouth.
P331	Do NOT induce vomiting.
P332	If skin irritation occurs:
P333	If skin irritation or rash occurs:
P334	Immerse in cool water/wrap in wet bandages.
P335	Brush off loose particles from skin.
P336	Thaw frosted parts with lukewarm water. Do not rub affected area.
P337	If eye irritation persists:
P338	Remove contact lenses, if present and easy to do. Continue rinsing.
P340	Remove victim to fresh air and keep at rest in a position comfortable for breathing.
P341	If breathing is difficult, remove victim to fresh air and keep at rest in a position comfortable for breathing.
P342	If experiencing respiratory symptoms:
P350	Gently wash with plenty of soap and water.
P351	Rinse cautiously with water for several minutes.
P352	Wash with plenty of soap and water.
P353	Rinse skin with water/shower.
P360	Rinse immediately contaminated clothing and skin with plenty of water before removing clothes.
P361	Remove/Take off immediately all contaminated clothing.
P362	Take off contaminated clothing and wash before reuse.

continues on next page

Identifier	Statement
P363	Wash contaminated clothing before reuse.
P370	In case of fire:
P371	In case of major fire and large quantities:
P372	Explosion risk in case of fire.
P373	DO NOT fight fire when fire reaches explosives.
P374	Fight fire with normal precautions from a reasonable distance.
P375	Fight fire remotely due to the risk of explosion.
P376	Stop leak if safe to do so.
P377	Leaking gas fire: Do not extinguish, unless leak can be stopped safely.
P378	Use ... for extinction.
P380	Evacuate area.
P381	Eliminate all ignition sources if safe to do so.
P390	Absorb spillage to prevent material damage.
P391	Collect spillage.
P301 + P310	IF SWALLOWED: Immediately call a POISON CENTER or doctor/physician.
P301 + P312	IF SWALLOWED: Call a POISON CENTER or doctor/physician if you feel unwell.
P301 + P330 + P331	IF SWALLOWED: rinse mouth. Do NOT induce vomiting.
P302 + P334	IF ON SKIN: Immerse in cool water/wrap in wet bandages.
P302 + P350	IF ON SKIN: Gently wash with plenty of soap and water.
P302 + P352	IF ON SKIN: Wash with plenty of soap and water.
P303 + P361 + P353	IF ON SKIN (or hair): Remove/Take off immediately all contaminated clothing. Rinse skin with water/shower.
P304 + P340	IF INHALED: Remove victim to fresh air and keep at rest in a position comfortable for breathing.
P304 + P341	IF INHALED: If breathing is difficult, remove victim to fresh air and keep at rest in a position comfortable for breathing.
P305 + P351 + P338	IF IN EYES: Rinse cautiously with water for several minutes. Remove contact lenses, if present and easy to do. Continue rinsing.
P306 + P360	IF ON CLOTHING: Rinse immediately contaminated clothing and skin with plenty of water before removing clothes.
P307 + P311	IF exposed: Call a POISON CENTER or doctor/physician.
P308 + P313	IF exposed or concerned: Get medical advice/attention.
P309 + P311	IF exposed or if you feel unwell: Call a POISON CENTER or doctor/physician.

continues on next page

Identifier	Statement
P332 + P313	If skin irritation occurs: Get medical advice/attention.
P333 + P313	If skin irritation or rash occurs: Get medical advice/attention.
P335 + P334	Brush off loose particles from skin. Immerse in cool water/wrap in wet bandages.
P337 + P313	If eye irritation persists: Get medical advice/attention.
P342 + P311	If experiencing respiratory symptoms: Call a POISON CENTER or doctor/physician.
P370 + P376	In case of fire: Stop leak if safe to do so.
P370 + P378	In case of fire: Use ... for extinction.
P370 + P380	In case of fire: Evacuate area.
P370 + P380 + P375	In case of fire: Evacuate area. Fight fire remotely due to the risk of explosion.
P371 + P380 + P375	In case of major fire and large quantities: Evacuate area. Fight fire remotely due to the risk of explosion.
P401	Store ...
P402	Store in a dry place.
P403	Store in a well-ventilated place.
P404	Store in a closed container.
P405	Store locked up.
P406	Store in corrosive resistant/... container with a resistant inner liner.
P407	Maintain air gap between stacks/pallets.
P410	Protect from sunlight.
P411	Store at temperatures not exceeding °C/°F.
P412	Store at temperatures not exceeding 50 °C/122 °F.
P413	Store bulk masses greater than kg/lbs at temperatures not exceeding °C/°F.
P420	Store away from other materials.
P422	Store contents under ...
P402 + P404	Store in a dry place. Store in a closed container.
P403 + P233	Store in a well-ventilated place. Keep container tightly closed.
P403 + P235	Store in a well-ventilated place. Keep cool.
P410 + P403	Protect from sunlight. Store in a well-ventilated place.
P410 + P412	Protect from sunlight. Do not expose to temperatures exceeding 50 °C/122 °F.
P411 + P235	Store at temperatures not exceeding °C/°F . Keep cool.
P501	Dispose of contents/container to ...

continues on next page

Identifier	Statement
------------	-----------

Part V.

Appendix

Suggestions and Bug Reports

Feedback on [CHEMMACROS](#), [CHEMFORMULA](#) and [GHSYSTEM](#) is highly appreciated and welcome! If you have suggestions for macros, missing features etc., please don't hesitate to contact me. If you recognize any errors, be it chemical ones, wrong documentation and the like, I'd be grateful about a short email⁴³.

If you find any bugs, it would be best, if you'd send me a minimal example, with which I can reproduce the bug. You can also submit an issue on <https://bitbucket.org/cgnieder/chemmacros/> instead.

Many thanks to all the people, who already provided me with feedback!

⁴³ contact@mychemistry.eu

Index

Section titles are indicated **bold**, packages sans serif, commands `\brown`, keys `green` and modules (only `CHEMMACROS`) `red`.

Symbols

<code>\-</code>	8 f.
<code>\^</code>	8 f.
<code>\ </code>	8 f.

A

<code>\abinitio</code>	10
absolute configuration	10
Acid/Base	11 f.
<code>adduct-space</code>	38 f.
<code>align</code>	22
<code>angle</code>	27, 29
<code>\anti</code>	9
<code>append</code>	12
APPENDIX	65
<code>\aq</code>	26
<code>arg</code>	24
<code>arrow-offset</code>	43
<code>arrow-ratio</code>	43
<code>arrow-style</code>	43
<code>arrow-tips</code>	43 f.
<code>arrow-yshift</code>	43
Arrows	41–44
customization	43
labels	42
types	41
<code>\atm</code>	11
<code>\atmosphere</code>	11
<code>atoms</code>	27

B

<code>\ba</code>	7, 31
<code>babel</code>	55
<code>back-atoms</code>	27
<code>bm</code>	3
<code>bond-length</code>	38
<code>booktabs</code>	56
<code>bpchem</code>	3, 5
<code>bpchem</code>	3, 5, 8 f.

C

<code>C-temperature</code>	52
Cahn-Ingold-Prelog	9
<code>\cal</code>	11
<code>\calory</code>	11
<code>\centering</code>	32
<code>\ch</code>	26, 30 f., 33 f., 36, 40 ff., 44, 46, 48
<code>charge-hshift</code>	38
<code>charge-style</code>	38
charges	
<code>append</code>	12
<code>chemfig</code>	8, 14 f.
CHEMFORMULA	33–49
CHEMMACROS	7–32
<code>chemmacros</code>	37
<code>\chemsetup</code>	6, 31, 34
<code>chemstyle</code>	7, 10 f.
<code>\cip</code>	9
<code>circled</code>	5, 8
<code>circletype</code>	5
<code>\cis</code>	9
<code>cis/trans</code>	9
<code>\cmc</code>	11
<code>color</code>	29
Commands for mhchem	22 f.
<code>compound-sep</code>	43
Compounds	36–39
adducts	36
bonds	38
length	39
charges	37
shift	38
commands	37
customization	38
subscripts	36
shift	39
superscripts	37
<code>cool</code>	5, 9

D

<code>decimal-marker</code>	13, 34
<code>\DeclareParticle</code>	8
<code>\delm</code>	14

<code>\delp</code>	14
<code>delta</code>	19 f.
<code>detect-bold</code>	5
<code>\Dfi</code>	9
<code>dist</code>	17
<code>dots</code>	52
E	
<code>\E</code>	5, 9
<code>effect</code>	51
<code>\El</code>	7
<code>\el</code>	7
<code>elpair</code>	8
<code>\Ent</code>	9
<code>\Enthalpy</code>	18 f., 32
<code>\Entropy</code>	18, 32
<code>environ</code>	3
Escaped Input	40 f.
<code>math</code>	41
<code>text</code>	40
<code>expl3</code>	3
<code>explicit-sign</code>	13
<code>exponent</code>	19 f.
<code>exposure</code>	51
<code>EZ</code>	5, 9
F	
<code>F-temperature</code>	52
<code>\fdelm</code>	14
<code>\fdelp</code>	14
<code>fill-in</code>	51
<code>Fischer</code>	9
<code>\fmch</code>	12, 31
<code>\fminus</code>	7
<code>font-family</code>	46
<code>font-series</code>	47
<code>font-shape</code>	47
<code>font-spec</code>	47
<code>fontsize</code>	22
<code>fontspec</code>	47
<code>format</code>	22, 46
Format and Font	45 ff.
<code>\fpch</code>	12, 31
<code>\fplus</code>	5, 7
<code>frac-style</code>	35
<code>\fscrm</code>	15

<code>\fscrp</code>	15
<code>\fsscrm</code>	15
<code>\fsscrp</code>	15

G

<code>\gas</code>	26
<code>german</code>	5, 11, 25, 55
<code>\ghs</code>	50
<code>ghs</code>	5
<code>\ghs*</code>	50
<code>\ghslistall</code>	56
<code>\ghspic</code>	53
GHSYSTEM	50–65
combined statements	52
get statements	50
statements with gaps	52
statements with placeholders	51
<code>\Gibbs</code>	18, 32
<code>graphicx</code>	3

H

<code>half</code>	29
Hazard and Precautionary Statements	50–53
<code>\hertz</code>	32
<code>hide</code>	50
<code>\Hpl</code>	7
<code>\Ht0</code>	7
<code>\Hyd</code>	7

I

<code>ifpdf</code>	3
<code>includegraphics</code>	53
<code>\insitu</code>	10
Installation	4
Ion Charges	12 f.
<code>\iupac</code>	8 f.
IUPAC Names	8 ff.
Cahn-Ingold-Prelog	9
cis/trans	9
Fischer	9
ortho/meta/para	10
syn/anti	9
tert	9
zusammen/entgegen	9

K

<code>\Ka</code>	11
------------------	----

<code>\Kb</code>	11	<code>name-width</code>	45
<code>kg-mass</code>	52	Names	44 f.
<code>\Kw</code>	12	customization.....	45
L		syntax.....	44
<code>l3kernel</code>	3	<code>\newman</code>	27, 32
<code>l3keys2e</code>	3	newman	
<code>l3packages</code>	3	angle.....	27
<code>label-offset</code>	43	atoms.....	27
<code>label-style</code>	43	back-atoms.....	27
<code>language</code>	56	ring.....	27
Latin Phrases	10 f.	scale.....	27
<code>lbs-mass</code>	52	Newman Projections	27 f.
<code>\Lfi</code>	9	<code>\newreaction</code>	24, 32
<code>\liquid</code>	26	<code>ngerman</code>	5
<code>list-entry</code>	25	<code>nicefrac</code>	3, 35
<code>list-name</code>	25	<code>\NMR</code>	5, 21, 32
List of All Statements	56–65	nmr	
<code>\listofreactions</code>	25	nucleus.....	22
Loading the Bundle	4	unit.....	21
<code>longtable</code>	3, 56	<code>\NMR*</code>	21
<code>\lqd</code>	5, 26, 31	<code>\normal</code>	11
M		<code>\ntr</code>	7
<code>math-space</code>	41	<code>\Nu</code>	5, 7, 31
Math Environments	48	<code>Nu</code>	5, 7
<code>mathspec</code>	5, 7	<code>\Nuc</code>	7
<code>mathtools</code>	3, 24	<code>nucleus</code>	22
<code>\mch</code>	12, 31	O	
<code>\mech</code>	15 f.	<code>opacity</code>	29
<code>\mega</code>	32	option	
<code>\meta</code>	10	bpchem.....	5
<code>method</code>	3, 5, 8, 22, 30	circled.....	5
<code>mhchem</code>	3, 8, 22, 24, 33, 41	circletype.....	5
<code>\mhName</code>	22, 32	detect-bold.....	5
mhName		EZ.....	5
align.....	22	german.....	5, 11, 25, 55
fontsize.....	22	ghs.....	5
format.....	22	method.....	5, 22, 30
width.....	22	ngerman.....	5
<code>\Molar</code>	11	Nu.....	5, 7
<code>\moLar</code>	11	strict.....	5
<code>\molar</code>	11	synchronize.....	5, 31
<code>\MolMass</code>	11	version.....	5 f.
N		xspace.....	6
<code>name-format</code>	45	Option Overview (chemmacros)	31 f.
		<code>\orbital</code>	28, 32

orbital	
angle	29
color	29
half	29
opacity	29
overlay	29
phase	29
scale	29
Orbitals	28 ff.
organs	51
\ortho	10
ortho/meta/para	10
overlay	29
\OX	16, 32
\ox	13 f., 31
ox	
decimal-marker	13
explicit-sign	13
parse	13
pos	13
roman	13
\ox*	13
Oxidation Numbers	13 f.
P	
\p	12
Package Options	4 ff.
\para	10
parse	13
Partial Charges	14 f.
particle	
elpair	8
Particles, Ions and Symbols	7 f.
own	8
predefined	7
\pch	12, 31
\pH	11
phase	29
Phases	26 f.
pic-type	55
Pictograms	53 ff.
\pKa	5, 11 f.
\pKb	12
plus-space	40
\pOH	11
polyglossia	55
pos	13
PRELIMINARIES	3–6
\prt	7
R	
\R	7
\Rcip	9
\Rconf	10
\reaction	24
reaction	
list-entry	25
list-name	25
reaction (env.)	23
reaction* (env.)	23
Reaction Environments	23–26
\reactionlistname	25
Reaction Mechanisms	15 f.
reactions (env.)	23
reactions* (env.)	23
\redox	16, 32
redox	
dist	17
sep	17
Redox Reactions	16 ff.
\renewstate	20, 32
\renewtagform	24
ring	27
roman	13
S	
scale	27, 29, 53
\Scip	9
\Sconf	10
\scrm	15
\scrp	15
sep	17
\setnewstate	19 f., 32
Setup	6
siunitx	3, 11, 18 f., 21
\sld	5, 26, 31
\solid	26
space	50
Special Types	40
Spectroscopy	21 f.
\standardstate	7, 32
star	24

<code>\State</code>	20, 32
<code>state</code>	
<code>delta</code>	20
<code>exponent</code>	20
<code>subscript-left</code>	20
Stereo Descriptors and Nomenclature ..	9 f.
<code>stoich-space</code>	35
Stoichiometric Factors	34 ff.
<code>nicefrac</code>	35
<code>space</code>	35
<code>xfrac</code>	35
<code>strict</code>	5, 8
<code>subscript</code>	19 f.
<code>subscript-left</code>	20
<code>subscript-style</code>	38
<code>subscript-vshift</code>	38 f.
<code>substance</code>	51
<code>\syn</code>	9
<code>synchronize</code>	5, 31
T	
<code>table-caption</code>	56
<code>table-caption-short</code>	56
<code>table-foot-rule</code>	56
<code>table-head-number</code>	56
<code>table-head-rule</code>	56
<code>table-head-text</code>	56
<code>table-label</code>	56
<code>table-last-foot-rule</code>	57
<code>table-next-page</code>	56
<code>table-row-sep</code>	56
<code>table-rules</code>	56
<code>table-top-head-rule</code>	56
<code>tabu</code>	3
<code>tert</code>	9
<code>\tert</code>	9
<code>text</code>	52
Thermodynamics	18–21
<code>TikZ</code>	3, 7, 10, 16, 27, 29, 41, 43
<code>tikz</code>	3
<code>\tiny</code>	32
<code>\torr</code>	11
<code>\trans</code>	9
<code>\transitionstatesymbol</code>	7

U	
<code>unit</code>	19, 21
Units	11
V	
<code>version</code>	5 f.
W	
<code>\water</code>	7
<code>width</code>	22
X	
<code>xfrac</code>	3, 14, 35
<code>xparse</code>	3
<code>xspace</code>	3, 6
<code>xspace</code>	3
Z	
<code>\Z</code>	9
<code>\Zus</code>	9
<code>zusammen/entgegen</code>	9