

Covalent links in **ACEDRG**

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Molecular Biology

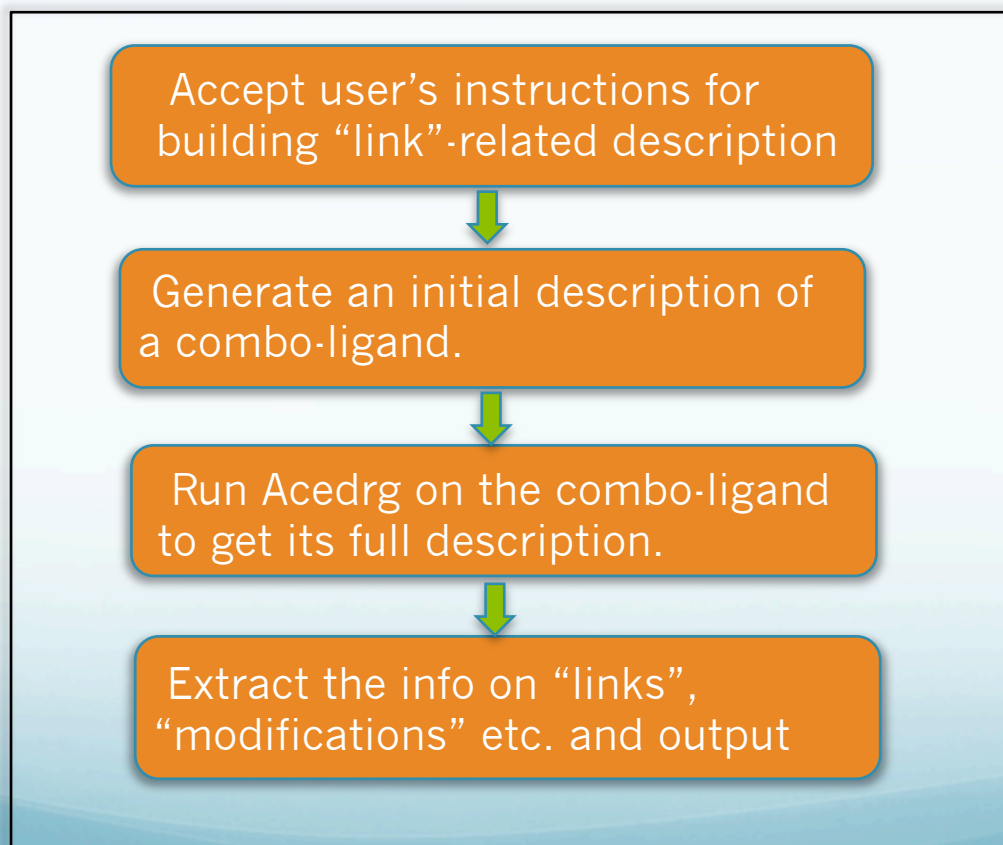
Covalent Links in Acedrg

❖ Contents

- A few cases of covalent links in Acedrg

Covalent Links in Acedrg

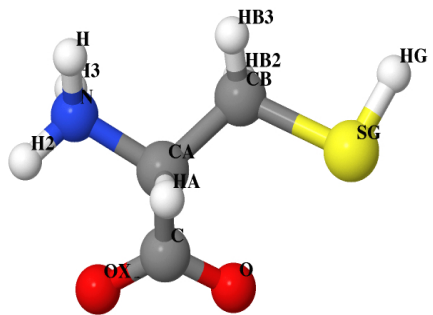
Flowchart for “link” in Acedrg



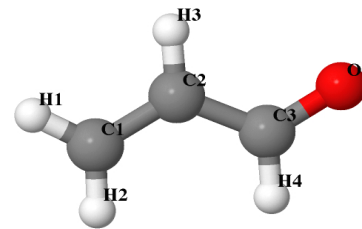
Covalent Links in Acedrg

A Few Cases on Covalent Links in Acedrg

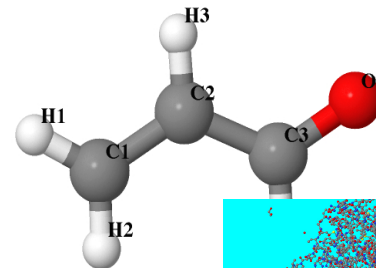
Case 1



CYS



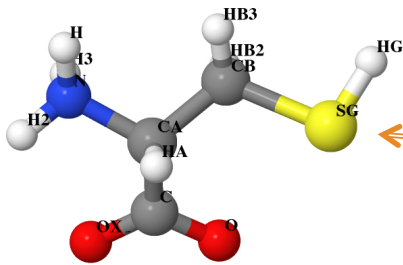
Monomer "TMP"



Real Structure

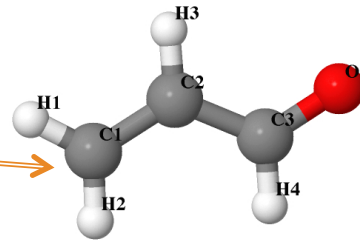
Covalent Links in Acedrg

Case 1: Scenario 1



CYS

Jmol



Monomer "TMP"

Jmol

No change in the bond between
C1 and C2 in "TMP"

Covalent Links in Acedrg

Case 1: Scenario 1

Acedrg command-line

```
acedrg -L inFiles/instruct_case1_sce1.txt -o case1_CYS_TMP_1
```

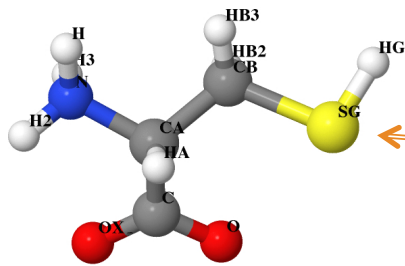
- It uses an instruction file: **instruct_case1_sce1.txt**
- It generates the dictionary file: **case1_CYS_TMP_1_link.cif**

instruct_case1_sce1.txt

```
LINK: RES-NAME-1 CYS ATOM-NAME-1 SG RES-NAME-2 TMP FILE-2 inFiles/case2_comp1.cif ATOM-NAME-2 C1
```

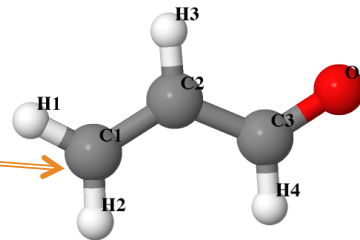
Covalent Links in Acedrg

Case 1: Scenario 1



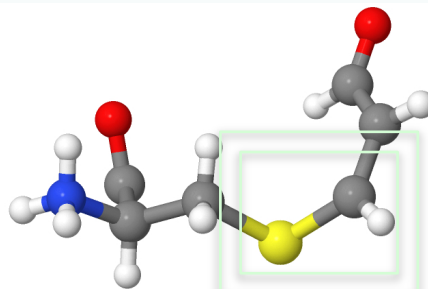
Jmol

CYS



Jmol

Monomer "TMP"



Jmol

The Linked molecule

Covalent Links in Acedrg

Case 1: Scenario 1

case1_CYS_TMP_1_link.cif

List Section

data_link_list

loop_

_chem_link.id

_chem_link.comp_id_1

_chem_link.mod_id_1

_chem_link.group_comp_1

_chem_link.comp_id_2

_chem_link.mod_id_2

_chem_link.group_comp_2

_chem_link.name

CYS-TMP	CYS	CYSmod1	L-PEPTIDE	TMP	TMPmod1
non-polymer	CYS-TMP				

Covalent Links in Acedrg

Case 1: Scenario 1

case1_CYS_TMP_1_link.cif

Component Section

data_comp_CYS

#

loop_

_chem_comp_atom.comp_id

_chem_comp_atom.atom_id

_chem_comp_atom.type_symbol

_chem_comp_atom.type_energy

_chem_comp_atom.charge

_chem_comp_atom.x

_chem_comp_atom.y

_chem_comp_atom.z

CYS	N	N	NT3	1	1.551	14.790	38.686
CYS	CA	C	CH1	0	0.616	13.682	39.054
CYS	C	C	C	0	1.049	13.053	40.388

Covalent Links in Acedrg

Case 1: Scenario 1

case1_CYS_TMP_1_link.cif

Component Section

data_comp_CYS

#

loop_

_chem_comp_atom.comp_id

_chem_comp_atom.atom_id

_chem_comp_atom.type_symbol

_chem_comp_atom.type_energy

_chem_comp_atom.charge

_chem_comp_atom.x

_chem_comp_atom.y

_chem_comp_atom.z

CYS	N	N	NT3	1	1.551	14.790	38.686
CYS	CA	C	CH1	0	0.616	13.682	39.054
CYS	C	C	C	0	1.049	13.053	40.388

Covalent Links in Acedrg

case1_CYS_TMP_1_link.cif

Modification Section

```
data_mod_TMPmod1
loop_
  _chem_mod_atom.mod_id
  _chem_mod_atom.function
  _chem_mod_atom.atom_id
  _chem_mod_atom.new_atom_id
  _chem_mod_atom.new_type_symbol
  _chem_mod_atom.new_type_energy
  _chem_mod_atom.new_charge
TMPmod1      delete      H2      .      H      H      0
TMPmod1      change      C1      .      C      C1     0
loop_
  _chem_mod_bond.mod_id
  _chem_mod_bond.function
  _chem_mod_bond.atom_id_1
  _chem_mod_bond.atom_id_2
  _chem_mod_bond.new_type
  _chem_mod_bond.new_value_dist
  _chem_mod_bond.new_value_dist_esd
TMPmod1      delete      C1      H2      single  .      .
```

Covalent Links in Acedrg

case1_CYS_TMP_1_link.cif

Link Section

data_link_CYS-TMP

loop_

_chem_link_bond.link_id

_chem_link_bond.atom_1_comp_id

_chem_link_bond.atom_id_1

_chem_link_bond.atom_2_comp_id

_chem_link_bond.atom_id_2

_chem_link_bond.type

_chem_link_bond.value_dist

_chem_link_bond.value_dist_esd

CYS-TMP	1	SG	2	C1	SINGLE	1.732	0.010
---------	---	----	---	----	--------	-------	-------

loop_

_chem_link_angle.link_id

_chem_link_angle.atom_1_comp_id

_chem_link_angle.atom_id_1

_chem_link_angle.atom_2_comp_id

_chem_link_angle.atom_id_2

_chem_link_angle.atom_3_comp_id

_chem_link_angle.atom_id_3

_chem_link_angle.value_angle

_chem_link_angle.value_angle_esd

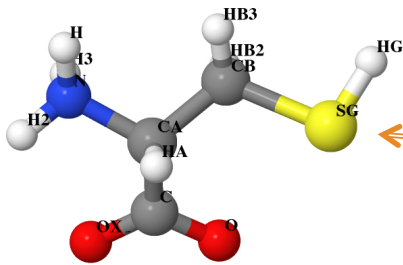
CYS-TMP	1	CB	1	SG	2	C1	100.987	2.18
---------	---	----	---	----	---	----	---------	------

CYS-TMP	2	C2	2	C1	1	SG	122.125	3.00
---------	---	----	---	----	---	----	---------	------

CYS-TMP	1	SG	2	C1	2	H1	117.563	1.50
---------	---	----	---	----	---	----	---------	------

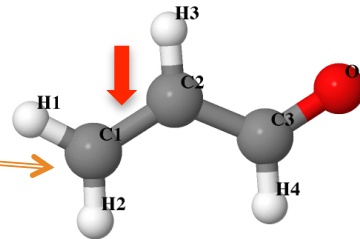
Covalent Links in Acedrg

Case 1: Scenario 1



CYS

Jmol



Monomer "TMP"

Jmol

The bond between C1 and C2 in
"TMP" becomes "single"

Covalent Links in Acedrg

Case 1: Scenario 1

Acedrg command-line

```
acedrg -L inFiles/instruct_case1_sce2.txt -o case1_CYS_TMP_2
```

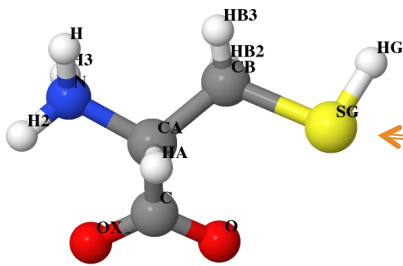
- It uses an instruction file: **instruct_case1_sce2.txt**
- It generates the dictionary file: **case1_CYS_TMP_2_link.cif**

instruct_case1_sce2.txt

```
LINK: RES-NAME-1 CYS ATOM-NAME-1 SG RES-NAME-2 TMP FILE-2 inFiles/case2_comp1.cif ATOM-NAME-2 C1  
CHANGE BOND C1 C2 SINGLE 2
```

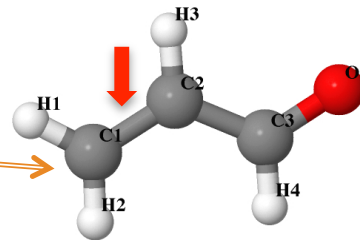
Covalent Links in Acedrg

Case 1: Scenario 2



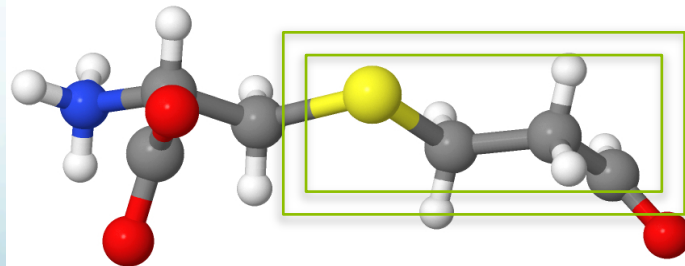
Jmol

CYS



Jmol

Monomer "TMP"

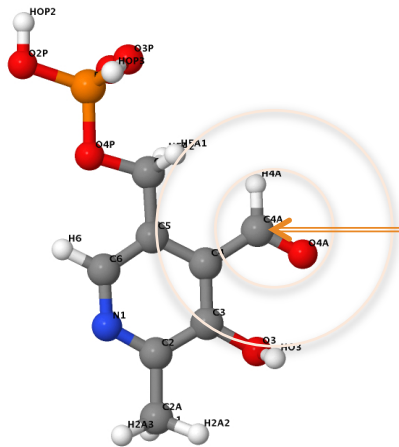


Jmol

The Linked molecule

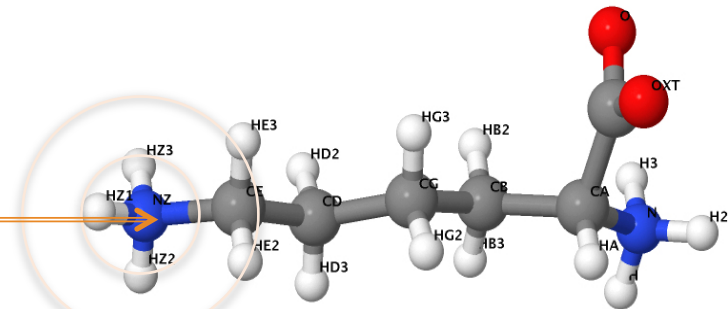
Covalent Links in Acedrg

Case 2



Jmol

PLP



Jmol

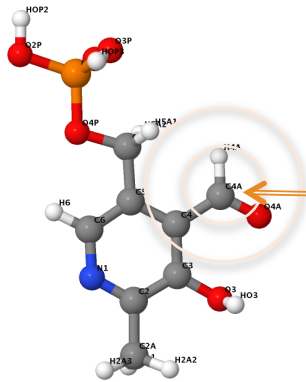
LYS

`instruct_LYS_PLP.txt`

```
LINK: RES-NAME-1 LYS ATOM-NAME-1 NZ RES-NAME-2 PLP ATOM-NAME-2 C4A  
DELETE ATOM O4A 2
```

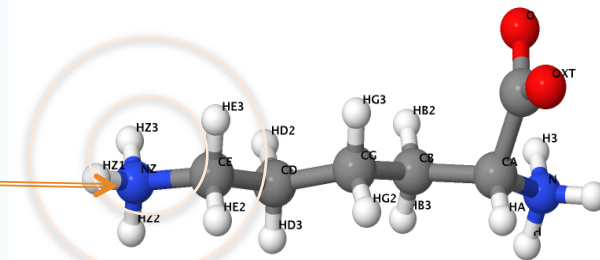

Covalent Links in Acedrg

Case 2



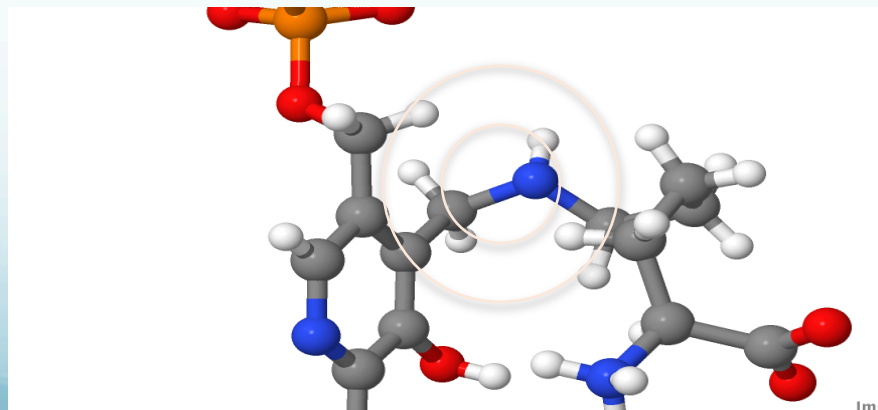
Jmol

PLP



Jmol

LYS



Jmol