

```
In [1]: import pickle
from rdkit import Chem
from rdkit.Chem import Draw
from rdkit.Chem.Draw import IPythonConsole
from rdkit.Chem import rdFMCS
from rdkit.Chem.Draw import rdDepictor

from IPython.display import display
import matplotlib.pyplot as plt
from IPython.display import HTML
import pandas as pd

IPythonConsole.ipython_useSVG=True
rdDepictor.SetPreferCoordGen(True)
#IPythonConsole.drawOptions.minFontSize=20
```

```
In [2]: with open('predictions/w_logs.pkl', 'rb') as file: w_te_data = pickle.load(fi
with open('predictions/wo_logs.pkl', 'rb') as file: wo_te_data = pickle.load(
original = pd.read_csv('predictions/chem_departm_output_wo_tie_embedding/outp
```

```
In [3]: def view_difference(mol1, mol2):
    mcs = rdFMCS.FindMCS([mol1, mol2])
    mcs_mol = Chem.MolFromSmarts(mcs.smartsString)
    match1 = mol1.GetSubstructMatch(mcs_mol)
    target_atm1 = []
    for atom in mol1.GetAtoms():
        if atom.GetIdx() not in match1:
            target_atm1.append(atom.GetIdx())
    match2 = mol2.GetSubstructMatch(mcs_mol)
    target_atm2 = []
    for atom in mol2.GetAtoms():
        if atom.GetIdx() not in match2:
            target_atm2.append(atom.GetIdx())
    return Draw.MolsToGridImage([mol1, mol2], highlightAtomLists=[target_atm1,
```

Generation

Notes:

- Predict the next fragment when probability $p > 0.5$
- The logic takes top-5 attachments from combinations of top-5 motifs and its possible configs. E.g., motif C1=CC=CC=C1 has 2 possible configs, C1=[CH:1]C=C[CH:2]=C1 or C1=[CH:1]C=CC=C1. The first config could be connected to other motifs that the connections are marked by :X, X is a number. The second config is the end motif that couldn't be connected to other motifs. Atoms marked by different mark numbers are connected together. No two atoms with same mark numbers are used for connection.
- For every attachment, it's checked for validity:
 - If the to-connect motif and to-be-connected (aka predicted motif) share common atoms for connections.
 - No self-loop.
 - If all atoms in the to-be-connected motif exist in the to-connect motif, no need to attach them.
- To view prediction logs of other molecules, subtract 2 from the molecule's index in Excel file.

In [4]:

```

def view(data, i, _original):
    print('Original: {}'.format(_original[i]))
    display(Draw.MolsToGridImage([Chem.MolFromSmiles(_original[i])]))

    sample = data[i]
    # step 0
    step_f0 = sample[0]
    print('*****Sample {}th*****'.format(i))
    print('-----Step-0-----')
    print('Root motif: {}'.format(step_f0['root']))
    print('Top 5 root motif configs:', '\n'.join([str(x) for x in step_f0['top-5-root-attachments']]))

    # display
    mol = Chem.MolFromSmiles(step_f0['top-5-root-attachments'][0][0])
    print('Displaying partial graph (aka molecule): {}'.format(step_f0['partial-graph']))
    display(Draw.MolsToGridImage([mol]))
    print('-----')

    # the remaining steps
    for i, step_f in enumerate(sample[1:]):
        print('-----Step-{}-----'.format(i + 1))
        if 'Generate fragment' in step_f:
            print('Generate next fragment p = {}'.format(step_f['Generate fragment p']))
        else:
            print('Skip, current fragment has no next fragment to be attached')
            continue

        if 'top-5-inter-cands' in step_f:
            print('Top 5 next motifs to attach:')
            for fragment in step_f['top-5-inter-cands']:
                print('Molecule {} and its specific config {} w/ p={}'.format(fragment[0], fragment[1], step_f['p']))
                display(Draw.MolsToGridImage([Chem.MolFromSmiles(fragment[1])]))
                print('-----')
                if 'Attaching Fragment' in step_f:
                    frag = step_f['Attaching Fragment']

                    sub_mol = Chem.MolFromSmiles(step_f['partial-graph'])
                    print('Attaching fragment {} of config {}'.format(frag[0], frag[1]))
                    print('Latest partial graph: {}'.format(step_f['partial-graph']))
                    print('Latest graph (left) vs graph in last step (right)')
                    display(view_difference(sub_mol, mol))
                    mol = sub_mol
                    print('-----')
                else:
                    print("Skip, the best next fragment to be attached to the current fragment is {}".format(frag[1]))

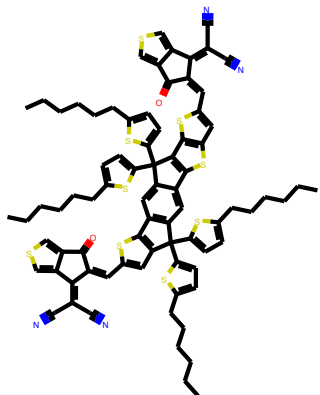
```

In [10]:

```
view(w_te_data, 12, original)
```

Original: CCCCCCC1=CC=C(S1)C2(C(S3)=CC=C3CCCCC)C(C(SC(/C=C(C4=O)/C(C5=CSC=C45

```
)=C(C#N)\C#N)=C6)=C6S7)=C7C8=CC9=C(C(SC(/C=C(C%10=O)/C(C%11=CSC=C%10%11)=C(C#N)\C#N)=C%12)=C%12C9(C(S%13)=CC=C%13CCCCC)C(S%14)=CC=C%14CCCCC)C=C28
```



*****Sample 12th*****

-----Step-0-----

Root motif: CC(=O)[O-]

Top 5 root motif configs: ('O=C([O-:1])[CH3:2]', tensor(13.9317))

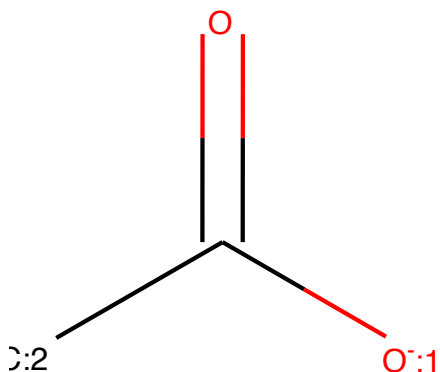
('O=C([O-])[CH3:1]', tensor(-12.5431))

('CC(=O)[O-:1]', tensor(-16.1744))

('[O:1]=[CH2:2]', tensor(-980.1329))

('C(#C[CH3:2])[CH3:1]', tensor(-980.1487))

Displaying partial graph (aka molecule): CC(=O)[O-]



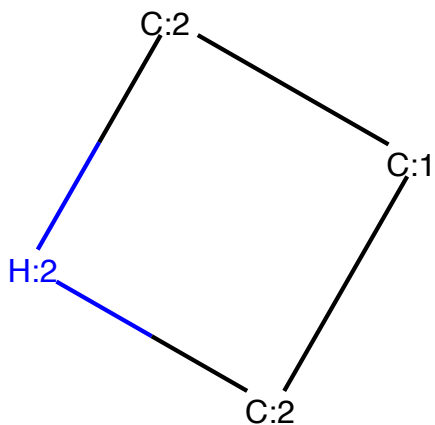
-----Step-1-----

Generate next fragment p = 1.0

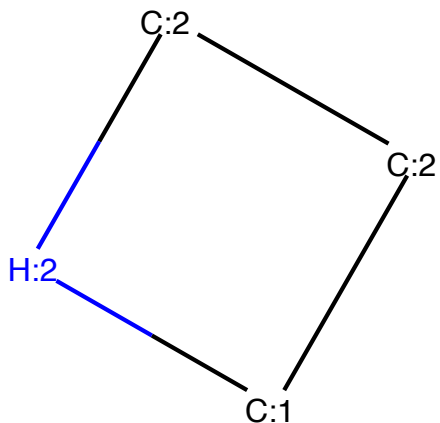
Top 5 next motifs to attach:

Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-1.0707383155822754

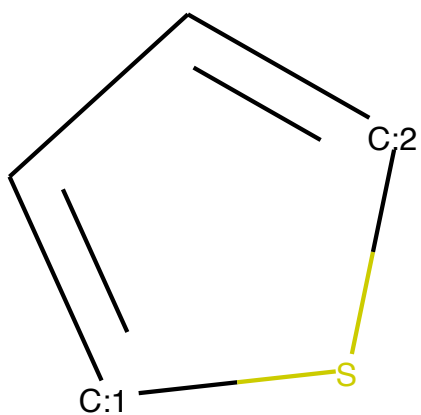
 Molecule C1CNC1 and its specific config [CH2:1]1[CH2:2][NH:2][CH2:2]1 w/ p=-1.
 16451895236969



 Molecule C1CNC1 and its specific config [CH2:1]1[CH2:2][CH2:2][NH:2]1 w/ p=-1.
 808347225189209

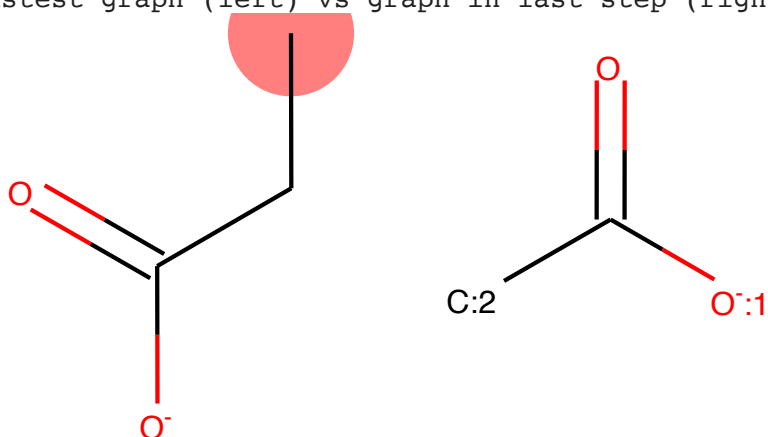


 Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-2.6687560
 081481934



 Molecule CN and its specific config [CH3:1][NH2:2] w/ p=-3.1203184127807617

 Attaching fragment [CH3:1][CH3:2] of config ['C[CH3:1]']
 Latest partial graph: CCC(=O)[O-]
 Lastest graph (left) vs graph in last step (right)

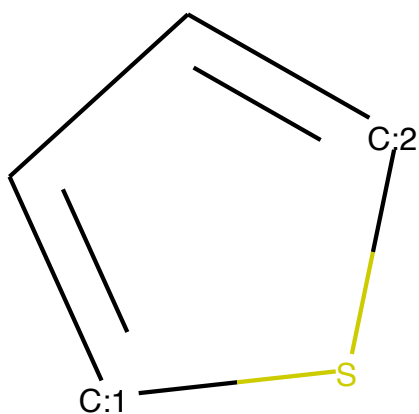


-----Step-2-----

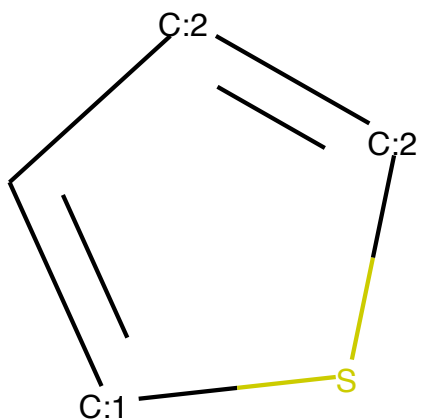
Generate next fragment p = 1.0

Top 5 next motifs to attach:

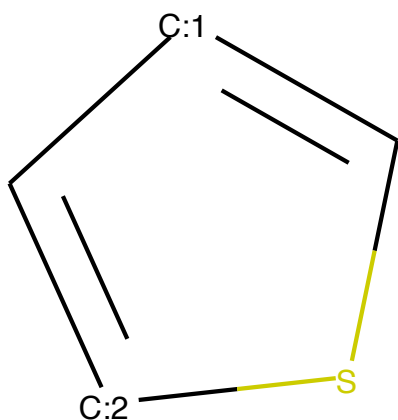
Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-0.0152112
 70190775394



 Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=[CH:2]1 w/ p=-4.91
 3881301879883

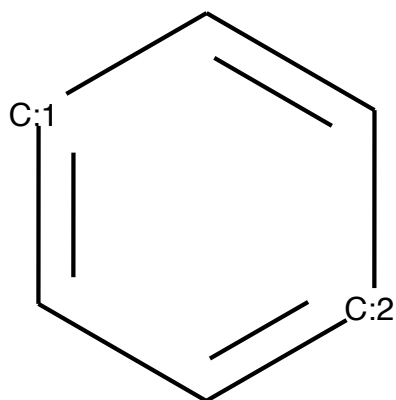


Molecule C1=CSC=C1 and its specific config C1=[CH:1]C=[CH:2]S1 w/ p=-5.474489212036133

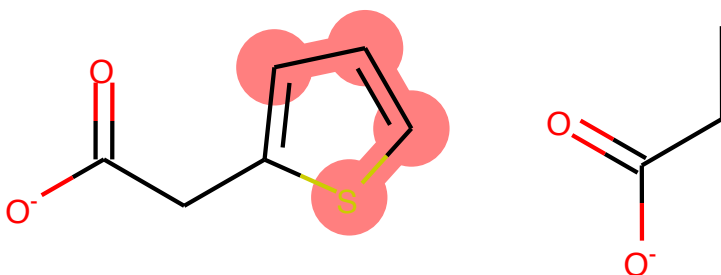


Molecule C and its specific config C w/ p=-5.725157260894775

Molecule C1=CC=CC=C1 and its specific config C1=[CH:1]C=C[CH:2]=C1 w/ p=-8.778680801391602



 Attaching fragment C1=[CH:1]S[CH:2]=C1 of config ['C1:C:S:[CH:1]:C:1']
 Latest partial graph: O=C([O-])Cc1cccs1
 Latest graph (left) vs graph in last step (right)



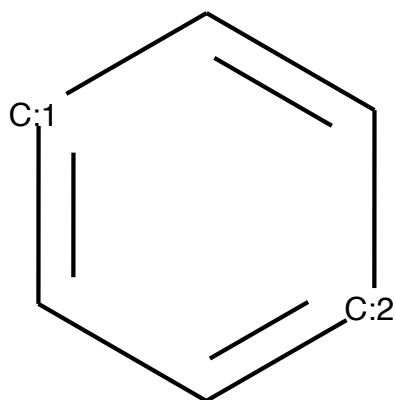
 -----Step-3-----

Generate next fragment p = 1.0

Top 5 next motifs to attach:

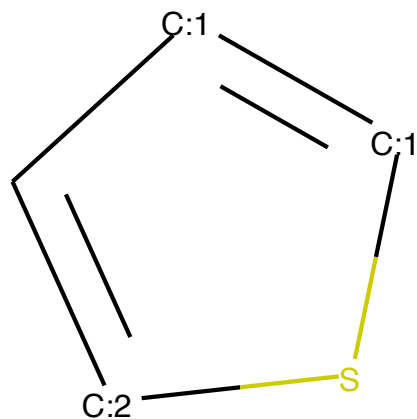
Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-1.811964830267243e-05

 Molecule C1=CC=CC=C1 and its specific config C1=[CH:1]C=C[CH:2]=C1 w/ p=-11.43
 4046745300293

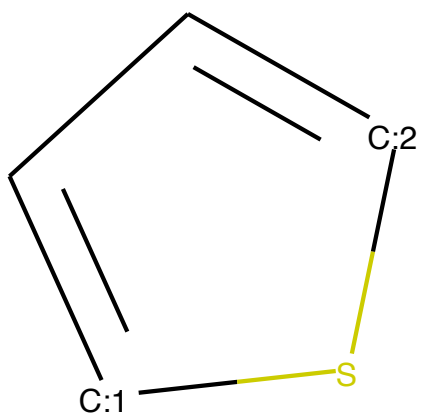


Molecule C and its specific config C w/ p=-11.994294166564941

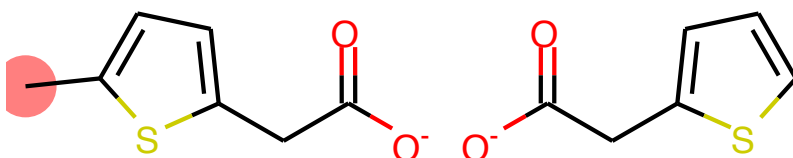
Molecule C1=CSC=C1 and its specific config C1=[CH:2]S[CH:1]=[CH:1]1 w/ p=-13.705684661865234



Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-17.18779754638672



 Attaching fragment [CH3:1][CH3:2] of config ['C[CH3:1]']
 Latest partial graph: Cc1ccc(CC(=O)[O-])s1
 Latest graph (left) vs graph in last step (right)

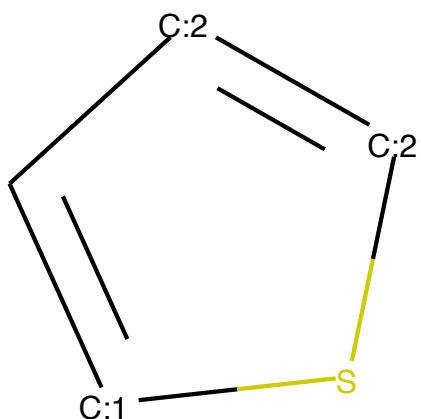


-----Step-4-----

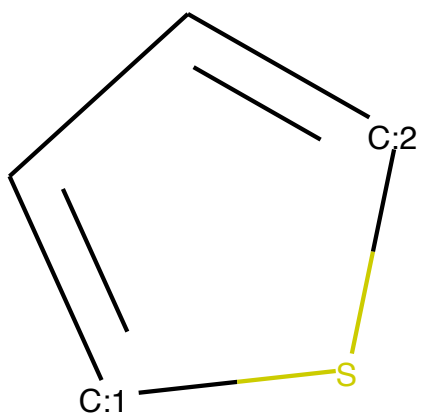
Generate next fragment p = 1.0

Top 5 next motifs to attach:

Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=[CH:2]1 w/ p=-0.00
 9146904572844505

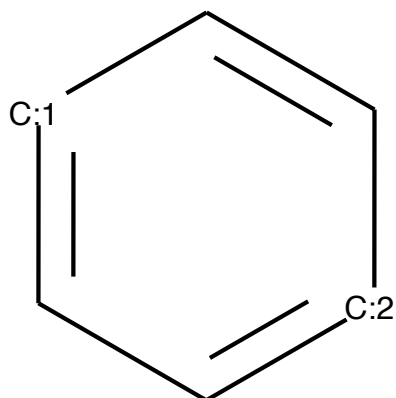


 Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-4.7870278
 35845947

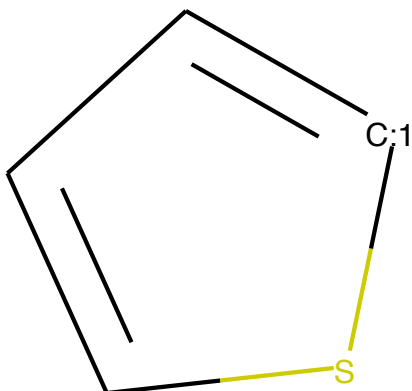


Molecule C and its specific config C w/ p=-7.912520408630371

Molecule C1=CC=CC=C1 and its specific config C1=[CH:1]C=C[CH:2]=C1 w/ p=-8.354876518249512



Molecule C1=CSC=C1 and its specific config C1=CS[CH:1]=C1 w/ p=-8.974496841430664



 Attaching fragment C1=[CH:1]S[CH:2]=[CH:2]1 of config ['C1:C:S:[CH:1]:C:1']
 Latest partial graph: O=C([O-])Cc1ccc(-c2cccs2)s1
 Latest graph (left) vs graph in last step (right)



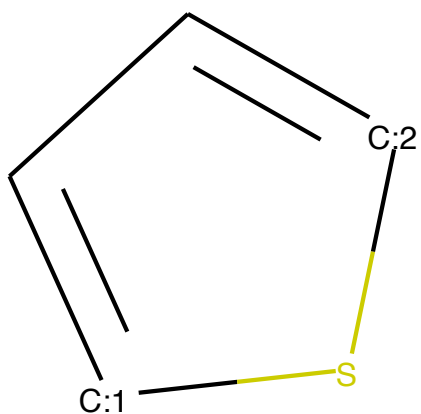
 -----Step-5-----

Generate next fragment p = 0.9997887015342712

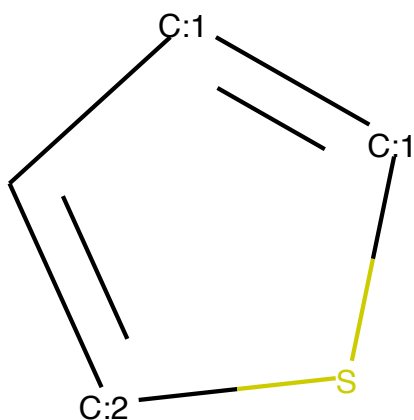
Top 5 next motifs to attach:

Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-0.5695070624351501

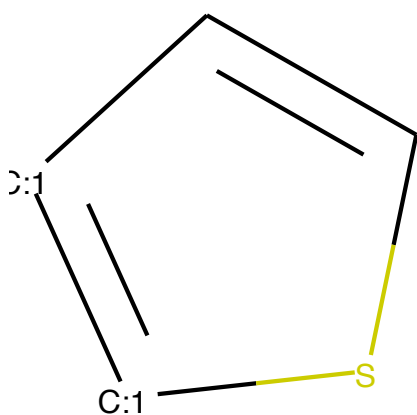
 Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-1.1753448
 247909546



Molecule C1=CSC=C1 and its specific config C1=[CH:2]S[CH:1]=[CH:1]1 w/ p=-2.7043516635894775

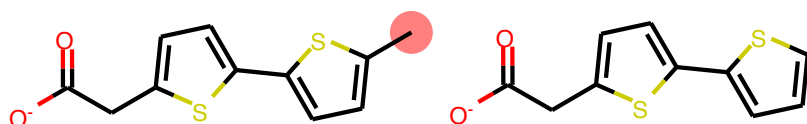


Molecule C1=CSC=C1 and its specific config C1=C[CH:1]=[CH:1]S1 w/ p=-3.261600971221924



Molecule C[SiH3] and its specific config [CH3:1][SiH3:2] w/ p=-4.336895942687988

 Attaching fragment [CH3:1][CH3:2] of config ['C[CH3:1]']
 Latest partial graph: Cc1ccc(-c2ccc(CC(=O)[O-])s2)s1
 Lastest graph (left) vs graph in last step (right)

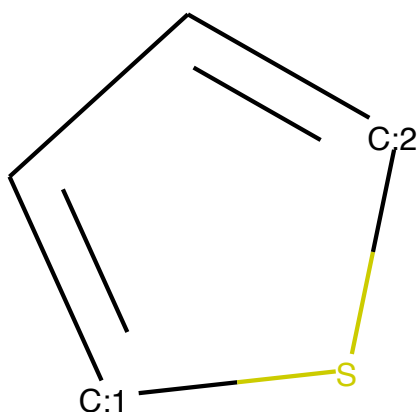


-----Step-6-----

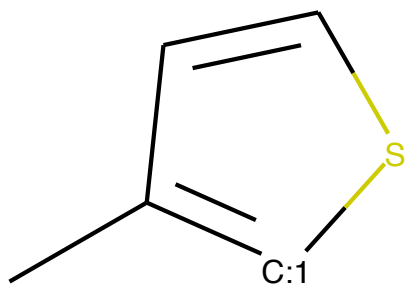
Generate next fragment p = 1.0

Top 5 next motifs to attach:

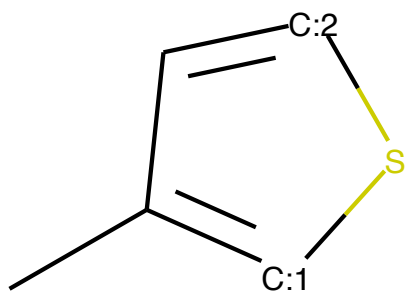
Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-0.0103656
 60302340984



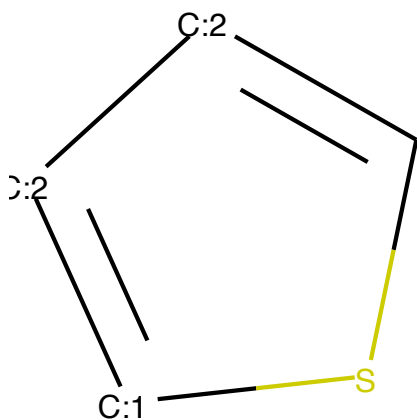
 Molecule CC1=CSC=C1 and its specific config CC1=[CH:1]SC=C1 w/ p=-4.7038173675
 53711



 Molecule CC1=CSC=C1 and its specific config CC1=[CH:1]S[CH:2]=C1 w/ p=-6.87516
 5939331055

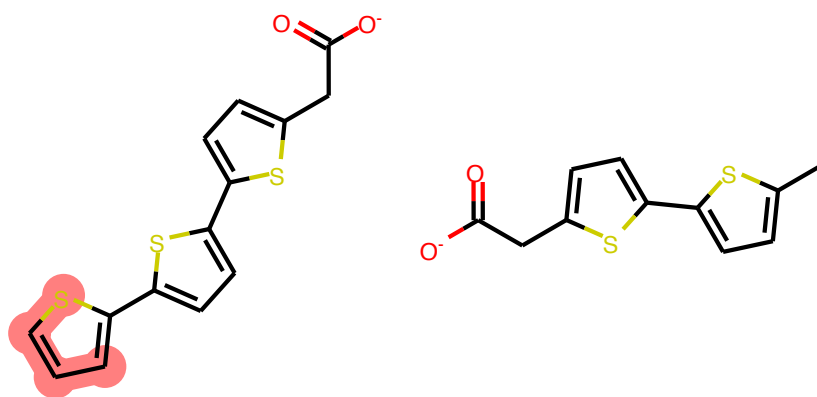


 Molecule C1=CSC=C1 and its specific config C1=[CH:2][CH:2]=[CH:1]S1 w/ p=-9.22
 6217269897461



 Molecule C and its specific config C w/ p=-9.870923042297363

 Attaching fragment C1=[CH:1]S[CH:2]=C1 of config ['C1:C:S:[CH:1]:C:1']
 Latest partial graph: O=C([O-])Cc1ccc(-c2ccc(-c3cccs3)s2)s1
 Lastest graph (left) vs graph in last step (right)



-----Step-7-----

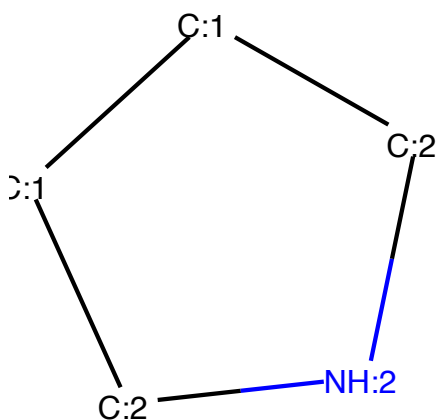
Generate next fragment p = 0.9998983144760132

Top 5 next motifs to attach:

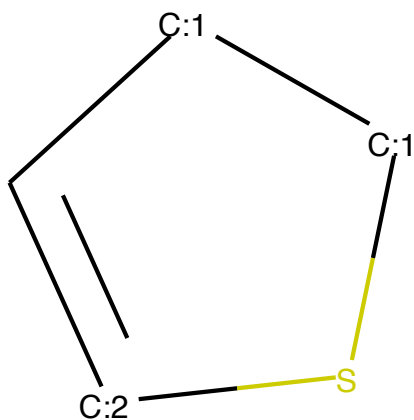
Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-0.0063044242560863495

 Molecule CC and its specific config C[CH3:1] w/ p=-5.355673313140869

 Molecule C1CCNC1 and its specific config [CH2:1]1[CH2:1][CH2:2][NH:2][CH2:2]1
 w/ p=-7.019346714019775

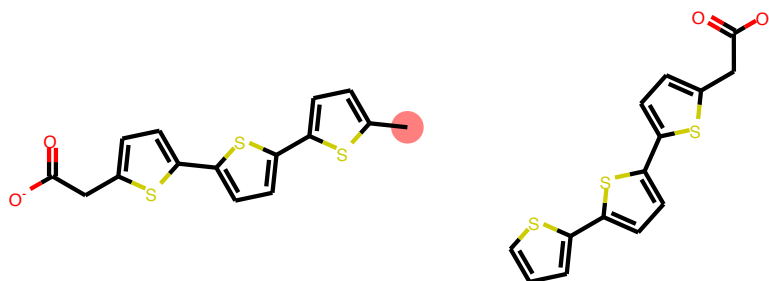


 Molecule C1=CSCC1 and its specific config C1=[CH:2]S[CH2:1][CH2:1]1 w/ p=-7.96
 6949939727783

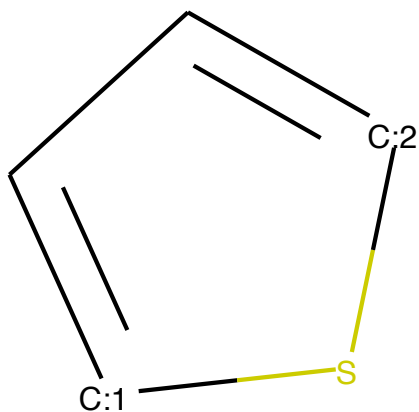


 Molecule C=O and its specific config O=[CH2:1] w/ p=-9.378978729248047

 Attaching fragment [CH3:1][CH3:2] of config ['C[CH3:1]']
 Latest partial graph: Cc1ccc(-c2ccc(-c3ccc(CC(=O)[O-])s3)s2)s1
 Lastest graph (left) vs graph in last step (right)

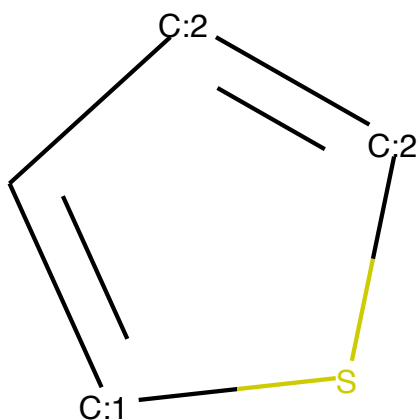


 -----Step-8-----
 Generate next fragment p = 0.9967923760414124
 Top 5 next motifs to attach:
 Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-0.1238685
 6973171234

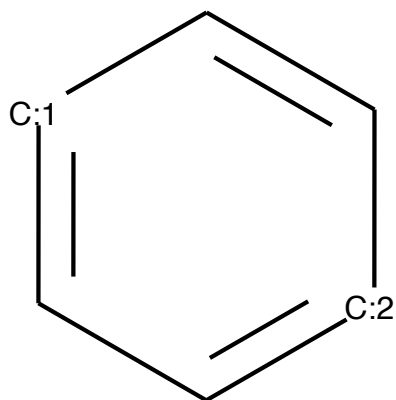


 Molecule C and its specific config C w/ p=-2.2312333583831787

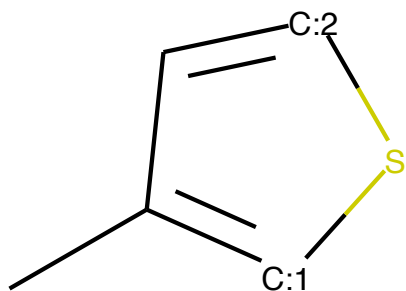
Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=[CH:2]1 w/ p=-5.37
5985622406006



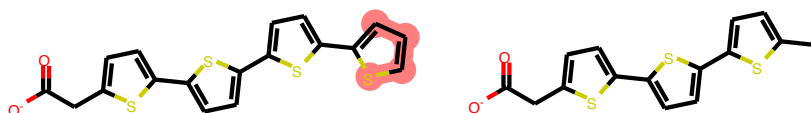
Molecule C1=CC=CC=C1 and its specific config C1=[CH:1]C=C[CH:2]=C1 w/ p=-6.157
557010650635



Molecule CC1=CSC=C1 and its specific config CC1=[CH:1]S[CH:2]=C1 w/ p=-6.57070
3983306885

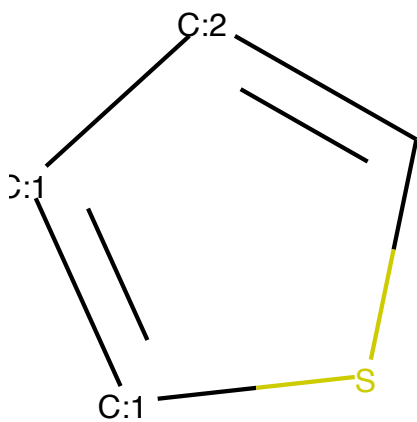


 Attaching fragment C1=[CH:1]S[CH:2]=C1 of config ['C1:C:S:[CH:1]:C:1']
 Latest partial graph: O=C([O-])Cclccc(-c2ccc(-c3ccc(-c4cccs4)s3)s2)s1
 Lastest graph (left) vs graph in last step (right)



 -----Step-9-----
 Generate next fragment p = 0.9993383288383484
 Top 5 next motifs to attach:
 Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-0.0006313720368780196

 Molecule C1=CSC=C1 and its specific config C1=[CH:2][CH:1]=[CH:1]S1 w/ p=-7.51
 4693737030029

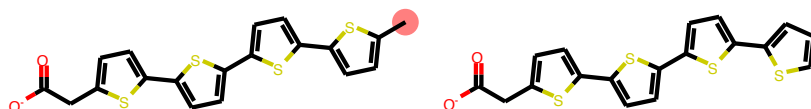


Molecule CS and its specific config [CH3:1][SH:2] w/ p=-11.040324211120605

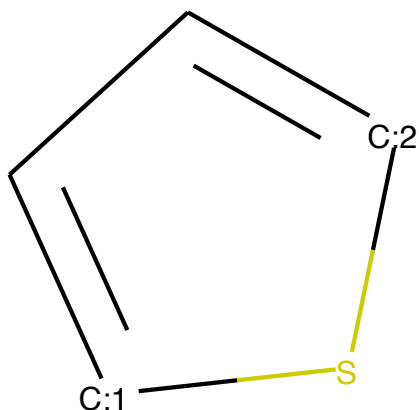
Molecule C=O and its specific config O=[CH2:1] w/ p=-11.106101036071777

Molecule C=N and its specific config [CH2:1]=[NH:2] w/ p=-11.327986717224121

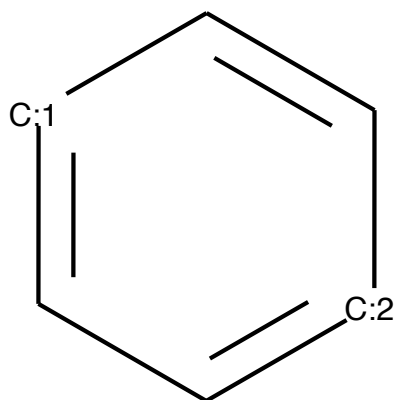
 Attaching fragment [CH3:1][CH3:2] of config ['C[CH3:1]']
 Latest partial graph: Cc1ccc(-c2ccc(-c3ccc(-c4ccc(CC(=O)[O-])s4)s3)s2)s1
 Lastest graph (left) vs graph in last step (right)



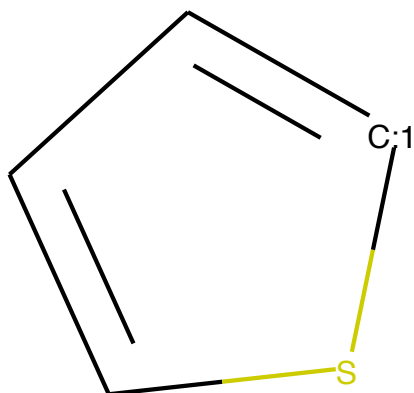
 -----Step-10-----
 Generate next fragment p = 1.0
 Top 5 next motifs to attach:
 Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-0.0962555
 2594661713



 Molecule C1=CC=CC=C1 and its specific config C1=[CH:1]C=C[CH:2]=C1 w/ p=-2.943
 636894226074



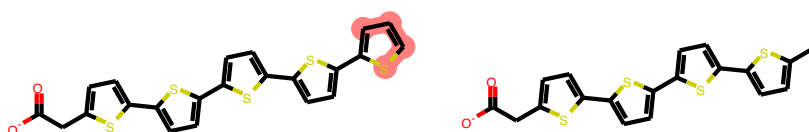
Molecule C1=CSC=C1 and its specific config C1=CS[CH:1]=C1 w/ p=-3.4166290760040283



Molecule C and its specific config C w/ p=-5.316184997558594

Molecule [SiH4] and its specific config [SiH4] w/ p=-6.750178337097168

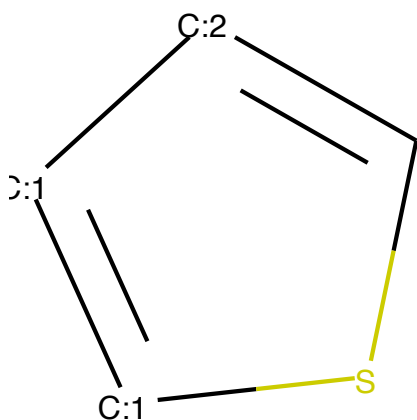
Attaching fragment C1=[CH:1]S[CH:2]=C1 of config ['C1:C:S:[CH:1]:C:1']
Latest partial graph: O=C([O-])Cc1ccc(-c2ccc(-c3ccc(-c4ccc(-c5cccs5)s4)s3)s2)s1
Lastest graph (left) vs graph in last step (right)



-----Step-11-----
Generate next fragment p = 0.8303309679031372
Top 5 next motifs to attach:
Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-0.041510120034217834

Molecule CS and its specific config [CH3:1][SH:2] w/ p=-3.2143540382385254

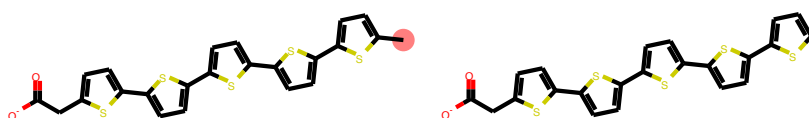
Molecule C1=CSC=C1 and its specific config C1=[CH:2][CH:1]=[CH:1]S1 w/ $p=-7.846226692199707$



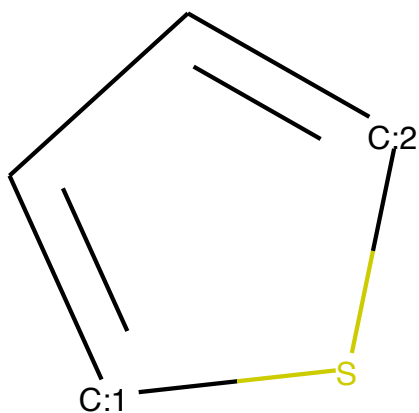
Molecule C=N and its specific config [CH2:1]=[NH:2] w/ $p=-10.278705596923828$

Molecule C=O and its specific config O=[CH2:1] w/ $p=-10.48699951171875$

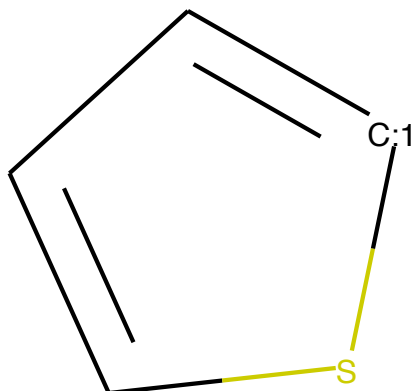
 Attaching fragment [CH3:1][CH3:2] of config ['C[CH3:1]']
 Latest partial graph: Cc1ccc(-c2ccc(-c3ccc(-c4ccc(-c5ccc(CC(=O)[O-])s5)s4)s3)s2)s1
 Lastest graph (left) vs graph in last step (right)



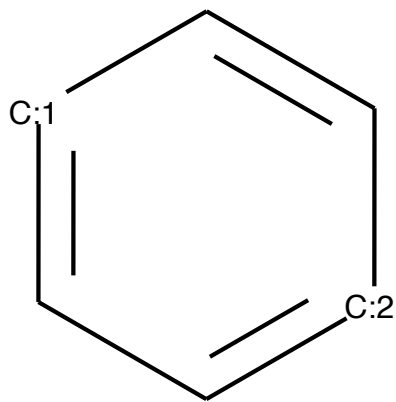
 -----Step-12-----
 Generate next fragment p = 1.0
 Top 5 next motifs to attach:
 Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-0.5011690258979797



 Molecule C1=CSC=C1 and its specific config C1=CS[CH:1]=C1 w/ p=-1.3847014904022217



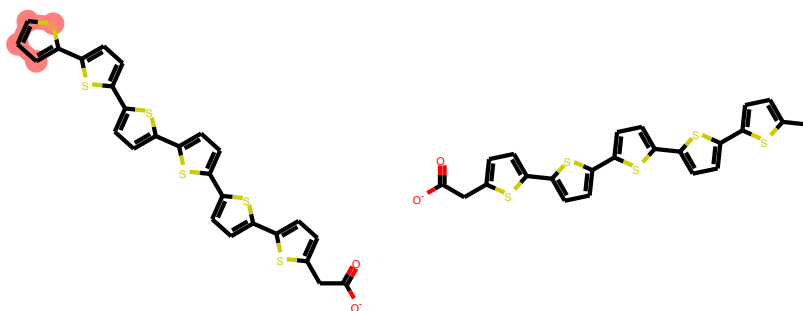
 Molecule C1=CC=CC=C1 and its specific config C1=[CH:1]C=C[CH:2]=C1 w/ p=-2.0013294219970703



 Molecule C and its specific config C w/ p=-4.850032329559326

 Molecule C#N and its specific config N#[CH:1] w/ p=-7.40615701675415

Attaching fragment C1=[CH:1]S[CH:2]=C1 of config ['C1:C:S:[CH:1]:C:1']
Latest partial graph: O=C([O-])Cclccc(-c2ccc(-c3ccc(-c4ccc(-c5ccc(-c6cccs6)s5)s4)s3)s2)s1
Lastest graph (left) vs graph in last step (right)

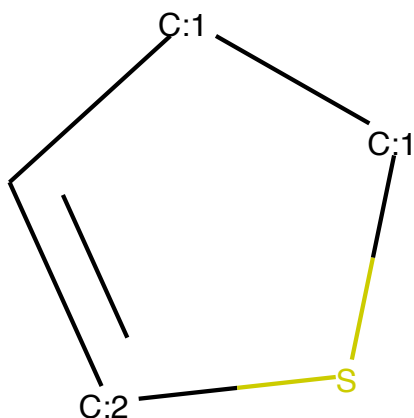


-----Step-13-----
Generate next fragment p = 0.31631457805633545
-----Step-14-----
Generate next fragment p = 2.1378728306289908e-15
-----Step-15-----
Generate next fragment p = 6.014240057083953e-07
-----Step-16-----
Generate next fragment p = 1.5512888011919718e-13
-----Step-17-----
Generate next fragment p = 0.9976533055305481
Top 5 next motifs to attach:
Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-0.0001510267611593008

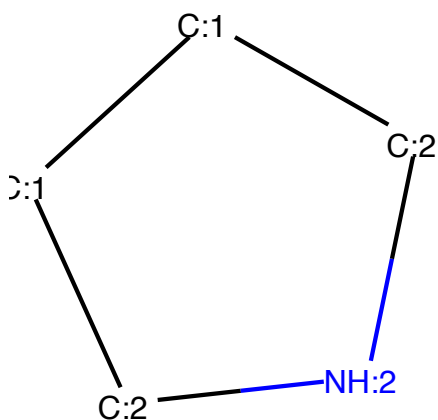
Molecule C=N and its specific config [CH2:1]=[NH:2] w/ p=-9.659462928771973

Molecule S and its specific config S w/ p=-9.775293350219727

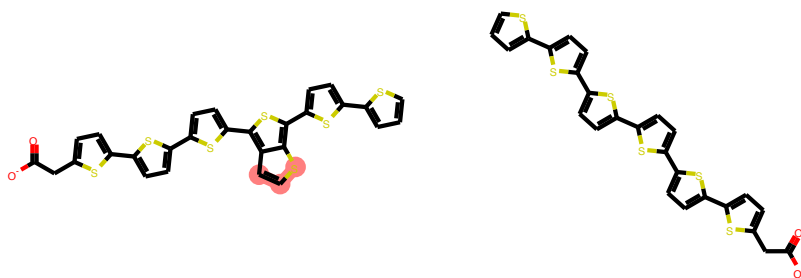
Molecule C1=CSCC1 and its specific config C1=[CH:2]S[CH2:1][CH2:1]1 w/ p=-10.877622604370117



 Molecule C1CCNC1 and its specific config [CH2:1]1[CH2:1][CH2:2][NH:2][CH2:2]1
 w/ p=-11.646870613098145



 Attaching fragment C1=[CH:2]S[CH2:1][CH2:1]1 of config ['C1=CS[CH2:1]C1', 'C1=C[CH2:1]CS1']
 Latest partial graph: O=C([O-])Cc1ccc(-c2ccc(-c3ccc(-c4sc(-c5ccc(-c6cccs6)s5)c5sccc45)s3)s2)s1
 Latest graph (left) vs graph in last step (right)



-----Step-18-----
 Generate next fragment p = 0.98321932554245
 Top 5 next motifs to attach:
 Molecule CS and its specific config [CH3:1][SH:2] w/ p=-3.814624506048858e-05

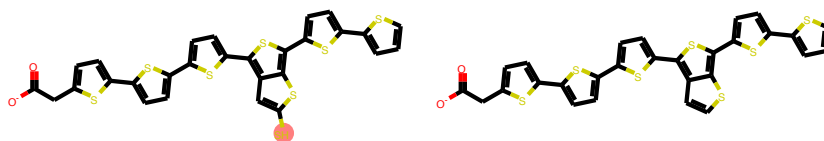
Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-10.231485366821289

Molecule [SiH4] and its specific config [SiH4] w/ p=-14.029769897460938

Molecule C=O and its specific config O=[CH2:1] w/ p=-14.399831771850586

 Molecule [CH2-]C and its specific config [CH3:1][CH2-:2] w/ p=-14.918670654296875

 Attaching fragment [CH3:1][SH:2] of config ['S[CH3:1]']
 Latest partial graph: O=C([O-])Cc1ccc(-c2ccc(-c3ccc(-c4sc(-c5ccc(-c6cccs6)s5)c5sc(S)cc45)s3)s2)s1
 Lastest graph (left) vs graph in last step (right)

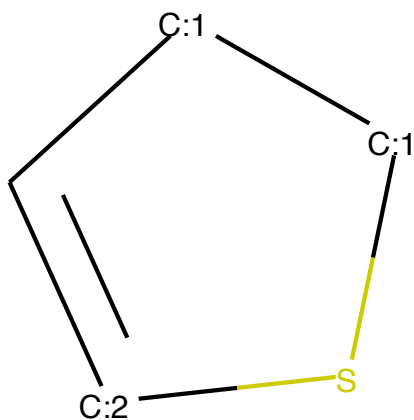


 -----Step-19-----
 Generate next fragment p = 1.0
 Top 5 next motifs to attach:
 Molecule S and its specific config S w/ p=-0.9460978507995605

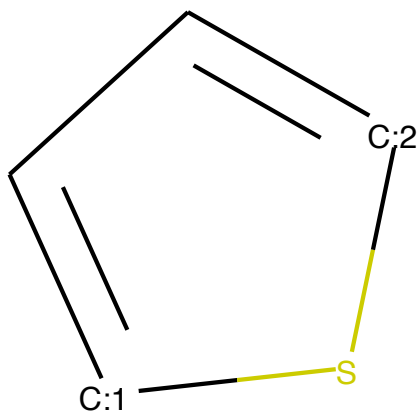
Molecule CS and its specific config C[SH:1] w/ p=-1.0368585586547852

Molecule C and its specific config C w/ p=-1.455458164215088

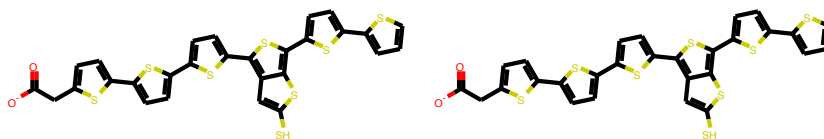
Molecule C1=CSCC1 and its specific config C1=[CH:2]S[CH2:1][CH2:1]1 w/ p=-3.820958375930786



 Molecule C1=CSC=C1 and its specific config C1=[CH:1]S[CH:2]=C1 w/ p=-6.773167610168457



 Attaching fragment S of config ['[SH2:1]']
 Latest partial graph: O=C([O-])Cc1ccc(-c2ccc(-c3ccc(-c4sc(-c5ccc(-c6cccs6)s5)c5sc(S)cc45)s3)s2)s1
 Latest graph (left) vs graph in last step (right)



-----Step-20-----

Generate next fragment p = 1.0

Top 5 next motifs to attach:

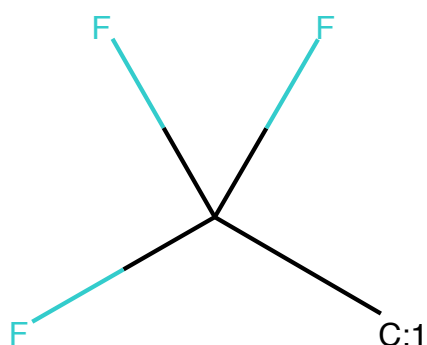
Molecule CS and its specific config C[SH:1] w/ p=-2.7656173188006505e-05

Molecule C[SiH3] and its specific config C[SiH3:1] w/ p=-10.856226921081543

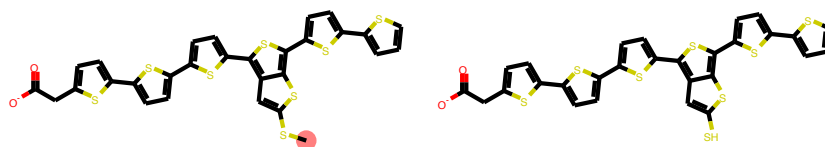
Molecule CC#N and its specific config N#C[CH3:1] w/ p=-11.877511024475098

Molecule O=S and its specific config O=[S:1] w/ p=-14.073086738586426

 Molecule CC(F)(F)F and its specific config FC(F)(F)[CH3:1] w/ p=-14.895258903503418



 Attaching fragment C[SH:1] of config ['C[SH:1]']
 Latest partial graph: CSc1cc2c(-c3ccc(-c4ccc(-c5ccc(CC(=O)[O-])s5)s4)s3)sc(-c3ccc(-c4cccs4)s3)c2s1
 Lastest graph (left) vs graph in last step (right)



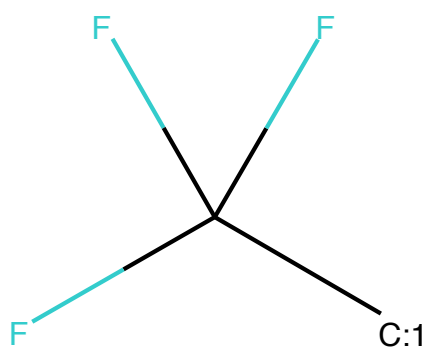
 -----Step-21-----
 Generate next fragment p = 7.399842871791407e-08
 -----Step-22-----
 Generate next fragment p = 1.0
 Top 5 next motifs to attach:
 Molecule O=S and its specific config O=[S:1] w/ p=-4.2199197196168825e-05

Molecule CS and its specific config C[SH:1] w/ p=-10.079389572143555

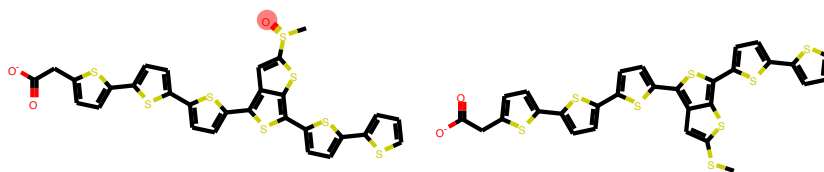
Molecule C[SiH3] and its specific config C[SiH3:1] w/ p=-15.358508110046387

Molecule [SiH4] and its specific config [SiH4] w/ p=-17.474271774291992

 Molecule CC(F)(F)F and its specific config FC(F)(F)[CH3:1] w/ p=-19.0468444824
 21875



 Attaching fragment O=[S:1] of config ['O=[S:1]']
 Latest partial graph: CS(=O)c1cc2c(-c3ccc(-c4ccc(-c5ccc(CC(=O)[O-])s5)s4)s3)sc(-c3ccc(-c4cccs4)s3)c2s1
 Lastest graph (left) vs graph in last step (right)



 -----Step-23-----
 Generate next fragment p = 0.0
 -----Step-24-----
 Generate next fragment p = 1.0
 Top 5 next motifs to attach:
 Molecule O=S and its specific config O=[S:1] w/ p=0.0

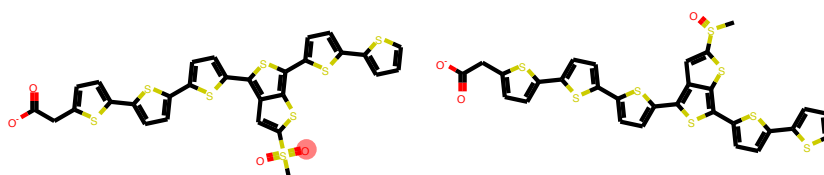
Molecule CC#N and its specific config N#C[CH3:1] w/ p=-19.532554626464844

Molecule CS and its specific config C[SH:1] w/ p=-20.424989700317383

Molecule C[SiH3] and its specific config C[SiH3:1] w/ p=-21.05328941345215

Molecule CN and its specific config N[CH3:1] w/ p=-22.404373168945312

Attaching fragment O=[S:1] of config ['O=[S:1]']
Latest partial graph: CS(=O)(=O)c1cc2c(-c3ccc(-c4ccc(-c5ccc(CC(=O)[O-])s5)s4)s3)sc(-c3ccc(-c4cccs4)s3)c2s1
Lastest graph (left) vs graph in last step (right)



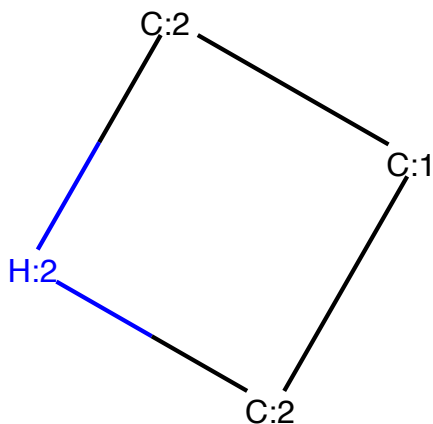

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-----Step-25-----  
Generate next fragment p = 1.737397008713094e-36  
-----Step-26-----  
Generate next fragment p = 5.188417517842936e-08  
-----Step-27-----  
Generate next fragment p = 1.4229140250709093e-15  
-----Step-28-----  
Generate next fragment p = 4.507595943096902e-15  
-----Step-29-----  
Generate next fragment p = 2.1907074678612082e-13  
-----Step-30-----  
Generate next fragment p = 2.852038053189957e-24  
-----Step-31-----  
Generate next fragment p = 2.4798725917207776e-06  
-----Step-32-----  
Generate next fragment p = 1.7128844150028954e-20  
-----Step-33-----  
Generate next fragment p = 0.24527804553508759  
-----Step-34-----  
Generate next fragment p = 3.1638699495767303e-27  
-----Step-35-----  
Generate next fragment p = 9.040126514037183e-08  
-----Step-36-----  
Generate next fragment p = 6.324027771182904e-23  
-----Step-37-----  
Generate next fragment p = 0.9880258440971375  
Top 5 next motifs to attach:  
Molecule CC and its specific config [CH3:1][CH3:2] w/ p=-0.0001016803507809527  
2
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Molecule CN and its specific config [NH2:1][CH3:2] w/ p=-9.30424690246582
```

Molecule CN and its specific config [CH3:1][NH2:2] w/ p=-11.479131698608398

Molecule C=O and its specific config O=[CH2:1] w/ p=-15.407944679260254

Molecule C1CNC1 and its specific config [CH2:1]1[CH2:2][NH:2][CH2:2]1 w/ p=-17.559106826782227



Skip, the best next fragment to be attached to the current fragment does not yield a valid sub-molecule . Go back to the previous fragment.

-----Step-38-----

Skip, current fragment has no next fragment to be attached. Go back to the previous fragment.

-----Step-39-----

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Skip, current fragment has no next fragment to be attached. Go back to the previous fragment.
-----Step-100-----
Skip, current fragment has no next fragment to be attached. Go back to the previous fragment.

In []: