

In [27]: `pwd`

Out[27]: `'C:\\Users\\L03121898\\Anaconda3\\envs\\Cadi22\\demo_chemi'`

In [28]: `ls`

El volumen de la unidad C no tiene etiqueta.

El número de serie del volumen es: C636-ABF1

Directorio de C:\\Users\\L03121898\\Anaconda3\\envs\\Cadi22\\demo\_chemi

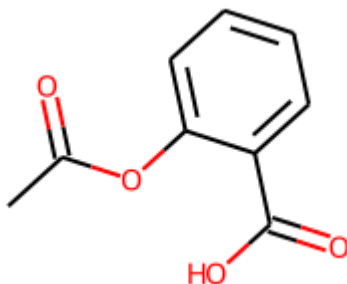
```
10/01/2022 07:31 p.m. <DIR> .
10/01/2022 07:31 p.m. <DIR> ..
10/01/2022 04:11 p.m. <DIR> .ipynb_checkpoints
10/01/2022 12:17 p.m. 1,273 aspirin.mol
10/01/2022 12:17 p.m. 3,641 aspirin.sdf
10/01/2022 12:17 p.m. 15,740,199 erk2.csv
10/01/2022 12:17 p.m. 24,914 export.sdf
10/01/2022 12:17 p.m. 203 imatinib.smi
10/01/2022 12:17 p.m. 4,300 sample.csv
10/01/2022 12:17 p.m. 2,666 sample.mol
10/01/2022 07:31 p.m. 56,384 Untitled.ipynb
      8 archivos 15,833,580 bytes
      3 dirs 80,171,159,552 bytes libres
```

In [11]: `from rdkit import Chem`  
`from rdkit.Chem import Draw`

In [12]: `x=Chem.MolFromSmiles("CC(=O)OC1=CC=CC=C1C(=O)O")`

In [13]: `x`

Out[13]:



In [26]:

8(37)23-8-6-22(7-9-23)20-36-16-14-35(2)15-17-36;1-5(2,3)4/h3-13,18-19H,14-17,2

In [14]: `path="C:\\Users\\L03121898\\Anaconda3\\envs\\Cadi22\\demo_chemi\\"`

x1

In [12]: `x1`

Out[12]:



```
In [15]: x2=Chem.SDMolSupplier(path+"aspirin.sdf")
x3=Chem.SmilesMolSupplier("imatinib.smi", delimiter="\t", titleLine=True)
x4= Chem.MolFromInchi("InChI=1S/C29H31N7O.CH4O3S/c1-21-5-10-25(18-27(21)34-29-
```

```
In [24]: x2=Chem.SmilesMolSupplier(path+"imatinib.smi",delimiter="\t", titleLine=True)
```

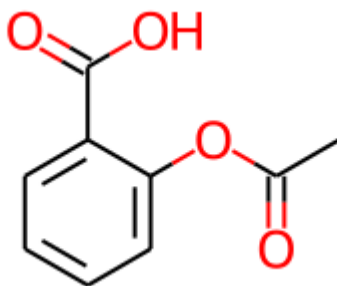
In [25]: `x2`Out[25]: `<rdkit.Chem.rdmolfiles.SmilesMolSupplier at 0x12ef0c663b0>`

```
In [23]: [mol.GetNumAtoms() for mol in x2 if mol is not None]
```

Out[23]: `[]`

```
In [18]: Chem.MolFromMolFile ("aspirin.mol")
```

Out[18]:



```
In [16]: x1=Chem.MolFromSmiles("C1=CC=CC=C1")
x2=Chem.SDMolSupplier(path+"aspirin.sdf")
x3=Chem.SmilesMolSupplier(path+"imatinib.smi", delimiter="\t", titleLine=True)
x4=Chem.MolFromInchi("InChI=1S/C29H31N7O.CH4O3S/c1-21-5-10-25(18-27(21)34-29-
```

In [17]:  x4

Out[17]:

