

## NAME

TopologicalAtomTorsionsFingerprints.pl - Generate topological atom torsions fingerprints for SD files

## SYNOPSIS

TopologicalAtomTorsionsFingerprints.pl SDFfile(s)...

```
TopologicalAtomTorsionsFingerprints.pl [-a, --AtomIdentifierType AtomicInvariantsAtomTypes] [
--AtomicInvariantsToUse "AtomicInvariant,AtomicInvariant..."] [--FunctionalClassesToUse
"FunctionalClass1,FunctionalClass2..."] [--CompoundID DataFieldName or LabelPrefixString] [--CompoundIDLabel text]
[--CompoundIDMode] [--DataFields "FieldLabel1,FieldLabel2,..."] [-d, --DataFieldsMode All | Common | Specify |
CompoundID] [-f, --Filter Yes | No] [--FingerprintsLabel text] [-h, --help] [-k, --KeepLargestComponent Yes |
No] [--OutDelim comma | tab | semicolon] [--output SD | text | both] [-o, --overwrite] [-q, --quote Yes | No] [-r,
--root RootName] [-v, --VectorStringFormat] [-w, --WorkingDir dirname] SDFfile(s)...
```

## DESCRIPTION

Generate topological atom torsions fingerprints [ Ref 58, Ref 72 ] for *SDFfile(s)* and create appropriate SD or CSV/TSV text file(s) containing fingerprints vector strings corresponding to molecular fingerprints.

Multiple SDFfile names are separated by spaces. The valid file extensions are *.sdf* and *.sd*. All other file names are ignored. All the SD files in a current directory can be specified either by *\*.sdf* or the current directory name.

The current release of MayaChemTools supports generation of topological atom torsions fingerprints corresponding to following *-a, --AtomIdentifierTypes*:

```
AtomicInvariantsAtomTypes, DREIDINGAtomTypes, EStateAtomTypes,
FunctionalClassAtomTypes, MMFF94AtomTypes, SLogPAtomTypes,
SYBYLAtomTypes, TPSAAtomTypes, UFFAtomTypes
```

Based on the values specified for *-a, --AtomIdentifierType* and *--AtomicInvariantsToUse*, initial atom types are assigned to all non-hydrogen atoms in a molecule. All unique atom torsions are identified and an atom torsion identifier is generated; the format of atom torsion identifier is:

```
<AtomType1>-<AtomType2>-<AtomType3>-<AtomType4>
```

```
AtomType1, AtomType2, AtomType3, AtomType4: Assigned atom types
```

```
where AtomType1 <= AtomType2 <= AtomType3 <= AtomType4
```

The atom torsion identifiers for all unique atom torsions corresponding to non-hydrogen atoms constitute topological atom torsions fingerprints of the molecule.

The current release of MayaChemTools generates the following types of topological atom torsions fingerprints vector strings:

```
FingerprintsVector;TopologicalAtomTorsions:AtomicInvariantsAtomTypes;11;
NumericalValues;IDsAndValuesString;C.X1.B01.H3-C.X3.B04-C.X2.B03.H1-
N.X2.B02.H1 C.X1.B01.H3-C.X3.B04-C.X3.B04-C.X2.B03.H1 C.X1.B01.H3-
C.X3.B04-C.X3.B04-O.X1.B02 C.X2.B03.H1-C.X2.B03.H1-C.X3.B04-C.X3.B04
C.X2.B03.H1-C.X2.B03.H1-C.X3.B04-O.X1.B02...;
1 1 1 1 1 1 1 1 1 1
```

```
FingerprintsVector;TopologicalAtomTorsions:AtomicInvariantsAtomTypes;11;
NumericalValues;IDsAndValuesPairsString;C.X1.B01.H3-C.X3.B04-C.X2.B03.H1-
N.X2.B02.H1 1 C.X1.B01.H3-C.X3.B04-C.X3.B04-C.X2.B03.H1 1 C.X1.B01.H3-
C.X3.B04-C.X3.B04-O.X1.B02 1 C.X2.B03.H1-C.X2.B03.H1-C.X3.B04-
C.X3.B04 1 C.X2.B03.H1-C.X2.B03.H1-C.X3.B04-O.X1.B02 1 C.X2.B03.H1-
C.X2.B03.H1-N.X2.B02.H1-C.X2.B03.H1 1 C.X2.B03.H1-C.X3.B04-C.X3.B04-
C.X2.B03.H1 1...
```

```
FingerprintsVector;TopologicalAtomTorsions:DREIDINGAtomTypes;12;Numerical
Values;IDsAndValuesString;C_2-C_3-C_3-C_3 C_2-C_3-C_3-O_3 C_2-N_3-C_3-C
_3 C_3-C_2-N_3-C_3 C_3-C_3-C_2-O_2 C_3-C_3-C_2-O_3 C_3-C_3-C_3-C_3-C_3-
```

```
C_3-C_3-N_3 C_3-C_3-C_3-O_3 C_3-N_3-C_2-N_2 C_3-N_3-C_2-N_3 C_3-N_3-C_2
-O_2;2 1 4 1 2 2 15 7 3 1 1 1
```

```
FingerprintsVector;TopologicalAtomTorsions:EStateAtomTypes;23;Numerical
Values;IDsAndValuesString;dNH-dssC-ssNH-sssCH dO-dssC-ssNH-sssCH dO-dss
C-sssCH-ssCH2 dO-dssC-sssCH-sssCH dssC-ssNH-sssCH-ssCH2 dssC-ssNH-sssCH
-sssCH dssC-sssCH-ssCH2-sssCH dssC-sssCH-sssCH-sOH dssC-sssCH-sssCH-...
;1 1 1 1 1 3 1 1 1 1 2 2 1 1 1 1 2 2 5 1 4 2 4
```

```
FingerprintsVector;TopologicalAtomTorsions:FunctionalClassAtomTypes;11;
NumericalValues;IDsAndValuesString;HBA-NI-None-None HBA-None-HBD-None
HBA.HBD-NI-None-None HBA.HBD-None-None-NI HBA.HBD-None-None-None HBD-
None-HBD-None HBD-None-None-None HBD.PI-None-HBD-None NI-None-None-None
None-HBD-None-None None-None-None-None;2 1 2 1 3 1 7 1 2 5 15
```

```
FingerprintsVector;TopologicalAtomTorsions:MMFF94AtomTypes;14;Numerical
Values;IDsAndValuesString;C=ON-NC=O-CR-CR CGD-NC=N-CR-CR COO-CR-CR-CR
COO-CR-CR-OR CR-C=ON-NC=O-CR CR-CR-COO-O=CO CR-CR-COO-OC=O CR-CR-CR-CR
CR-CR-CR-NC=N CR-CR-CR-NC=O CR-CR-CR-OR CR-NC=N-CGD-N=C CR-NC=N-CGD-NC=N
CR-NC=O-C=ON-O=CN;2 2 2 1 1 2 2 15 3 4 3 1 1 1
```

```
FingerprintsVector;TopologicalAtomTorsions:SLogPAtomTypes;25;Numerical
Values;IDsAndValuesString;C1-C1-C2-C1 C1-C1-C2-CS C1-C2-C5-O2 C1-C2-C5-
O9 C1-C2-CS-C2 C1-C2-CS-N2 C1-C2-CS-O2 C1-C5-N2-CS C1-CS-C2-CS C1-CS-N2
-C5 C2-C1-CS-C2 C2-C1-CS-N2 C2-CS-C2-C5 C2-CS-C2-CS C2-CS-N2-C5 C5-C...
;2 2 1 1 3 2 1 1 2 1 1 1 1 4 3 1 1 1 1 1 4 2 1 1 1
```

```
FingerprintsVector;TopologicalAtomTorsions:SYBYLAtomTypes;12;Numerical
Values;IDsAndValuesString;C.2-C.3-C.3-C.3 C.2-C.3-C.3-O.3 C.2-N.am-C.3-
C.3 C.3-C.2-N.am-C.3 C.3-C.3-C.2-O.co2 C.3-C.3-C.3-C.3 C.3-C.3-C.3-N.am
C.3-C.3-C.3-N.pl3 C.3-C.3-C.3-O.3 C.3-C.3-N.pl3-C.cat C.3-N.am-C.2-O.2
C.3-N.pl3-C.cat-N.pl3;2 1 2 1 4 15 4 3 3 2 1 2
```

```
FingerprintsVector;TopologicalAtomTorsions:TPSAAAtomTypes;8;Numerical
Values;IDsAndValuesString;N10-None-N7-None N7-None-None-None N9-None-N7
-None None-N7-None-None None-N7-None-O3 None-None-None-None None-None-
None-O3 None-None-None-O4;1 7 1 5 1 17 2 6
```

```
FingerprintsVector;TopologicalAtomTorsions:UFFAtomTypes;12;Numerical
Values;IDsAndValuesString;C_2-C_3-C_3-C_3 C_2-C_3-C_3-O_3 C_2-N_3-C_3-C
_3 C_3-C_2-N_3-C_3 C_3-C_3-C_2-O_2 C_3-C_3-C_2-O_3 C_3-C_3-C_3-C_3 C_3-
C_3-C_3-N_3 C_3-C_3-C_3-O_3 C_3-N_3-C_2-N_2 C_3-N_3-C_2-N_3 C_3-N_3-C_2
-O_2;2 1 4 1 2 2 15 7 3 1 1 1
```

## OPTIONS

-a, --AtomIdentifierType *AtomicInvariantsAtomTypes* | *DREIDINGAtomTypes* | *EStateAtomTypes* | *FunctionalClassAtomTypes* | *MMFF94AtomTypes* | *SLogPAtomTypes* | *SYBYLAtomTypes* | *TPSAAAtomTypes* | *UFFAtomTypes*

Specify atom identifier type to use for assignment of initial atom identifier to non-hydrogen atoms during calculation of topological atom torsions fingerprints. Possible values in the current release are: *AtomicInvariantsAtomTypes*, *DREIDINGAtomTypes*, *EStateAtomTypes*, *FunctionalClassAtomTypes*, *MMFF94AtomTypes*, *SLogPAtomTypes*, *SYBYLAtomTypes*, *TPSAAAtomTypes*, *UFFAtomTypes*. Default value: *AtomicInvariantsAtomTypes*.

--AtomicInvariantsToUse "*AtomicInvariant,AtomicInvariant...*"

This value is used during *AtomicInvariantsAtomTypes* value of a, --AtomIdentifierType option. It's a list of comma separated valid atomic invariant atom types.

Possible values for atomic invariants are: *AS, X, BO, LBO, SB, DB, TB, H, Ar, RA, FC, MN, SM*. Default value: *AS,X,BO,H,FC*.

The atomic invariants abbreviations correspond to:

AS = Atom symbol corresponding to element symbol

X<n> = Number of non-hydrogen atom neighbors or heavy atoms  
 BO<n> = Sum of bond orders to non-hydrogen atom neighbors or heavy atoms  
 LBO<n> = Largest bond order of non-hydrogen atom neighbors or heavy atoms  
 SB<n> = Number of single bonds to non-hydrogen atom neighbors or heavy atoms  
 DB<n> = Number of double bonds to non-hydrogen atom neighbors or heavy atoms  
 TB<n> = Number of triple bonds to non-hydrogen atom neighbors or heavy atoms  
 H<n> = Number of implicit and explicit hydrogens for atom  
 Ar = Aromatic annotation indicating whether atom is aromatic  
 RA = Ring atom annotation indicating whether atom is a ring  
 FC<+n/-n> = Formal charge assigned to atom  
 MN<n> = Mass number indicating isotope other than most abundant isotope  
 SM<n> = Spin multiplicity of atom. Possible values: 1 (singlet), 2 (doublet) or 3 (triplet)

Atom type generated by AtomTypes::AtomicInvariantsAtomTypes class corresponds to:

AS.X<n>.BO<n>.LBO<n>.<SB><n>.<DB><n>.<TB><n>.H<n>.Ar.RA.FC<+n/-n>.MN<n>.SM<n>

Except for AS which is a required atomic invariant in atom types, all other atomic invariants are optional. Atom type specification doesn't include atomic invariants with zero or undefined values.

In addition to usage of abbreviations for specifying atomic invariants, the following descriptive words are also allowed:

X : NumOfNonHydrogenAtomNeighbors or NumOfHeavyAtomNeighbors  
 BO : SumOfBondOrdersToNonHydrogenAtoms or SumOfBondOrdersToHeavyAtoms  
 LBO : LargestBondOrderToNonHydrogenAtoms or LargestBondOrderToHeavyAtoms  
 SB : NumOfSingleBondsToNonHydrogenAtoms or NumOfSingleBondsToHeavyAtoms  
 DB : NumOfDoubleBondsToNonHydrogenAtoms or NumOfDoubleBondsToHeavyAtoms  
 TB : NumOfTripleBondsToNonHydrogenAtoms or NumOfTripleBondsToHeavyAtoms  
 H : NumOfImplicitAndExplicitHydrogens  
 Ar : Aromatic  
 RA : RingAtom  
 FC : FormalCharge  
 MN : MassNumber  
 SM : SpinMultiplicity

AtomTypes::AtomicInvariantsAtomTypes module is used to assign atomic invariant atom types.

--FunctionalClassesToUse "*FunctionalClass1,FunctionalClass2...*"

This value is used during *FunctionalClassAtomTypes* value of a, --AtomIdentifierType option. It's a list of comma separated valid functional classes.

Possible values for atom functional classes are: *Ar, CA, H, HBA, HBD, Hal, NI, PI, RA*. Default value [ Ref 24 ]: *HBD,HBA,PI,NI,Ar,Hal*.

The functional class abbreviations correspond to:

HBD: HydrogenBondDonor  
 HBA: HydrogenBondAcceptor  
 PI : PositivelyIonizable  
 NI : NegativelyIonizable  
 Ar : Aromatic  
 Hal : Halogen  
 H : Hydrophobic  
 RA : RingAtom  
 CA : ChainAtom

Functional class atom type specification for an atom corresponds to:

Ar.CA.H.HBA.HBD.Hal.NI.PI.RA

AtomTypes::FunctionalClassAtomTypes module is used to assign functional class atom types. It uses following definitions [ Ref 60-61, Ref 65-66 ]:

HydrogenBondDonor: NH, NH2, OH

HydrogenBondAcceptor: N[!H], O  
 PositivelyIonizable: +, NH2  
 NegativelyIonizable: -, C(=O)OH, S(=O)OH, P(=O)OH

--CompoundID *DataFieldName* or *LabelPrefixString*

This value is --CompoundID mode specific and indicates how compound ID is generated.

For *DataField* value of --CompoundID mode option, it corresponds to datafield label name whose value is used as compound ID; otherwise, it's a prefix string used for generating compound IDs like *LabelPrefixString*<Number>. Default value, *Cmpd*, generates compound IDs which look like *Cmpd*<Number>.

Examples for *DataField* value of --CompoundID mode:

MolID  
 ExtReg

Examples for *LabelPrefix* or *MolNameOrLabelPrefix* value of --CompoundID mode:

Compound

The value specified above generates compound IDs which correspond to *Compound*<Number> instead of default value of *Cmpd*<Number>.

--CompoundIDLabel *text*

Specify compound ID column label for CSV/TSV text file(s) used during *CompoundID* value of --DataFieldsMode option. Default value: *CompoundID*.

--CompoundIDMode *DataField* | *MolName* | *LabelPrefix* | *MolNameOrLabelPrefix*

Specify how to generate compound IDs and write to CSV/TSV text file(s) along with generated fingerprints for *text* | *both* values of --output option: use a *SDFFile(s)* datafield value; use molname line from *SDFFile(s)*; generate a sequential ID with specific prefix; use combination of both *MolName* and *LabelPrefix* with usage of *LabelPrefix* values for empty molname lines.

Possible values: *DataField* | *MolName* | *LabelPrefix* | *MolNameOrLabelPrefix*. Default value: *LabelPrefix*.

For *MolNameAndLabelPrefix* value of --CompoundID mode, molname line in *SDFFile(s)* takes precedence over sequential compound IDs generated using *LabelPrefix* and only empty molname values are replaced with sequential compound IDs.

This is only used for *CompoundID* value of --DataFieldsMode option.

--DataFields "*FieldLabel1,FieldLabel2,...*"

Comma delimited list of *SDFFile(s)* data fields to extract and write to CSV/TSV text file(s) along with generated fingerprints for *text* | *both* values of --output option.

This is only used for *Specify* value of --DataFieldsMode option.

Examples:

Extreg  
 MolID,CompoundName

-d, --DataFieldsMode *All* | *Common* | *Specify* | *CompoundID*

Specify how data fields in *SDFFile(s)* are transferred to output CSV/TSV text file(s) along with generated fingerprints for *text* | *both* values of --output option: transfer all SD data field; transfer SD data files common to all compounds; extract specified data fields; generate a compound ID using molname line, a compound prefix, or a combination of both. Possible values: *All* | *Common* | *specify* | *CompoundID*. Default value: *CompoundID*.

-f, --Filter *Yes* | *No*

Specify whether to check and filter compound data in *SDFFile(s)*. Possible values: *Yes* or *No*. Default value: *Yes*.

By default, compound data is checked before calculating fingerprints and compounds containing atom data corresponding to non-element symbols or no atom data are ignored.

--FingerprintsLabel *text*

SD data label or text file column label to use for fingerprints string in output SD or CSV/TSV text file(s) specified by --output. Default value: *TopologicalAtomTorsionsFingerprints*.

-h, --help

Print this help message.

-k, --KeepLargestComponent *Yes / No*

Generate fingerprints for only the largest component in molecule. Possible values: *Yes or No*. Default value: *Yes*.

For molecules containing multiple connected components, fingerprints can be generated in two different ways: use all connected components or just the largest connected component. By default, all atoms except for the largest connected component are deleted before generation of fingerprints.

--OutDelim *comma | tab | semicolon*

Delimiter for output CSV/TSV text file(s). Possible values: *comma, tab, or semicolon* Default value: *comma*.

--output *SD | text | both*

Type of output files to generate. Possible values: *SD, text, or both*. Default value: *text*

-o, --overwrite

Overwrite existing files.

-q, --quote *Yes / No*

Put quote around column values in output CSV/TSV text file(s). Possible values: *Yes or No*. Default value: *Yes*

-r, --root *RootName*

New file name is generated using the root: <Root>.<Ext>. Default for new file names: <SDFFileName><TopologicalAtomTorsionsFP>.<Ext>. The file type determines <Ext> value. The sdf, csv, and tsv <Ext> values are used for SD, comma/semicolon, and tab delimited text files, respectively. This option is ignored for multiple input files.

-v, --VectorStringFormat *IDsAndValuesString | IDsAndValuesPairsString | ValuesAndIDsString | ValuesAndIDsPairsString*

Format of fingerprints vector string data in output SD or CSV/TSV text file(s) specified by --output used during *PathLengthCount* value of -m, --mode option. Possible values: *IDsAndValuesString | IDsAndValuesPairsString | ValuesAndIDsString | ValuesAndIDsPairsString*. Default value: *IDsAndValuesString*.

Examples:

```
FingerprintsVector;TopologicalAtomTorsions:AtomicInvariantsAtomTypes;11;
NumericalValues;IDsAndValuesString;C.X1.B01.H3-C.X3.B04-C.X2.B03.H1-
N.X2.B02.H1 C.X1.B01.H3-C.X3.B04-C.X3.B04-C.X2.B03.H1 C.X1.B01.H3-
C.X3.B04-C.X3.B04-O.X1.B02 C.X2.B03.H1-C.X2.B03.H1-C.X3.B04-C.X3.B04
C.X2.B03.H1-C.X2.B03.H1-C.X3.B04-O.X1.B02...;
1 1 1 1 1 1 1 1 1 1 1
```

```
FingerprintsVector;TopologicalAtomTorsions:AtomicInvariantsAtomTypes;11;
NumericalValues;IDsAndValuesPairsString;C.X1.B01.H3-C.X3.B04-C.X2.B03.H1-
N.X2.B02.H1 1 C.X1.B01.H3-C.X3.B04-C.X3.B04-C.X2.B03.H1 1 C.X1.B01.H3-
C.X3.B04-C.X3.B04-O.X1.B02 1 C.X2.B03.H1-C.X2.B03.H1-C.X3.B04-
C.X3.B04 1 C.X2.B03.H1-C.X2.B03.H1-C.X3.B04-O.X1.B02 1 C.X2.B03.H1-
C.X2.B03.H1-N.X2.B02.H1-C.X2.B03.H1 1 C.X2.B03.H1-C.X3.B04-C.X3.B04-
C.X2.B03.H1 1...
```

-w, --WorkingDir *DirName*

Location of working directory. Default value: current directory.

## EXAMPLES

To generate topological atom torsions fingerprints using atomic invariants atom types in *IDsAndValuesString* format and create a *SampleTATFP.csv* file containing sequential compound IDs along with fingerprints vector strings data, type:

---

```
% TopologicalAtomTorsionsFingerprints.pl -r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using atomic invariants atom types in IDsAndValuesString format and create both SampleTATFP.csv and SampleTATFP.sdf files containing sequential compound IDs in CSV file along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl --output both -r SampleTATFP
-o Sample.sdf
```

To generate topological atom torsions fingerprints using atomic invariants atom types in IDsAndValuesPairsString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl --VectorStringFormat
IDsAndValuesPairsString -r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using DREIDING atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a DREIDINGAtomTypes
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using E-state atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a EStateAtomTypes
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using functional class atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a FunctionalClassAtomTypes
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using MMFF94 atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a MMFF94AtomTypes
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using SLogP atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a SLogPAtomTypes
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using SYBYL atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a SYBYLAtomTypes
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using TPSA atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a TPSAAtomTypes
```

---

```
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using UFF atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a UFFAtomTypes
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using only AS,X atomic invariants atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing sequential compound IDs along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a AtomicInvariantsAtomTypes
--AtomicInvariantsToUse "AS,X" -r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using atomic invariants atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing compoundID from molecule name line along with fingerprints vector strings, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a AtomicInvariantsAtomTypes
--DataFieldsMode CompoundID -CompoundIDMode MolName
-r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using atomic invariants atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing compound IDs using specified data field along with fingerprints vector strings, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a AtomicInvariantsAtomTypes
--DataFieldsMode CompoundID -CompoundIDMode DataField --CompoundID
Mol_ID -r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using atomic invariants atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing compound ID using combination of molecule name line and an explicit compound prefix along with fingerprints vector strings data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a AtomicInvariantsAtomTypes
--DataFieldsMode CompoundID -CompoundIDMode MolnameOrLabelPrefix
--CompoundID Cmpd --CompoundIDLabel MolID -r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using atomic invariants atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing specific data fields columns along with fingerprints vector strings, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a AtomicInvariantsAtomTypes
--DataFieldsMode Specify --DataFields Mol_ID -r SampleTATFP
-o Sample.sdf
```

To generate topological atom torsions fingerprints using atomic invariants atom types in IDsAndValuesString format and create a SampleTATFP.csv file containing common data fields columns along with fingerprints vector strings, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a AtomicInvariantsAtomTypes
--DataFieldsMode Common -r SampleTATFP -o Sample.sdf
```

To generate topological atom torsions fingerprints using atomic invariants atom types in IDsAndValuesString format and create both SampleTATFP.csv and SampleTATFP.sdf files containing all data fields columns in CSV file along with fingerprints data, type:

```
% TopologicalAtomTorsionsFingerprints.pl -a AtomicInvariantsAtomTypes
--DataFieldsMode All --output both -r SampleTATFP
-o Sample.sdf
```

#### AUTHOR

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#### SEE ALSO

InfoFingerprintsSDFFiles.pl, InfoFingerprintsTextFiles.pl, SimilarityMatrixSDFFiles.pl, SimilarityMatrixTextFiles.pl, AtomNeighborhoodsFingerprints.pl, ExtendedConnectivityFingerprints.pl, MACCSKeysFingerprints.pl, PathLengthFingerprints.pl, TopologicalAtomPairsFingerprints.pl, TopologicalPharmacophoreAtomPairsFingerprints.pl, TopologicalPharmacophoreAtomTripletsFingerprints.pl

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