

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, TPSAAAtomTypes, 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 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 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 --CompoundIDMode specific and indicates how compound ID is generated.

For *DataField* value of --CompoundIDMode 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 --CompoundIDMode:

```
MolID
ExtReg
```

Examples for *LabelPrefix* or *MolNameOrLabelPrefix* value of --CompoundIDMode:

```
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 `--CompoundIDMode`, *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.BO1.H3-C.X3.BO4-C.X2.BO3.H1-
N.X2.BO2.H1 C.X1.BO1.H3-C.X3.BO4-C.X3.BO4-C.X2.BO3.H1 C.X1.BO1.H3-
C.X3.BO4-C.X3.BO4-O.X1.BO2 C.X2.BO3.H1-C.X2.BO3.H1-C.X3.BO4-C.X3.BO4
C.X2.BO3.H1-C.X2.BO3.H1-C.X3.BO4-O.X1.BO2...;
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
```

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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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