NAME

ElementalAnalysisSDFiles.pl - Perform elemental analysis using formula data field in SDFile(s)

SYNOPSIS

ElementalAnalysisSDFiles.pl SDFile(s)...

ElementalAnalysisSDFiles.pl [-d, --detail infolevel] [--fast] [--formulafield SD data field name] [-f, --formulamode *DataField | StructureData*] [--formulaout yes or no] [-m, --mode All | "ElementalAnalysis, [MolecularWeight, ExactMass]"] [-h, --help] [-o, --overwrite] [-r, --root rootname] [-v --valuefieldnames Name, Label, [Name, Label,...]] [-w, --workingdir dirname] SDFile(s)...

DESCRIPTION

Perform elemental analysis using molecular formula specified by a data field name or generated from structure data in *SDFile(s)*.

In addition to straightforward molecular formulas - H2O, HCI, C3H7O2N - other supported variations are: Ca3(PO4)2, [PCI4]+, [Fe(CN)6]4-, C37H42N2O6+2, Na2CO3.10H2O, 8H2S.46H2O, and so on. Charges are simply ignored. Isotope symbols in formulas specification, including D and T, are not supported.

The file names are separated by space. 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.

OPTIONS

-d, --detail infolevel

Level of information to print about compound records being ignored. Default: 1. Possible values: 1, 2 or 3.

--fast

In this mode, the formula data field specified using -f, --formulafield option is assumed to contain valid molecular formula data and initial formula validation check is skipped.

--formulafield SD data field name

SDFile(s) data field name containing molecular formulas used for performing elemental analysis during DataField value of -f, --formulamode option. Default value: SD data field containing the word formula in its name.

This option is ignore during StructureData value of -f, --formulamode option.

-f, --formulamode DataField | StructureData

Specify source of molecular formula used for performing elemental analysis: retrieve formula using *SDFile(s)* data field name or generate formula from structure. Possible values: *DataField or StructureData*. Default value: *DataField*.

--formulaout yes or no

Specify whether to write out formula to SD file during *StructureData* value of -f, --formulamode option. Possible values: *Yes or No.* Default: *No.*

-m, --mode All | "ElementalAnalysis, [MolecularWeight, ExactMass]"

Specify what values to calculate using molecular formula data field or structure data from *SDFile(s)*: calculate all supported values or specify a comma delimited list of values. Possible values: *All | "Elemental Analysis, [Molecular Weight, Exact Mass]"*. Default: *All*

-h, --help

Print this help message.

-o, --overwrite

Overwrite existing files.

-p, --precision *number*

Precision of calculated values in the output file. Default: up to 2 decimal places. Valid values: positive integers.

-r, --root rootname

New SD file name is generated using the root: <Root>.<Ext>. Default new file name: <InitialSDFileName>ElementalAnalysis.<Ext>. This option is ignored for multiple input files.

-v --valuefieldnames Name, Label, [Name, Label, ...]

Specify SD data field names to use for calculated values. In general, it's a comma delimited list of value name and SD field name pairs. Supported value names: *ElementalAnalysis, MolecularWeight, ExactMass, and MolecularFormula*. Default labels: *ElementalAnalysis, MolecularWeight, ExactMass, and MolecularFormula*.

MolecularFormula label is only used during StructureData value of -f, --formulamode option.

-w, --workingdir dirname

Location of working directory. Default: current directory.

FXAMPLES

To perform elemental analysis, calculate molecular weight and exact mass using SD field name value with the word Formula in its name and generate a new SD file NewSample1.sdf, type:

% ElementalAnalysisSDFiles.pl -o -r NewSample1 Sample1.sdf

To perform elemental analysis, calculate molecular weight and exact mass using structure data in SD file and generate a new SD file NewSample1.sdf, type:

```
% ElementalAnalysisSDFiles.pl --formulamode StructureData -o
-r NewSamplel Sample1.sdf
```

To perform elemental analysis using formulas in SD field name Formula, use field name Analysis for calculated data, and generate a new SD file NewSample1.sdf, type:

```
% ElementalAnalysisSDFiles.pl --m ElementalAnalysis --formulafield
Formula --valuefieldnames "ElementalAnalysis,Analysis" -o
-r NewSamplel Samplel.sdf
```

To calculate molecular weight, using formulas in SD field name Formula, with four decimal precision and generate a new SD file NewSample1.sdf, type

```
% ElementalAnalysisSDFiles.pl --m MolecularWeight --formulafield
Formula --precision 4 -o -r NewSamplel Samplel.sdf
```

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SEE ALSO

AnalyzeSDFilesData.pl, InfoSDFiles.pl, ExtractFromSDFiles.pl

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