NAME

SDFileUtil

SYNOPSIS

```
use SDFileUtil ;
use SDFileUtil qw(:all);
```

DESCRIPTION

SDFileUtil module provides the following functions:

GenerateCmpdAtomAliasPropertyLines, GenerateCmpdAtomLine, GenerateCmpdBondLine, GenerateCmpdChargePropertyLines, GenerateCmpdCommentsLine, GenerateCmpdCountsLine, GenerateCmpdDataHeaderLabelsAndValuesLines, GenerateCmpdIsotopePropertyLines, GenerateCmpdMiscInfoLine, GenerateCmpdMolNameLine, GenerateCmpdRadicalPropertyLines, GenerateEmptyCtabBlockLines, GenerateEmptyCtabBlockLines, GenerateMiscLineDateStamp, GetAllAndCommonCmpdDataHeaderLabels, GetCmpdDataHeaderLabels, GetCmpdDataHeaderLabelsAndValues, GetCmpdFragments, GetCtabLinesCount, GetInvalidAtomNumbers, GetUnknownAtoms, InternalBondOrderToMDLBondType, InternalBondStereochemistryToMDLBondStereo, InternalChargeToMDLCharge, InternalSpinMultiplicityToMDLRadical, IsCmpd2D, IsCmpd3D, MDLBondStereoToInternalBondStereochemistry, MDLBondTypeToInternalBondOrder, MDLChargeToInternalCharge, MDLRadicalToInternalSpinMultiplicity, ParseCmpdAtomAliasPropertyLine, ParseCmpdAtomLine, ParseCmpdCountsLine, ParseCmpdBondLine, ParseCmpdCountsLine, ParseCmpdRadicalPropertyLine, ParseCmpdRadicalPropertyLine, ReadCmpdString, RemoveCmpdDataHeaderLabelAndValue, WashCmpd

METHODS

GenerateCmpdAtomAliasPropertyLines

```
@Lines = GenerateCmpdAtomAliasPropertyLines($AliasValuePairsRef);
```

Returns a formatted atom alias property lines corresponding to successive pairs of atom number and alias values specified by a refernce to an array. Two lines are generate for each atom number and alias value pairs: First line - A <AtomNum>; Second line: <AtomAlias>.

GenerateCmpdAtomLine

Returns a formatted atom data line containing all the input values.

${\tt GenerateCmpdBondLine}$

Returns a formatted bond data line containing all the input values.

${\tt GenerateCmpdChargePropertyLines}$

```
@Lines = GenerateCmpdChargePropertyLines($ChargeValuePairsRef);
```

Returns a formatted M CHG property lines corresponding to successive pairs of atom number and charge values specified by a refernce to an array.

GenerateCmpdCommentsLine

```
$Line = GenerateCmpdCommentsLine($Comments);
```

Returns a formatted comments data line.

GenerateCmpdCountsLine

Returns a formatted line containing all the input values. The default values of 999 and V2000 are used for

PropertyCount and Version.

GenerateCmpdDataHeaderLabelsAndValuesLines

Returns formatted data lines containing header label and values lines corresponding to all data header labels in array reference *DataHeaderLabelsRef* with values in hash reference *DataHeaderLabelsAndValuesRef*. By default, data header labels are not sorted and correspond to the label order in array reference *DataHeaderLabelsRef*.

GenerateCmpdIsotopePropertyLines

```
@Lines = GenerateCmpdIsotopePropertyLines($IsotopeValuePairsRef);
```

Returns a formatted M ISO property lines corresponding to successive pairs of atom number and isotope values specified by a refernce to an array.

GenerateCmpdMiscInfoLine

Returns a formatted line containing specified user initial, program name, date and code. Default values are: *ProgramName - MayaChem; UserInitial - NULL; Code - 2D.*

GenerateCmpdMoINameLine

```
$Line = GenerateCmpdMolNameLine($MolName);
```

Returns a formatted molecule name data line.

GenerateCmpdRadicalPropertyLines

```
@Lines = GenerateCmpdRadicalPropertyLines($RadicalValuePairsRef);
```

Returns a formatted M CHG property lines corresponding to successive pairs of atom number and multiplicity values specified by a refernce to an array.

GenerateEmptyCtabBlockLines

```
$Lines = GenerateCmpdMiscInfoLine([$Date]);
```

Returns formatted lines representing empty CTAB block.

GenerateMiscLineDateStamp

```
$Line = GenerateMiscLineDateStamp();
```

Returns date stamp for misc line.

${\sf GetAllAndCommonCmpdDataHeaderLabels}$

```
($CmpdCount, $DataFieldLabelsArrayRef,
  $CommonDataFieldLabelsArrayRef) =
   GetAllAndCommonCmpdDataHeaderLabels(\*SDFILE);
```

Returns number of comopunds, a reference to an array containing all unique data header label and a reference to an array containing common data field labels for all compounds in SD file.

GetCmpdDataHeaderLabels

```
(@Labels) = GetCmpdDataHeaderLabels(\@CmpdLines);
```

Returns an array containg data header labels for a compound

GetCmpdDataHeaderLabelsAndValues

```
(%DataValues) = GetCmpdDataHeaderLabelsAndValues(\@CmpdLines);
```

Returns a hash conating data header labes and values for a compound.

GetCmpdFragments

```
($FragmentCount, $FragmentString) = GetCmpdFragments(\@CmpLines);
```

Figures out the number of disconnected fragments and return their values along with the atom numbers in a string delimited by new line character. Fragment data in FragmentString is sorted on based on its size.

GetCtabLinesCount

```
$CtabLinesCount = GetCtabLinesCount(\@CmpdLines);
```

Returns number of lines present between the 4th line and the line containg "M END".

GetInvalidAtomNumbers

```
($InvalidAtomNumbersCount, $InvalidAtomNumbers, $InvalidAtomNumberLines) =
GetInvalidAtomNumbers(\@CmpdLines);
```

Returns a list of values containing information about invalid atom numbers present in block or atom property lines.

GetUnknownAtoms

```
($UnknownAtomCount, $UnknownAtoms, $UnknownAtomLines) =
GetUnknownAtoms(\@CmpdLines);
```

Returns a list of values containing information about atoms which contain special element symbols not present in the periodic table.

InternalBondOrderToMDLBondType

```
$MDLBondType = InternalBondOrderToMDLBondType($InternalBondOrder);
```

Returns value of MDLBondType corresponding to InternalBondOrder.

 ${\tt InternalBondOrder} \quad {\tt MDLBondType}$

1	1
2	2
3	3
1.5	4

Internal Bond Stere ochem is try ToMDL Bond Stere o

Returns value of MDLBondStereo corresponding to InternalBondStereo using following mapping:

InternalBondStereo MDLBondStereo

Up	1
UpOrDown	4
Down	6
CisOrTrans	3
Other	0

Internal Charge To MDL Charge

```
$MDLCharge = InternalChargeToMDLCharge($InternalCharge);
```

Returns value of $\emph{MDLCharge}$ corresponding to $\emph{InternalCharge}$ using following mapping:

InternalCharge MDLCharge

3	1
2	2
1	3

-1 5 -2 6 -3 7

InternalSpinMultiplicityToMDLRadical

Returns value of MDLRadical corresponding to Internal SpinMultiplicity. These value are equivalent.

MDLBondStereoToInternalBondType

```
$InternalBondType = MDLBondStereoToInternalBondType($MDLBondStereo);
```

Returns value of *InternalBondType* corresponding to *MDLBondStereo* using mapping shown for InternalBondTypeToMDLBondStereo function.

IsCmpd2D

```
$Status = IsCmpd2D();
```

Returns 1 or 0 based on whether z-coordinate of any atom is non-zero.

IsCmpd3D

```
$Status = IsCmpd3D();
```

Returns 1 or 0 based on whether z-coordinate of any atom is non-zero.

MDLBondStereoToInternalBondStereochemistry

Returns value of $\it InternalBondStereo$ corresponding to $\it MDLBondStereo$ using mapping shown for InternalBondStereochemistryToMDLBondStereo function.

MDLBondTypeToInternalBondOrder

```
$InternalBondOrder = MDLBondTypeToInternalBondOrder($MDLBondType);
```

Returns value of *InternalBondOrder* corresponding to *MDLBondType* using mapping shown for InternalBondOrderToMDLBondType function.

MDLChargeToInternalCharge

```
$InternalCharge = MDLChargeToInternalCharge($MDLCharge);
```

Returns value of *\$InternalCharge* corresponding to *MDLCharge* using mapping shown for InternalChargeToMDLCharge function.

MDLRadicalToInternalSpinMultiplicity

Returns value of InternalSpinMultiplicity corresponding to MDLRadical. These value are equivalent.

${\tt ParseCmpdAtomAliasPropertyLine}$

Parses atom alias properly lines in CTAB generic properties block and returns an array with successive pairs of values corresponding to atom number and its alias.

ParseCmpdAtomLine

```
($AtomSymbol, $AtomX, $AtomY, $AtomZ, $MassDifference, $Charge,
  $StereoParity) = ParseCmpdAtomLine($AtomDataLine);
```

Parses compound data line containing atom information and returns a list of values.

ParseCmpdBondLine

```
($FirstAtomNum, $SecondAtomNum, $BondType) =
ParseCmpdBondLine($BondDataLine);
```

Parses compound data line containing bond information and returns a list of values.

ParseCmpdCommentsLine

```
$Comments = ParseCmpdCommentsLine($CommentsDataLine);
```

Returns the comment string.

ParseCmpdChargePropertyLine

Parses charge properly line in CTAB generic properties block and returns an array with successive pairs of values corresponding to atom number and its charge.

ParseCmpdCountsLine

```
($AtomCount, $BondCount, $ChiralFlag, $PropertyCount, $Version) =
ParseCmpdCountsLine(\@CountDataLines);
```

Returns a list of values containing count information.

ParseCmpdMiscInfoLine

```
($UserInitial, $ProgramName, $Date, $Code, $ScalingFactor1, $ScalingFactor2,
$Energy, $RegistryNum) = ParseCmpdMiscInfoLine($Line);
```

Returns a list of values containing miscellaneous information.

ParseCmpdI sotopePropertyLine

Parses isotopic properly line in CTAB generic properties block and returns an array with successive pairs of values corresponding to atom number and absolute mass of atom isotope.

ParseCmpdMoINameLine

```
$MolName = ParseCmpdMolNameLine($Line);
```

Returns a string containing molecule name.

ParseCmpdRadicalPropertyLine

Parses radical properly line in CTAB generic properties block and returns an array with successive pairs of values corresponding to atom number and radical number value.

RemoveCmpdDataHeaderLabelAndValue

Returns a NewCmpdString after removing DataHeaderLabel along with its value from CmpdString.

ReadCmpdString

```
$CmpdString = ReadCmpdString(\*SDFILEHANDLE);
```

Returns a string containing all the data lines for the next available compound in an already open file indicated by SDFILEHANDLE. A NULL string is returned on EOF.

WashCmpd

```
($FragmentCount, $Fragments, $WashedCmpdString) =
WashCmpd(\@CmpdLines);
```

Figures out the number of disconnected fragments and return their values along with the atom numbers in a string delimited by new line character. Fragment data in FragmentString is sorted on based on its size.

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SEE ALSO

TextUtil.pm

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