### NAME

AnalyzeTextFilesData.pl - Analyze numerical coulmn data in TextFile(s)

### **SYNOPSIS**

AnalyzeTextFilesData.pl TextFile(s)...

AnalyzeTextFilesData.pl [-c, --colmode colnum | collabel] [--columns "colnum,[colnum,...]" |
"collabel,[collabel,...]" | All] [--columnpairs "colnum,colnum,[colnum,colnum]..." |
"collabel,collabel,[collabel,collabel]..." | AllPairs] [-d, --detail infolevel] [-f, --fast] [--frequencybins number |
"number,number,[number,...]"] [-h, --help] [--indelim comma | semicolon] [--klargest number] [--ksmallest
number] [-m, --mode DescriptiveStatisticsBasic | DescriptiveStatisticsAll | All | "function1, [function2,...]"] [-o,
--overwrite] [--outdelim comma | tab | semicolon] [-p, --precision number] [-q, --quote yes | no] [-r, --root
rootname] [--trimfraction number] [-w, --workingdir dirname] TextFiles(s)...

### **DESCRIPTION**

Anaylze numerical column data in *TextFile(s)* using a combination of various statistical functions; Non-numerical values are simply ignored. For *Correlation, RSquare, and Covariance* analysis, the count of valid values in specified column pair must be same; otherwise, column pair is ignored. The file names are separated by space. The valid file extensions are .csv and .tsv for comma/semicolon and tab delimited text files respectively. All other file names are ignored. All the text files in a current directory can be specified by \*.csv, \*.tsv, or the current directory name. The --indelim option determines the format of *TextFile(s)*. Any file which doesn't correspond to the format indicated by --indelim option is ignored.

#### **OPTIONS**

#### -c, --colmode colnum | collabel

Specify how columns are identified in TextFile(s): using column number or column label. Possible values: colnum or collabel. Default value: colnum.

### --columns "colnum,[colnum,...]" | "collabel,[collabel]..." | All

This value is mode specific. It's a list of comma delimited columns to use for data analysis. Default value: First column.

This value is ignored during *Correlation/Pearson Correlation* and *Covariance* data analysis; -coulmnparis option is used instead.

For colnum value of -c, --colmode option, input values format is: colnum,colnum,.... Example:

1,3,5

For *collabel* value of -c, --colmode option, input values format is: *collabel,collabel,...* Example: ALogP,MolWeight,EC50

--columnpairs "colnum,colnum,[colnum,colnum,...]" | "collabel,collabel,[collabel,collabel,...]" | AllPairs

This value is mode specific and is only used for *Correlation, PearsonCorrelation, or Covariance* value of -m, --mode option. It is a comma delimited list of column pairs to use for data analysis during *Correlation* and *Covariance* calculations. Default value: *First column, Second column*.

For *colnum* value of -c, --colmode option, input values format is: *colnum,colnum,[colnum,colnum]....* Example: 1,3,5,6,1,6

For *collabel* value of -c, --colmode option, input values format is: *collabel,collabel,collabel,collabelj...* Example: MolWeight, EC50, NumN+O, PSA

For AllPairs value of --columnparis option, all column pairs are used for Correlation and Covariance calculations.

## -d, --detail infolevel

Level of information to print about column values being ignored. Default: 1. Possible values: 1, 2, 3, or 4.

#### -f, --fast

In this mode, all the columns specified for analysis are assumed to contain numerical data and no checking is performed before analysis. By default, only numerical data is used for analysis; other types of column

data is ignored.

### --frequencybins number | "number,number,[number,...]"

Specify number of bins or bin range to use for frequency analysis. Default value: 10

Number of bins value along with the smallest and largest value for a column is used to group the column values into different groups.

The bin range list is used to group values for a column into different groups; It must contain values in ascending order. Examples:

```
10,20,30
0.1,0.2,0.3,0.4,0.5
```

The frequency value calculated for a specific bin corresponds to all the column values which are greater than the previous bin value and less than or equal to the current bin value.

### -h, --help

Print this help message.

### --indelim comma | semicolon

Input delimiter for CSV *TextFile(s)*. Possible values: *comma or semicolon*. Default value: *comma*. For TSV files, this option is ignored and *tab* is used as a delimiter.

#### --klargest number

Kth largest value to find by KLargest function. Default value: 2 Valid values: positive integers.

#### --ksmallest number

Kth smallest value to find by KSmallest function. Default value: 2. Valid values: positive integers.

#### -m, --mode DescriptiveStatisticsBasic | DescriptiveStatisticsAll | All | "function1, [function2,...]"

Specify how to analyze data in TextFile(s): calculate basic or all descriptive statistics; or use a comma delimited list of supported statistical functions. Possible values: DescriptiveStatisticsBasic | DescriptiveStatisticsAll | "function1,[function2]...". Default value: DescriptiveStatisticsBasic

DescriptiveStatisticsBasic includes these functions: Count, Maximum, Minimum, Mean, Median, Sum, StandardDeviation, StandardError, Variance.

DescriptiveStatisticsAll, in addition to DescriptiveStatisticsBasic functions, includes: GeometricMean, Frequency, HarmonicMean, KLargest, KSmallest, Kurtosis, Mode, RSquare, Skewness, TrimMean.

All uses complete list of supported functions: Average, AverageDeviation, Correlation, Count, Covariance, GeometricMean, Frequency, HarmonicMean, KLargest, KSmallest, Kurtosis, Maximum, Minimum, Mean, Median, Mode, RSquare, Skewness, Sum, SumOfSquares, StandardDeviation, StandardDeviationN, StandardError, StandardScores, StandardScoresN, TrimMean, Variance, VarianceN. The function names ending with N calculate corresponding values assuming an entire population instead of a population sample.

Here are the formulas for these functions:

Average: See Mean

AverageDeviation: SUM( ABS(x[i] - Xmean) ) / n

Correlation: See Pearson Correlation

Covariance: SUM((x[i] - Xmean)(y[i] - Ymean))/n

GeometricMean: NthROOT( PRODUCT(x[i]) )

HarmonicMean: 1 / (SUM(1/x[i]) / n)

Mean: SUM(x[i])/n

Median: Xsorted[(n - 1)/2 + 1] for even values of n; (Xsorted[n/2] + Xsorted[n/2 + 1])/2 for odd values of

Kurtosis:  $[ \{n(n + 1)/(n - 1)(n - 2)(n - 3)\} SUM\{ ((x[i] - Xmean)/STDDEV)^4 \} ] - \{3((n - 1)^2)\}/\{(n - 2)(n-3)\}$ 

PearsonCorrelation: SUM( (x[i] - Xmean)(y[i] - Ymean)) / SQRT( SUM(  $(x[i] - Xmean)^2$ ) (SUM(  $(y[i] - Ymean)^2$ )))

RSquare: PearsonCorrelation^2

```
Skewness: \{n/(n-1)(n-2)\}\ SUM\{(x[i]-Xmean)/STDDEV)^3\} StandardDeviation: SQRT ( SUM( (x[i]-Mean)^2) / (n-1) ) StandardDeviationN: SQRT ( SUM( (x[i]-Mean)^2) / (n-1) ) StandardError: StandardDeviation / SQRT( (n-1) ) StandardScore: (x[i]-Mean) / (n-1) StandardScoreN: (x[i]-Mean) / (n-1) Variance: SUM( (x[i]-Xmean)^2 / (n-1) ) VarianceN: SUM( (x[i]-Xmean)^2 / (n-1)
```

### -o, --overwrite

Overwrite existing files.

### --outdelim comma | tab | semicolon

Output text file delimiter. Possible values: comma, tab, or semicolon Default value: comma.

#### -p, --precision number

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

## -q, --quote yes | no

Put quotes around column values in output text file. Possible values: yes or no. Default value: yes.

#### -r, --root rootname

New text file name is generated using the root: <Root>.<Ext>. Default new file name: <InitialTextFileName><Mode>.<Ext>. Based on the specified analysis, <Mode> corresponds to one of these values: DescriptiveStatisticsBasic, DescriptiveStatisticsAll, AllStatistics, SpecifiedStatistics, Covariance, Correlation, Frequency, or StandardScores. The csv, and tsv <Ext> values are used for comma/semicolon, and tab delimited text files respectively. This option is ignored for multiple input files.

### --trimfraction number

Fraction of data to exclude from the top and bottom of the data set during TrimMean calculation. Default value: 0.1. Valid values: > 0 and < 1.

### -w --workingdir text

Location of working directory. Default: current directory.

### **EXAMPLES**

To calculate basic statistics for data in first column and generate a NewSample1DescriptiveStatisticsBasic.csv file, type:

```
% AnalyzeTextFilesData.pl -o -r NewSample1 Sample1.csv
```

To calculate basic statistics for data in third column and generate a NewSample1DescriptiveStatisticsBasic.csv file, type:

```
% AnalyzeTextFilesData.pl --columns 3 -o -r NewSample1 Sample1.csv
```

To calculate basic statistics for data in MolWeight column and generate a NewSample1DescriptiveStatisticsBasic.csv file, type:

```
% AnalyzeTextFilesData.pl -colmode collabel --columns MolWeight -o
-r NewSamplel Samplel.csv
```

To calculate all available statistics for data in third column and all column pairs, and generate NewSample1DescriptiveStatisticsAll.csv, NewSample1CorrelationMatrix.csv, NewSample1CorrelationMatrix.csv, and NewSample1MolWeightFrequencyAnalysis.csv files, type:

```
% AnalyzeTextFilesData.pl -m DescriptiveStatisticsAll --columns 3 -o
--columnpairs AllPairs -r NewSamplel Sample1.csv
```

To compute frequency distribution of data in third column into five bins and generate NewSample1MolWeightFrequencyAnalysis.csv, type:

```
% AnalyzeTextFilesData.pl -m Frequency --frequencybins 5 --columns 3
-o -r NewSamplel Samplel.csv
```

To compute frequency distribution of data in third column into specified bin range values, and generate NewSample1MolWeightFrequencyAnalysis.csv, type:

```
% AnalyzeTextFilesData.pl -m Frequency --frequencybins "100,200,400"
--columns 3 -o -r NewSamplel Samplel.csv
```

To calculate all available statistics for data in all columns and column pairs, type:

```
% AnalyzeTextFilesData.pl -m All --columns All --columnpairs
AllPairs -o -r NewSamplel Sample1.csv
```

#### **AUTHOR**

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# SEE ALSO

JoinTextFiles.pl, MergeTextFilesWithSD.pl, ModifyTextFilesFormat.pl, SplitTextFiles.pl, TextFilesToHTML.pl

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