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 specifed 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,collabel]...* 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 n.

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: $SQRT (SUM((x[i] - Mean)^2) / n)$

StandardError: $StandardDeviation / SQRT(n)$

StandardScore: $(x[i] - Mean) / (n - 1)$

StandardScoreN: $(x[i] - Mean) / n$

Variance: $SUM((x[i] - Xmean)^2 / (n - 1))$

VarianceN: $SUM((x[i] - Xmean)^2 / n)$

-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 NewSample1 Sample1.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 NewSample1 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 NewSample1 Sample1.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 NewSample1 Sample1.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 NewSample1 Sample1.csv
```

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SEE ALSO

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

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