**spearman** — Spearman’s and Kendall’s correlations

### Description

`spearman` displays Spearman’s rank correlation coefficients for all pairs of variables in `varlist` or, if `varlist` is not specified, for all the variables in the dataset. When there are two variables, an exact \( p \)-value can be calculated optionally using permutations.

`ktau` displays Kendall’s rank correlation coefficients between the variables in `varlist` or, if `varlist` is not specified, for all the variables in the dataset. `ktau` is intended for use on small- and moderate-sized datasets; it requires considerable computation time for larger datasets.

### Quick start

**Spearman’s rank correlation coefficient with approximate \( p \)-value for \( v1 \) and \( v2 \)**

`spearman v1 v2`

As above, but report an exact \( p \)-value calculated using Monte Carlo permutations

`spearman v1 v2, exact`

As above, but perform 100,000 Monte Carlo permutations rather than the default of 10,000, and set the random-number seed for reproducibility

`spearman v1 v2, exact(montecarlo, reps(100000) rseed(1234))`

Display Spearman’s rank correlation coefficients in a matrix for all pairs of \( v1 \), \( v2 \), and \( v3 \)

`spearman v1 v2 v3`

Display \( p \)-values as well as correlation coefficients

`spearman v1 v2 v3, stats(rho p)`

Same as above, but perform Bonferroni’s adjustment to \( p \)-values

`spearman v1 v2 v3, stats(rho p) bonferroni`

Kendall’s rank correlation coefficients, scores, and standard errors of the scores for pairs of \( v1 \), \( v2 \), and \( v3 \)

`ktau v1 v2 v3, stats(taua taub score se)`

Same as above, but use pairwise instead of casewise deletion

`ktau v1 v2 v3, stats(taua taub score se) pw`

### Menu

**spearman**

Statistics > Nonparametric analysis > Tests of hypotheses > Spearman’s rank correlation

**ktau**

Statistics > Nonparametric analysis > Tests of hypotheses > Kendall’s rank correlation
## Syntax

### Spearman’s rank correlation coefficients

```
spearman [varlist] [if] [in] [ , spearman_options ]
```

### Kendall’s rank correlation coefficients

```
ktau [varlist] [if] [in] [ , ktau_options ]
```

### spearman_options

<table>
<thead>
<tr>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>stat(list)</td>
</tr>
<tr>
<td>print(#)</td>
</tr>
<tr>
<td>star(#)</td>
</tr>
<tr>
<td>bonferroni</td>
</tr>
<tr>
<td>sidak</td>
</tr>
<tr>
<td>pw</td>
</tr>
<tr>
<td>matrix</td>
</tr>
<tr>
<td>exact( exact_specs )</td>
</tr>
</tbody>
</table>

### ktau_options

<table>
<thead>
<tr>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>stat(list)</td>
</tr>
<tr>
<td>print(#)</td>
</tr>
<tr>
<td>star(#)</td>
</tr>
<tr>
<td>bonferroni</td>
</tr>
<tr>
<td>sidak</td>
</tr>
<tr>
<td>pw</td>
</tr>
<tr>
<td>matrix</td>
</tr>
</tbody>
</table>

by and collect are allowed with `spearman` and `ktau`; see [U] 11.1.10 Prefix commands.

where the elements of `spearman_list` may be

<table>
<thead>
<tr>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>rho</td>
</tr>
<tr>
<td>obs</td>
</tr>
<tr>
<td>p</td>
</tr>
</tbody>
</table>

and the elements of `ktau_list` may be

<table>
<thead>
<tr>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>taua</td>
</tr>
<tr>
<td>taub</td>
</tr>
<tr>
<td>score</td>
</tr>
<tr>
<td>se</td>
</tr>
<tr>
<td>obs</td>
</tr>
<tr>
<td>p</td>
</tr>
</tbody>
</table>
Options for spearman

`stats(spearman_list)` specifies the statistics to be displayed in the matrix of output. `stats(rho)` is the default. Up to three statistics may be specified; `stats(rho obs p)` would display the correlation coefficient, number of observations, and p-value. If `varlist` contains only two variables, all statistics are shown in tabular form, and `stats()`, `print()`, and `star()` have no effect unless the `matrix` option is specified.

`print(#)` specifies the p-value cutoff for correlation coefficients to be printed. Correlation coefficients with larger p-values are left blank in the matrix. Typing `spearman, print(.10)` would display only those correlation coefficients that have p-values less than or equal to 0.10.

`star(#)` specifies the p-value cutoff for correlation coefficients to be marked with a star. Typing `spearman, star(.05)` would star all correlation coefficients that have p-values less than or equal to 0.05.

`bonferroni` makes the Bonferroni adjustment to p-values. This adjustment affects displayed p-values and the `print()` and `star()` options. Thus, `spearman, print(.05) bonferroni` prints coefficients with Bonferroni-adjusted p-values of 0.05 or less.

`sidak` makes the Šidák adjustment to p-values. This adjustment affects displayed p-values and the `print()` and `star()` options. Thus, `spearman, print(.05) sidak` prints coefficients with Šidák-adjusted p-values of 0.05 or less.

`pw` specifies that correlations be calculated using pairwise deletion of observations with missing values. By default, `spearman` uses casewise deletion, where observations are ignored if any of the variables in `varlist` are missing.

`matrix` forces `spearman` to display the statistics as a matrix, even if `varlist` contains only two variables. `matrix` is implied if more than two variables are specified.

`exact` and `exact(exact_specs)` specify that an exact p-value be reported. This option is available only when `varlist` contains only two variables.

`exact` specifies that an exact p-value from a Monte Carlo permutation test be reported. `exact` is a synonym for `exact(montecarlo)`.

`exact(montecarlo[, options] | enumerate[, options])` specifies that an exact p-value be reported in addition to the approximate p-value. Specifying `exact(montecarlo)` does a Monte Carlo permutation test. Specifying `exact(enumerate)` does an enumeration of all possible permutations. Because the number of all possible permutations is typically extremely large, enumeration is feasible only for very small datasets. The number of permutations will be displayed, and you can click on `Break` to stop the computation. The exact p-value is computed by `permute`.

`exact(montecarlo[, options])` allows options `show`, `reps(#)`, `rseed(#)`, `saving(filename[, sav_options])`, `level(#)`, `dots(#)`, `nodots`, and `eps(#)`.

`exact(enumerate[, options])` allows options `show`, `saving(filename[, sav_options])`, `dots(#)`, `nodots`, and `eps(#)`.

See `Options in [R] permute`.

`exact(montecarlo[, options])` allows options `show`, `saving(filename[, sav_options])`, `dots(#)`, `nodots`, and `eps(#)`.

See `Options in [R] permute`.
Options for ktau

_stats(ktau_list) specifies the statistics to be displayed in the matrix of output. _stats(taua) is the default. Up to six statistics may be specified; _stats(taua taub score se obs p) would display the correlation coefficients $\tau_a$ and $\tau_b$, score, standard error of score, number of observations, and $p$-value. If _varlist contains only two variables, all statistics are shown in tabular form and _stats(), _print(), and _star() have no effect unless the _matrix option is specified.

_print(#) specifies the $p$-value cutoff for correlation coefficients to be printed. Correlation coefficients with larger $p$-values are left blank in the matrix. Typing _ktau, _print(.10) would display only those correlation coefficients that have $p$-values less than or equal to 0.10.

_star(#) specifies the $p$-value cutoff for correlation coefficients to be marked with a star. Typing _ktau, _star(.05) would star all correlation coefficients that have $p$-values less than or equal to 0.05.

_bonferroni makes the Bonferroni adjustment to $p$-values. This adjustment affects displayed $p$-values and the _print() and _star() options. Thus, _ktau, _print(.05) _bonferroni prints coefficients with Bonferroni-adjusted $p$-values of 0.05 or less.

_sidak makes the Šidák adjustment to $p$-values. This adjustment affects displayed $p$-values and the _print() and _star() options. Thus, _ktau, _print(.05) _sidak prints coefficients with Šidák-adjusted $p$-values of 0.05 or less.

_pw specifies that correlations be calculated using pairwise deletion of observations with missing values. By default, _ktau uses casewise deletion, where observations are ignored if any of the variables in _varlist are missing.

_matrix forces _ktau to display the statistics as a matrix, even if _varlist contains only two variables. _matrix is implied if more than two variables are specified.

Remarks and examples

Example 1

We wish to calculate the correlation coefficients among marriage rate (_mrgrate), divorce rate (_divorce_rate), and median age (_medage) in state data. We can calculate the standard Pearson correlation coefficients and $p$-values by typing

```
use https://www.stata-press.com/data/r18/states2
.pwcorr _mrgrate _divorce_rate _medage, sig
```

<table>
<thead>
<tr>
<th></th>
<th>_mrgrate</th>
<th>_divorce_rate</th>
<th>_medage</th>
</tr>
</thead>
<tbody>
<tr>
<td>_mrgrate</td>
<td>1.0000</td>
<td></td>
<td></td>
</tr>
<tr>
<td>_divorce_rate</td>
<td>0.7895</td>
<td>1.0000</td>
<td>0.0000</td>
</tr>
<tr>
<td>_medage</td>
<td>0.0011</td>
<td>-0.1526</td>
<td>1.0000</td>
</tr>
<tr>
<td></td>
<td>0.9941</td>
<td>0.2900</td>
<td></td>
</tr>
</tbody>
</table>
We can calculate Spearman’s rank correlation coefficients by typing

```
.spearman mrgrate divorce_rate medage, stats(rho p)
```

Number of observations = 50

<table>
<thead>
<tr>
<th>Key</th>
<th>Mrgrate</th>
<th>Divorce_rate</th>
<th>Medage</th>
</tr>
</thead>
<tbody>
<tr>
<td>rho</td>
<td>1.0000</td>
<td>0.6933 1.0000</td>
<td>-0.4869 -0.2455 1.0000</td>
</tr>
<tr>
<td>p-value</td>
<td>0.0000</td>
<td>0.0004 0.0857</td>
<td></td>
</tr>
</tbody>
</table>

The large difference in the results is caused by one observation. Nevada’s marriage rate is almost 10 times higher than the state with the next-highest marriage rate. An important feature of the Spearman rank correlation is its reduced sensitivity to extreme values compared with the Pearson correlation.

We can calculate Kendall’s rank correlations by typing

```
.ktau mrgrate divorce_rate medage, stats(taua taub p)
```

Number of observations = 50

<table>
<thead>
<tr>
<th>Key</th>
<th>Mrgrate</th>
<th>Divorce_rate</th>
<th>Medage</th>
</tr>
</thead>
<tbody>
<tr>
<td>Tau_a</td>
<td>0.9829</td>
<td>0.5110 0.9804</td>
<td>-0.3486 -0.1698 0.9845</td>
</tr>
<tr>
<td>Tau_b</td>
<td>1.0000</td>
<td>0.5206 1.0000</td>
<td>-0.3544 -0.1728 1.0000</td>
</tr>
<tr>
<td>p-value</td>
<td>0.0004</td>
<td>0.0004 0.0828</td>
<td></td>
</tr>
</tbody>
</table>

There are tied values for variables `mrgrate`, `divorce_rate`, and `medage`, so average ranks are used for the tied values. As a result, $\tau_a < 1$ on the diagonal (see Methods and formulas for the definition of $\tau_a$).

According to Conover (1999, 323), “Spearman’s $\rho$ tends to be larger than Kendall’s $\tau$ in absolute value. However, as a test of significance, there is no strong reason to prefer one over the other because both will produce nearly identical results in most cases.”
Newson (2000a, 2000b, 2000c, 2001, 2003, 2005, 2006) introduces confidence intervals for Kendall’s \( \tau_a \). The community-contributed `somersd` command provides these confidence intervals along with additional rank statistics such as Somers’ \( D \) and Harrell’s \( C \) and their corresponding confidence intervals.

See Seed (2001) for confidence intervals for Spearman’s rank correlation.

### Example 2

We illustrate `spearman` and `ktau` with the auto data, which contains some missing values.

```stata
. use https://www.stata-press.com/data/r18/auto
   (1978 automobile data)
. spearman mpg rep78
Number of observations = 69
Spearman’s rho = 0.3098
Test of H0: mpg and rep78 are independent
   Prob = 0.0098
```

Because we specified two variables, `spearman` displayed the sample size, correlation, and \( p \)-value in tabular form. To obtain just the correlation coefficient displayed in matrix form, we type

```stata
. spearman mpg rep78, stats(rho) matrix
Number of observations = 69

<table>
<thead>
<tr>
<th></th>
<th>mpg</th>
<th>rep78</th>
</tr>
</thead>
<tbody>
<tr>
<td>mpg</td>
<td>1.0000</td>
<td></td>
</tr>
<tr>
<td>rep78</td>
<td>0.3098</td>
<td>1.0000</td>
</tr>
</tbody>
</table>
```

We can specify the `pw` option with `spearman` and `ktau` so that all nonmissing observations between a pair of variables when calculating their correlation coefficient are used. In the output below, some correlations are based on 74 observations, whereas others are based on 69 because 5 observations contain a missing value for `rep78`.

```stata
. spearman mpg price rep78, pw stats(rho obs p) star(0.01)
Number of observations:
    min = 69
    avg = 71
    max = 74

<table>
<thead>
<tr>
<th></th>
<th>rho</th>
<th>Number of obs</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>mpg</td>
<td>1.0000</td>
<td>74</td>
<td></td>
</tr>
<tr>
<td>price</td>
<td>-0.5419*</td>
<td>74</td>
<td>0.0000</td>
</tr>
<tr>
<td>rep78</td>
<td>0.3098*</td>
<td>69</td>
<td>0.0098</td>
</tr>
</tbody>
</table>
```
The Bonferroni and Sidak options provide adjusted \( p \)-values:

```
.ktau mpg price rep78, stats(taua taub score se p) bonferroni
```

Number of observations = 69

<table>
<thead>
<tr>
<th></th>
<th>mpg</th>
<th>price</th>
<th>rep78</th>
</tr>
</thead>
<tbody>
<tr>
<td>tau_a</td>
<td>0.9471</td>
<td>1.0000</td>
<td>2222.0000</td>
</tr>
<tr>
<td>tau_b</td>
<td>191.8600</td>
<td></td>
<td></td>
</tr>
<tr>
<td>score</td>
<td></td>
<td>-0.3973</td>
<td>1.0000</td>
</tr>
<tr>
<td>se of score</td>
<td>-0.4082</td>
<td>-932.0000</td>
<td>2346.0000</td>
</tr>
<tr>
<td>p-value</td>
<td>0.0000</td>
<td>192.4561</td>
<td>193.0682</td>
</tr>
</tbody>
</table>

Example 3

We continue with the auto data and show an example of `spearman` with the `exact` option.

```
.set seed 1234
.spearman mpg gear_ratio if foreign == 1, exact
```

Permutations (10,000): ........1,000.........2,000.........3,000.........4,000.
> ........5,000.........6,000.........7,000.........8,000.........9,000.........
> 10,000 done

Number of observations = 22
Spearman's rho = 0.4881

Test of H0: mpg and gear_ratio are independent
Prob = 0.0222
Exact prob = 0.0214 (10,000 Monte Carlo permutations)

By default, `exact` does a Monte Carlo permutation test with 10,000 permutations. Because it is a Monte Carlo test, we set the random-number generator seed before running `spearman` for reproducibility.

The exact \( p \)-value from the Monte Carlo permutation test is 0.0214, which is close to the approximate \( p \)-value of 0.0222. Note that the approximate \( p \)-value is not based on the normal distribution or the \( t \) distribution. It is calculated using a beta distribution fit to the first four moments of the null distribution of Spearman's rank correlation, and these four moments are calculated exactly for any value of \( N \), the number of observations. See Methods and formulas below.
Let’s increase the number of Monte Carlo permutations to 1,000,000 to see how close the approximate p-value is to the exact p-value. We specify dots(10000) to see a dot every 10,000th permutation, rather than the default of every 1,000th permutation. This time we set the random-number generator seed using a suboption. The exact p-value is computed by `permute`, and we can see `permute`’s output by specifying the show suboption.

```
. spearman mpg gear_ratio if foreign == 1,
    > exact(montecarlo, reps(1000000) dots(10000) rseed(1234) show)
```

Permutations (1,000,000): ........100,000.........200,000.........300,000......
> ...400,000.........500,000.........600,000.........700,000.........800,000....
> .....900,000.........1,000,000 done

Monte Carlo permutation results

<table>
<thead>
<tr>
<th>T</th>
<th>T(obs)</th>
<th>Test</th>
<th>c</th>
<th>n</th>
<th>p</th>
<th>SE(p)</th>
<th>[95% CI(p)]</th>
</tr>
</thead>
<tbody>
<tr>
<td>rho</td>
<td>0.4880642</td>
<td>lower</td>
<td>988530</td>
<td>1000000</td>
<td>.9885</td>
<td>.0001</td>
<td>.9883 .9887</td>
</tr>
<tr>
<td></td>
<td></td>
<td>upper</td>
<td>115110</td>
<td>1000000</td>
<td>.0115</td>
<td>.0001</td>
<td>.0113 .0117</td>
</tr>
<tr>
<td>two-sided</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>.0230</td>
<td>.0001 .0227 .0233</td>
</tr>
</tbody>
</table>

Notes: For lower one-sided test, c = #{T <= T(obs)} and p = p_lower = c/n.
For upper one-sided test, c = #{T >= T(obs)} and p = p_upper = c/n.
For two-sided test, p = 2*min(p_lower, p_upper); SE and CI approximate.

Number of observations = 22
Spearman’s rho = 0.4881
Test of H0: mpg and gear_ratio are independent
Prob = 0.0222
Exact prob = 0.0230 (1,000,000 Monte Carlo permutations)

With this increase in the number of permutations, the exact p-value is now calculated as 0.0230. From the table produced by `permute`, we see that this p-value has a confidence interval of [0.0227, 0.0233], accounting for the Monte Carlo error. The approximate p-value of 0.0222 falls outside this confidence interval, but it is still very close. Not bad for N = 22.

Example 4

For very small sample sizes, an exact p-value can be computed by enumerating the full permutation distribution. Here is an example using the auto data with sample size N = 11.

```
. spearman mpg gear_ratio if foreign == 1 & mpg <= 24,
    > exact(enumerate, dots(10000))
```

(enumerating all 831,600 possible permutations)

Permutations (831,600): ........100,000.........200,000.........300,000......
> ...400,000.........500,000.........600,000.........700,000.........800,000....
> one

Number of observations = 11
Spearman’s rho = 0.5636
Test of H0: mpg and gear_ratio are independent
Prob = 0.0722
Exact prob = 0.0747 (enumerated all 831,600 permutations)

The exact p-value is 0.0747. The approximate p-value is 0.0722, which is quite close to the exact p-value in this case, even with only 11 observations.
For $N = 11$, the permutation distribution consists of $11! = 39,916,800$ permutations. However, the output above says that there are only $831,600$ permutations. This is because the values of mpg in this sample are not all unique.

```
. tabulate mpg if foreign == 1 & mpg <= 24

<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>14</td>
<td>1</td>
<td>9.09</td>
<td>9.09</td>
</tr>
<tr>
<td>17</td>
<td>2</td>
<td>18.18</td>
<td>27.27</td>
</tr>
<tr>
<td>18</td>
<td>2</td>
<td>18.18</td>
<td>45.45</td>
</tr>
<tr>
<td>21</td>
<td>2</td>
<td>18.18</td>
<td>63.64</td>
</tr>
<tr>
<td>23</td>
<td>3</td>
<td>27.27</td>
<td>90.91</td>
</tr>
<tr>
<td>24</td>
<td>1</td>
<td>9.09</td>
<td>100.00</td>
</tr>
</tbody>
</table>

Total     11  100.00
```

The multiplicities in the values of mpg yield permutations that give identical results, and we need to enumerate only the permutations that are distinct. From the values of mpg, we see that each distinct permutation has a multiplicity of $2!2!2!3! = 48$, and $39,916,800/48 = 831,600$, which reduces considerably the number of permutations that need to be computed.

`permute`, which computes the permutations, permutes the first variable of the two in the `spearman varlist` command. So to best exploit this, the variable that produces the most multiplicities in the permutations should be placed first in `varlist` when doing an enumeration.

Charles Edward Spearman (1863–1945) was a British psychologist who made contributions to correlation, factor analysis, test reliability, and psychometrics. After several years’ military service, he obtained a PhD in experimental psychology at Leipzig and became a professor at University College London, where he sustained a long program of work on the interpretation of intelligence tests. Ironically, the rank correlation version bearing his name is not the formula he advocated.

Maurice George Kendall (1907–1983) was a British statistician who contributed to rank correlation, time series, multivariate analysis, among other topics, and wrote many statistical texts. Most notably, perhaps, his advanced survey of the theory of statistics went through several editions, later ones with Alan Stuart; the baton has since passed to others. Kendall was employed in turn as a government and business statistician, as a professor at the London School of Economics, as a consultant, and as director of the World Fertility Survey. He was knighted in 1974.
### Stored results

`spearman` stores the following in `r()`:

**Scalars**
- `r(N)` number of observations (last variable pair)
- `r(rho)` \( \rho \) (last variable pair)
- `r(p)` two-sided \( p \)-value (last variable pair)
- `r(p_l)` lower one-sided \( p \)-value (last variable pair)
- `r(p_u)` upper one-sided \( p \)-value (last variable pair)
- `r(p_exact)` two-sided exact \( p \)-value
- `r(p_l_exact)` lower one-sided exact \( p \)-value
- `r(p_u_exact)` upper one-sided exact \( p \)-value
- `r(n_perm)` number of permutations performed

**Macros**
- `r(exact)` "montecarlo" or "enumerate"
- `r(rngstate)` random-number state used for Monte Carlo permutations

**Matrices**
- `r(Nobs)` number of observations
- `r(Rho)` \( \rho \)
- `r(P)` two-sided \( p \)-value

If `exact(..., show)` is specified, the stored results from `permute` are returned as well; see Stored results in [R] `permute`.

`ktau` stores the following in `r()`:

**Scalars**
- `r(N)` number of observations (last variable pair)
- `r(tau_a)` \( \tau_a \) (last variable pair)
- `r(tau_b)` \( \tau_b \) (last variable pair)
- `r(score)` Kendall’s score (last variable pair)
- `r(se_score)` standard error of score (last variable pair)
- `r(p)` two-sided \( p \)-value (last variable pair)

**Matrices**
- `r(Nobs)` number of observations
- `r(Tau_a)` \( \tau_a \)
- `r(Tau_b)` \( \tau_b \)
- `r(Score)` Kendall’s score
- `r(Se_Score)` standard error of score
- `r(P)` two-sided \( p \)-value

### Methods and formulas

Methods and formulas are presented under the following headings:

- **Spearman’s rank correlation**
- **Exact \( p \)-values**
- **Kendall’s tau**

### Spearman’s rank correlation

Spearman’s (1904) rank correlation is calculated as Pearson’s correlation computed on the ranks (averaged for ties) (Conover 1999, 314–315). Ranks are as calculated by `egen`; see [D] `egen`.
If \( x_i \) and \( y_i \), where \( i = 1, 2, \ldots, n \), are the ranks of one variable pair, and \( n \) is the number of observations, then Spearman’s rank correlation is

\[
\rho = \frac{\sum_i (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_i (x_i - \bar{x})^2 \sum_i (y_i - \bar{y})^2}}
\]

where \( \bar{x} = (\sum_i x_i)/n \) is the mean of \( x \) and \( \bar{y} \) is defined similarly.

Under the null hypothesis of independence (or, more generally, exchangeability), the distribution of \( \rho \) is given by all the possible permutations of \( x_i \) with \( y_i \) fixed. For the permutation distribution, an equivalent statistic to \( \rho \) is

\[
T = \sum_i x_i y_i
\]

The moments of \( T \) for the permutation distribution can be computed exactly. Its mean is

\[
E(T) = \frac{1}{n} \{\sum_i x_i\} \{\sum_i y_i\}
\]

Assume now that the ranks, \( x_i \) and \( y_i \), are adjusted so that their means are zero. With this assumption, we have (Stuart, Ord, and Arnold 1999, eqs. 27.42 and 27.43)

\[
E(T^2) = \frac{1}{n-1} \{\sum_i x_i^2\} \{\sum_i y_i^2\}
\]

\[
E(T^3) = \frac{1}{n} \{\sum_i x_i^3\} \{\sum_i y_i^3\}
+ \frac{3}{n(n-1)} \{\sum_{i \neq j} x_i^2 x_j\} \{\sum_{i \neq j} y_i^2 y_j\}
+ \frac{36}{n(n-1)(n-2)} \{\sum_{i < j < k} x_i x_j x_k\} \{\sum_{i < j < k} y_i y_j y_k\}
\]

\[
E(T^4) = \frac{1}{n} \{\sum_i x_i^4\} \{\sum_i y_i^4\}
+ \frac{4}{n(n-1)} \{\sum_{i \neq j} x_i^3 x_j\} \{\sum_{i \neq j} y_i^3 y_j\}
+ \frac{12}{n(n-1)} \{\sum_{i < j} x_i^2 x_j^2\} \{\sum_{i < j} y_i^2 y_j^2\}
+ \frac{24}{n(n-1)(n-2)} \{\sum_{i \neq j, k; j < k} x_i^2 x_j x_k\} \{\sum_{i \neq j, k; j < k} y_i^2 y_j y_k\}
+ \frac{576}{n(n-1)(n-2)(n-3)} \{\sum_{i < j < k < l} x_i x_j x_k x_l\} \{\sum_{i < j < k < l} y_i y_j y_k y_l\}
\]

Note that Stuart, Ord, and Arnold (1999, eq. 27.43) express \( E(T^3) \) and \( E(T^4) \) in terms of \( k \) statistics, but that formulation is exactly equivalent to the equations given above.
An approximate $p$-value for Spearman’s rank correlation is calculated by fitting a four-parameter beta distribution to the first four moments of $T$. (See Lord [1965] and Hanson [1991] for a description of the technique of fitting a four-parameter beta distribution to moments of another distribution.) The four-parameter beta distribution with domain $[l, u]$ is

$$f(x) = \frac{(-l + x)^{\alpha - 1}(u - x)^{\beta - 1}}{(u - l)^{\alpha + \beta - 1}B(\alpha, \beta)}$$

where $0 \leq l < u \leq 1$ and $B(\alpha, \beta)$ is the beta function with $\alpha > 0$ and $\beta > 0$.

The parameters for the beta distribution are calculated as follows. Let $m$ be the mean of $T$, $v$ its variance, $g_3$ its skewness, and $g_4$ its kurtosis (see Methods and Formulas in [R] summarize). Define

$$r = \frac{6(g_4 - g_3^2 - 1)}{6 + 3g_3^2 - 2g_4}$$

and

$$d = 1 - \frac{24(r + 1)}{(r + 2)(r + 3)g_4 - 3(r - 6)(r + 1)}$$

Let

$$a = \frac{r(1 - \sqrt{d})}{2}$$
$$b = \frac{r(1 + \sqrt{d})}{2}$$

If $g_3 > 0$, then $\alpha = a$ and $\beta = b$; otherwise, $\alpha = b$ and $\beta = a$. The domain boundaries are given by

$$l = m - \alpha \sqrt{\frac{v(\alpha + \beta + 1)}{\alpha \beta}}$$
$$u = m + \beta \sqrt{\frac{v(\alpha + \beta + 1)}{\alpha \beta}}$$

To calculate the $p$-values, we first scale the observed value of $T$ to the domain of beta:

$$s = \frac{T_{obs} - l}{u - l}$$

Then the lower and upper one-sided $p$-values are given by

$$p_{lower} = \text{ibeta}(\alpha, \beta, s)$$
$$p_{upper} = \text{ibetatail}(\alpha, \beta, s)$$

where \text{ibeta} is Stata’s two-parameter cumulative beta distribution and \text{ibetatail} is the function for its upper tail; see [FN] Statistical functions. The two-sided $p$-value is given by

$$p = \min\{1, 2 \min(p_{lower}, p_{upper})\}$$
Exact p-values

Exact p-values for Spearman’s rank correlation are computed by permute. For details on the permutation computation, see [R] permute.

Kendall’s tau

Kendall’s $\tau$ is calculated in the following manner. For any two pairs of ranks $(x_i, y_i)$ and $(x_j, y_j)$, $1 \leq i, j \leq n$, define them as concordant if

$$(x_i - x_j)(y_i - y_j) > 0$$

and discordant if this product is less than zero.

Kendall’s (1938; also see Kendall and Gibbons [1990] or Bland [2015], 187–188) score $S$ is defined as $C - D$, where $C$ ($D$) is the number of concordant (discordant) pairs. Let $N = n(n-1)/2$ be the total number of pairs, so $\tau_a$ is given by

$$\tau_a = \frac{S}{N}$$

and $\tau_b$ is given by

$$\tau_b = \frac{S}{\sqrt{N - U} \sqrt{N - V}}$$

where

$$U = \sum_{i=1}^{N_1} u_i (u_i - 1)/2$$

$$V = \sum_{j=1}^{N_2} v_j (v_j - 1)/2$$

and where $N_1$ is the number of sets of tied $x$ values, $u_i$ is the number of tied $x$ values in the $i$th set, $N_2$ is the number of sets of tied $y$ values, and $v_j$ is the number of tied $y$ values in the $j$th set.

Under the null hypothesis of independence, the variance of $S$ is exactly (Kendall and Gibbons 1990, 66)

$$\text{Var}(S) = \frac{1}{18} \left\{ n(n-1)(2n+5) - \sum_{i=1}^{N_1} u_i (u_i - 1)(2u_i + 5) - \sum_{j=1}^{N_2} v_j (v_j - 1)(2v_j + 5) \right\}$$

$$+ \frac{1}{9n(n-1)(n-2)} \left\{ \sum_{i=1}^{N_1} u_i (u_i - 1)(u_i - 2) \right\} \left\{ \sum_{j=1}^{N_2} v_j (v_j - 1)(v_j - 2) \right\}$$

$$+ \frac{1}{2n(n-1)} \left\{ \sum_{i=1}^{N_1} u_i (u_i - 1) \right\} \left\{ \sum_{j=1}^{N_2} v_j (v_j - 1) \right\}$$

Using a normal approximation with a continuity correction,

$$z = \frac{|S| - 1}{\sqrt{\text{Var}(S)}}$$
For the hypothesis of independence, the statistics $S$, $\tau_a$, and $\tau_b$ produce equivalent tests and give the same $p$-value.

For Kendall’s $\tau$, the normal approximation is surprisingly accurate for sample sizes as small as 8, at least for calculating $p$-values under the null hypothesis for continuous variables. See Kendall and Gibbons [1990, chap. 4], who also present some tables for calculating exact $p$-values for $n < 10$.

Let $v$ be the number of variables specified so that $k = v(v - 1)/2$ correlation coefficients are to be estimated. If bonferroni is specified, the adjusted $p$-value is $p' = \min(1, kp)$. If sidak is specified, $p' = \min\{1, 1 - (1 - p)^n\}$. See Methods and formulas in [R] oneway for a more complete description of the logic behind these adjustments.

Early work on rank correlation is surveyed by Kruskal (1958).

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References


Also see

[R] `correlate` — Correlations of variables
[R] `nptrend` — Tests for trend across ordered groups
[R] `permute` — Permutation tests