

Pearson correlation coefficient

In statistics, the **Pearson correlation coefficient** (**PCC**, pronounced /ˈpiərsən/), also referred to as **Pearson's *r***, the **Pearson product-moment correlation coefficient** (**PPMCC**) or the **bivariate correlation**,^[1] is a measure of the linear correlation between two variables *X* and *Y*. According to the Cauchy–Schwarz inequality it has a value between +1 and −1, where 1 is total positive linear correlation, 0 is no linear correlation, and −1 is total negative linear correlation. It is widely used in the sciences. It was developed by Karl Pearson from a related idea introduced by Francis Galton in the 1880s and for which the mathematical formula was derived and published by Auguste Bravais in 1844.^{[2][3][4][5][6]} The naming of the coefficient is thus an example of Stigler's Law.

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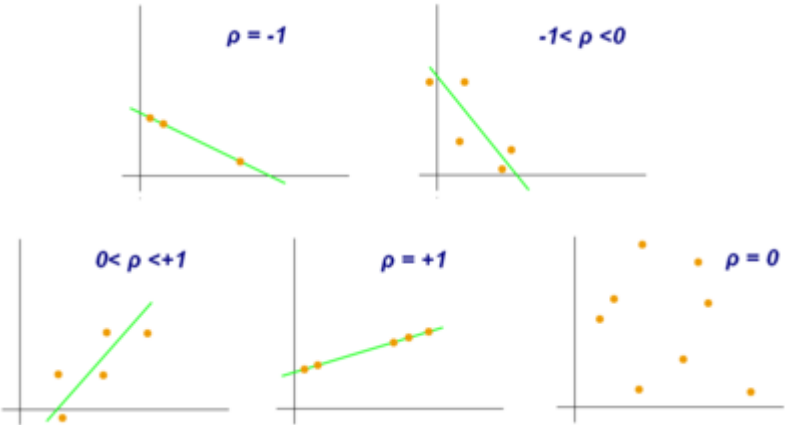
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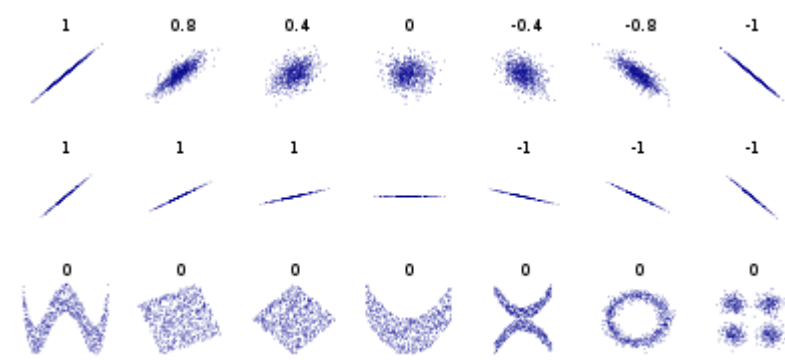
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Examples of scatter diagrams with different values of correlation coefficient (*ρ*)



Several sets of (*x*, *y*) points, with the correlation coefficient of *x* and *y* for each set. Note that the correlation reflects the non-linearity and direction of a linear relationship (top row), but not the slope of that relationship (middle), nor many aspects of nonlinear relationships (bottom). N.B.: the figure in the center has a slope of 0 but in that case the correlation coefficient is undefined because the variance of *Y* is zero.

Definition

Pearson's correlation coefficient is the covariance of the two variables divided by the product of their standard deviations. The form of the definition involves a "product moment", that is, the mean (the first moment about the origin) of the product of the mean-adjusted random variables; hence the modifier *product-moment* in the name.

For a population

Pearson's correlation coefficient when applied to a population is commonly represented by the Greek letter *ρ* (rho) and may be referred to as the *population correlation coefficient* or the *population Pearson correlation coefficient*. Given a pair of random variables (***X***, ***Y***), the formula for *ρ*^[7] is:

$$\rho_{X,Y} = \frac{\mathbf{cov}(X,Y)}{\sigma_X\sigma_Y}$$

(Eq.1)

where:

- **cov** is the covariance
- ***σ_X*** is the standard deviation of ***X***
- ***σ_Y*** is the standard deviation of ***Y***

The formula for ***ρ*** can be expressed in terms of mean and expectation. Since

$$\mathbf{cov}(X,Y) = \mathbf{E}[(X-\mu_X)(Y-\mu_Y)],^{[7]}$$

the formula for ***ρ*** can also be written as

$$\rho_{X,Y} = \frac{\mathbf{E}[(X-\mu_X)(Y-\mu_Y)]}{\sigma_X\sigma_Y}$$

(Eq.2)

where:

- ***σ_Y*** and ***σ_X*** are defined as above
- ***μ_X*** is the mean of ***X***
- ***μ_Y*** is the mean of ***Y***
- **E** is the expectation.

The formula for ***ρ*** can be expressed in terms of uncentered moments. Since

- ***μ_X*** = **E**[***X***]
- ***μ_Y*** = **E**[***Y***]
- ***σ_X²*** = **E**[(***X*** − **E**[***X***])²] = **E**[***X***²] − [**E**[***X***]]²
- ***σ_Y²*** = **E**[(***Y*** − **E**[***Y***])²] = **E**[***Y***²] − [**E**[***Y***]]²
- **E**[(***X*** − ***μ_X***)(***Y*** − ***μ_Y***)] = **E**[(***X*** − **E**[***X***])(***Y*** − **E**[***Y***])] = **E**[***XY***] − **E**[***X***] **E**[***Y***],

the formula for ***ρ*** can also be written as

$$\rho_{X,Y} = \frac{\mathbf{E}[XY] - \mathbf{E}[X] \mathbf{E}[Y]}{\sqrt{\mathbf{E}[X^2] - [\mathbf{E}[X]]^2} \sqrt{\mathbf{E}[Y^2] - [\mathbf{E}[Y]]^2}}.$$

For a sample

Pearson's correlation coefficient when applied to a sample is commonly represented by r_{xy} and may be referred to as the *sample correlation coefficient* or the *sample Pearson correlation coefficient*. We can obtain a formula for r_{xy} by substituting estimates of the covariances and variances based on a sample into the formula above. Given paired data $\{(x_1, y_1), \dots, (x_n, y_n)\}$ consisting of n pairs, r_{xy} is defined as:

$$r_{xy} = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_{i=1}^n (x_i - \bar{x})^2} \sqrt{\sum_{i=1}^n (y_i - \bar{y})^2}}$$

(Eq.3)

where:

- n is sample size
- x_i, y_i are the individual sample points indexed with i
- $\bar{x} = \frac{1}{n} \sum_{i=1}^n x_i$ (the sample mean); and analogously for \bar{y}

Rearranging gives us this formula for r_{xy} :

$$r_{xy} = \frac{n \sum x_i y_i - \sum x_i \sum y_i}{\sqrt{n \sum x_i^2 - (\sum x_i)^2} \sqrt{n \sum y_i^2 - (\sum y_i)^2}}.$$

where:

- n, x_i, y_i are defined as above
- This formula suggests a convenient single-pass algorithm for calculating sample correlations, but, depending on the numbers involved, it can sometimes be numerically unstable.

Rearranging again gives us this^[7] formula for r_{xy} :

$$r_{xy} = \frac{\sum x_i y_i - n \bar{x} \bar{y}}{\sqrt{(\sum x_i^2 - n \bar{x}^2)} \sqrt{(\sum y_i^2 - n \bar{y}^2)}}.$$

where:

- $n, x_i, y_i, \bar{x}, \bar{y}$ are defined as above

An equivalent expression gives the formula for r_{xy} as the mean of the products of the standard scores as follows:

$$r_{xy} = \frac{1}{n-1} \sum_{i=1}^n \left(\frac{x_i - \bar{x}}{s_x} \right) \left(\frac{y_i - \bar{y}}{s_y} \right)$$

where

- $n, x_i, y_i, \bar{x}, \bar{y}$ are defined as above, and s_x, s_y are defined below
- $\left(\frac{x_i - \bar{x}}{s_x} \right)$ is the standard score (and analogously for the standard score of y)

Alternative formulae for r_{xy} are also available. One can use the following formula for r_{xy} :

r

xy

=

∑

x

i

y

i

−
n
x
¯

y
¯

(
n
−
1
)

s

x

s

y

where:

- n*, *x_i*, *y_i*, *x̄*, *ȳ* are defined as above and:
- s_x* =

1

n
−
1

∑

i
=
1

n

(

x

i

−
x
¯

)

2

 (the sample standard deviation); and analogously for *s_y*

Practical issues

Under heavy noise conditions, extracting the correlation coefficient between two sets of stochastic variables is nontrivial, in particular where Canonical Correlation Analysis reports degraded correlation values due to the heavy noise contributions. A generalization of the approach is given elsewhere.^[8]

In case of missing data, Garren derived the maximum likelihood estimator.^[9]

Mathematical properties

The absolute values of both the sample and population Pearson correlation coefficients are less than or equal to 1. Correlations equal to 1 or −1 correspond to data points lying exactly on a line (in the case of the sample correlation), or to a bivariate distribution entirely supported on a line (in the case of the population correlation). The Pearson correlation coefficient is symmetric: corr(*X*,*Y*) = corr(*Y*,*X*).

A key mathematical property of the Pearson correlation coefficient is that it is invariant under separate changes in location and scale in the two variables. That is, we may transform *X* to *a* + *bX* and transform *Y* to *c* + *dY*, where *a*, *b*, *c*, and *d* are constants with *b*, *d* > 0, without changing the correlation coefficient. (This holds for both the population and sample Pearson correlation coefficients.) Note that more general linear transformations do change the correlation: see § Decorrelation of n random variables for an application of this.

Interpretation

The correlation coefficient ranges from −1 to 1. A value of 1 implies that a linear equation describes the relationship between *X* and *Y* perfectly, with all data points lying on a line for which *Y* increases as *X* increases. A value of −1 implies that all data points lie on a line for which *Y* decreases as *X* increases. A value of 0 implies that there is no linear correlation between the variables.

More generally, note that (*X_i* −

X
¯
)(*Y_i* −

Y
¯
) is positive if and only if *X_i* and *Y_i* lie on the same side of their respective means. Thus the correlation coefficient is positive if *X_i* and *Y_i* tend to be simultaneously greater than, or simultaneously less than, their respective means. The correlation coefficient is negative (anti-correlation) if *X_i* and *Y_i* tend to lie on opposite sides of their respective means. Moreover, the stronger is either tendency, the larger is the absolute value of the correlation coefficient.

Rogers and Nicewander^[10] cataloged thirteen ways of interpreting correlation:

- Function of raw scores and means
- Standardized covariance
- Standardized slope of the regression line
- Geometric mean of the two regression slopes
- Square root of the ratio of two variances
- Mean cross-product of standardized variables
- Function of the angle between two standardized regression lines
- Function of the angle between two variable vectors
- Rescaled variance of the difference between standardized scores
- Estimated from the balloon rule
- Related to the bivariate ellipses of isoconcentration
- Function of test statistics from designed experiments
- Ratio of two means

Geometric interpretation

For uncentered data, there is a relation between the correlation coefficient and the angle *φ* between the two regression lines, *y* = *g*_{*X*}(*x*) and *x* = *g*_{*Y*}(*y*), obtained by regressing *y* on *x* and *x* on *y* respectively. (Here *φ* is measured counterclockwise within the first quadrant formed around the lines' intersection point if *r* > 0, or counterclockwise from the fourth to the second quadrant if *r* < 0.) One can show^[11] that if the standard deviations are equal, then *r* = sec *φ* − tan *φ*, where sec and tan are trigonometric functions.

For centered data (i.e., data which have been shifted by the sample means of their respective variables so as to have an average of zero for each variable), the correlation coefficient can also be viewed as the cosine of the angle *θ* between the two observed vectors in *N*-dimensional space (for *N* observations of each variable)^{[12]:ch. 5} (as illustrated for a special case in the next paragraph).

Both the uncentered (non-Pearson-compliant) and centered correlation coefficients can be determined for a dataset. As an example, suppose five countries are found to have gross national products of 1, 2, 3, 5, and 8 billion dollars, respectively. Suppose these same five countries (in the same order) are found to have 11%, 12%, 13%, 15%, and 18% poverty. Then let **x** and **y** be ordered 5-element vectors containing the above data: **x** = (1, 2, 3, 5, 8) and **y** = (0.11, 0.12, 0.13, 0.15, 0.18).

By the usual procedure for finding the angle *θ* between two vectors (see dot product), the *uncentered* correlation coefficient is:

$$\cos \theta = \frac{\mathbf{x} \cdot \mathbf{y}}{\|\mathbf{x}\| \|\mathbf{y}\|} = \frac{2.93}{\sqrt{103}\sqrt{0.0983}} = 0.920814711.$$

This uncentred correlation coefficient is identical with the cosine similarity. Note that the above data were deliberately chosen to be perfectly correlated: *y* = 0.10 + 0.01 *x*. The Pearson correlation coefficient must therefore be exactly one. Centering the data (shifting **x** by E(**x**) = 3.8 and **y** by E(**y**) = 0.138) yields **x** = (−2.8, −1.8, −0.8, 1.2, 4.2) and **y** = (−0.028, −0.018, −0.008, 0.012, 0.042), from which

$$\cos \theta = \frac{\mathbf{x} \cdot \mathbf{y}}{\|\mathbf{x}\| \|\mathbf{y}\|} = \frac{0.308}{\sqrt{30.8}\sqrt{0.00308}} = 1 = \rho_{xy},$$

as expected.

Interpretation of the size of a correlation

Several authors have offered guidelines for the interpretation of a correlation coefficient.^{[13][14]} However, all such criteria are in some ways arbitrary.^[14] The interpretation of a correlation coefficient depends on the context and purposes. A correlation of 0.8 may be very low if one is verifying a physical law using high-quality instruments, but may be regarded as very high in the social sciences where there may be a greater contribution from complicating factors.

Inference

Statistical inference based on Pearson's correlation coefficient often focuses on one of the following two aims:

- One aim is to test the null hypothesis that the true correlation coefficient *ρ* is equal to 0, based on the value of the sample correlation coefficient *r*.
- The other aim is to derive a confidence interval that, on repeated sampling, has a given probability of containing *ρ*.

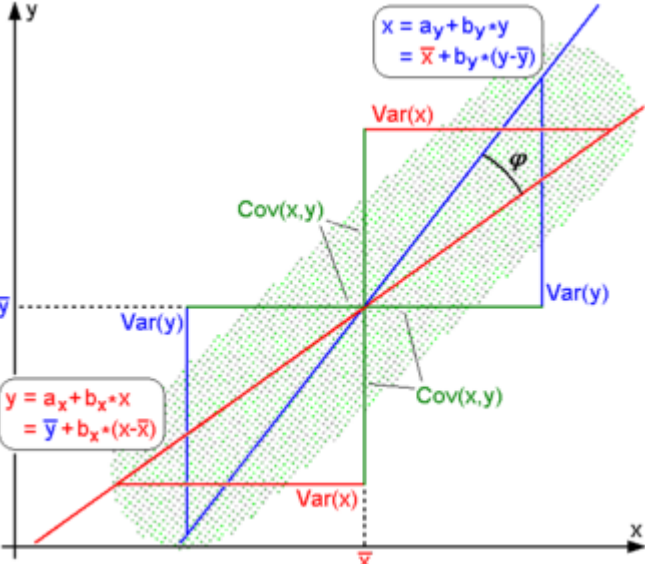
We discuss methods of achieving one or both of these aims below.

Using a permutation test

Permutation tests provide a direct approach to performing hypothesis tests and constructing confidence intervals. A permutation test for Pearson's correlation coefficient involves the following two steps:

- Using the original paired data (*x*_{*i*}, *y*_{*i*}), randomly redefine the pairs to create a new data set (*x*_{*i*}, *y*_{*i*}'), where the *i*′ are a permutation of the set {1,...,*n*}. The permutation *i*′ is selected randomly, with equal probabilities placed on all *n*! possible permutations. This is equivalent to drawing the *i*′ randomly without replacement from the set {1, ..., *n*}. In bootstrapping, a closely related approach, the *i* and the *i*′ are equal and drawn with replacement from {1, ..., *n*};
- Construct a correlation coefficient *r* from the randomized data.

To perform the permutation test, repeat steps (1) and (2) a large number of times. The p-value for the permutation test is the proportion of the *r* values generated in step (2) that are larger than the Pearson correlation coefficient that was calculated from the original data. Here "larger" can mean either that the value is larger in magnitude, or larger in signed value, depending on whether a two-sided or one-sided test is desired.



Regression lines for *y* = *g*_{*X*}(*x*) [**red**] and *x* = *g*_{*Y*}(*y*) [**blue**]

Using a bootstrap

The bootstrap can be used to construct confidence intervals for Pearson's correlation coefficient. In the "non-parametric" bootstrap, *n* pairs (*x*_{*i*}, *y*_{*i*}) are resampled "with replacement" from the observed set of *n* pairs, and the correlation coefficient *r* is calculated based on the resampled data. This process is repeated a large number of times, and the empirical distribution of the resampled *r* values are used to approximate the sampling distribution of the statistic. A 95% confidence interval for *ρ* can be defined as the interval spanning from the 2.5th to the 97.5th percentile of the resampled *r* values.

Testing using Student's *t*-distribution

For pairs from an uncorrelated bivariate normal distribution, the sampling distribution of a certain function of Pearson's correlation coefficient follows Student's *t*-distribution with degrees of freedom *n* − 2. Specifically, if the underlying variables are white and have a bivariate normal distribution, the variable

$$t = r \sqrt{\frac{n-2}{1-r^2}}$$

has a Student's *t*-distribution in the null case (zero correlation).^[15] This holds approximately in case of non-normal observed values if sample sizes are large enough.^[16] For determining the critical values for *r* the inverse function is needed:

$$r = \frac{t}{\sqrt{n-2+t^2}}.$$

Alternatively, large sample, asymptotic approaches can be used.

Another early paper^[17] provides graphs and tables for general values of *ρ*, for small sample sizes, and discusses computational approaches.

In the case where the underlying variables are not white, the sampling distribution of Pearson's correlation coefficient follows a Student's *t*-distribution, but the degrees of freedom are reduced.^[18]

Using the exact distribution

For data that follow a bivariate normal distribution, the exact density function *f*(*r*) for the sample correlation coefficient *r* of a normal bivariate is^[19] ^[20] ^[21]

$$f(r) = \frac{(n-2) \, \Gamma(n-1) (1-\rho^2)^{\frac{n-1}{2}} (1-r^2)^{\frac{n-4}{2}}}{\sqrt{2\pi} \, \Gamma\left(n-\frac{1}{2}\right) (1-\rho r)^{n-\frac{3}{2}}} \, {}_2\mathbf{F}_1\left(\frac{1}{2}, \frac{1}{2}; \frac{2n-1}{2}; \frac{\rho r+1}{2}\right)$$

where **Γ** is the gamma function and **{}_2\mathbf{F}_1(a, b; c; z)** is the Gaussian hypergeometric function.

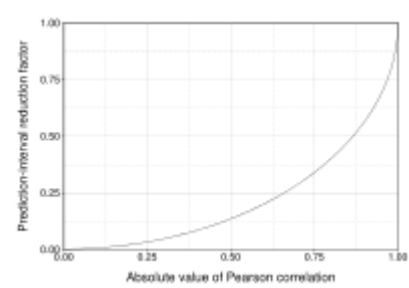
In the special case when *ρ* = 0, the exact density function *f*(*r*) can be written as:

$$f(r) = \frac{(1-r^2)^{\frac{n-4}{2}}}{\mathbf{B}\left(\frac{1}{2}, \frac{n-2}{2}\right)},$$

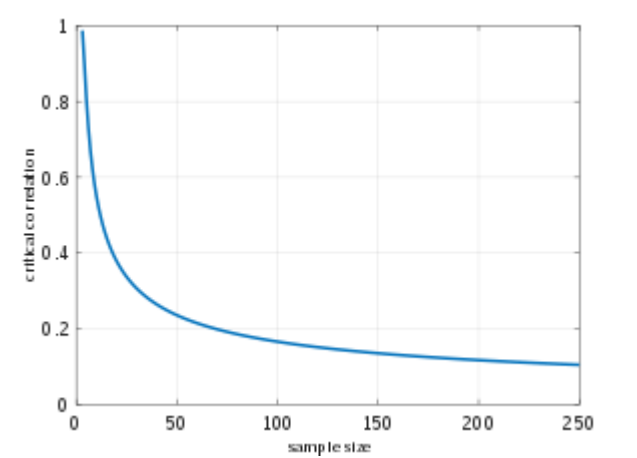
where **B** is the beta function, which is one way of writing the density of a Student's t-distribution, as above.

Using the Fisher transformation

In practice, confidence intervals and hypothesis tests relating to ρ are usually carried out using the Fisher transformation, ***F***:



This figure gives a sense of how the usefulness of a Pearson correlation for predicting values varies with its magnitude. Given jointly normal *X*, *Y* with correlation *ρ*, $1 - \sqrt{1 - \rho^2}$ (plotted here as a function of *ρ*) is the factor by which a given prediction interval for *Y* may be reduced given the corresponding value of *X*. For example, if *ρ* = .5, then the 95% prediction interval of *Y*|*X* will be about 13% smaller than the 95% prediction interval of *Y*.



Critical values of Pearson's correlation coefficient that must be exceeded to be considered significantly nonzero at the 0.05 level.

$$F(r) = \frac{1}{2} \ln \frac{1+r}{1-r} = \mathbf{arctanh}(r).$$

where n is the sample size. $F(r)$ approximately follows a normal distribution with

$$\mathbf{mean} = F(\rho) = \mathbf{arctanh}(\rho) \quad \text{and standard error} = \mathbf{SE} = \frac{1}{\sqrt{n-3}}.$$

The approximation error is lowest for a large sample size \boldsymbol{n} and small \boldsymbol{r} and $\boldsymbol{\rho_0}$ and increases otherwise.

Using the approximation, a z-score is

$$z = \frac{\boldsymbol{x} - \mathbf{mean}}{\mathbf{SE}} = [F(r) - F(\rho_0)]\sqrt{n-3}$$

under the null hypothesis of that $\boldsymbol{\rho} = \boldsymbol{\rho_0}$, given the assumption that the sample pairs are independent and identically distributed and follow a bivariate normal distribution. Thus an approximate p-value can be obtained from a normal probability table. For example, if $z = 2.2$ is observed and a two-sided p-value is desired to test the null hypothesis that $\boldsymbol{\rho} = \mathbf{0}$, the p-value is $2 \cdot \Phi(-2.2) = 0.028$, where Φ is the standard normal cumulative distribution function.

To obtain a confidence interval for ρ , we first compute a confidence interval for $F(\rho)$:

$$100(1-\alpha)\% \mathbf{CI} : \mathbf{arctanh}(\rho) \in [\mathbf{arctanh}(r) \pm z_{\alpha/2}SE]$$

The inverse Fisher transformation brings the interval back to the correlation scale.

$$100(1-\alpha)\% \mathbf{CI} : \boldsymbol{\rho} \in [\mathbf{tanh(arctanh}(r) - z_{\alpha/2}SE), \mathbf{tanh(arctanh}(r) + z_{\alpha/2}SE)]$$

For example, suppose we observe $r = 0.3$ with a sample size of $n=50$, and we wish to obtain a 95% confidence interval for ρ . The transformed value is $\arctanh(r) = 0.30952$, so the confidence interval on the transformed scale is $0.30952 \pm 1.96/\sqrt{47}$, or (0.023624, 0.595415). Converting back to the correlation scale yields (0.024, 0.534).

In least squares regression analysis

The square of the sample correlation coefficient is typically denoted r^2 and is a special case of the coefficient of determination. In this case, it estimates the fraction of the variance in Y that is explained by X in a simple linear regression. So if we have the observed dataset $\boldsymbol{Y_1, \ldots, Y_n}$ and the fitted dataset $\hat{\boldsymbol{Y_1, \ldots, Y_n}}$ then as a starting point the total variation in the Y_i around their average value can be decomposed as follows

$$\sum_i (Y_i - \bar{Y})^2 = \sum_i (Y_i - \hat{Y}_i)^2 + \sum_i (\hat{Y}_i - \bar{Y})^2,$$

where the $\hat{\boldsymbol{Y_i}}$ are the fitted values from the regression analysis. This can be rearranged to give

$$1 = \frac{\sum_i (Y_i - \hat{Y}_i)^2}{\sum_i (Y_i - \bar{Y})^2} + \frac{\sum_i (\hat{Y}_i - \bar{Y})^2}{\sum_i (Y_i - \bar{Y})^2}.$$

The two summands above are the fraction of variance in Y that is explained by X (right) and that is unexplained by X (left).

Next, we apply a property of least square regression models, that the sample covariance between $\hat{\boldsymbol{Y_i}}$ and $\boldsymbol{Y_i} - \hat{\boldsymbol{Y_i}}$ is zero. Thus, the sample correlation coefficient between the observed and fitted response values in the regression can be written (calculation is under expectation, assumes Gaussian statistics)

$$\begin{aligned}
r(Y, \hat{Y}) &= \frac{\sum_i (Y_i - \bar{Y})(\hat{Y}_i - \bar{Y})}{\sqrt{\sum_i (Y_i - \bar{Y})^2 \cdot \sum_i (\hat{Y}_i - \bar{Y})^2}} \\
&= \frac{\sum_i (Y_i - \hat{Y}_i + \hat{Y}_i - \bar{Y})(\hat{Y}_i - \bar{Y})}{\sqrt{\sum_i (Y_i - \bar{Y})^2 \cdot \sum_i (\hat{Y}_i - \bar{Y})^2}} \\
&= \frac{\sum_i [(Y_i - \hat{Y}_i)(\hat{Y}_i - \bar{Y}) + (\hat{Y}_i - \bar{Y})^2]}{\sqrt{\sum_i (Y_i - \bar{Y})^2 \cdot \sum_i (\hat{Y}_i - \bar{Y})^2}} \\
&= \frac{\sum_i (\hat{Y}_i - \bar{Y})^2}{\sqrt{\sum_i (Y_i - \bar{Y})^2 \cdot \sum_i (\hat{Y}_i - \bar{Y})^2}} \\
&= \sqrt{\frac{\sum_i (\hat{Y}_i - \bar{Y})^2}{\sum_i (Y_i - \bar{Y})^2}}.
\end{aligned}$$

Thus

$$r(Y, \hat{Y})^2 = \frac{\sum_i (\hat{Y}_i - \bar{Y})^2}{\sum_i (Y_i - \bar{Y})^2}$$

where

- $r(Y, \hat{Y})^2$ is the proportion of variance in Y explained by a linear function of X .

That equation can be written as:

$$r(Y, \hat{Y})^2 = \frac{SS_{\text{reg}}}{SS_{\text{tot}}}$$

where

- SS_{reg} is the regression sum of squares, also called the explained sum of squares
- SS_{tot} is the total sum of squares (proportional to the variance of the data)
- $SS_{\text{reg}} = \sum_i (\hat{Y}_i - \bar{Y})^2$
- $SS_{\text{tot}} = \sum_i (Y_i - \bar{Y})^2$

Sensitivity to the data distribution

Existence

$$(2) \qquad r = \mathbf{E}[r] = r_{\text{adj}} - \frac{r_{\text{adj}}(1 - r_{\text{adj}}^2)}{2n}.$$

An approximate solution to equation (2) is:

$$(3) \qquad r_{\text{adj}} = r \left[1 + \frac{1 - r^2}{2n} \right],$$

where in (3):

- r, n are defined as above,
- r_{adj} is a suboptimal estimator,
- r_{adj} can also be obtained by maximizing $\log(f(r))$,
- r_{adj} has minimum variance for large values of n ,
- r_{adj} has a bias of order $1/(n - 1)$.

Another proposed^[7] adjusted correlation coefficient is:

$$r_{\text{adj}} = \sqrt{1 - \frac{(1 - r^2)(n - 1)}{(n - 2)}}.$$

Note that $r_{\text{adj}} \approx r$ for large values of n .

Weighted correlation coefficient

Suppose observations to be correlated have differing degrees of importance that can be expressed with a weight vector w . To calculate the correlation between vectors x and y with the weight vector w (all of length n),^{[29][30]}

- Weighted mean:

$$\mathbf{m}(x; w) = \frac{\sum_i w_i x_i}{\sum_i w_i}.$$

- Weighted covariance

$$\mathbf{cov}(x, y; w) = \frac{\sum_i w_i \cdot (x_i - \mathbf{m}(x; w))(y_i - \mathbf{m}(y; w))}{\sum_i w_i}.$$

- Weighted correlation

$$\mathbf{corr}(x, y; w) = \frac{\mathbf{cov}(x, y; w)}{\sqrt{\mathbf{cov}(x, x; w) \mathbf{cov}(y, y; w)}}.$$

Reflective correlation coefficient

The reflective correlation is a variant of Pearson's correlation in which the data are not centered around their mean values. The population reflective correlation is

$$\mathbf{corr}_r(X,Y) = \frac{\mathbf{E}[XY]}{\sqrt{\mathbf{E}\,X^2\cdot\mathbf{E}\,Y^2}}.$$

The reflective correlation is symmetric, but it is not invariant under translation:

$$\mathbf{corr}_r(X,Y) = \mathbf{corr}_r(Y,X) = \mathbf{corr}_r(X,bY) \neq \mathbf{corr}_r(X,a+bY), \quad a \neq 0, b > 0.$$

The sample reflective correlation is equivalent to cosine similarity:

$$rr_{xy} = \frac{\sum x_i y_i}{\sqrt{(\sum x_i^2)(\sum y_i^2)}}.$$

The weighted version of the sample reflective correlation is

$$rr_{xy,w} = \frac{\sum w_i x_i y_i}{\sqrt{(\sum w_i x_i^2)(\sum w_i y_i^2)}}.$$

Scaled correlation coefficient

Scaled correlation is a variant of Pearson's correlation in which the range of the data is restricted intentionally and in a controlled manner to reveal correlations between fast components in time series.^[31] Scaled correlation is defined as average correlation across short segments of data.

Let ***K*** be the number of segments that can fit into the total length of the signal ***T*** for a given scale ***s***:

$$K = \mathbf{round}\left(\frac{T}{s}\right).$$

The scaled correlation across the entire signals ***r̄_s*** is then computed as

$$\bar{r}_s = \frac{1}{K} \sum_{k=1}^K r_k,$$

where ***r_k*** is Pearson's coefficient of correlation for segment ***k***.

By choosing the parameter ***s***, the range of values is reduced and the correlations on long time scale are filtered out, only the correlations on short time scales being revealed. Thus, the contributions of slow components are removed and those of fast components are retained.

Pearson's distance

A distance metric for two variables X and Y known as *Pearson's distance* can be defined from their correlation coefficient as^[32]

$$d_{X,Y} = 1 - \rho_{X,Y}.$$

Considering that the Pearson correlation coefficient falls between [−1, 1], the Pearson distance lies in [0, 2]. The Pearson distance has been used in cluster analysis and data detection for communications and storage with unknown gain and offset^[33]

Circular correlation coefficient

For variables

X

=

{

x

1

,...,

x

n

}

 and

Y

=

{

y

1

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y

n

}

 that are defined on the unit circle [0, 2*π*), it is possible to define a circular analog of Pearson's coefficient.^[34] This is done by transforming data points in X and Y with a sine function such that the correlation coefficient is given as:

$$r_{\mathrm{circular}} = \frac{\sum_{i=1}^n \sin(x_i - \bar{x}) \sin(y_i - \bar{y})}{\sqrt{\sum_{i=1}^n \sin(x_i - \bar{x})^2} \sqrt{\sum_{i=1}^n \sin(y_i - \bar{y})^2}}$$

where ***x
¯*** and ***y
¯*** are the circular means of *X* and *Y*. This measure can be useful in fields like meteorology where the angular direction of data is important.

Partial correlation

If a population or data-set is characterized by more than two variables, a partial correlation coefficient measures the strength of dependence between a pair of variables that is not accounted for by the way in which they both change in response to variations in a selected subset of the other variables.

Decorrelation of *n* random variables

It is always possible to remove the correlations between all pairs of an arbitrary number of random variables by using a data transformation, even if the relationship between the variables is nonlinear. A presentation of this result for population distributions is given by Cox & Hinkley.^[35]

A corresponding result exists for reducing the sample correlations to zero. Suppose a vector of *n* random variables is observed *m* times. Let *X* be a matrix where **X

i
,
j** is the *j*th variable of observation *i*. Let **Z

m
,
m** be an *m* by *m* square matrix with every element 1. Then *D* is the data transformed so every random variable has zero mean, and *T* is the data transformed so all variables have zero mean and zero correlation with all other variables – the sample correlation matrix of *T* will be the identity matrix. This has to be further divided by the standard deviation to get unit variance. The transformed variables will be uncorrelated, even though they may not be independent.

$$D = X - \frac{1}{m}Z_{m,m}X$$

$$T = D(D^{\mathsf{T}}D)^{-\frac{1}{2}},$$

where an exponent of −1/2 represents the matrix square root of the inverse of a matrix. The correlation matrix of *T* will be the identity matrix. If a new data observation *x* is a row vector of *n* elements, then the same transform can be applied to *x* to get the transformed vectors *d* and *t*:

$$d = x - \frac{1}{m}Z_{1,m}X,$$

$$t = d(D^{\mathsf{T}}D)^{-\frac{1}{2}}.$$

This decorrelation is related to principal components analysis for multivariate data.

Software Implementations

- R's statistics base-package implements the test cor.test(x, y, method = "pearson") (http://stat.ethz.ch/R-manual/R-patched/library/stats/html/cor.test.html) in its "stats" package (also cor(x, y, method = "pearson") will work, but without returning the p-value). Since pearson is the default, the method argument could also be omitted.

See also

- Anscombe's quartet
- Association (statistics)
- Coefficient of colligation