

# Package ‘jfa’

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**Title** Bayesian and Classical Audit Sampling

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**Description** Implements the audit sampling workflow as discussed in Derks et al. (2019) <[doi:10.31234/osf.io/9f6ub](https://doi.org/10.31234/osf.io/9f6ub)>. The package makes it easy for an auditor to plan a statistical sample, select the sample from the population, and evaluate the misstatement in the sample compliant with the International Standards on Auditing. Next to classical audit sampling methodology, the package implements Bayesian equivalents of these methods whose statistical underpinnings are described in Derks et al. (2021) <[doi:10.1111/ijau.12240](https://doi.org/10.1111/ijau.12240)> and Derks et al. (2021) <[doi:10.31234/osf.io/kzqp5](https://doi.org/10.31234/osf.io/kzqp5)>.

**BugReports** <https://github.com/koenderks/jfa/issues>

**URL** <https://koenderks.github.io/jfa/>, <https://github.com/koenderks/jfa>

**Imports** extraDistr, graphics, stats

**Suggests** kableExtra, knitr, MUS, rmarkdown, testthat

**Language** en-US

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** true

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**VignetteBuilder** knitr

## R topics documented:

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## Description

jfa is an R package for statistical audit sampling. The package provides functions for planning, performing, evaluating, and reporting an audit sample. Specifically, these functions implement standard audit sampling techniques for calculating sample sizes, selecting items from a population, and evaluating the misstatement from a data sample or from summary statistics. Additionally, the jfa package allows the user to create a prior probability distribution to perform Bayesian audit sampling using these functions.

The package and its intended workflow are also implemented with a graphical user interface in the Audit module of **JASP**, a free and open-source statistical software program.

For documentation on jfa itself, including the manual and user guide for the package, worked examples, and other tutorial information visit the [package website](#).

## Reference tables

Below you can find several links to reference tables that contain statistical sample sizes, upper limits, and Bayes factors. These tables are created using the `planning()` and `evaluation()` functions provided in the package. See the corresponding help files for more information about these functions and how to replicate this output.

### *Sample sizes*

- [Sample sizes based on the Poisson distribution](#)
- [Sample sizes based on the binomial distribution](#)
- [Sample sizes based on the hypergeometric distribution](#)

### *Upper limits*

- [Upper limits based on the Poisson distribution](#)
- [Upper limits based on the binomial distribution](#)
- [Upper limits based on the hypergeometric distribution](#)

### *One-sided p values*

- [One sided p values based on the Poisson distribution](#)
- [One sided p values based on the binomial distribution](#)
- [One sided p values based on the hypergeometric distribution](#)

### *Bayes factors*

- [Impartial Bayes factors based on the gamma distribution](#)
- [Impartial Bayes factors based on the beta distribution](#)
- [Impartial Bayes factors based on the beta-binomial distribution](#)

## Author(s)

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Please use the citation provided by R when citing this package. A BibTex entry is available from `citation('jfa')`.

## See Also

Useful links:

- The [cheat sheet](#) for a quick overview of the intended workflow.
- The [vignettes](#) for worked examples.
- The [issue page](#) to submit a bug report or feature request.

## Examples

```
# Load the jfa package
library(jfa)

# Load the BuildIt population
data('BuildIt')

#####
### Example 1: Classical audit sampling ###
#####

# Stage 1: Planning
stage1 <- planning(materiality = 0.03, expected = 0.01,
                    likelihood = 'poisson', conf.level = 0.95)
summary(stage1)

# Stage 2: Selection
stage2 <- selection(data = BuildIt, size = stage1,
                    units = 'values', values = 'bookValue',
                    method = 'interval', start = 1)
summary(stage2)

# Stage 3: Execution
sample <- stage2[['sample']]

# Stage 4: Evaluation
stage4 <- evaluation(materiality = 0.03, method = 'stringer',
                     conf.level = 0.95, data = sample,
                     values = 'bookValue', values.audit = 'auditValue')
summary(stage4)

#####
### Example 2: Bayesian audit sampling using a non-informed prior ###
#####

# Create the prior distribution
prior <- auditPrior(method = 'default', likelihood = 'poisson')
summary(prior)

# Stage 1: Planning
stage1 <- planning(materiality = 0.03, expected = 0.01,
```

```

                                likelihood = 'poisson', conf.level = 0.95, prior = prior)
summary(stage1)

# Stage 2: Selection
stage2 <- selection(data = BuildIt, size = stage1,
                    units = 'values', values = 'bookValue',
                    method = 'interval', start = 1)
summary(stage2)

# Stage 3: Execution
sample <- stage2[['sample']]

# Stage 4: Evaluation
stage4 <- evaluation(materiality = 0.03, conf.level = 0.95, data = sample,
                     values = 'bookValue', values.audit = 'auditValue',
                     prior = prior)
summary(stage4)

#####
### Example 3: Bayesian audit sampling using an informed prior ###
#####

# Create the prior distribution
prior <- auditPrior(method = 'arm', likelihood = 'poisson',
                    expected = 0.01, materiality = 0.03, cr = 0.6, ir = 1)
summary(prior)

# Stage 1: Planning
stage1 <- planning(materiality = 0.03, expected = 0.01,
                   likelihood = 'poisson', conf.level = 0.95, prior = prior)
summary(stage1)

# Stage 2: Selection
stage2 <- selection(data = BuildIt, size = stage1,
                    units = 'values', values = 'bookValue',
                    method = 'interval', start = 1)
summary(stage2)

# Stage 3: Execution
sample <- stage2[['sample']]

# Stage 4: Evaluation
stage4 <- evaluation(materiality = 0.03, conf.level = 0.95, data = sample,
                     values = 'bookValue', values.audit = 'auditValue',
                     prior = prior)
summary(stage4)

```

---

auditPrior

---

*Prior Distributions for Audit Sampling*


---

## Description

auditPrior is used to create a prior distribution for Bayesian audit sampling. The interface allows a complete customization of the prior distribution as well as a formal translation of pre-existing audit information into a prior distribution. auditPrior returns an object of class jfaPrior that can

be subsequently used in the `planning()` and `evaluation()` functions via their prior argument. Objects with class `jfaPrior` can be used with associated `summary()` and `plot()` methods.

For more details on how to use this function, see the package vignette: `vignette('jfa', package = 'jfa')`

## Usage

```
auditPrior(method = 'default', likelihood = c('poisson', 'binomial', 'hypergeometric'),
  N.units = NULL, alpha = NULL, beta = NULL, materiality = NULL, expected = 0,
  ir = NULL, cr = NULL, ub = NULL, p.hmin = NULL, x = NULL,
  n = NULL, factor = NULL, conf.level = 0.95)
```

## Arguments

|                          |  |
|--------------------------|--|
| <code>method</code>      | a character specifying the method by which the prior distribution is constructed. Possible options are <code>default</code> , <code>strict</code> , <code>impartial</code> , <code>param</code> , <code>arm</code> , <code>bram</code> , <code>hyp</code> , <code>sample</code> , and <code>factor</code> . See the details section for more information.              |
| <code>likelihood</code>  | a character specifying the likelihood for updating the prior distribution. Possible options are <code>poisson</code> (default) for a conjugate gamma prior distribution, <code>binomial</code> for a conjugate beta prior distribution, or <code>hypergeometric</code> for a conjugate beta-binomial prior distribution. See the details section for more information. |
| <code>N.units</code>     | a numeric value larger than 0 specifying the total number of units in the population. Only used for the <code>hypergeometric</code> likelihood.  |
| <code>alpha</code>       | a numeric value specifying the $\alpha$ parameter of the prior distribution. Only used for method <code>param</code> .   |
| <code>beta</code>        | a numeric value specifying the $\beta$ parameter of the prior distribution. Only used for method <code>param</code> .  |
| <code>materiality</code> | a numeric value between 0 and 1 specifying the performance materiality (i.e., the maximum misstatement) as a fraction of the total number of units in the population. Only used for methods <code>impartial</code> , <code>arm</code> , and <code>hyp</code> .   |
| <code>expected</code>    | a numeric value between 0 and 1 specifying the expected errors in the sample relative to the total sample size. Only used for methods <code>impartial</code> , <code>arm</code> , <code>bram</code> , and <code>hyp</code> .   |
| <code>ir</code>          | a numeric value between 0 and 1 specifying the inherent risk in the audit risk model. Defaults to 1 for 100% risk. Only used for method <code>arm</code> .   |
| <code>cr</code>          | a numeric value between 0 and 1 specifying the internal control risk in the audit risk model. Defaults to 1 for 100% risk. Only used for method <code>arm</code> .   |
| <code>ub</code>          | a numeric value between 0 and 1 specifying the <code>conf.level</code> -% upper bound for the prior distribution as a fraction of the total number of units in the population. Only used for method <code>bram</code> .  |
| <code>p.hmin</code>      | a numeric value between 0 and 1 specifying the prior probability of the hypothesis of tolerable misstatement ( $H1: \theta < \text{materiality}$ ). Only used for method <code>hyp</code> .  |
| <code>x</code>           | a numeric value larger than, or equal to, 0 specifying the sum of proportional errors (taints) in a prior sample. Only used for methods <code>sample</code> and <code>factor</code> .  |
| <code>n</code>           | a numeric value larger than 0 specifying the sample size of a prior sample. Only used for methods <code>sample</code> and <code>factor</code> .  |

|            |   |
|------------|---|
| factor     | a numeric value between 0 and 1 specifying the weight of the prior sample. Only used for method factor. |
| conf.level | a numeric value between 0 and 1 specifying the confidence level. Defaults to 0.95 for 95% confidence.   |

## Details

To perform Bayesian audit sampling you must assign a prior probability distribution to the parameter in the model, i.e., the population misstatement  $\theta$ . The prior distribution can incorporate pre-existing audit information about  $\theta$  before seeing a sample, which consequently allows for a more efficient or more accurate estimate of  $\theta$ . However, the default priors used in `jfa` are purposely indifferent towards the individual values of  $\theta$ . For these priors, the posterior distribution is based solely on the data from the sample, i.e., the likelihood. Note that the default prior is a conservative choice of prior since it assumes all possible misstatement is equally likely before seeing information from a data sample. It is therefore strongly recommended to construct an informed prior distribution based on pre-existing audit information if possible.

This section elaborates on the available options for the method argument.

- **default**: This method produces a  $\text{gamma}(1, 1)$ ,  $\text{beta}(1, 1)$ , or  $\text{beta-binomial}(N, 1, 1)$  prior distribution. These priors are indifferent towards the possible values of the misstatement.
- **strict**: This method produces an improper  $\text{gamma}(1, 0)$ ,  $\text{beta}(1, 0)$ , or  $\text{beta-binomial}(N, 1, 0)$  prior distribution. These prior distributions exactly match sample sizes and upper limits from classical methods.
- **impartial**: This method produces an impartial prior distribution. These prior distributions assume that tolerable misstatement ( $\theta < \text{materiality}$ ) and intolerable misstatement ( $\theta > \text{materiality}$ ) are equally likely.
- **param**: This method produces a  $\text{gamma}(\alpha, \beta)$ ,  $\text{beta}(\alpha, \beta)$ , or  $\text{beta-binomial}(N, \alpha, \beta)$  prior distribution.
- **hyp**: This method translates an assessment of the prior probability for tolerable misstatement ( $\theta < \text{materiality}$ ) to a prior distribution.
- **arm**: This method translates an assessment of inherent risk and internal control risk (Audit Risk Model, Derks et al., 2021) to a prior distribution.
- **bram**: This method translates an assessment of the expected most likely error and x-% upper bound to a prior distribution.
- **sample**: This method translates sampling results from an earlier sample to a prior distribution.
- **factor**: This method translates and weighs sampling results from an earlier sample to a prior distribution.

This section elaborates on the available likelihoods and corresponding prior distributions for the likelihood argument.

- **poisson**: The Poisson distribution is an approximation of the binomial distribution. The Poisson distribution is defined as:

$$f(\theta, n) = \frac{\lambda^\theta e^{-\lambda}}{\theta!}$$

The conjugate  $\text{gamma}(\alpha, \beta)$  prior has probability density function:

$$p(\theta; \alpha, \beta) = \frac{\beta^\alpha \theta^{\alpha-1} e^{-\beta\theta}}{\Gamma(\alpha)}$$

- **binomial:** The binomial distribution is an approximation of the hypergeometric distribution. The binomial distribution is defined as:

$$f(\theta, n, x) = \binom{n}{x} \theta^x (1 - \theta)^{n-x}$$

The conjugate  $beta(\alpha, \beta)$  prior has probability density function:

$$p(\theta; \alpha, \beta) = \frac{1}{B(\alpha, \beta)} \theta^{\alpha-1} (1 - \theta)^{\beta-1}$$

- **hypergeometric:** The hypergeometric distribution is defined as:

$$f(x, n, K, N) = \frac{\binom{K}{x} \binom{N-K}{n-x}}{\binom{N}{n}}$$

The conjugate  $beta\text{-binomial}(\alpha, \beta)$  prior (Dyer and Pierce, 1993) has probability mass function:

$$f(x, n, \alpha, \beta) = \binom{n}{x} \frac{B(x + \alpha, n - x + \beta)}{B(\alpha, \beta)}$$

## Value

An object of class `jfaPrior` containing:

|                          |   |
|--------------------------|---|
| <code>prior</code>       | a string describing the functional form of the prior distribution.  |
| <code>description</code> | a list containing a description of the prior distribution, including the parameters of the prior distribution and the implicit sample on which the prior distribution is based.   |
| <code>statistics</code>  | a list containing statistics of the prior distribution, including the mean, mode, median, and upper bound of the prior distribution.  |
| <code>specifics</code>   | a list containing specifics of the prior distribution that vary depending on the method.  |
| <code>hypotheses</code>  | if <code>materiality</code> is specified, a list containing information about the hypotheses, including prior probabilities and odds for the hypothesis of tolerable misstatement (H1) and the hypothesis of intolerable misstatement (H0). |
| <code>method</code>      | a character indicating the method by which the prior distribution is constructed.   |
| <code>likelihood</code>  | a character indicating the likelihood of the data.  |
| <code>materiality</code> | if <code>materiality</code> is specified, a numeric value between 0 and 1 indicating the materiality used to construct the prior distribution.  |
| <code>expected</code>    | a numeric value larger than, or equal to, 0 indicating the input for the number of expected errors.   |
| <code>conf.level</code>  | a numeric value between 0 and 1 indicating the confidence level used.   |
| <code>N.units</code>     | if <code>N.units</code> is specified, the number of units in the population.  |

## Author(s)

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## References

Derks, K., de Swart, J., Wagenmakers, E.-J., Wille, J., & Wetzels, R. (2021). JASP for audit: Bayesian tools for the auditing practice. *Journal of Open Source Software*, 6(68), 2733.

Derks, K., de Swart, J., van Batenburg, P., Wagenmakers, E.-J., & Wetzels, R. (2021). Priors in a Bayesian audit: How integration of existing information into the prior distribution can improve audit transparency and efficiency. *International Journal of Auditing*, 25(3), 621-636.

## See Also

[planning selection evaluation report](#)

## Examples

```
# Default uniform beta(1, 1) prior distribution
auditPrior(method = "default", likelihood = "binomial")

# Translate inherent risk (ir) and control risk (cr) to a gamma prior distribution
auditPrior(method = "arm", expected = 0.025, materiality = 0.05, ir = 1, cr = 0.6)

# Impartial beta prior distribution (equal prior probabilities)
auditPrior(method = "impartial", likelihood = "binomial", materiality = 0.05)
```

---

BuildIt

*BuildIt Construction Financial Statements*

---

## Description

Fictional data from a construction company in the United States, containing 3500 observations identification numbers, book values, and audit values. The audit values are added for illustrative purposes, as these would need to be assessed by the auditor in the execution stage of the audit.

## Usage

```
data(BuildIt)
```

## Format

A data frame with 3500 rows and 3 variables.

**ID** unique record identification number.

**bookValue** book value in US dollars (\$14.47–\$2,224.40).

**auditValue** true value in US dollars (\$14.47–\$2,224.40).

## References

Derks, K., de Swart, J., Wagenmakers, E.-J., Wille, J., & Wetzels, R. (2019). JASP for audit: Bayesian tools for the auditing practice.

## Examples

```
data(BuildIt)
```



---

carrier*Carrier Company Financial Statements*

---

### Description

Fictional data from a carrier company in Europe, containing 202 ledger items across 10 company entities.

### Usage

```
data(carrier)
```

### Format

A data frame with 202 rows and 12 variables.

**description** description of the ledger item.

**entity1** recorded values for entity 1, in US dollars.

**entity2** recorded values for entity 2, in US dollars.

**entity3** recorded values for entity 3, in US dollars.

**entity4** recorded values for entity 4, in US dollars.

**entity5** recorded values for entity 5, in US dollars.

**entity6** recorded values for entity 6, in US dollars.

**entity7** recorded values for entity 7, in US dollars.

**entity8** recorded values for entity 8, in US dollars.

**entity9** recorded values for entity 9, in US dollars.

**entity10** recorded values for entity 10, in US dollars.

**total** total value, in US dollars.

### Source

<https://towardsdatascience.com/data-driven-audit-1-automated-sampling-using-python-52e83347add5>

### Examples

```
data(carrier)
```

evaluation

*Evaluate a statistical audit sample***Description**

This function takes a data frame (using `data`, `values`, and `values.audit`) or summary statistics (using `x` and `n`) and performs inference on the misstatement in the sample. The function returns an object of class `jfaEvaluation` which can be used with associated `summary()` and `plot()` methods.

For more details on how to use this function, see the package vignette: `vignette('jfa', package = 'jfa')`

**Usage**

```
evaluation(materiality = NULL, min.precision = NULL, method = 'poisson',
            alternative = c('less', 'two.sided', 'greater'), conf.level = 0.95,
            data = NULL, values = NULL, values.audit = NULL, times = NULL,
            x = NULL, n = NULL, N.units = NULL, N.items = NULL,
            r.delta = 2.7, m.type = 'accounts', cs.a = 1, cs.b = 3, cs.mu = 0.5,
            prior = FALSE)
```

**Arguments**

|                            |   |
|----------------------------|---|
| <code>materiality</code>   | a numeric value between 0 and 1 specifying the performance materiality (maximum tolerable error) as a fraction of the total size of the population. If specified, the function also returns the conclusion of the analysis with respect to the performance materiality. The value is discarded when direct, difference, quotient, or regression method is chosen.   |
| <code>min.precision</code> | a numeric value between 0 and 1 specifying the required minimum precision (upper bound minus most likely error) as a fraction of the total size of the population. If specified, the function also returns the conclusion of the analysis with respect to the required minimum precision.   |
| <code>method</code>        | a character specifying the method to be used in the evaluation. Possible options are <code>poisson</code> , <code>binomial</code> (default), <code>hypergeometric</code> , <code>mpu</code> , <code>stringer</code> , <code>stringer.meikle</code> , <code>stringer.lta</code> , <code>stringer.pvz</code> , <code>rohrbach</code> , <code>moment</code> , <code>direct</code> , <code>difference</code> , <code>quotient</code> , or <code>regression</code> . See the details section for more information. |
| <code>alternative</code>   | a character indicating the alternative hypothesis to be tested (and the type of interval to be produced). This must be one of <code>less</code> (default), <code>two.sided</code> , or <code>greater</code> . You can specify just the initial letter.  |
| <code>conf.level</code>    | a numeric value between 0 and 1 specifying the confidence level used in the evaluation. Defaults to 0.95 for 95% confidence.  |
| <code>data</code>          | a data frame containing the sample to be evaluated. The sample must at least contain a column of book values and a column of audit (true) values.   |
| <code>values</code>        | a character specifying name of a column in <code>data</code> containing the book values of the items.   |
| <code>values.audit</code>  | a character specifying name of a column in <code>data</code> containing the audit (true) values of the items.   |
| <code>times</code>         | a character specifying name of a column in <code>data</code> containing the number of times each item in the data should be counted in the evaluation (due to it being selected multiple times for the sample).   |

|         |   |
|---------|---|
| x       | a numeric value larger than 0 specifying the sum of errors found in the sample. If specified, overrides the data, values and values.audit arguments and assumes that the data come from summary statistics specified by both x and n.   |
| n       | an integer larger than 0 specifying the number of items in the sample. If specified, overrides the data, values and values.audit arguments and assumes that the data come from summary statistics specified by both x and n.  |
| N.units | an integer larger than 0 specifying the total number of sampling units in the population (i.e., the population size / value). Only required if method is one of 'hypergeometric', direct, difference, quotient, or regression.  |
| N.items | an integer larger than 0 specifying the total number of items in the population. Only required if method is one of direct, difference, quotient, or regression.   |
| r.delta | if method = 'rohrbach', a numeric value specifying $\Delta$ in Rohrbach's augmented variance bound (Rohrbach, 1993).  |
| m.type  | if method = 'moment', a character specifying the type of population (Dworin and Grimlund, 1984). Possible options are accounts and inventory. This argument affects the calculation of the central moments in the bound.  |
| cs.a    | if method = "coxsnell", a numeric value specifying the $\alpha$ parameter of the prior distribution on the mean taint. Defaults to 1 as recommended by Cox and Snell (1979).  |
| cs.b    | if method = "coxsnell", a numeric value specifying the $\beta$ parameter of the prior distribution on the mean taint. Defaults to 3 as recommended by Cox and Snell (1979).   |
| cs.mu   | if method = "coxsnell", a numeric value between 0 and 1 specifying the mean of the prior distribution on the mean taint. Defaults to 0.5 as recommended by Cox and Snell (1979).  |
| prior   | a logical specifying if a prior distribution must be used, or an object of class jfaPrior or jfaPosterior containing the prior distribution. Defaults to FALSE for frequentist planning. If TRUE, a minimal information prior is chosen by default. Chooses a conjugate gamma distribution for the Poisson likelihood, a conjugate beta distribution for the binomial likelihood, and a conjugate beta-binomial distribution for the hypergeometric likelihood. |

## Details

This section lists the available options for the methods argument.

- **poisson**: Evaluates the sample with the Poisson distribution. If combined with prior = TRUE, performs Bayesian evaluation using a *gamma* prior and posterior.
- **binomial**: Evaluates the sample with the binomial distribution. If combined with prior = TRUE, performs Bayesian evaluation using a *beta* prior and posterior.
- **hypergeometric**: Evaluates the sample with the hypergeometric distribution. If combined with prior = TRUE, performs Bayesian evaluation using a *beta-binomial* prior and posterior.
- **mpu**: Evaluates the sample with the mean-per-unit estimator.
- **stringer**: Evaluates the sample with the Stringer bound (Stringer, 1963).
- **stringer.meikle**: Evaluates the sample with the Stringer bound with Meikle's correction for understatements (Meikle, 1972).
- **stringer.lta**: Evaluates the sample with the Stringer bound with LTA correction for understatements (Leslie, Teitlebaum, and Anderson, 1979).

- `stringer.pvz`: Evaluates the sample with the Stringer bound with Pap and van Zuijlen's correction for understatements (Pap and van Zuijlen, 1996).
- `rohrbach`: Evaluates the sample with Rohrbach's augmented variance bound (Rohrbach, 1993).
- `moment`: Evaluates the sample with the modified moment bound (Dworin and Grimlund, 1984).
- `coxsnell`: Evaluates the sample with the Cox and Snell bound (Cox and Snell, 1979).
- `direct`: Evaluates the sample with the direct estimator (Touw and Hoogduin, 2011).
- `difference`: Evaluates the sample with the difference estimator (Touw and Hoogduin, 2011).
- `quotient`: Evaluates the sample with the quotient estimator (Touw and Hoogduin, 2011).
- `regression`: Evaluates the sample with the regression estimator (Touw and Hoogduin, 2011).

## Value

An object of class `jfaEvaluation` containing:

|                            |   |
|----------------------------|---|
| <code>conf.level</code>    | a numeric value between 0 and 1 indicating the confidence level used.   |
| <code>mle</code>           | a numeric value between 0 and 1 indicating the most likely error in the population as a fraction of its total size.   |
| <code>ub</code>            | a numeric value indicating the upper bound on the (probability of) misstatement.  |
| <code>lb</code>            | a numeric value indicating the lower bound of the interval around the (probability of) misstatement.  |
| <code>precision</code>     | a numeric value between 0 and 1 indicating the difference between the most likely error and the upper bound in the population as a fraction of the total population size. |
| <code>p.value</code>       | a numeric value indicating the one-sided p-value.   |
| <code>x</code>             | an integer larger than, or equal to, 0 indicating the number of items in the sample that contained an error.  |
| <code>t</code>             | a value larger than, or equal to, 0, indicating the sum of observed taints.   |
| <code>n</code>             | an integer larger than 0 indicating the sample size.  |
| <code>materiality</code>   | if <code>materiality</code> is specified, a numeric value between 0 and 1 indicating the performance materiality as a fraction of the total population size.              |
| <code>min.precision</code> | if <code>min.precision</code> is specified, a numeric value between 0 and 1 indicating the minimum required precision as a fraction of the total population size.         |
| <code>alternative</code>   | a character indicating the alternative hypothesis.  |
| <code>method</code>        | a character indicating the evaluation method.   |
| <code>N.units</code>       | if <code>N.units</code> is specified, in integer larger than 0 indicating the total number of units in the population.  |
| <code>N.items</code>       | if <code>N.items</code> is specified, in integer larger than 0 indicating the total number of items in the population.  |
| <code>K</code>             | if <code>method = 'hypergeometric'</code> , an integer indicating the assumed total errors in the population.   |
| <code>prior</code>         | an object of class <code>'jfaPrior'</code> that contains the prior distribution.  |
| <code>posterior</code>     | an object of class <code>'jfaPosterior'</code> that contains the posterior distribution.  |
| <code>data</code>          | a data frame containing the relevant columns from the data.   |
| <code>data.name</code>     | a character string giving the name of the data.   |

**Author(s)**

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**References**

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**See Also**

[auditPrior planning selection report](#)

**Examples**

```
data("BuildIt")

# Draw a sample of 100 monetary units from the population using
# fixed interval monetary unit sampling
sample <- selection(
  data = BuildIt, size = 100, units = "values",
  method = "interval", values = "bookValue"
)$sample

# Classical evaluation using the Stringer bound
evaluation(
  materiality = 0.05, method = "stringer", conf.level = 0.95,
  data = sample, values = "bookValue", values.audit = "auditValue"
)

# Classical evaluation using the Poisson likelihood
evaluation(
  materiality = 0.05, method = "poisson", conf.level = 0.95,
  data = sample, values = "bookValue", values.audit = "auditValue"
)
```

```
# Bayesian evaluation using a noninformative gamma prior distribution
evaluation(
  materiality = 0.05, method = "poisson", conf.level = 0.95,
  data = sample, values = "bookValue", values.audit = "auditValue",
  prior = TRUE
)

# Bayesian evaluation using an informed prior distribution
evaluation(
  materiality = 0.05, method = "poisson", conf.level = 0.95,
  data = sample, values = "bookValue", values.audit = "auditValue",
  prior = auditPrior(method = "param", alpha = 1, beta = 10)
)
```

---

jfa-methods

*Methods for jfa objects*


---

## Description

Methods defined for objects returned from the [auditPrior](#), [planning](#), [selection](#), and [evaluation](#) functions.

## Usage

```
## S3 method for class 'jfaPrior'
print(x, ...)

## S3 method for class 'jfaPosterior'
print(x, ...)

## S3 method for class 'jfaPlanning'
print(x, ...)

## S3 method for class 'jfaSelection'
print(x, ...)

## S3 method for class 'jfaEvaluation'
print(x, digits = getOption("digits"), ...)

## S3 method for class 'summary.jfaPrior'
print(x, digits = getOption("digits"), ...)

## S3 method for class 'summary.jfaPosterior'
print(x, digits = getOption("digits"), ...)

## S3 method for class 'summary.jfaPlanning'
print(x, digits = getOption("digits"), ...)

## S3 method for class 'summary.jfaSelection'
print(x, digits = getOption("digits"), ...)

## S3 method for class 'summary.jfaEvaluation'
```

```

print(x, digits = getOption("digits"), ...)

## S3 method for class 'jfaPrior'
summary(object, digits = getOption("digits"), ...)

## S3 method for class 'jfaPosterior'
summary(object, digits = getOption("digits"), ...)

## S3 method for class 'jfaPlanning'
summary(object, digits = getOption("digits"), ...)

## S3 method for class 'jfaSelection'
summary(object, digits = getOption("digits"), ...)

## S3 method for class 'jfaEvaluation'
summary(object, digits = getOption("digits"), ...)

## S3 method for class 'jfaPrior'
predict(object, n, lim = NULL, cumulative = FALSE, ...)

## S3 method for class 'jfaPosterior'
predict(object, n, lim = NULL, cumulative = FALSE, ...)

## S3 method for class 'jfaPrior'
plot(x, xlim = c(0, 1), ...)

## S3 method for class 'jfaPosterior'
plot(x, xlim = c(0, 1), ...)

## S3 method for class 'jfaPlanning'
plot(x, xlim = c(0, 1), ...)

## S3 method for class 'jfaSelection'
plot(x, ...)

## S3 method for class 'jfaEvaluation'
plot(x, xlim = c(0, 1), ...)

```

### Arguments

|            |  |
|------------|--|
| ...        | further arguments, currently ignored.  |
| digits     | an integer specifying the number of digits to which output should be rounded. Used in summary. |
| object, x  | an object of class jfaPrior, jfaPosterior, jfaPlanning, jfaSelection, or jfaEvaluation.        |
| n          | used in predict. Specifies the sample size for which predictions should be made.               |
| lim        | used in predict. Limits the number of errors for which predictions should be made.             |
| cumulative | used in predict. Specifies whether cumulative probabilities should be shown.                   |
| xlim       | used in plot. Specifies the x limits (x1, x2) of the plot.                                     |

**Value**

The summary methods return a `data.frame` which contains the input and output.

The print methods simply print and return nothing.

---

planning

*Plan a Statistical Audit Sample*


---

**Description**

This function calculates the minimum sample size for a statistical audit sample based on the Poisson, binomial, or hypergeometric likelihood. The function returns an object of class `jfaPlanning` which can be used with associated `summary()` and `plot()` methods.

For more details on how to use this function, see the package vignette: `vignette('jfa', package = 'jfa')`

**Usage**

```
planning(materiality = NULL, min.precision = NULL, expected = 0,
          likelihood = c('poisson', 'binomial', 'hypergeometric'),
          conf.level = 0.95, N.units = NULL, by = 1, max = 5000,
          prior = FALSE)
```

**Arguments**

|               |   |
|---------------|---|
| materiality   | a numeric value between 0 and 1 specifying the performance materiality (i.e., maximum upper limit) as a fraction of the total population size. Can be <code>NULL</code> , but <code>min.precision</code> should be specified in that case.  |
| min.precision | a numeric value between 0 and 1 specifying the minimum precision (i.e., upper bound minus most likely error) as a fraction of the total population size. Can be <code>NULL</code> , but <code>materiality</code> should be specified in that case.  |
| expected      | a numeric value between 0 and 1 specifying the expected errors in the sample relative to the total sample size, or a number ( $\geq 1$ ) that represents the number of expected errors in the sample. It is advised to set this value conservatively to minimize the probability of the observed errors exceeding the expected errors, which would imply that insufficient work has been done in the end. |
| likelihood    | a character specifying the likelihood assumed in the calculation. This can be either <code>poisson</code> (default) for the Poisson likelihood, <code>binomial</code> for the binomial likelihood, or <code>hypergeometric</code> for the hypergeometric likelihood. See the details section for more information about the available likelihoods.  |
| conf.level    | a numeric value between 0 and 1 specifying the confidence level used in the planning. Defaults to 0.95 for 95% confidence.  |
| N.units       | an integer larger than 0 specifying the total number of units or items in the population (i.e., the population size). Only required when <code>likelihood = 'hypergeometric'</code> .   |
| by            | an integer larger than 0 specifying the desired increment for the sample size calculation.  |
| max           | an integer larger than 0 specifying the maximum sample size that is considered in the calculation. Defaults to 5000 for efficiency. Increase this value if the sample size cannot be found due to it being too large (e.g., for a low materiality).   |



**prior** a logical specifying whether to use a prior distribution when planning, or an object of class `jfaPrior` or `jfaPosterior` containing the prior distribution. Defaults to `FALSE` for frequentist planning. If `TRUE`, a minimal information prior is chosen by default. Chooses a conjugate gamma distribution for the Poisson likelihood, a conjugate beta distribution for the binomial likelihood, and a conjugate beta-binomial distribution for the hypergeometric likelihood.

## Details

This section elaborates on the available likelihoods and corresponding prior distributions for the likelihood argument.

- **poisson**: The Poisson distribution is an approximation of the binomial distribution. The Poisson distribution is defined as:

$$f(\theta, n) = \frac{\lambda^\theta e^{-\lambda}}{\theta!}$$

The conjugate  $gamma(\alpha, \beta)$  prior has probability density function:

$$p(\theta; \alpha, \beta) = \frac{\beta^\alpha \theta^{\alpha-1} e^{-\beta\theta}}{\Gamma(\alpha)}$$

- **binomial**: The binomial distribution is an approximation of the hypergeometric distribution. The binomial distribution is defined as:

$$f(\theta, n, x) = \binom{n}{x} \theta^x (1 - \theta)^{n-x}$$

The conjugate  $beta(\alpha, \beta)$  prior has probability density function:

$$p(\theta; \alpha, \beta) = \frac{1}{B(\alpha, \beta)} \theta^{\alpha-1} (1 - \theta)^{\beta-1}$$

- **hypergeometric**: The hypergeometric distribution is defined as:

$$f(x, n, K, N) = \frac{\binom{K}{x} \binom{N-K}{n-x}}{\binom{N}{n}}$$

The conjugate  $beta-binomial(\alpha, \beta)$  prior (Dyer and Pierce, 1993) has probability mass function:

$$f(x, n, \alpha, \beta) = \binom{n}{x} \frac{B(x + \alpha, n - x + \beta)}{B(\alpha, \beta)}$$

## Value

An object of class `jfaPlanning` containing:

|                         |   |
|-------------------------|---|
| <code>conf.level</code> | a numeric value between 0 and 1 indicating the confidence level used.                                     |
| <code>x</code>          | a numeric value larger than, or equal to, 0 indicating the number of tolerable errors in the sample.      |
| <code>n</code>          | an integer larger than 0 indicating the required sample size.   |
| <code>ub</code>         | a numeric value between 0 and 1 indicating the expected upper bound if the sample goes according to plan. |

|               |   |
|---------------|---|
| precision     | a numeric value between 0 and 1 indicating the expected precision if the sample goes according to plan.   |
| p.value       | a numeric value indicating the one-sided p-value.   |
| K             | if likelihood = 'hypergeometric', an integer larger than 0 indicating the assumed population errors.  |
| N.units       | an integer larger than 0 indicating the population size (only returned if N is specified).  |
| materiality   | a numeric value between 0 and 1 indicating the specified materiality.   |
| min.precision | a numeric value between 0 and 1 indicating the minimum precision to be obtained.  |
| expected      | a numeric value larger than, or equal to, 0 indicating the expected errors input.   |
| likelihood    | a character indicating the specified likelihood.  |
| errorType     | a character indicating whether the expected errors were specified as a percentage or as an integer.   |
| iterations    | a numeric value indicating the number of iterations used.   |
| prior         | if a prior distribution is specified, an object of class <code>jfaPrior</code> that contains information about the prior distribution.                  |
| posterior     | if a prior distribution is specified, an object of class <code>jfaPosterior</code> that contains information about the expected posterior distribution. |

### Author(s)

Koen Derks, <k.derks@nyenrode.nl>

### References

- Derks, K., de Swart, J., van Batenburg, P., Wagenmakers, E.-J., & Wetzels, R. (2021). Priors in a Bayesian audit: How integration of existing information into the prior distribution can improve audit transparency and efficiency. *International Journal of Auditing*, 25(3), 621-636.
- Dyer, D. and Pierce, R.L. (1993). On the choice of the prior distribution in hypergeometric sampling. *Communications in Statistics - Theory and Methods*, 22(8), 2125 - 2146.

### See Also

[auditPrior selection evaluation report](#)

### Examples

```
# Classical planning using a Poisson likelihood
planning(materiality = 0.05, expected = 0.025, likelihood = "poisson")

# Bayesian planning using a noninformative gamma prior distribution
planning(
  materiality = 0.05, expected = 0.025, likelihood = "poisson",
  prior = TRUE
)

# Bayesian planning using an informed gamma prior distribution
planning(
  materiality = 0.05, expected = 0.025, likelihood = "poisson",
  prior = auditPrior(method = "impartial", materiality = 0.05)
)
```

---

|        |   |
|--------|---|
| report | <i>Create a statistical audit sampling report</i> |
|--------|---|

---

## Description

This function takes an object of class `jfaEvaluation` as returned by the `evaluation()` function automatically generates a html or pdf report containing the analysis results and their interpretation.

For more details on how to use this function, see the package vignette: `vignette('jfa', package = 'jfa')`

## Usage

```
report(object, file = 'report.html', format = c('html_document', 'pdf_document'))
```

## Arguments

|                     |   |
|---------------------|---|
| <code>object</code> | an object of class <code>jfaEvaluation</code> as returned by the <code>evaluation()</code> function.  |
| <code>file</code>   | a character specifying the name of the report (e.g. <code>report.html</code> ). By default, the report is created in your current working directory.  |
| <code>format</code> | a character specifying the output format of the report. Possible options are <code>html_document</code> (default) and <code>pdf_document</code> , but compiling to pdf format requires a local version of MikTeX. |

## Value

A html or pdf file containing a report of the evaluation.

## Author(s)

Koen Derks, <k.derks@nyenrode.nl>

## See Also

[auditPrior planning selection evaluation](#)

## Examples

```
data("BuildIt")

# Draw a sample of 100 monetary units from the population using
# fixed interval monetary unit sampling
sample <- selection(
  data = BuildIt, size = 100, method = "interval",
  units = "values", values = "bookValue"
)$sample

# Evaluate using the Stringer bound
result <- evaluation(
  conf.level = 0.95, materiality = 0.05, method = "stringer",
  data = sample, values = "bookValue", values.audit = "auditValue"
)
## Not run:
```

```
report(result)

## End(Not run)
```

---

selection

*Select a Statistical Audit Sample*


---

## Description

This function takes a data frame and performs statistical selection according to one of four algorithms: fixed interval sampling, cell sampling, random sampling, and modified sieve sampling. Selection is done on the level of two possible sampling units: items (records / rows) or monetary units. The function returns an object of class `jfaSelection` which can be used with associated `summary()` and `plot()` methods.

For more details on how to use this function, see the package vignette: `vignette('jfa', package = 'jfa')`

## Usage

```
selection(data, size, units = c('items', 'values'),
          method = c('interval', 'cell', 'random', 'sieve'), values = NULL,
          order = NULL, decreasing = FALSE, randomize = FALSE,
          replace = FALSE, start = 1)
```

## Arguments

|                         |  |
|-------------------------|--|
| <code>data</code>       | a data frame containing the population of items the auditor wishes to sample from.   |
| <code>size</code>       | an integer larger than 0 specifying the number of sampling units that need to be selected from the population. Can also be an object of class <code>jfaPlanning</code> .   |
| <code>units</code>      | a character specifying the sampling units used. Possible options are <code>items</code> (default) for selection on the level of items (rows) or <code>values</code> for selection on the level of monetary units.  |
| <code>method</code>     | a character specifying the sampling algorithm used. Possible options are <code>interval</code> (default) for fixed interval sampling, <code>cell</code> for cell sampling, <code>random</code> for random sampling, or <code>sieve</code> for modified sieve sampling. |
| <code>values</code>     | a character specifying the name of a column in <code>data</code> containing the book values of the items.  |
| <code>order</code>      | a character specifying the name of a column in <code>data</code> containing the ranks of the items. The items in the data are ordered according to these values in the order indicated by <code>decreasing</code> .  |
| <code>decreasing</code> | if <code>order</code> is specified, a logical specifying whether to order the items from smallest to largest. Defaults to <code>FALSE</code> .   |
| <code>randomize</code>  | a logical specifying whether the items in the data should be randomly shuffled before selection. Defaults to <code>FALSE</code> . Note that specifying if <code>randomize = TRUE</code> overrules <code>order</code> .   |
| <code>replace</code>    | if <code>method = 'random'</code> , a logical specifying whether sampling should be performed with replacement. Defaults to <code>FALSE</code> .   |
| <code>start</code>      | if <code>method = 'interval'</code> , an integer larger than 0 specifying the starting point of the algorithm.   |

## Details

The first part of this section elaborates on the two possible options for the `units` argument:

- `items`: In record sampling each item in the population is seen as a sampling unit. An item of \$5000 is therefore equally likely to be selected as an item of \$500.
- `values`: In monetary unit sampling each monetary unit in the population is seen as a sampling unit. An item of \$5000 is therefore ten times more likely to be selected as an item of \$500.

The second part of this section elaborates on the four possible options for the `method` argument:

- `interval`: In fixed interval sampling the sampling units in the population are divided into a number (equal to the sample size) of intervals. From each interval one sampling unit is selected according to a fixed starting point (specified by `start`).
- `cell`: In cell sampling the sampling units in the population are divided into a number (equal to the sample size) of intervals. From each interval one sampling unit is selected with equal probability.
- `random`: In random sampling each sampling unit in the population is drawn with equal probability.
- `sieve`: In modified sieve sampling each item in the population is selected proportional to its value (Hoogduin, Hall, & Tsay, 2010).

## Value

An object of class `jfaSelection` containing:

|                        |  |
|------------------------|--|
| <code>data</code>      | a data frame containing the input data.  |
| <code>sample</code>    | a data frame containing the selected sample of items.  |
| <code>n.req</code>     | an integer indicating the requested sample size.   |
| <code>n.units</code>   | an integer indicating the total number of obtained sampling units.                                   |
| <code>n.items</code>   | an integer indicating the total number of obtained sample items.                                     |
| <code>N.units</code>   | an integer indicating the total number of sampling units in the population.                          |
| <code>N.items</code>   | an integer indicating the total number of items in the population.                                   |
| <code>interval</code>  | if <code>method = 'interval'</code> , a numeric value indicating the size of the selection interval. |
| <code>units</code>     | a character indicating the sampling units that were used to create the selection.                    |
| <code>method</code>    | a character indicating the the algorithm that was used to create the selection.                      |
| <code>values</code>    | if <code>values</code> is specified, a character indicating the name of the book value column.       |
| <code>start</code>     | if <code>method = 'interval'</code> , an integer indicating the starting point in the interval.      |
| <code>data.name</code> | a character string giving the name of the data.  |

## Author(s)

Koen Derks, <k.derks@nyenrode.nl>

## References

- Hoogduin, L. A., Hall, T. W., & Tsay, J. J. (2010). Modified sieve sampling: A method for single- and multi-stage probability-proportional-to-size sampling. *Auditing: A Journal of Practice & Theory*, 29(1), 125-148.
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- Wampler, B., & McEacharn, M. (2005). Monetary-unit sampling using Microsoft Excel. *The CPA journal*, 75(5), 36.

## See Also

[auditPrior planning evaluation report](#)

## Examples

```
data("BuildIt")

# Select 100 items using random sampling
selection(data = BuildIt, size = 100, method = "random")

# Select 150 monetary units using fixed interval sampling
selection(
  data = BuildIt, size = 150, units = "values",
  method = "interval", values = "bookValue"
)
```

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