brr: Basic usage.

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The **brr** package performs Bayesian inference on the rate ratio $\phi = \frac{\lambda}{\mu}$ in the *two Poisson samples model* given by two independent observations

$$\begin{cases} x \sim \mathcal{P}(\lambda S) \\ y \sim \mathcal{P}(\mu T) \end{cases}$$

where λ and μ are the unknown incidence rates and S and T are the known observation-opportunity sizes, or, for short, the sample sizes. S and T are also called the times at risk when they represent some durations.

The **brr** package implements the *semi-conjugate family of prior distributions*. Precisely, for positive numbers a, b, c and d, to be set by the user, the following independent prior distributions are assigned on μ and $\phi = \lambda/\mu$:

$$\mu \sim \mathcal{G}(a,b) \quad \text{and} \quad \phi \sim \frac{T+b}{S} \times \mathcal{B}'(c,d),$$

Then the joint posterior on (μ, ϕ) is given by

$$(\mu \mid \phi, x, y) \sim \mathcal{G}(a + x + y, b + \phi S + T) \quad \text{and} \quad (\phi \mid x, y) \sim \frac{T + b}{S} \times \mathcal{B}'(c + x, a + d + y).$$

In particular:

- when a=c=0.5 and b=d=0, the prior is the reference prior, also called the *non-informative prior*;
- when a, b > 0, c = 0.5 and d = 0, the prior is the semi-reference prior, also called the *semi-informative prior*, that is to say the reference prior after the arbitrary Gamma prior distribution $\mathcal{G}(a, b)$ is assigned on μ .

Setting parameters with brr

Use the **Brr** function to set the prior parameters, the sample sizes, and the observed counts. One can proceed step by step, for example below we start by supplying the parameters a and b of the prior Gamma distribution on μ :

```
library(brr)
model <- Brr(a=2,b=3)
summary(model)</pre>
```

```
## Type of prior distribution: semi-informative prior
##
## *Prior distribution on \mu*: Gamma(a=2,b=3)
## | mode | mean | sd |
                           Q1
                                  Q2
## | 0.3333 | 0.6667 | 0.4714 | 0.3204 | 0.5594 | 0.8975 |
## +----+----+-----+
##
## *Prior distribution on \phi*: Non-informative prior
##
## *Sample sizes*
##
   S (treated group): not supplied yet
##
   T (control group): not supplied yet
##
## *Observed counts*
##
   x (treated group): not supplied yet
##
   y (control group): not supplied yet
##
## *Posterior distribution on \phi*:
   a, b, c, d, S, T, x and y must be supplied
##
```

Since c and d were not supplied, **brr** automatically considers the non-informative prior on ϕ . Equivalently the same **brr** object can be defined by **Brr(a=2, b=3, c=NULL, d=NULL)** or **Brr(a=2, b=3, c=0.5, d=0)**.

The **brr** object is a function which can be used to update itself with new parameters, for example:

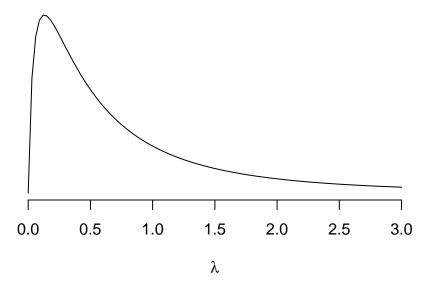
```
model <- model(c=3, d=3, S=100, T=100)
summary(model)
```

```
## Type of prior distribution: informative prior
##
## *Prior distribution on \mu*: Gamma(a=2,b=3)
##
## +----+
## | mode | mean |
               sd
                  - 1
                      Q1
                        - 1
                            Q2
                                  03
## +======+=====+======+=====+=====+=====+
## | 0.3333 | 0.6667 | 0.4714 | 0.3204 | 0.5594 | 0.8975 |
## *Prior distribution on \phi*: Beta2(c=3,d=3,scale=1.03)
##
## +----+
```

```
| mean | sd | Q1
                                | Q2
## +======+=====+=====+=====+=====+======+
## | 0.515
          | 1.545 | 1.995 | 0.578 | 1.03 | 1.836 |
## +----+
##
## *Sample sizes*
    S (treated group): 100
##
##
    T (control group): 100
##
## *Observed counts*
##
    x (treated group): not supplied yet
##
    y (control group): not supplied yet
##
## *Posterior distribution on \phi*:
##
    a, b, c, d, S, T, x and y must be supplied
```

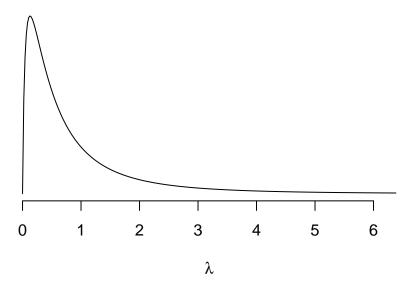
Now that a, b, c, d, S and T have been supplied, the user can play with all functions related to the prior distributions.

For example, **dprior** (model, "lambda", .) evaluates the density of the prior distribution of λ . This is convenient to generate a graphic with the **curve** function:



The **brr** package also provides a way to generate a plot with automatic aesthetics:

```
par(mar=c(4, 3, 1, 1))
plot(model, dprior(lambda))
```



If you are not pleased with the automatic bounds of the interval over which the curve is plotted, set your bounds in the **bounds** argument of the **plot** function. You can also set usual graphical parameters such as **lwd**, **lty**, ...

Posterior inference with brr

Posterior inference is available after, obviously, setting the observed counts x and y:

```
model <- model(x=14, y=20)
summary(model)</pre>
```

```
## Type of prior distribution: informative prior
##
## *Prior distribution on \u03c4*:
                    Gamma(a=2,b=3)
##
             - 1
                sd
                    Q1
                          Q2
                                  Q3
   mode
        mean
## | 0.3333 | 0.6667 | 0.4714 | 0.3204 | 0.5594 | 0.8975 |
##
## *Prior distribution on \( \phi^* : \) Beta2(c=3,d=3,scale=1.03)
   mode
         mean
                sd
                     Q1
                          Q2
## +======+=====+=====+=====+
       | 1.545 | 1.995 | 0.578 | 1.03 | 1.836 |
##
```

```
## *Sample sizes*
    S (treated group): 100
##
    T (control group): 100
##
## *Observed counts*
   x (treated group): 14
##
   y (control group): 20
##
## *Posterior distribution on \( \phi^* : \) Beta2(17,25,scale=1.03)
##
## +-----+
## | mode | mean | sd |
                          Q1
                             | Q2
                                        Q3
## +======+====+=====+====++====++=====+
## | 0.6338 | 0.7296 | 0.2363 | 0.5613 | 0.696 | 0.8605 |
##
## Pr('relative risk is greater than 1') = 0.876132340775555
```

Estimates are provided by the **coef** function:

coef (model)

```
## Estimates of ф
##
## mode : 0.6338462
## mean : 0.7295833
## median : 0.6959761
## intrinsic : 0.6966658
## intrinsic2 : 0.7033363
```

Posterior credibility intervals are provided by the **confint** function:

confint (model)

```
## 95%-credibility intervals about ¢
##
## +------+
## | interval | lwr | upr |
## +=======+
## | equi-tailed | 0.3679 | 1.284 |
## +------+
## | equi-tailed* | 0.3679 | 1.284 |
```

```
## | hpd | 0.3232 | 1.202 |

## +-----+

## | intrinsic | 0.3624 | 1.268 |

## +-----+

## | intrinsic2 | 0.3522 | 1.244 |

## +------+
```

Predictions are provided by the **predict** function after adding the sample sizes of the future experiment:

```
model <- model(Snew=500, Tnew=500)
predict(model)</pre>
```

```
## Predictions and 95%-credibility prediction intervals
##
## +----+
## | obs | size | median | lwr | upr
## +=====++=====++====++====++
## | xnew
        500
           - 1
              71
                    38
                         117
## +----+
## | ynew | 500
              102
                    62
                         156 |
           1
                  1
## +----+
```