The poweRlaw package: Examples

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The poweRlaw package provides an easy to use interface for fitting and visualising heavy tailed distributions, including power-laws. The fitting procedure follows the method detailed in Clauset *et al.*¹. This vignette gives examples of the fitting procedure.

¹ A. Clauset, C.R. Shalizi, and M.E.J. Newman. Power-law distributions in empirical data. *SIAM review*, 51(4):661–703, 2009

1 Discrete data: Moby Dick

The Moby Dick dataset contains the frequency of unique words in the the novel Moby Dick by Herman Melville. The data set can be downloaded from

http://tuvalu.santafe.edu/~aaronc/powerlaws/data.htm or loaded directly

```
data(moby)
```

To fit a discrete power-law to this data², we use the displ method

² The object moby is a simple R vector.

```
m_pl = displ*new(moby)
```

The resulting object, m_pl, is a displ³ object. It also inherits the discrete_distribution class. After creating the displ object, a typical first step would be to infer model parameters.⁴ We can estimate the lower threshold, via:

```
<sup>3</sup> displ: discrete power-law.
```

⁴ When the displ object is first created, the default parameter values are NULL and x_{min} is set to the minimum x-value.

For a given value x_{min} , the scaling parameter is estimated using its MLE:

$$\hat{\alpha} \simeq 1 + n \left[\sum_{i=1}^{n} \log \left(\frac{x_i}{x_{\min} - 0.5} \right) \right]^{-1}$$

This yields a threshold estimate of $x_{\min} = 7$ and scaling parameter $\alpha = 1.95$, which matches results found in Clauset et al. [2009]. Alternatively, we could perform a parameter scan for each value of x_{\min} :

```
estimate_xmin(m_pl, pars = seq(1.5, 2.5, 0.1))
```

The parameter scan method is more exact, but is slightly slower.

To fit a discrete log-normal distribution, we follows a similar procedure, except we begin by creating a dislnorm.⁵

⁵ dislnorm: discrete log-normal object

```
m_ln = dislnorm$new(moby)
est = estimate_xmin(m_ln)
```

which yields a lower threshold of $x_{min} = 3$ and parameters (-17.9, 4.87). A similar procedure is applied to fit the Poisson distribution; we create a distribution object using dispois, then fit as before.

The data CDF and lines of best fit can be easily plotted:

```
plot(m_pl)
lines(m_pl, col = 2)
lines(m_ln, col = 3)
lines(m_pois, col = 4)
```

to obtain figure 1. It clear that the Poisson distribution is not appropriate for this data set. However, the log-normal and power-law distribution both provide reasonable fits to the data.

Parameter uncertainty 1.1

To get a handle on the uncertainty in the parameter estimates, we use a bootstrapping procedure, via the bootstrap function. This procedure can be applied to any distribution object.⁶ Furthermore, the bootstrap procedure can utilize multiple CPU cores to speed up inference.7

```
##5000 bootstraps using two cores
bs = bootstrap(m_pl, no_of_sims=5000, threads=2)
```

By default, the bootstrap function will use the maximum likelihood estimate to estimate the parameter and check all values of x_{min} . When possible x_{\min} values are large, then it is recommend that the search space is reduced. For example, this function call

```
bootstrap(m_pl, xmins = seq(2, 20, 2))
```

will only calculate the Kolmogorov-Smirnoff statistics at values of x_{\min} equal to

$$2,4,6,\ldots,20$$
.

A similar argument exists for the parameters.⁸

The bootstrap function, returns bs_xmin object that has three components:

- 1. The goodness of fit statistic obtained from the Kolmogorov-Smirnoff test. This value should correspond to the value obtained from the estimate_xmin function.
- 2. A data frame containing the results for the bootstrap procedure.
- 3. The average simulation time, in seconds, for a single bootstrap.

The boostrap results can be explored in a variety way. First we can estimate the standard deviation of the parameter uncertainty, i.e.

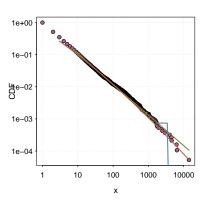


Figure 1: Data CDF of the Moby Dick data set. The fitted power-law (green line), log-normal (red line) and poisson (blue) distributions are also given.

⁶ For example, bootstrap(m_ln).

⁷ The output of this bootstrapping procedure can be obtained via data(bootstrap_moby).

⁸ For single parameter models, pars should be a vector. For the log-normal distribution, pars should be a matrix of values.

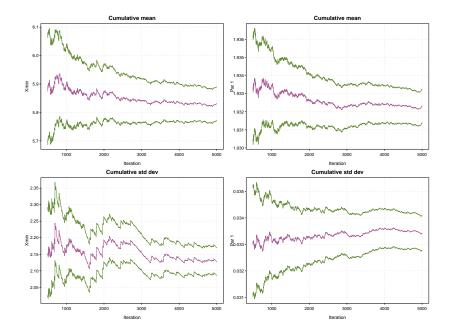


Figure 2: Results from the standard bootstrap procedure (for the power-law model) using the Moby Dick data set: bootstrap(m_pl). The top row shows the mean estimate of parameters x_{min} and α . The bottom row shows the estimate of standard deviation for each parameter. The dashed-lines give approximate 95% confidence intervals.

After 5000 iterations, the standard deviation of x_{min} and α is estimated to be 2.1 and 0.03 respectively.

```
sd(bs$bootstraps[, 2])
## [1] 2.127
sd(bs$bootstraps[, 3])
## [1] 0.03342
```

Alternatively, we can visualise the results using the plot function:

```
plot(bs, trim = 0.1)
```

to obtain figure 2. This top row of graphics in figure 2 give a 95% confidence interval for mean estimate of the parameters. The bottom row of graphics give a 95% confidence for the standard deviation of the parameters. The parameter trim in the plot function controls the percentage of samples displayed. When trim=0.1, we only display the final 90% of data.

We can also construct histograms.

```
hist(bs$bootstraps[, 2])
hist(bs$bootstraps[, 3])
```

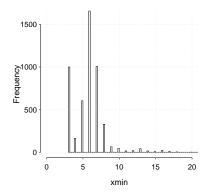
to get figure 3.

A similar bootstrap analysis can be obtained for the log-normal distribution

```
bs1 = bootstrap(m_ln)
```

in this case we would obtain uncertainty estimates for both of the log-normal parameters.

9 When trim=0, all iterations are displayed.



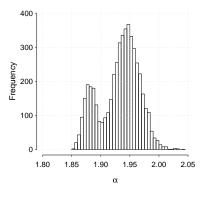


Figure 3: Characterising uncertainty in parameter values. (a) x_{min} uncertainty (standard deviation 2) (b) α uncertainty (std dev. 0.03)

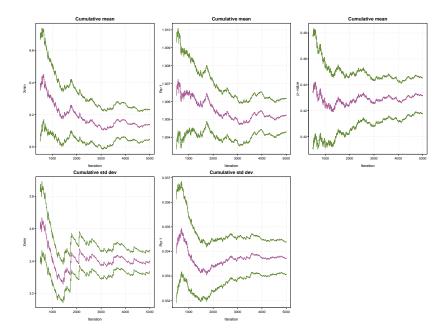


Figure 4: Results from the bootstrap procedure (for the power-law model) using the Moby Dick data set: bootstrap $_p(m_pl)$. The top row shows the mean estimate of parameters x_{\min} , α and the *p*-value. The bottom row shows the estimate of standard deviation for each parameter. The dashed-lines give approximate 95% confidence intervals.

Testing the power-law hypothesis

Since it is possible to fit a power-law distribution to any data set, it is appropriate to test whether it the observed data set actually follows a power-law. Clauset et al, suggest that this hypothesis is tested using a goodness-of-fit test, via a bootstrapping procedure. This test generates a p-value that can be used to quantify the plausibility of the hypothesis. If the *p*-value is large, than any difference between the empirical data and the model can be explained with statistical fluctuations. If $p \simeq 0$, then the model does not provide a plausible fit to the data and another distribution may be more appropriate. In this scenario,

 H_0 : data is generated from a power-law distribution.

 H_1 : data is not generated from a power-law distribution.

To test these hypothesis, we use the bootstrap_p function

The point estimate of the *p*-value is one of the elements of the bs_p object10

bs_p\$p ## [1] 0.4318

Alternatively we can plot the results

to obtain figure 4. The graph in the top right hand corner gives the cumulative estimate of the *p*-value; the final value of the purple line corresponds to bs_p\$p. Also given are approximate 95% confidence intervals.

10 Also given is the average time in seconds of a single bootstrap: $bs_p sim_time = 2.86$.

1.3 Investigating the effect in x_{\min}

The estimate of the scaling parameter, α , is typically highly correlated with the threshold limit, x_{min} . This relationship can be easily investigated with the poweRlaw package. First, we create a vector of thresholds to scan

```
xmins = 1:1500
```

then a vector to store the results

```
est_scan = 0 * xmins
```

Next, we loop over the x_{min} values and estimate the parameter value conditional on the x_{min} value:

```
for (i in seq_along(xmins)) {
    m_pl$setXmin(xmins[i])
    est = estimate_pars(m_pl, pars = seq(1.2, 2.8, 0.01))
    est_scan[i] = est*pars
}
```

The results are plotted figure 5. For this data set, as the lower threshold increases, so does the point estimate of α .

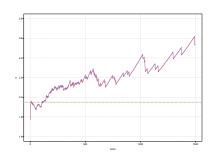


Figure 5: Estimated parameter values conditional on the threshold, x_{min} . The horizontal line corresponds to $\alpha = 1.95$.

Continuous data: electrical blackouts

In this example, we will investigate the numbers of customers affected in electrical blackouts in the United States between 1984 and 2002.¹¹ The data set can be downloaded from Clauset's website:12

```
blackouts = read.table("blackouts.txt")
```

Although the blackouts data set is discrete, since the values are large, it makes sense to treat the data as continuous. Continuous power-law objects take vectors as inputs, so

```
m_bl = conpl$new(blackouts$V1)
```

then we estimate the lower-bound via

```
est = estimate_xmin(m_bl)
```

This gives a point estimate of $x_{\min} = 230000$. We can then update distribution object:

```
m_bl$setXmin(est)
```

Using generic plot method

```
plot(m_bl)
lines(m_bl, col = 2, lwd = 2)
```

we get figure 6. To fit a discrete log-normal distribution we follow a similar procedure:

```
m_bl_ln = conlnorm$new(blackouts$V1)
est = estimate_xmin(m_bl_ln)
m_bl_ln$setXmin(est)
```

and add the line of best fit to the plot via

```
lines (m_bl_ln, col = 3, lwd = 2)
```

It is clear from figure 6 that the log-normal distribution provides a better fit to this data set.

11 M.E.J. Newman. Power laws, pareto distributions and zipf's law. Contemporary physics, 46(5):323-351, 2005

12 http://goo.gl/BsqnP

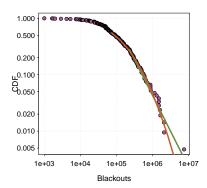


Figure 6: CDF plot of the blackout dataset with line of best fit. Since the minimum value of x is large, we fit a continuous power-law as this is more it efficient. The power-law fit is the green line, the discrete log-normal is the red

Multiple data sets: the American-Indian war

In a recent paper, Bohorquez et al. investigated insurgent attacks in Afghanistan, Iraq, Colombia, and Peru.¹³ Each time, the data resembled power laws. Friedman used the power-law nature of casualties to infer under-reporting in the American-Indian war. Briefly, by fitting a power-law distribution to the observed process, the latent, unobserved casualties can be inferred.¹⁴

The number of casualties observed in the American-Indian War can be obtained via

```
data(NativeAmerican)
data(USAmerican)
```

Each data set is a data frame with two columns. The first column is number of casualties recorded, the second the conflict date

```
head(NativeAmerican, 3)
     Cas
               Date
## 1
      18 1776-07-15
## 2
      26 1776-07-20
## 3
      13 1776-07-20
```

The records span around one hundred years, 1776 – 1890. The data is plotted in figure 7.

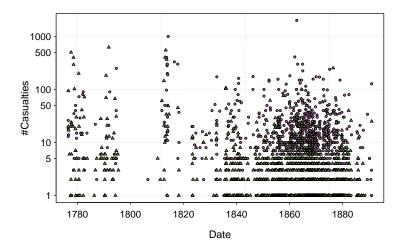
It is straightforward to fit a discrete power-law to this data set. First, we create discrete power-law objects:

```
m_na = displ$new(NativeAmerican$Cas)
m_us = displ$new(USAmerican$Cas)
```

then we estimate x_{min} for each data set:

```
est_na = estimate_xmin(m_na, pars = seq(1.5, 2.5, 0.001))
est_us = estimate_xmin(m_us, pars = seq(1.5, 2.5, 0.001))
```

and update the power-law objects



¹³ J. C. Bohorquez, S. Gourley, A. R. Dixon, M. Spagat, and N. F. Johnson. Common ecology quantifies human insurgency. Nature, 462(7275):911-914, 2009

Figure 7: Casualty record for the Indian-American war, 1776 - 1890. Native Americans casualties (purple circles) and US Americans casualties (green triangles). Data taken from Friedman.

¹⁴ J. A. Friedman. Using power laws to estimate conflict size. 2013

```
m_na$setXmin(est_na)
m_us$setXmin(est_us)
```

The resulting fitted distributions can be plotted on the same figure

```
plot(m_na)
lines(m_na)
## Don't create a new plot Just store the output
d = plot(m_us, draw = FALSE)
points(d$x, d$y, col = 2)
lines(m_us, col = 2)
```

The result is given in figure 8. The tails of the distributions appear to follow a power-law. This is consistent with the expectation that smaller-scale engagements are less likely to be recorded. However, for larger scale engagements, it is very likely that a record is made.

References

- J. C. Bohorquez, S. Gourley, A. R. Dixon, M. Spagat, and N. F. Johnson. Common ecology quantifies human insurgency. Nature, 462(7275):911-914, 2009.
- A. Clauset, C.R. Shalizi, and M.E.J. Newman. Power-law distributions in empirical data. SIAM review, 51(4):661-703, 2009.
- J. A. Friedman. Using power laws to estimate conflict size. 2013.
- M.E.J. Newman. Power laws, pareto distributions and zipf's law. Contemporary physics, 46(5):323-351, 2005.

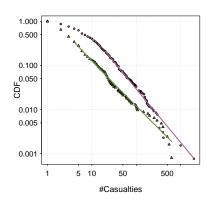


Figure 8: Plots of the CDFs for the Native American and US American casualties. The lines of best fit are also given.