dRepanalyzer report
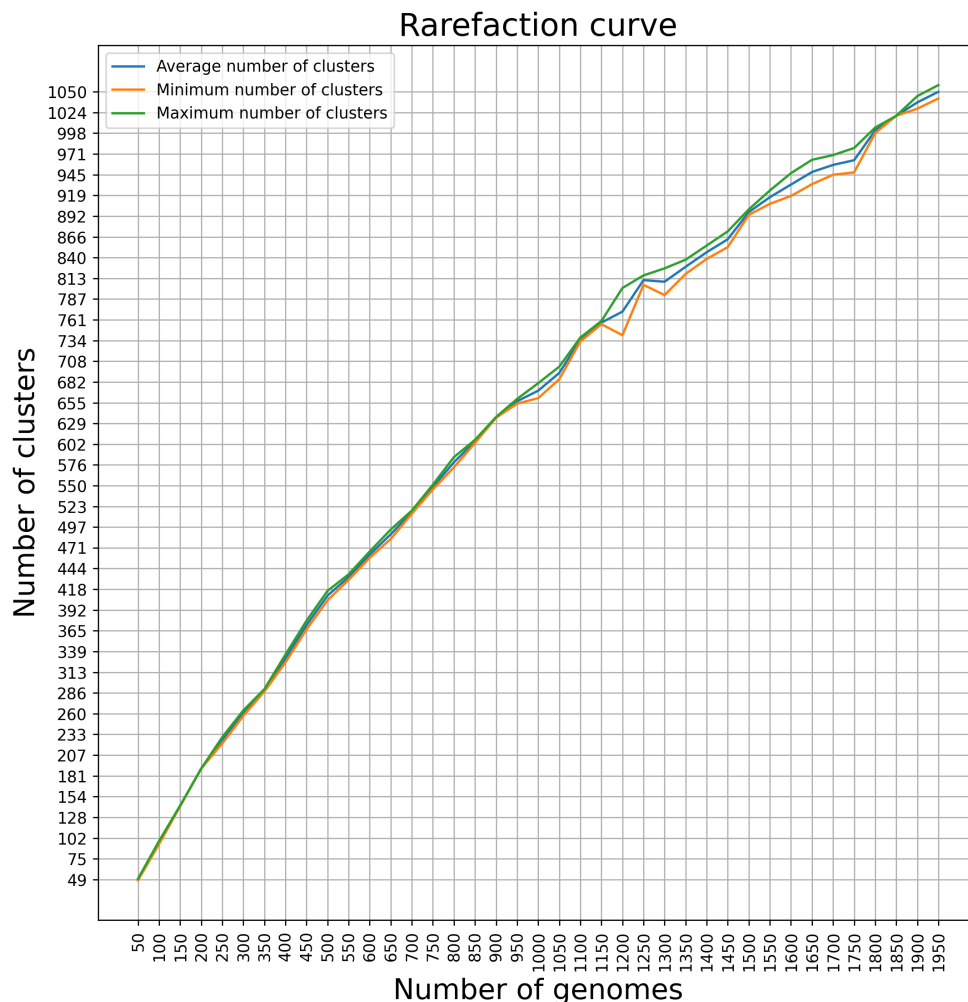
Date   07/06/2023 23:13:33

Input   Cdb_big.csv

Output   Cdb_big

Rarefaction plot

The rarefaction plot is produced by a resampling procedure starting from a minimum sample size N to the maximum sample size M. At each iteration the sample size increased by k and per each sample size the sampling was repeated RR times to compute the mean, the minimum and the maximum number of species detectable by a particular sample size. If filter is true, the cluster detected just one time gets taken out.

- Minimum sample size (N): 50

- Maximum sample size (M): 2000

- Growth iteration rate (k): 50

- Resample rate (RR): 2

- Filter under represented cluster: True