

POP\_TYPE.x MARINE FRESHWATER 1.00 -0.75 **-**Number of loci 10 100 1000 0.25 -0.00 -

> 100 1 log<sub>10</sub>(Total number of missing genotypes)

10

100

Mean % of missing genotypes

10

ECOTYPE.x BIG\_LAKE SEA 1.00 -0.75 -0.50 -0.25 -Mean % of missing genotypes Number of loci 10 SMALL\_LAKE RIVER 100 1000 0.75 -0.50 -0.25 0.00 -10 10 100 100  $log_{10}(Total\ number\ of\ missing\ genotypes)$