

Species tree error (RF distance)

- 1 alleles x 800 genes
- 2 alleles x 400 genes
- 5 alleles x 160 genes

0.4
0.3
0.2
0.1
0.0

(1.16e+05,3.55e+05]

(3.55e+05,4.49e+05]

(4.49e+05,5.73e+05]

(5.73e+05,1.18e+06]

Number of generations

