

Gene copy variation relative to outgroup

retained
(1:1)

39.2 %

40.8 %

6.3 %

4.2 %

lost
(1:0)

45.2 %

44.9 %

5.7 %

4.7 %

gained
(1:n)

15.5 %

14.2 %

88.0 %

91.1 %

Species

- Clusia_multiflora_H1
- Clusia_multiflora_H2
- Clusia_minor
- Clusia_rosea

0 2500 5000 7500 10000

Number of gene families