

phylometric = colless_like_index

phylometric = mean_pairwise_distance

phylometric = num_ancestors

phylometric = mean_evolutionary_distinctiveness

plain

weak selection

strong selection

spatial structure

weak 4 niche ecology

4 niche ecology

8 niche ecology

0

200000

400000

value

0

10

20

30

40

value

0

10000

20000

value

0

5

10

15

20

value