

phylogenetic

colless_like_index -

mean_evolutionary_distinctiveness -

mean_pairwise_distance -

sum_distance -

plain

weak selection

strong selection

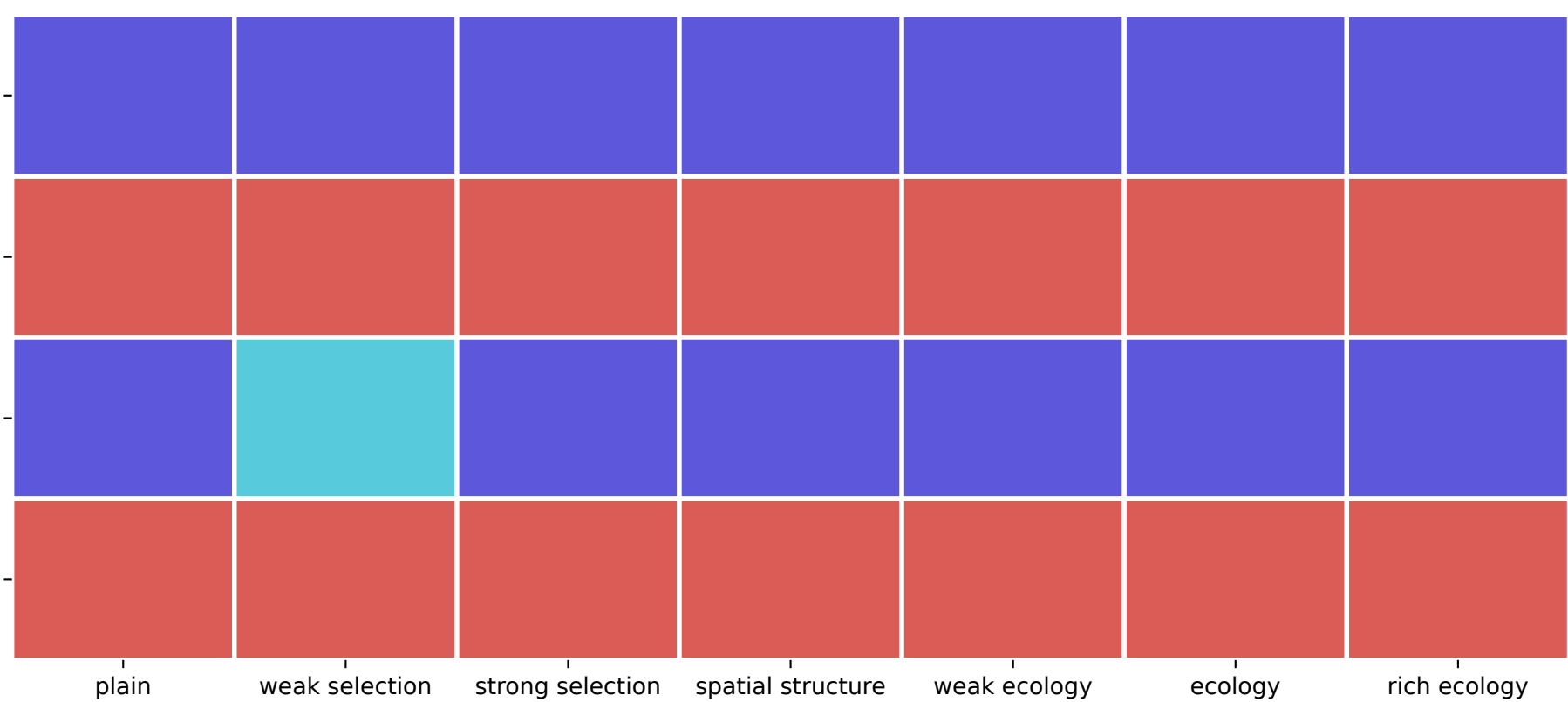
spatial structure
regime

weak ecology

ecology

rich ecology

reconstruction error insignificance threshold



33
10
3
1