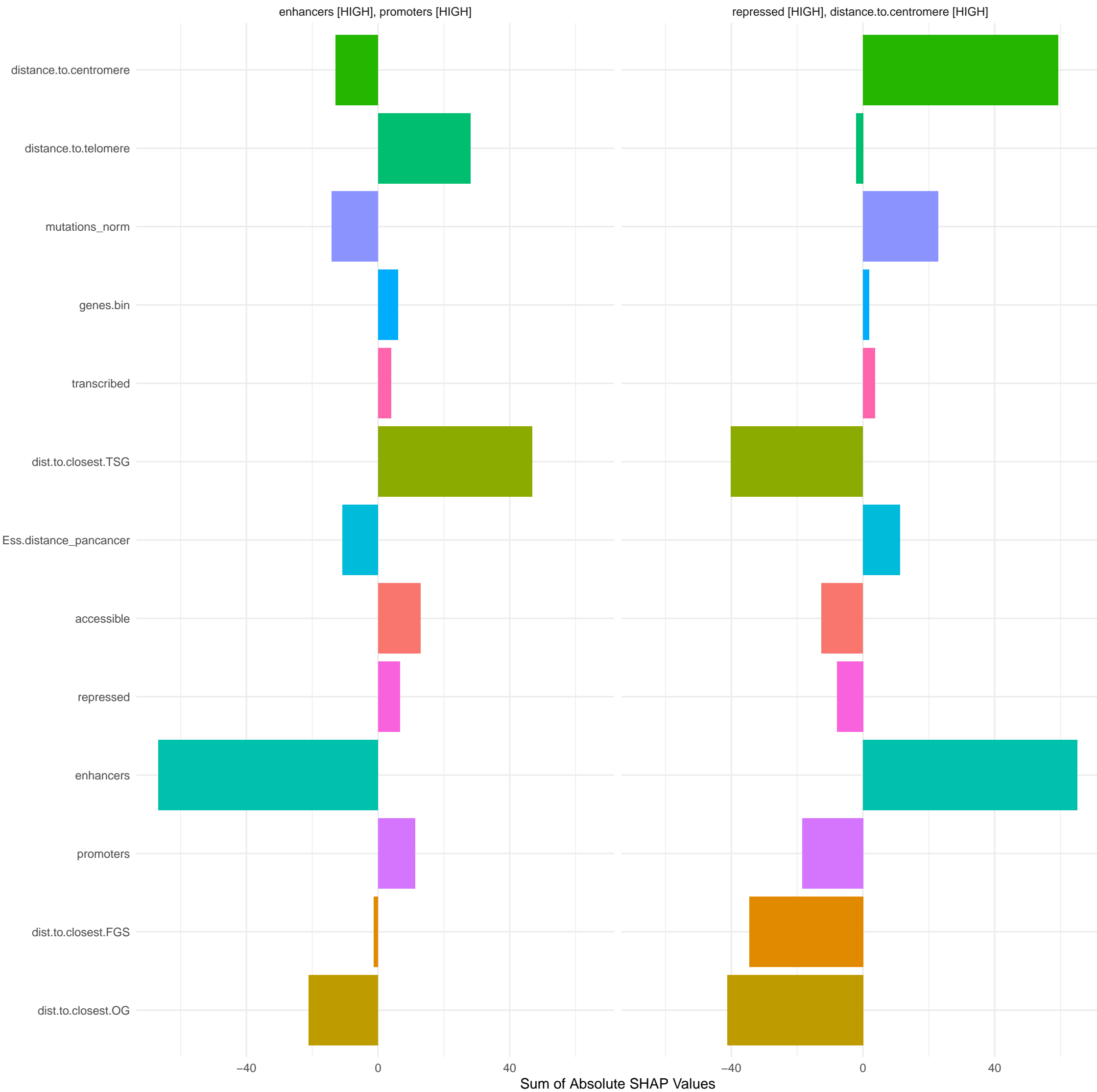


Clustering depth 1 – Ampl

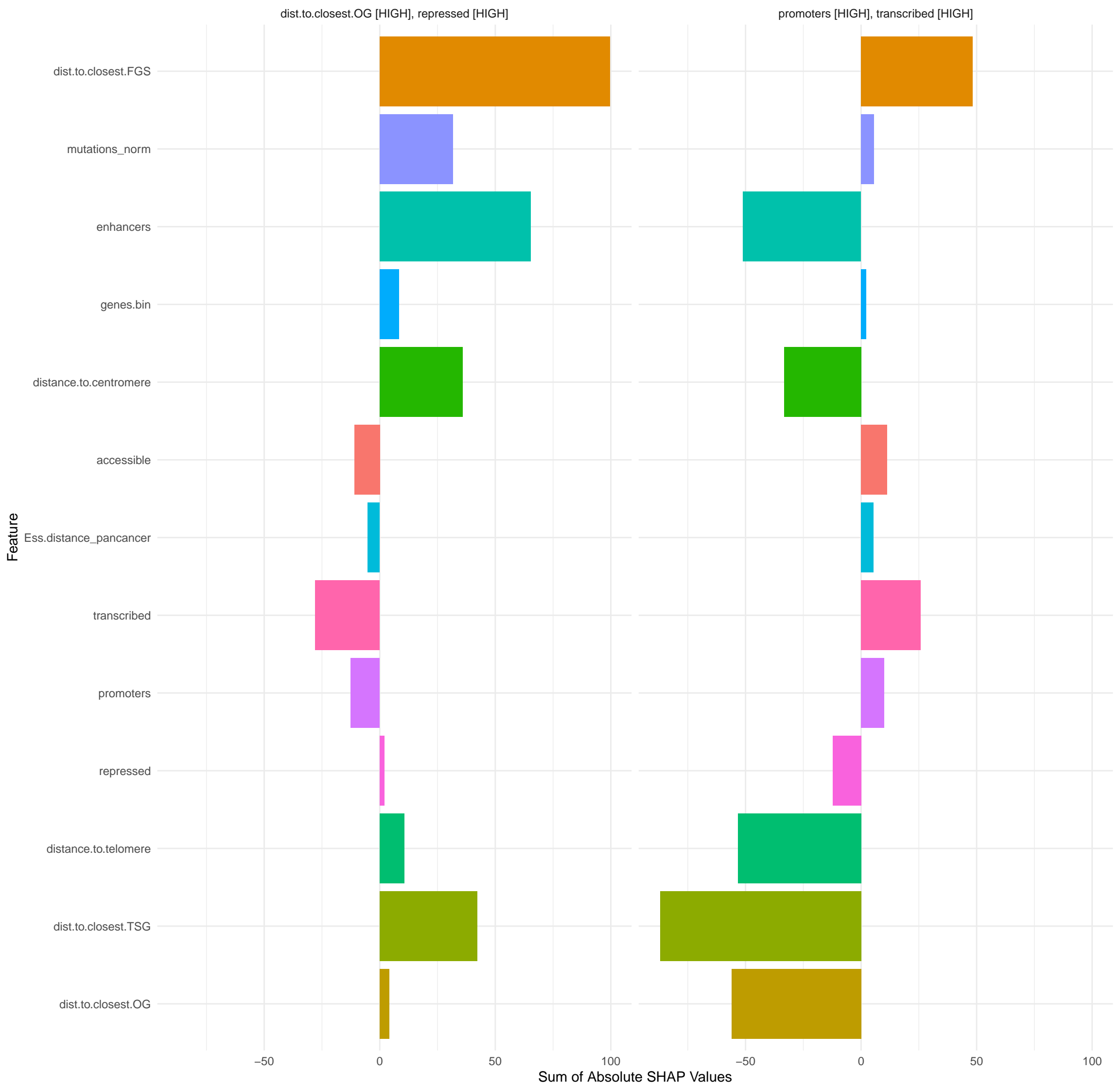
enhancers [HIGH], promoters [HIGH]

repressed [HIGH], distance.to.centromere [HIGH]

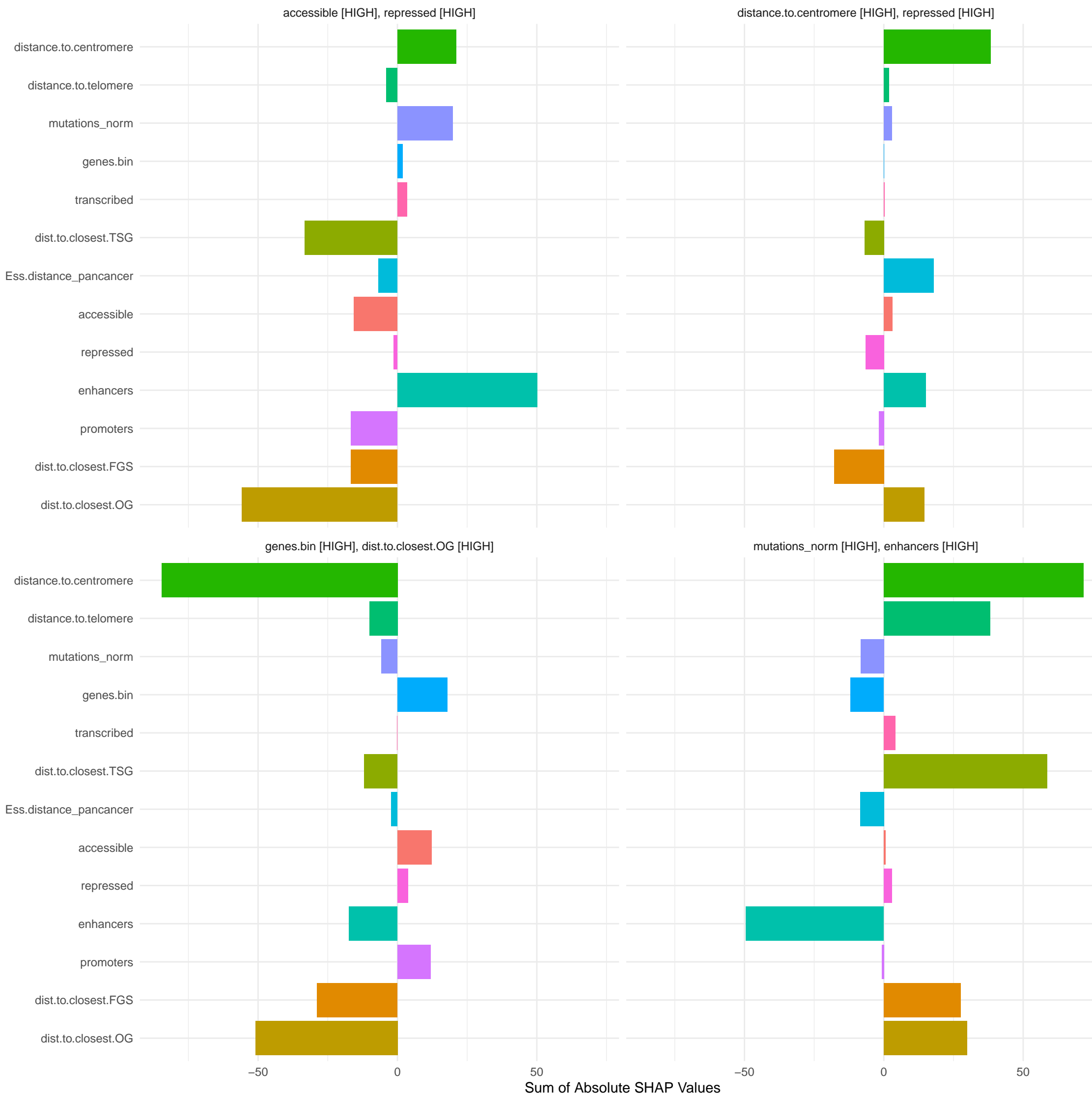
Feature



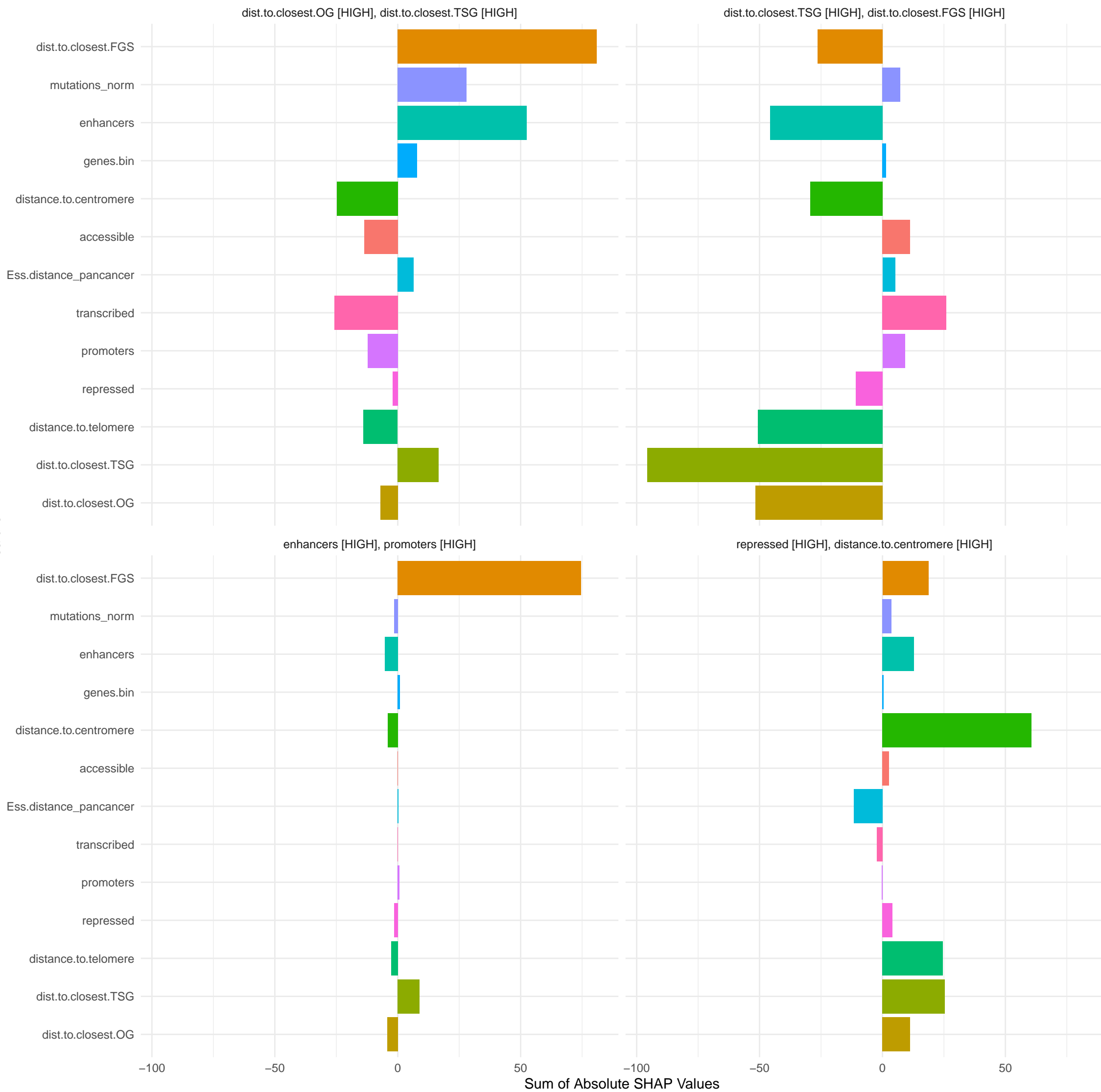
Clustering depth 1 – Del



Clustering depth 2 – Ampl



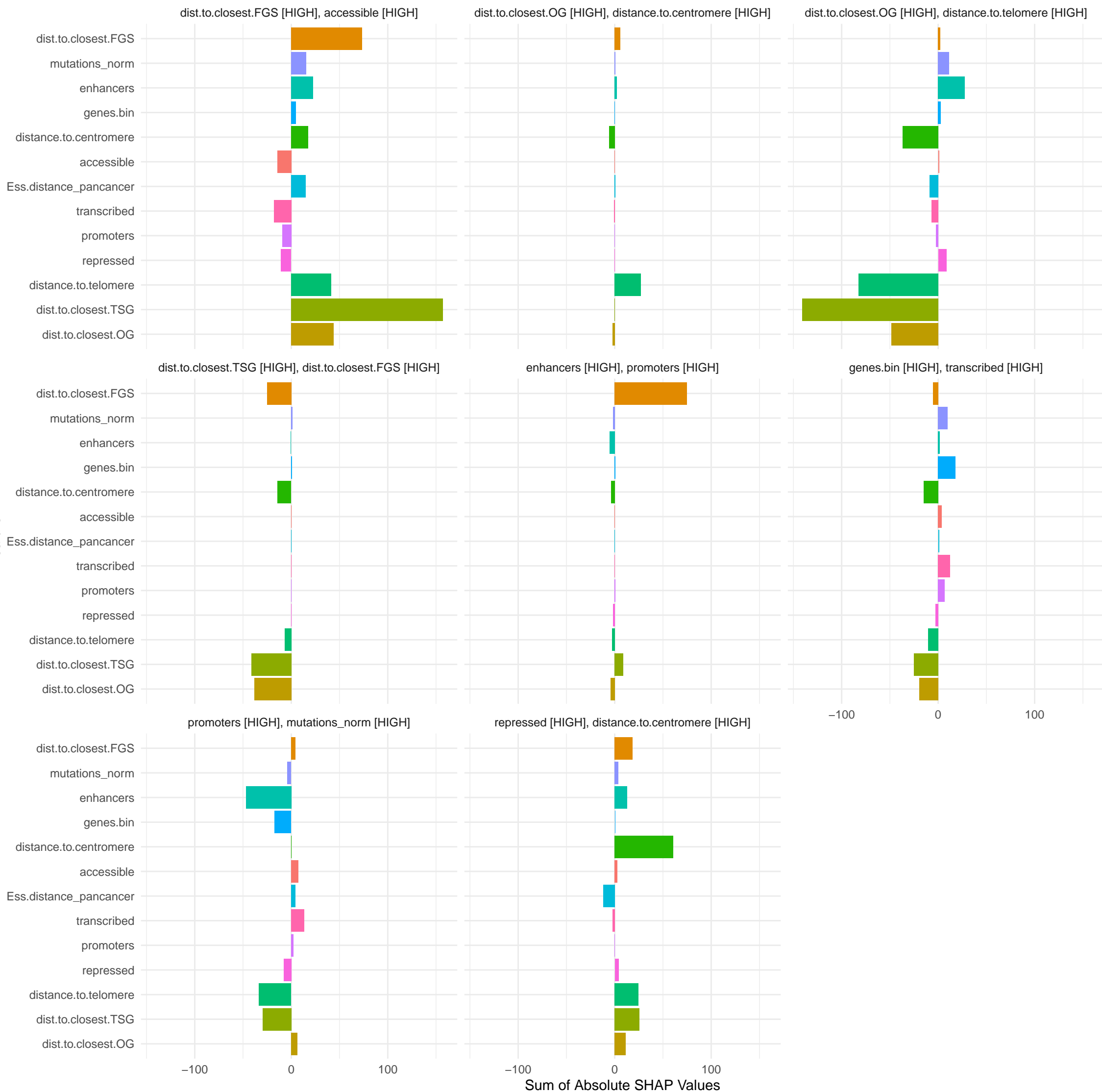
Clustering depth 2 – Del



Clustering depth 3 – Ampl

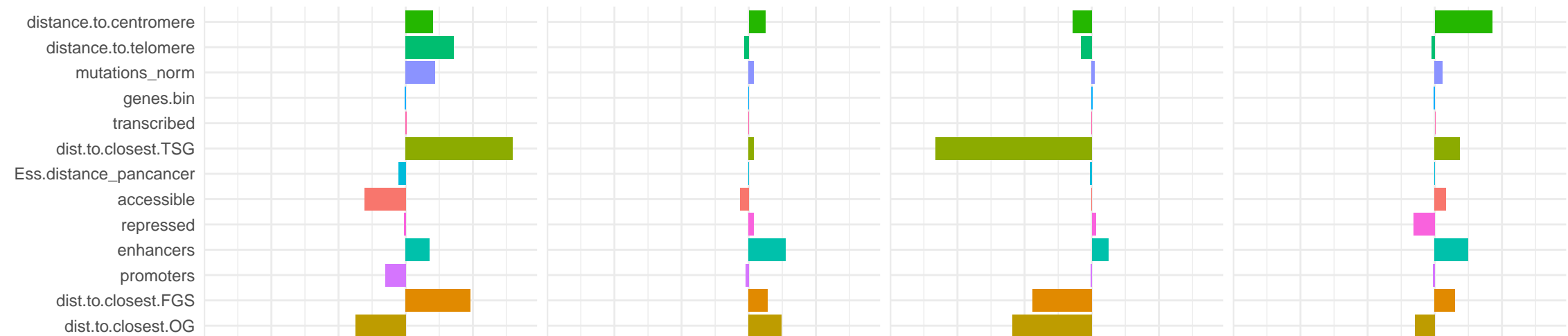


Clustering depth 3 – Del

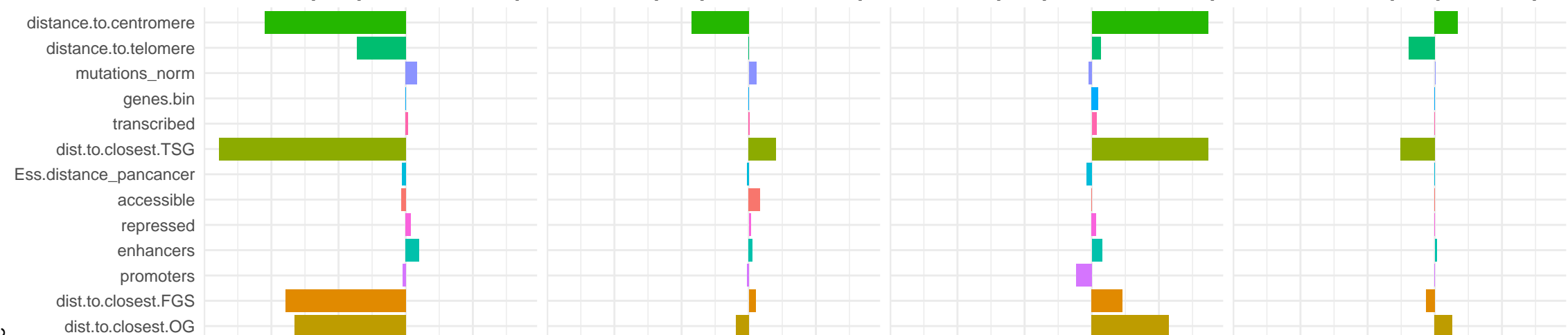


Clustering depth 4 – Ampl

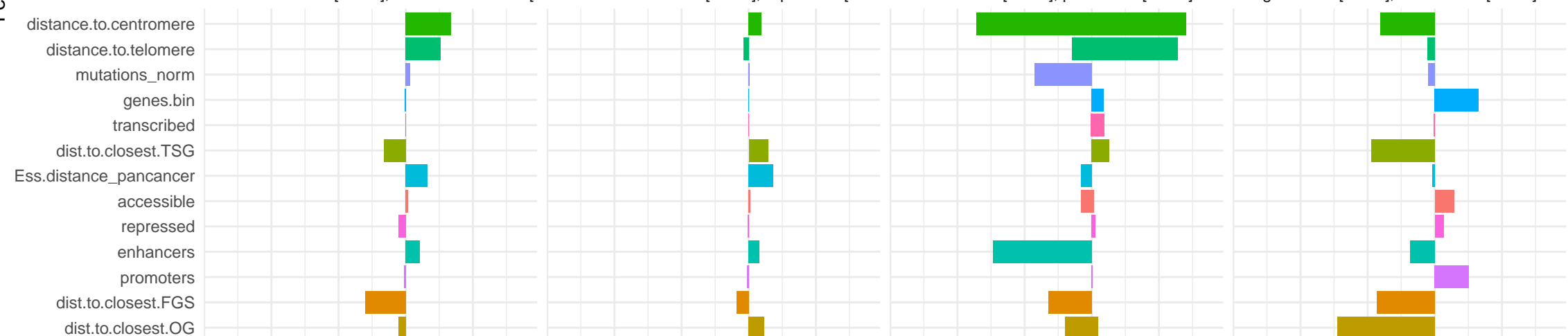
accessible [HIGH], dist.to.closest.FGS [HIGH] accessible [HIGH], distance.to.telomere [HIGH] accessible [HIGH], Ess.distance_pancancer [HIGH] closest.FGS [HIGH], distance.to.centromere [HIGH]



distance.to.closest.OG [HIGH], dist.to.closest.FGS [HIGH] distance.to.closest.OG [HIGH], dist.to.closest.TSG [HIGH] distance.to.closest.TSG [HIGH], distance.to.telomere [HIGH] distance.to.centromere [HIGH], accessible [HIGH]



distance.to.centromere [HIGH], dist.to.closest.OG [HIGH] distance.to.centromere [HIGH], repressed [HIGH] enhancers [HIGH], promoters [HIGH] genes.bin [HIGH], transcribed [HIGH]



mutations_norm [HIGH], accessible [HIGH] mutations_norm [HIGH], transcribed [HIGH] repressed [HIGH], distance.to.centromere [HIGH]



Sum of Absolute SHAP Values

Clustering depth 4 – Del

