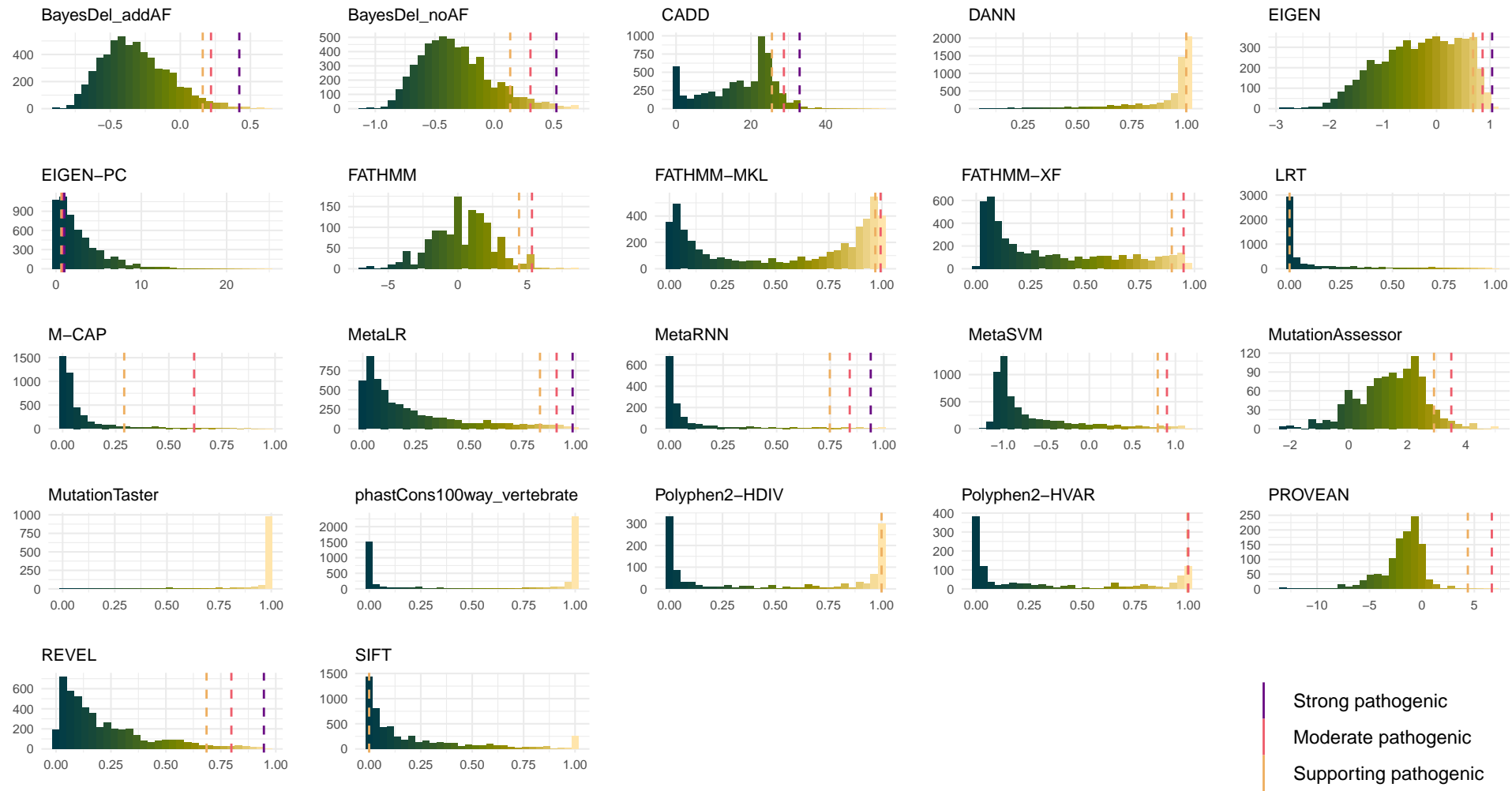


No. qualifying variants



in silico prediction score