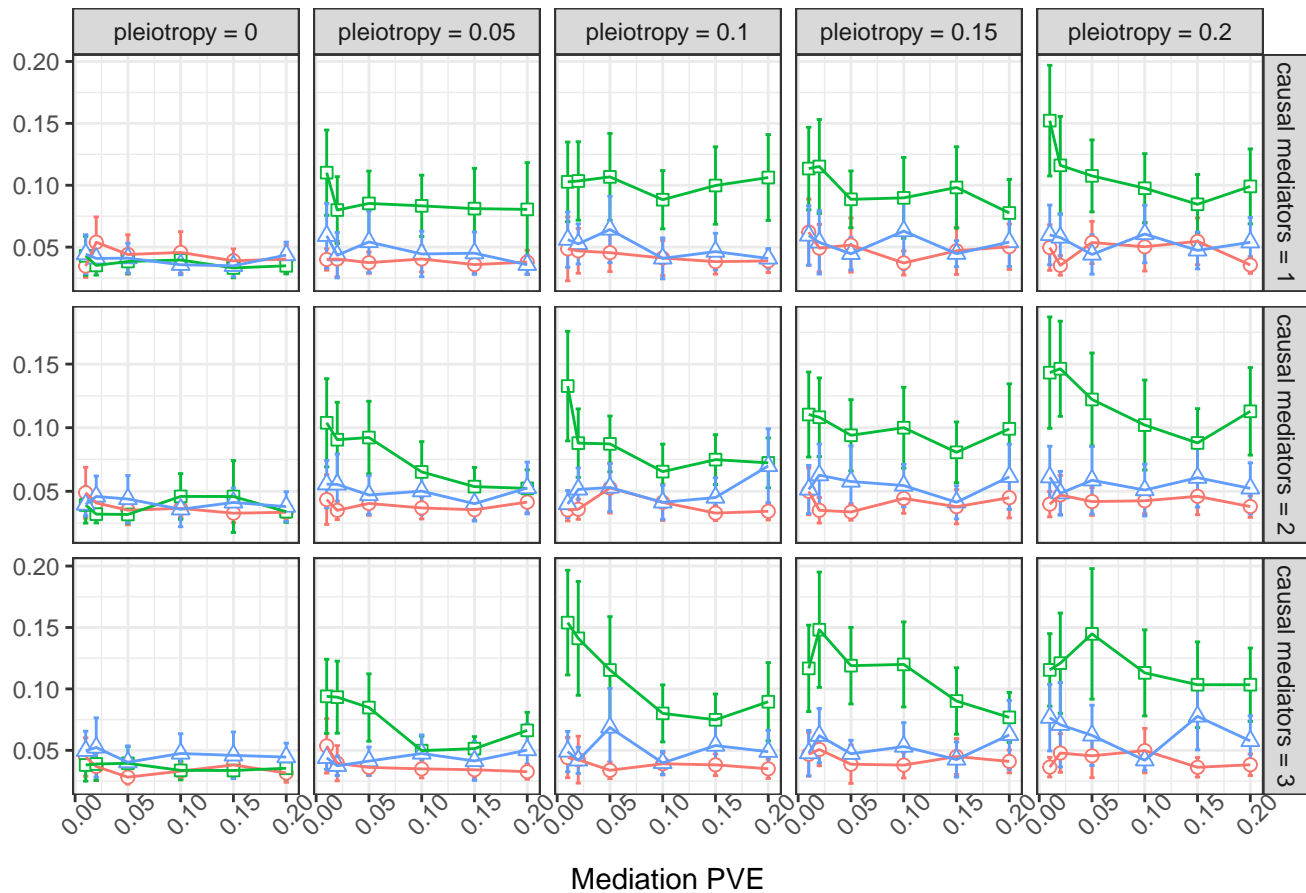


QTLs per gene = 3

Mis-classification AUPRC



zqtl
twas
egger