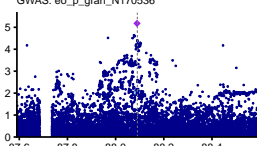
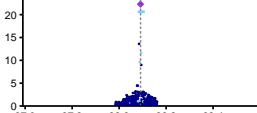


GWAS: eo_p_gran_N170536



Macrophage_H3K27ac: G3694

502;0.82797502813444



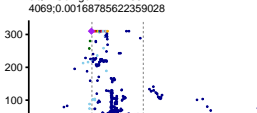
Macrophage_H3K27ac: G3694

502;0.828674305945468



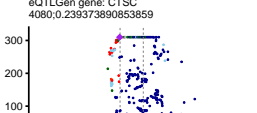
eQTLGen gene: RAB38

4069;0.00168785622359028



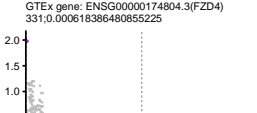
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4080;0.239373890853859



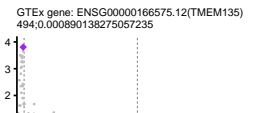
GTEx gene: ENSG00000174804.3(FZD4)

331;0.000618386480855225



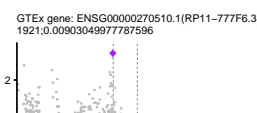
GTEx gene: ENSG00000166575.12(TMEM135)

494;0.000890138275057235



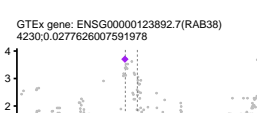
GTEx gene: ENSG00000270510.1(RP11-777F6.3)

1921;0.00903049977787596



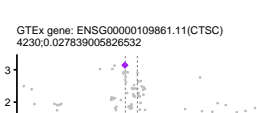
GTEx gene: ENSG00000123892.7(RAB38)

4230;0.0277626007591978



GTEx gene: ENSG00000109861.11(CTSC)

4230;0.027839005826532



GTEx gene: ENSG00000249489.1(GAPDHP70)

4230;0.0219463947510952

