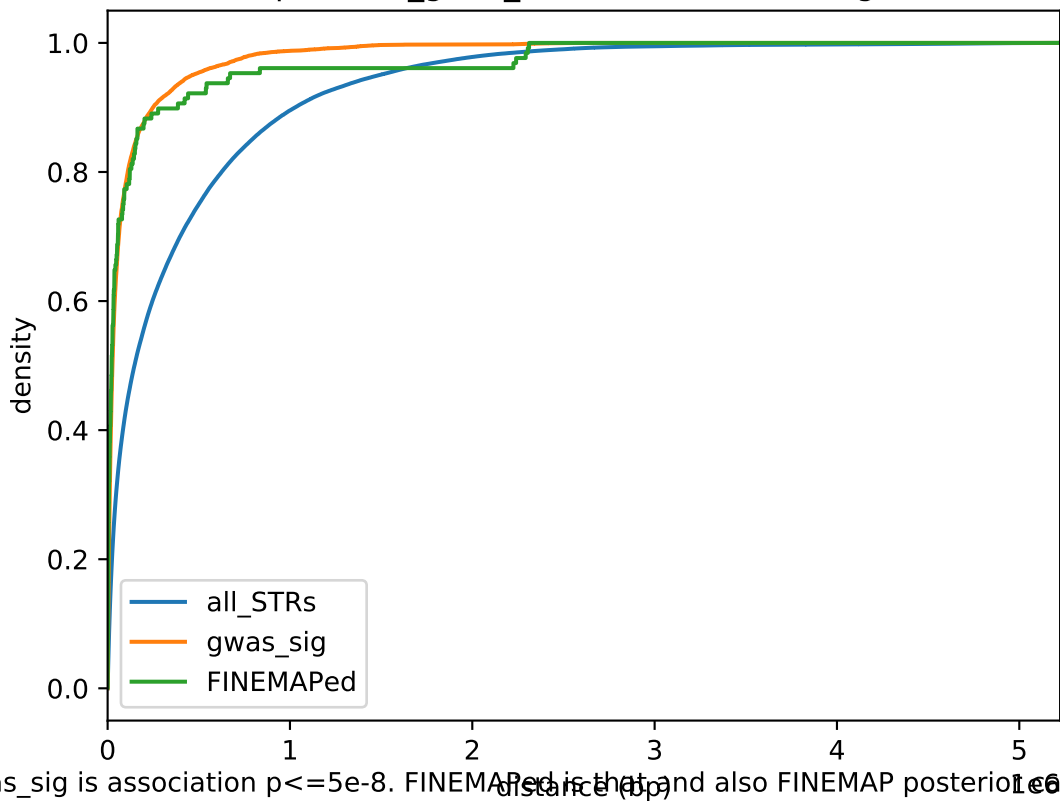


CDF of upstream_gene_dist from within intergenic STRs



gwas_sig is association $p \leq 5e-8$. FINEMAPed is that and also FINEMAP posterior ≥ 0.1