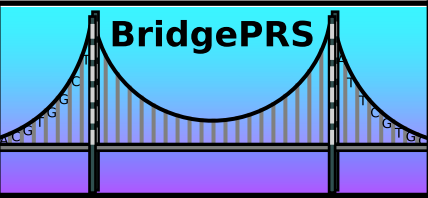


bridgePRS: Weighted-Combined Analysis

	Target	Model
Populations: (Reference, GWAS)	AFR, AFR	EUR, EUR
GWAS Sample Size:	20000	80000
GWAS Total SNPs: (Correlation: 0.09)	19685	19662
Phenotype: (type)	y (continuous)	NA
Phenotype Dist: (Mean,Std,Min,Max)	-1.74, 1.01, -4.7, 1.4	NA
PRS Dist: (Mean,Std,Min,Max)	-1.75, 0.15, -2.4, -1.2	NA
Result: SNP Weights (4812)	GWAS Corr= 0.43	GWAS Corr=0.44

