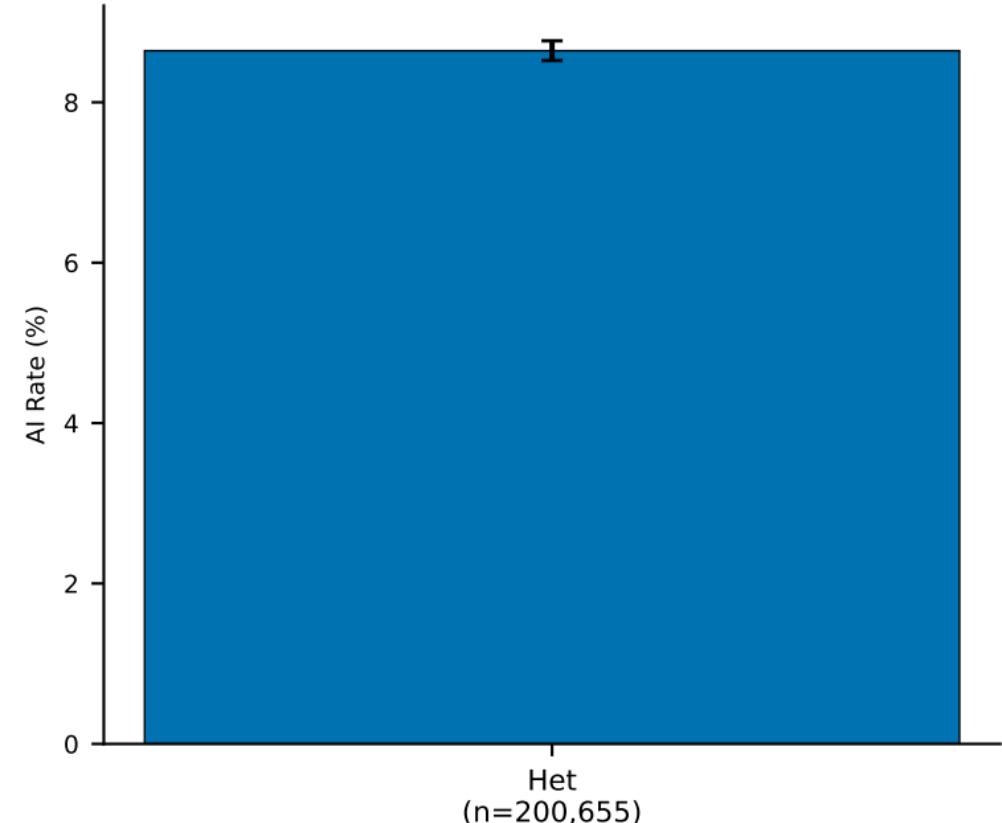


Corrected Het/Homo Validation
(Using SNV own genotype, not lead genotype)

ATAC-seq (caQTL)



RNA-seq (eQTL)

