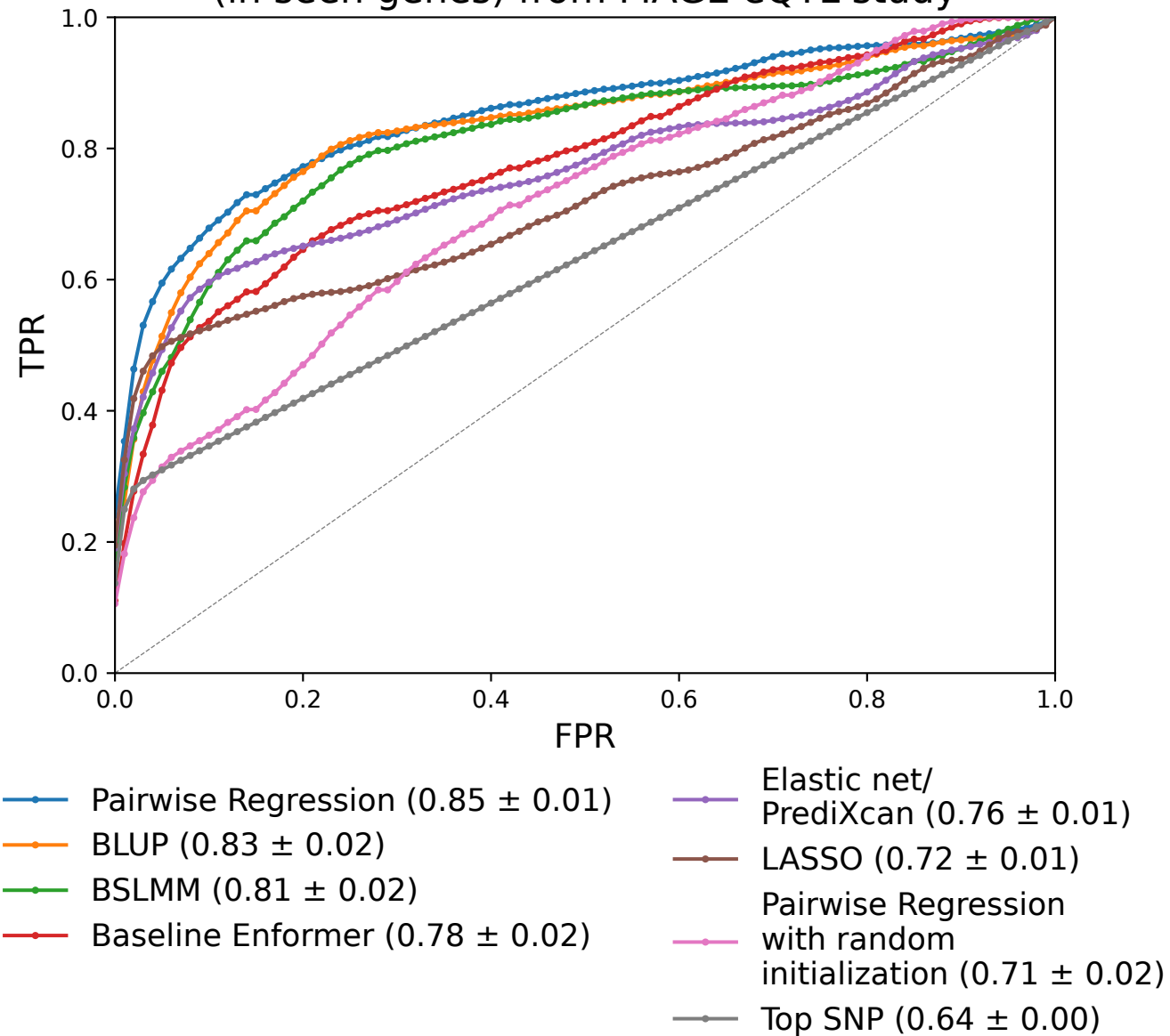


Fine-mapped vs. non-fine-mapped variants  
(in seen genes) from MAGE eQTL study



Enrichment of driver variants in  
LCL DNase I footprints

