DITTO Benchmarking for snRNA Receiver Operating Characteristic (ROC) curves Precision Recall (PRC) curves CADD = 0.76CADD = 0.231.0 CSCAPE = 0.73CSCAPE = 0.12ClinPred = 0.39ClinPred = 0.79DANN = 0.42DANN = 0.78 $DANN_Coding = 0.48$  $DANN_Coding = 0.80$  $DGI_score = 0.56$ DGI score = 0.830.8  $FATHMM_coding = 0.79$  $FATHMM_coding = 0.47$ Funseq2 = 0.19Funseq2 = 0.75Linsight = 0.25Linsight = 0.80LRT = 0.53LRT = 0.81Loftool = 0.52Loftool = 0.81Rate 9.0 0.6 MetaSVM = 0.38MetaSVM = 0.80--- MetaLR = 0.79 MetaLR = 0.39MutPred = 0.80MutPred = 0.50 $MutPred_indel = 0.50$  $MutPred_indel = 0.80$ — Mutation\_Assessor = 0.80  $Mutation_Assessor = 0.50$ 0.4  $Mutation_{Taster} = 0.50$  $Mutation_{Taster} = 0.80$ Provean = 0.80Provean = 0.50PHDSNPG = 0.55PHDSNPG = 0.82—— Revel = 0.50--- Revel = 0.80 --- SIFT = 0.50 --- SIFT = 0.80 0.2 VEST = 0.50---- VEST = 0.80 --- dbSCSNV\_ADA = 0.48 --- dbSCSNV\_ADA = 0.80 --- dbSCSNV\_RF = 0.48 --- dbSCSNV\_RF = 0.80 -- varity\_r = 0.45 - varity\_r = 0.79 --- varity\_er = 0.79 --- varity\_er = 0.45 0.0 DITTO = 1.00- DITTO = 1.00 0.0 0.2 0.0 0.2 0.6 8.0 0.4 0.6 8.0 1.0 1.0 False Positive Rate Recall