C) CATH/Gene3D

Search tools: AUC1000 scores

CSBLAST:	0.9100
PHMMER:	0.9032
HHSEARCH:	0.9009
NCBI−BLAST:	0.8784
USEARCH:	0.8613
FASTA:	0.8523
UBLAST:	0.8158

True positives

False positives