

C

TNBC

normal

tumor

average precision

average recall

software

- MonoVar default
- ...●... MonoVar no consensus
- ProSolo default
- ...●... ProSolo imputation^{-B}
- ...●... SCAN-SNV sensitive
- SCcaller default dbsnp
- ...●... SCcaller sensitive dbsnp
- SCIPhl default
- ...●... SCIPhl sensitive^{-B}

threshold used

- ▲ 1e-06
- × 1e-05
- * 1e-04
- 0.001
- ⊠ 0.005
- △ 0.01
- ◊ 0.02
- ⊗ 0.05
- + 0.1
- ⊕ 0.2
- ◇ 0.5

