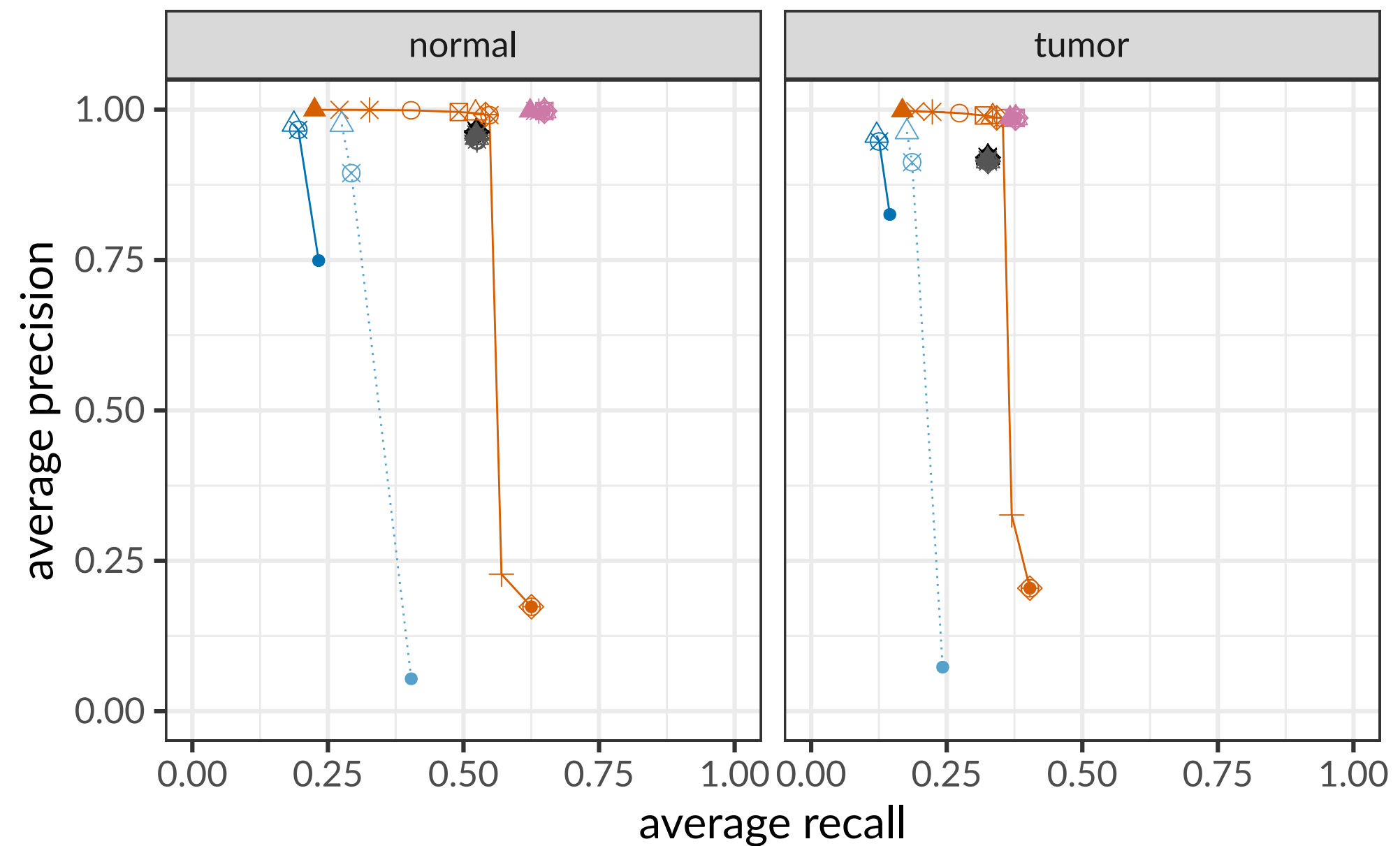


C

TNBC

normal

tumor



- 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