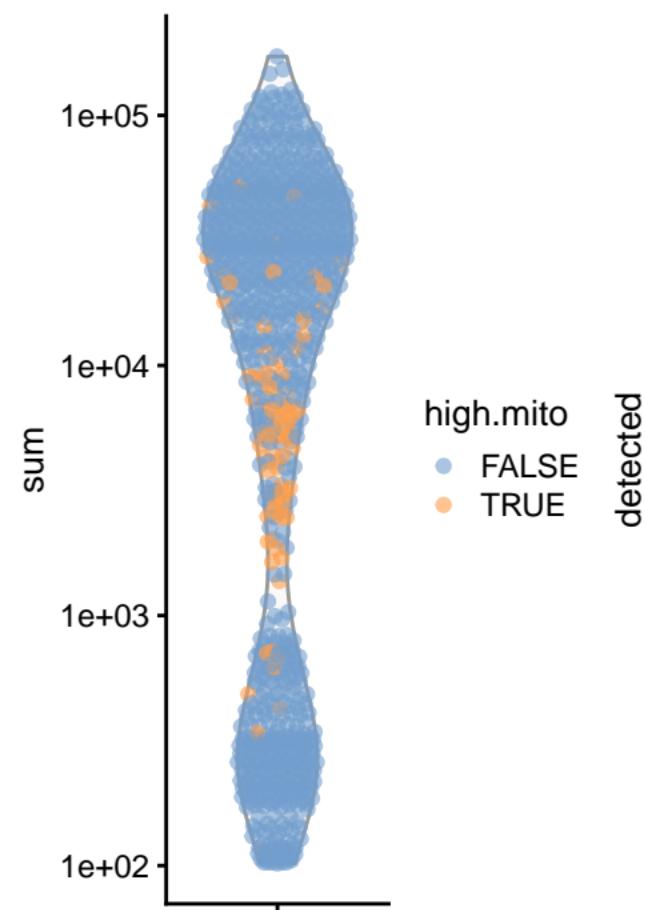
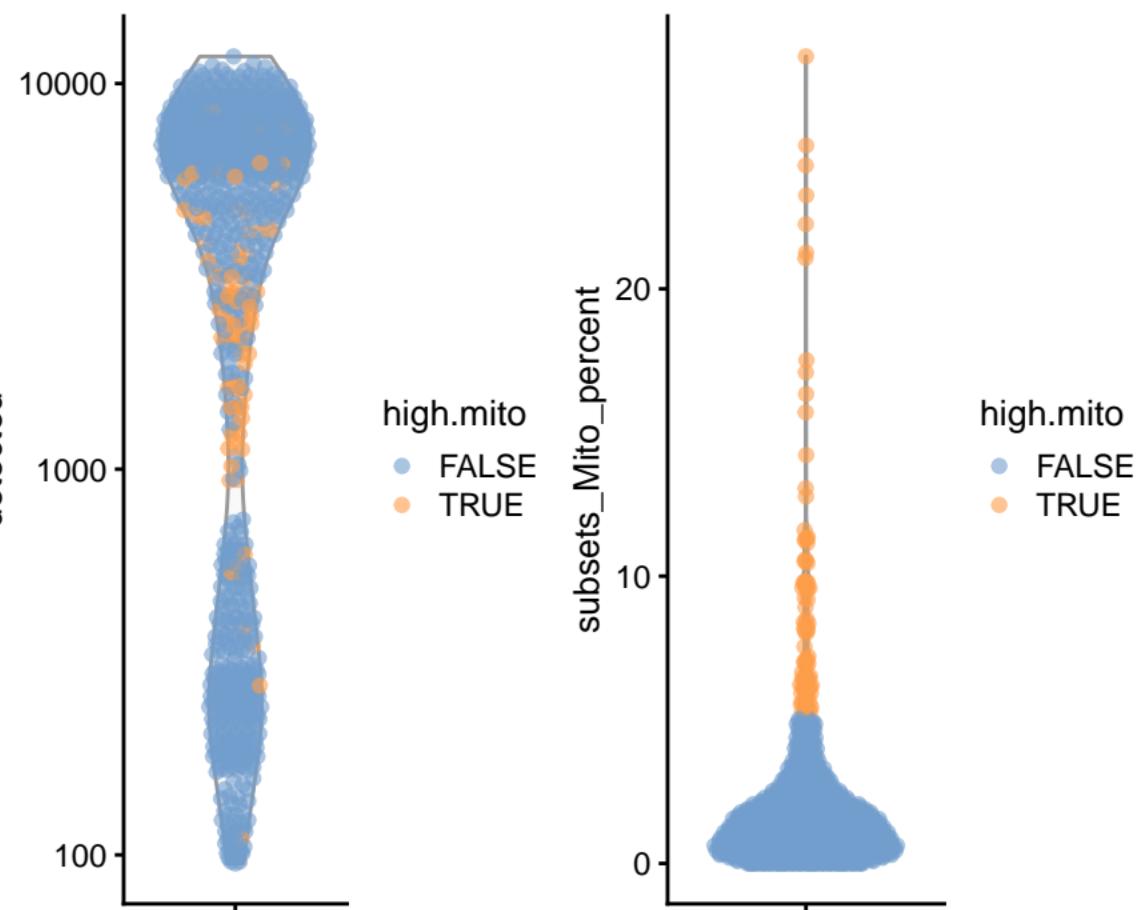


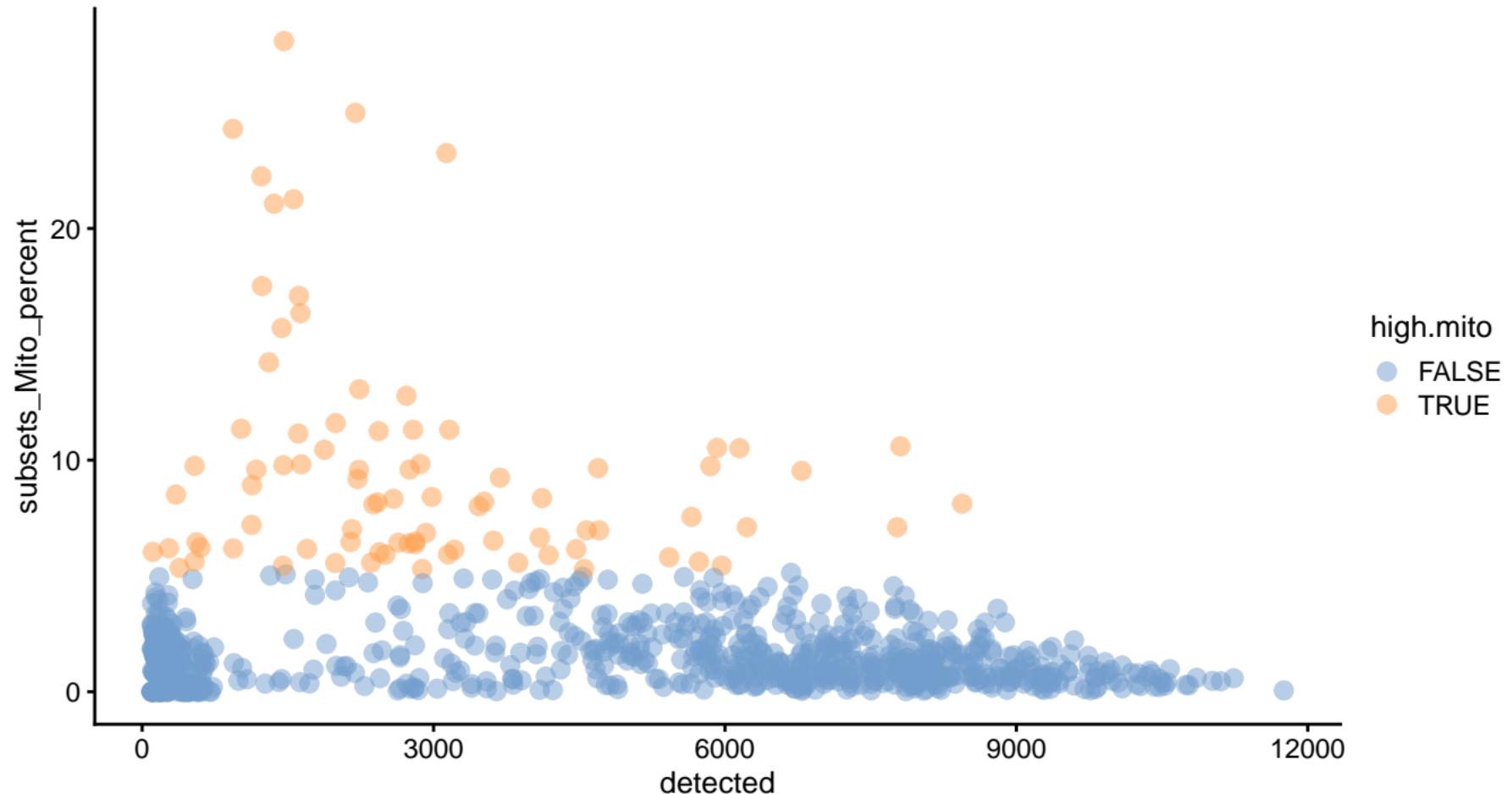
Total count: br5276.sacc.neun    Detected features



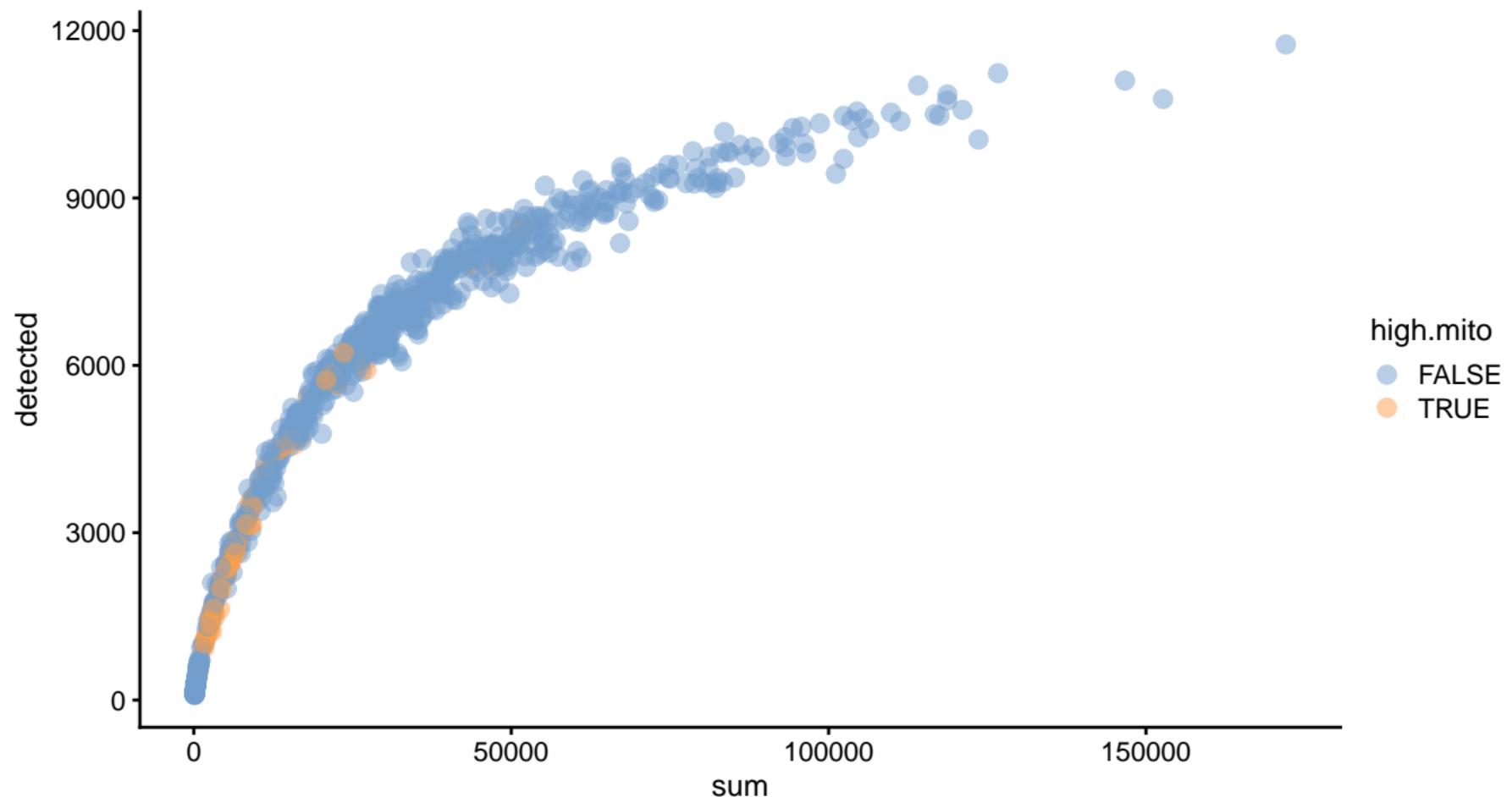
Mito % (cutoff = 5.191)

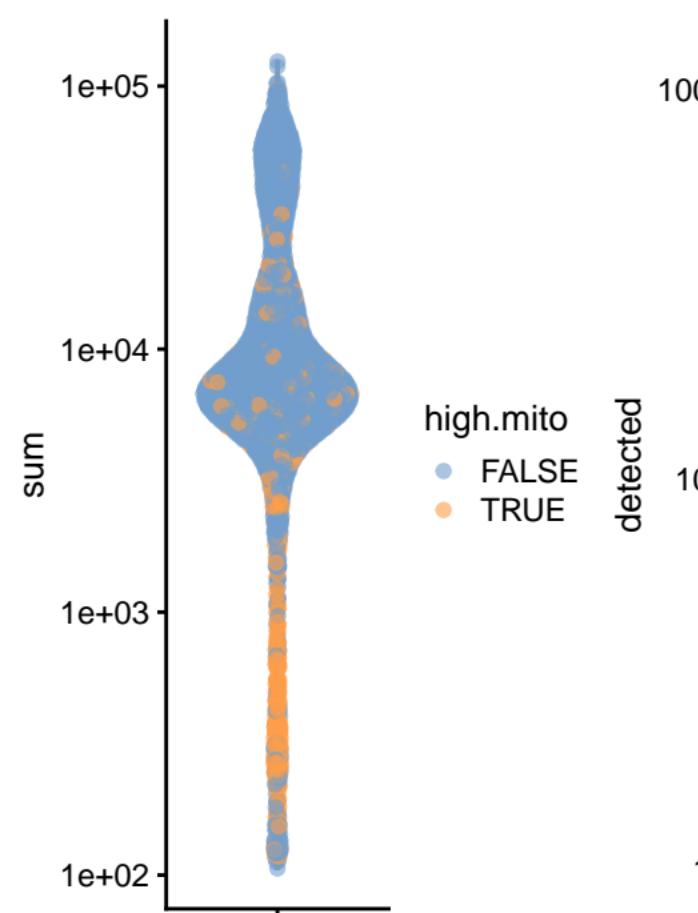
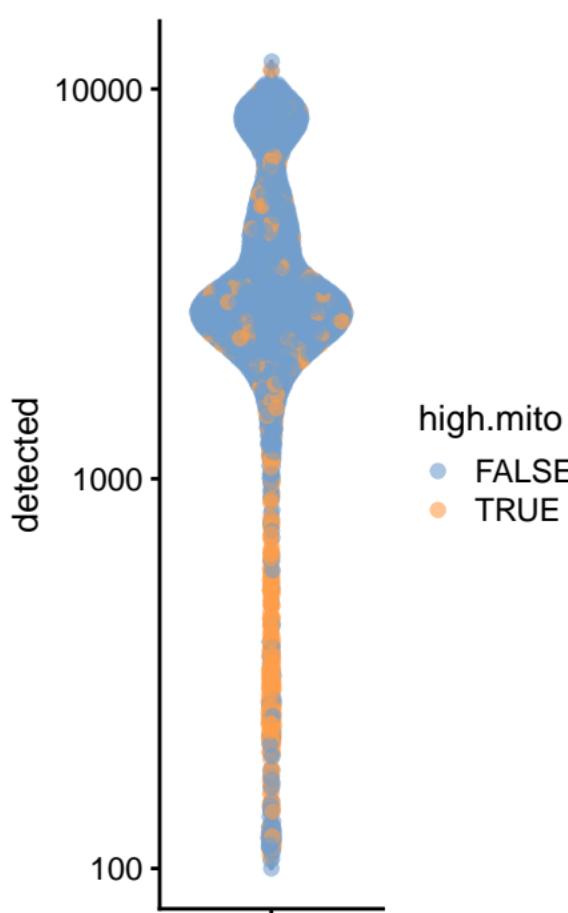
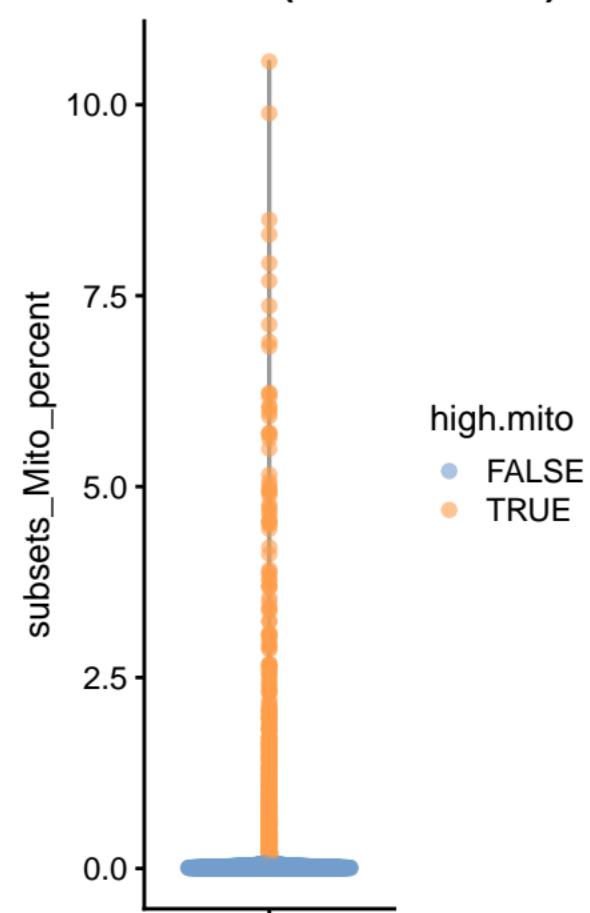


Sample: br5276.sacc.neun; pre-QC nNuclei: 926; nNuclei kept: 842 (0.91%)

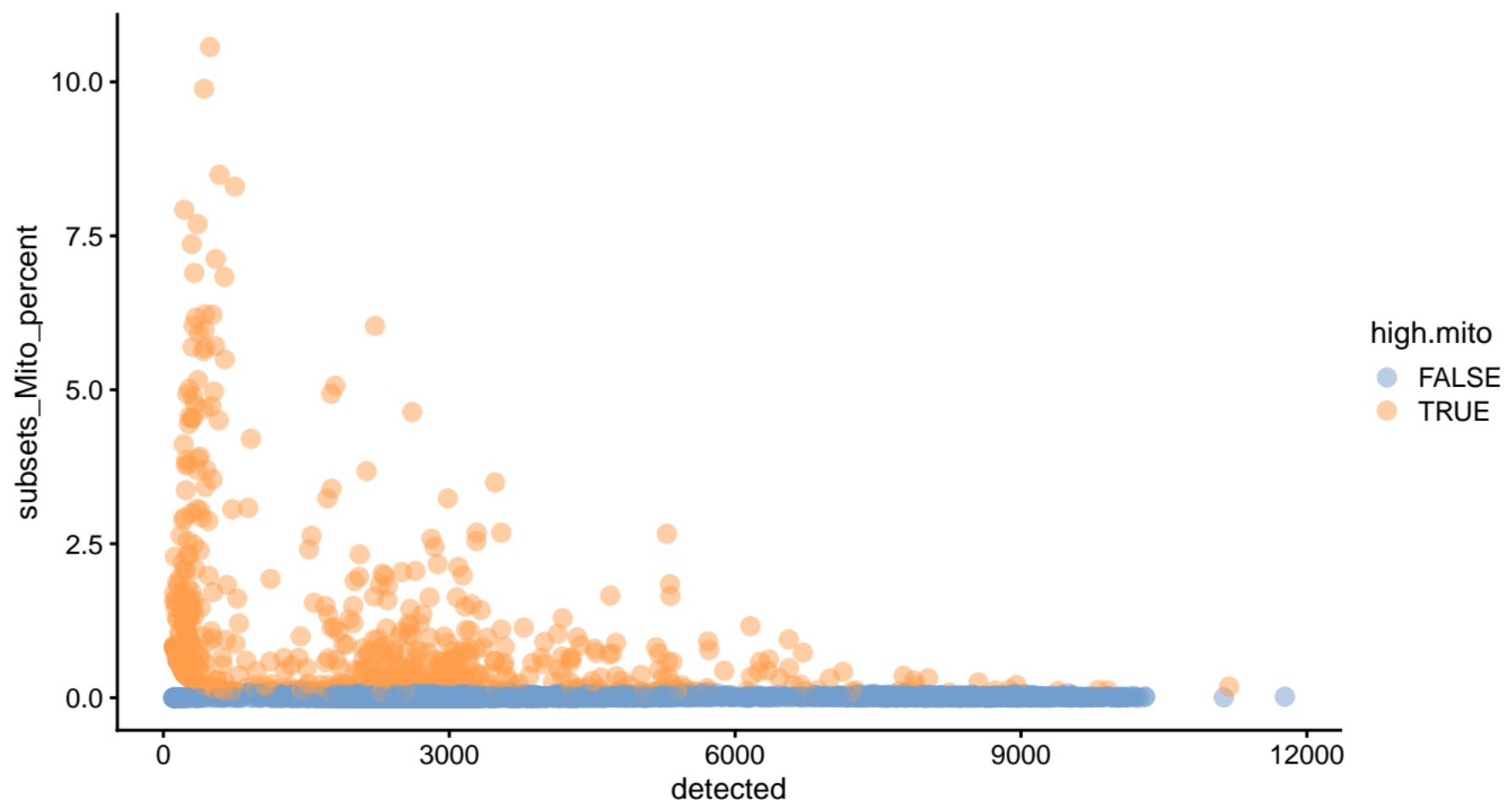


Sample: br5276.sacc.neun; pre-QC nNuclei: 926; nNuclei kept: 842 (0.91%)

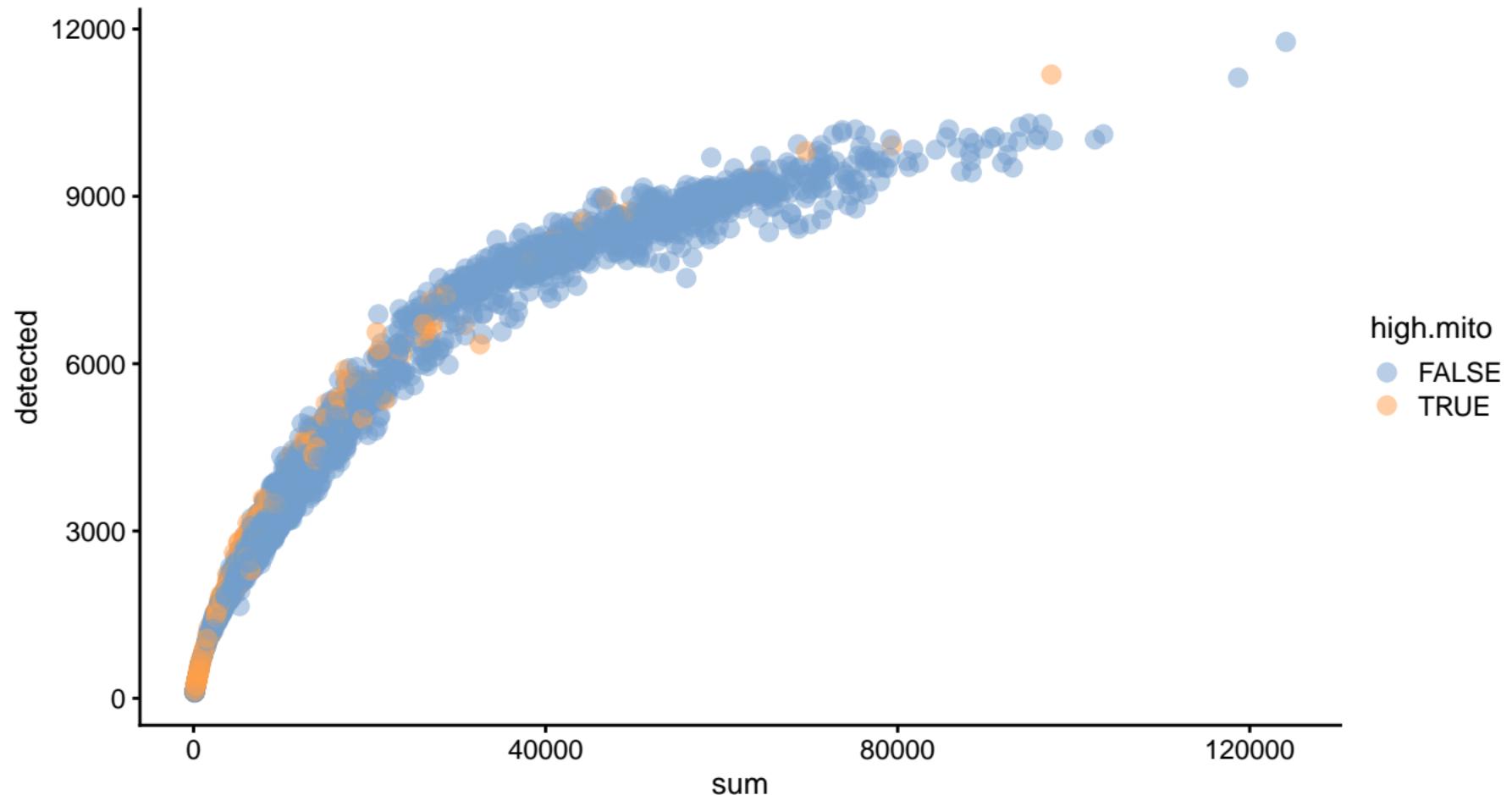


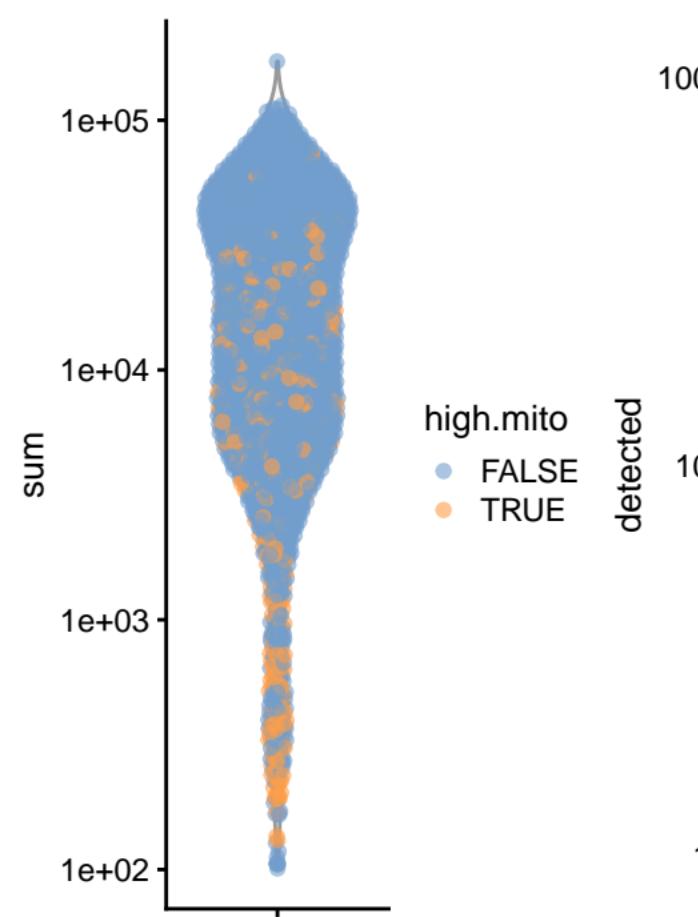
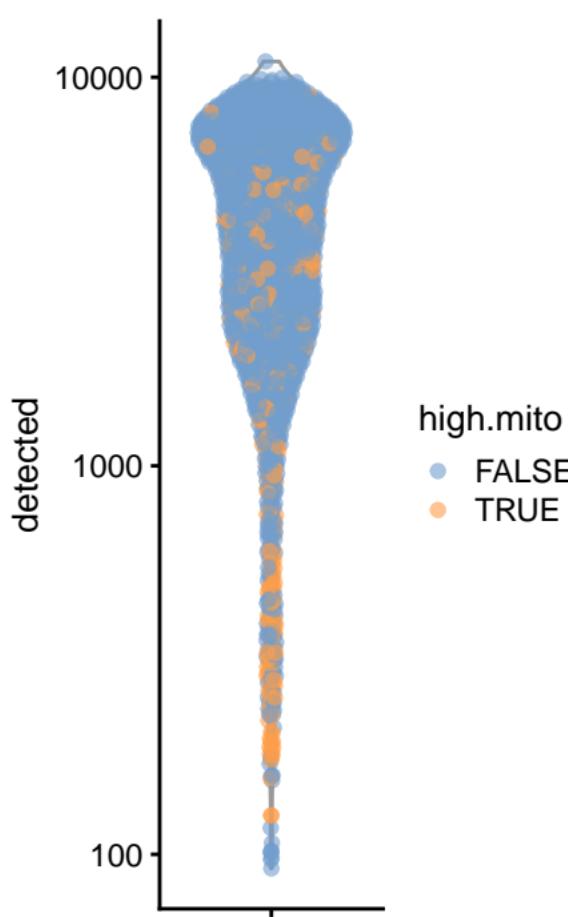
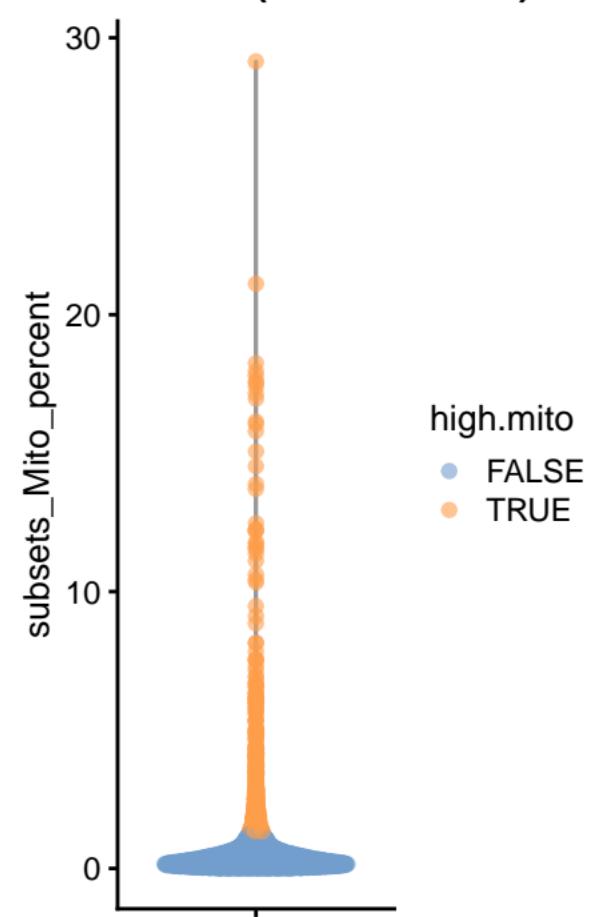
**Total count: br5400.nac****Detected features****Mito % (cutoff = 0.082)**

Sample: br5400.nac; pre-QC nNuclei: 4859; nNuclei kept: 4246 (0.87%)

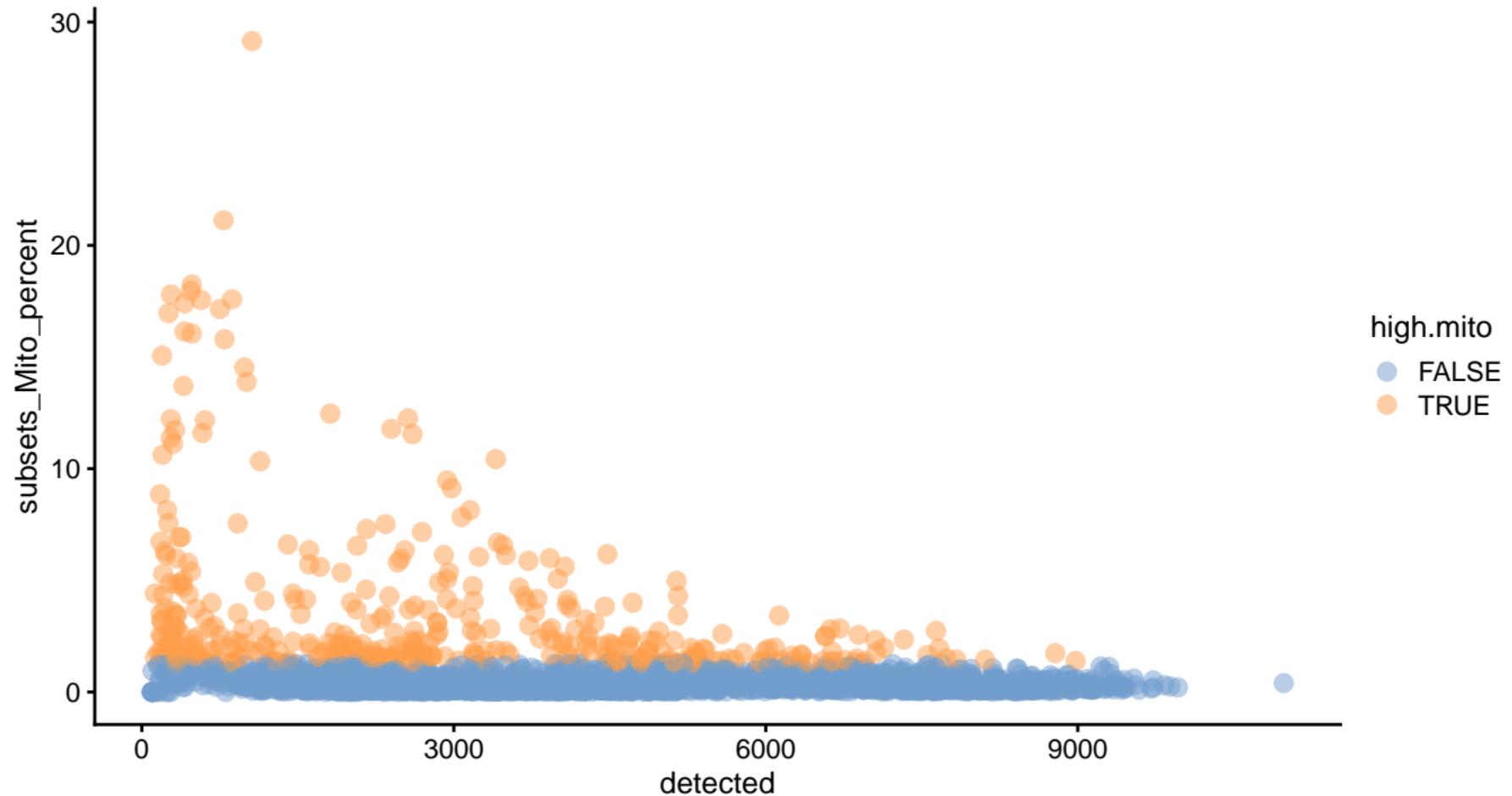


Sample: br5400.nac; pre-QC nNuclei: 4859; nNuclei kept: 4246 (0.87%)

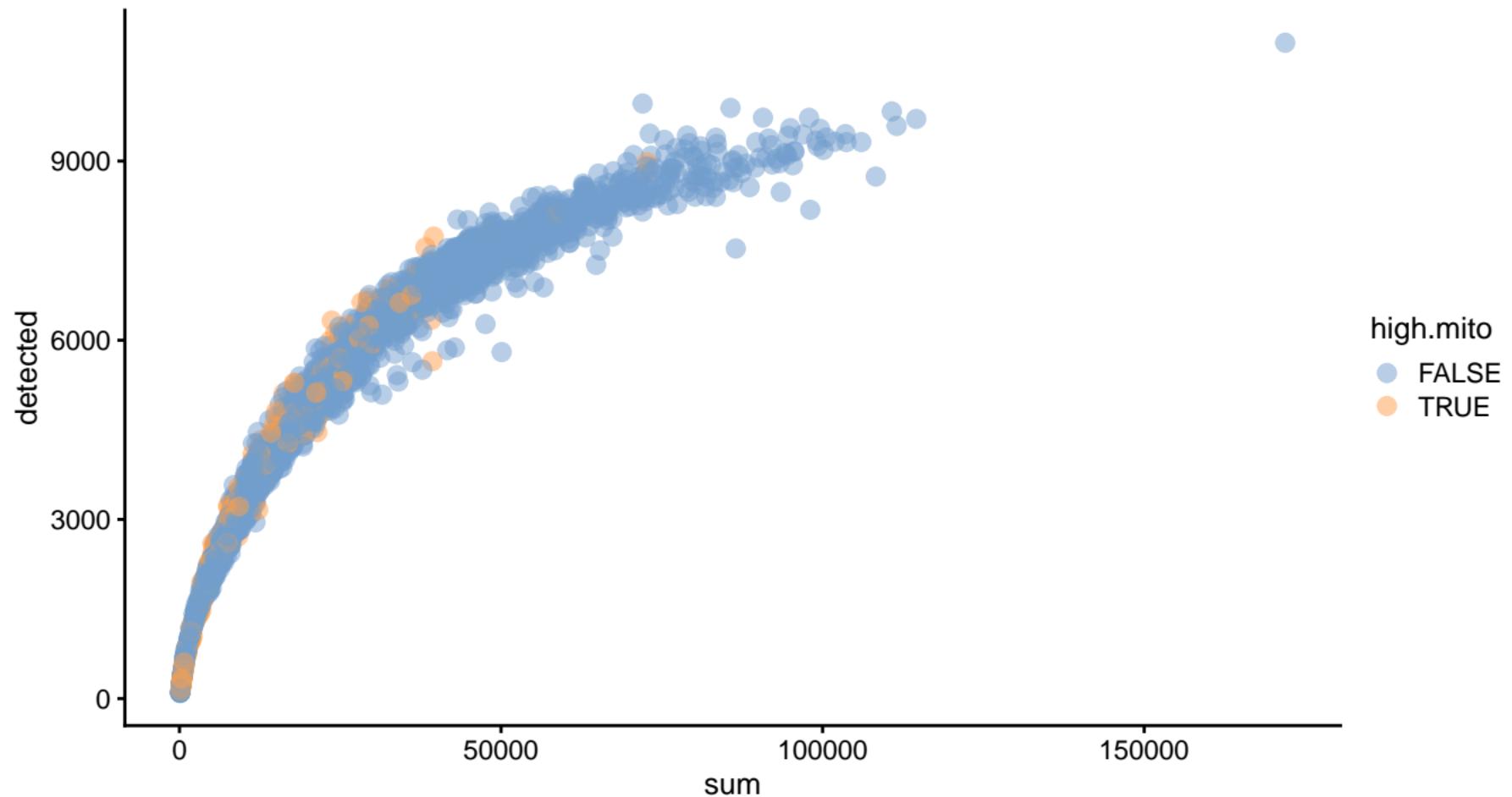


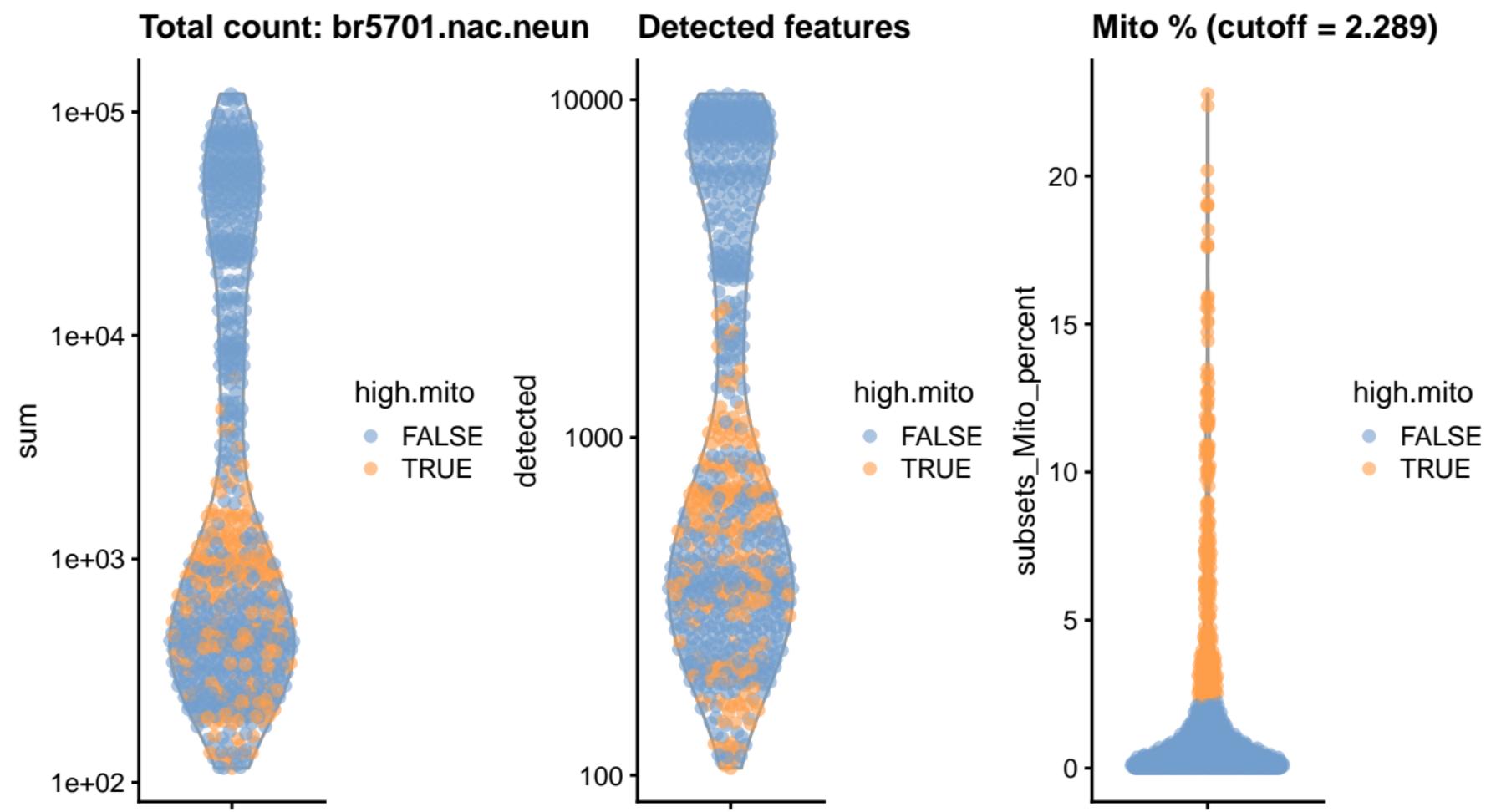
**Total count: br5276.nac****Detected features****Mito % (cutoff = 1.291)**

Sample: br5276.nac; pre-QC nNuclei: 3012; nNuclei kept: 2638 (0.88%)

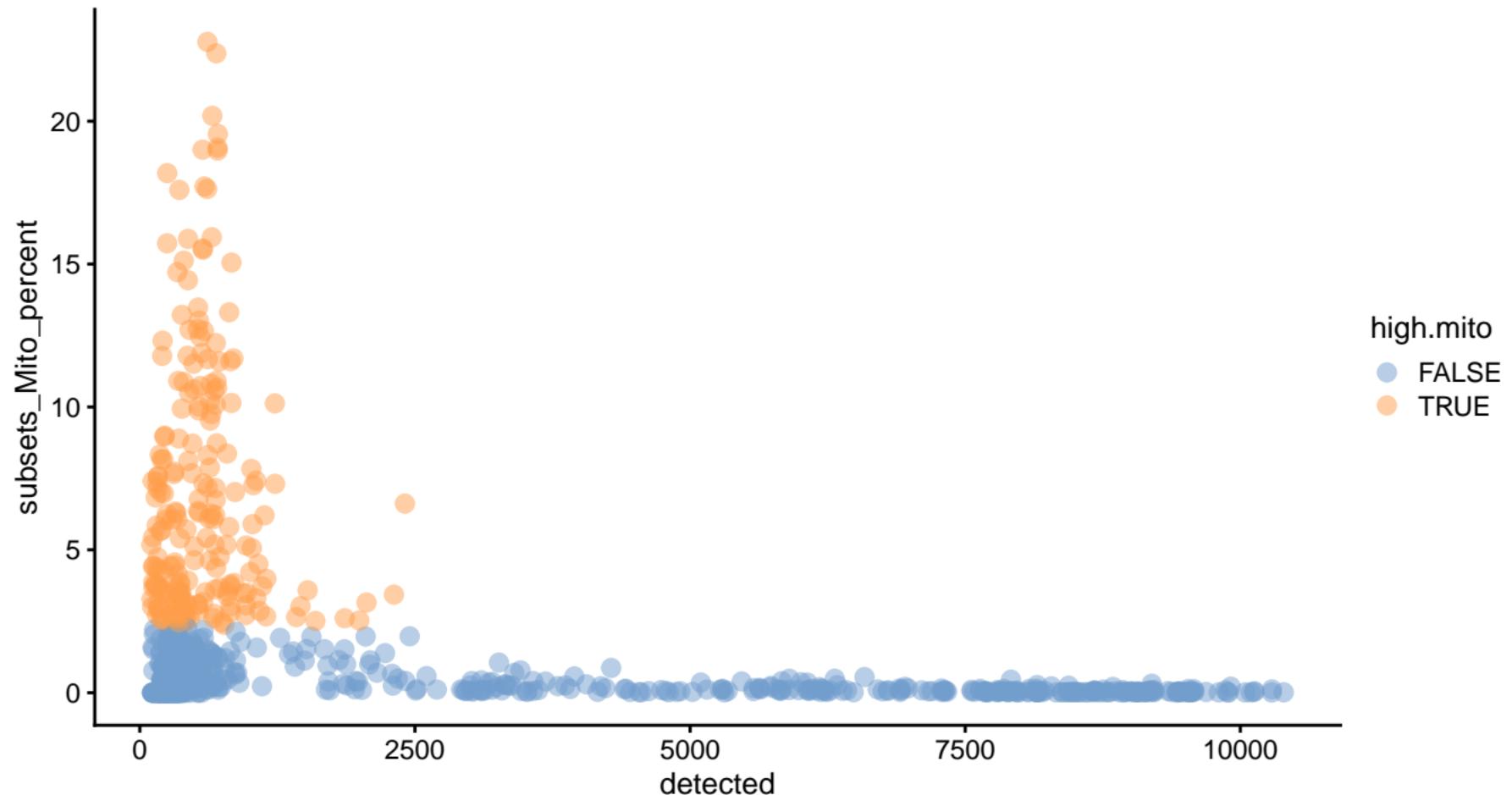


Sample: br5276.nac; pre-QC nNuclei: 3012; nNuclei kept: 2638 (0.88%)

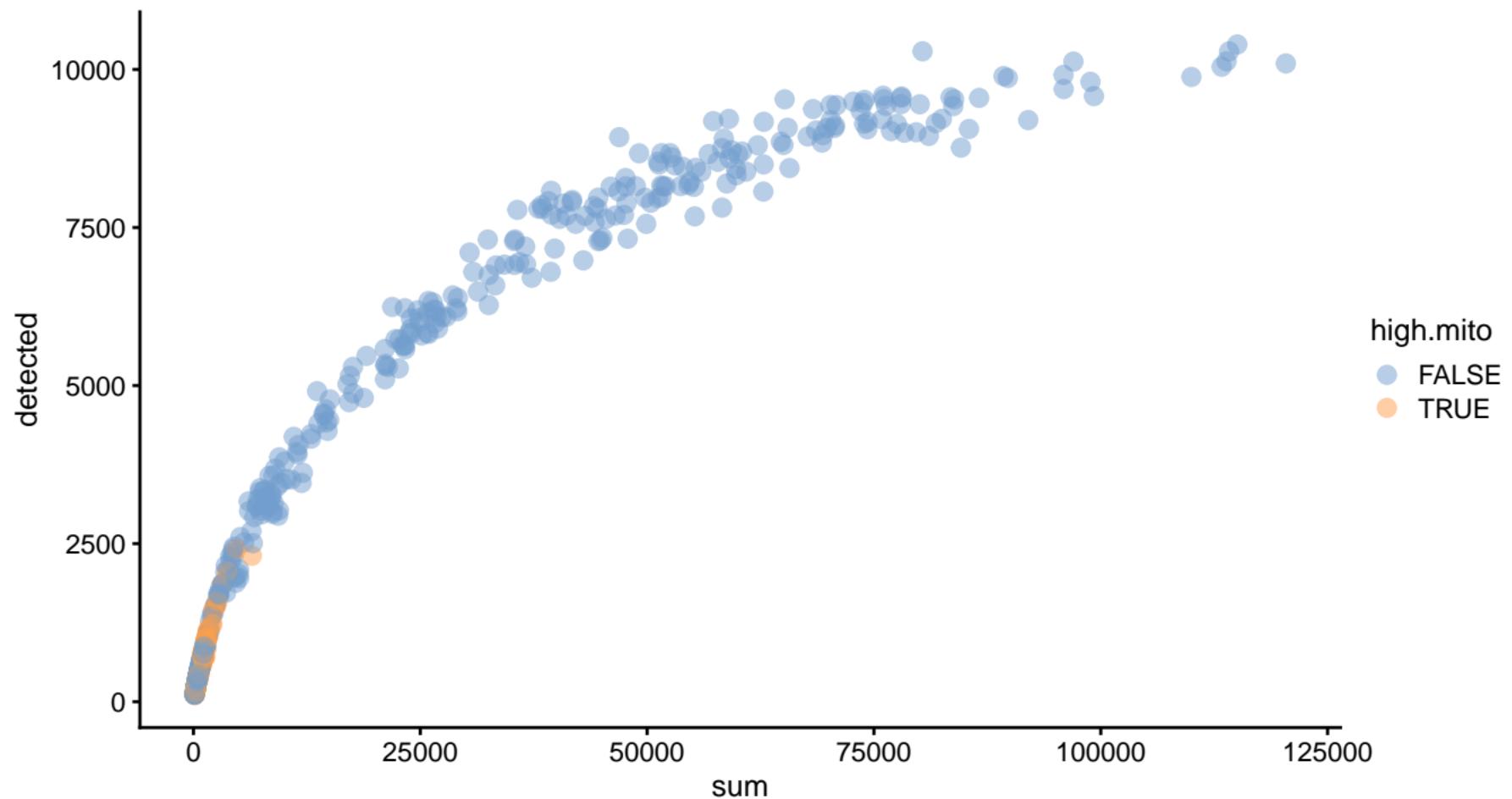


**Total count: br5701.nac.neun****Detected features****Mito % (cutoff = 2.289)**

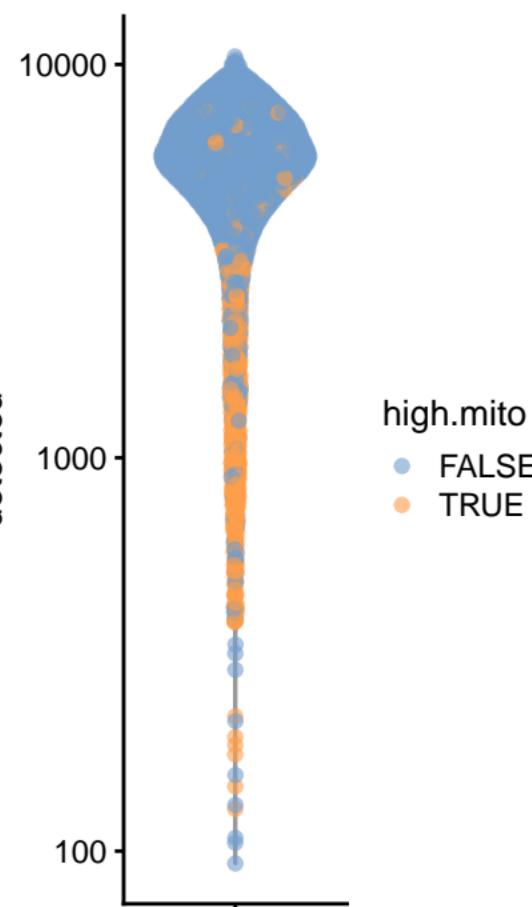
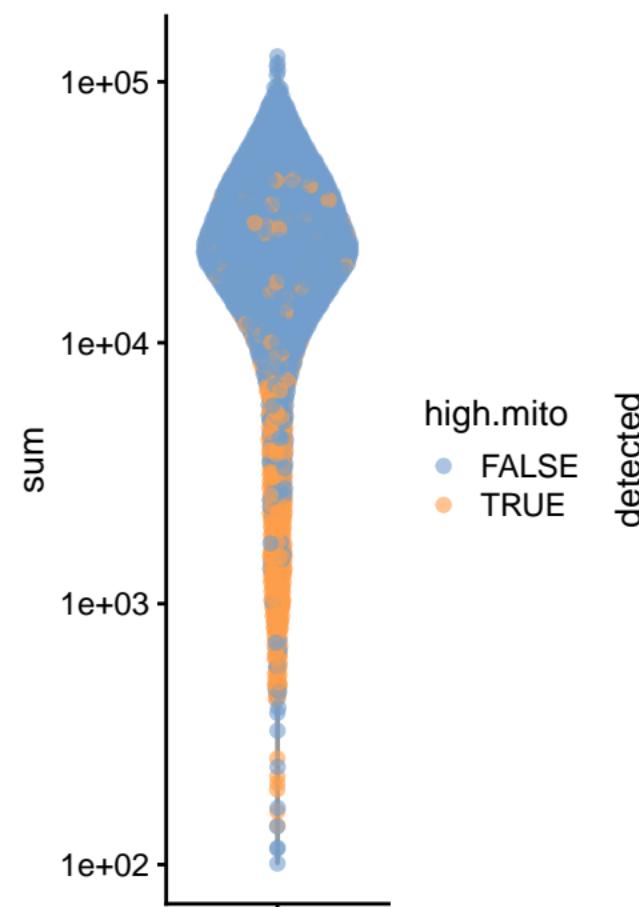
Sample: br5701.nac.neun; pre-QC nNuclei: 814; nNuclei kept: 599 (0.74%)



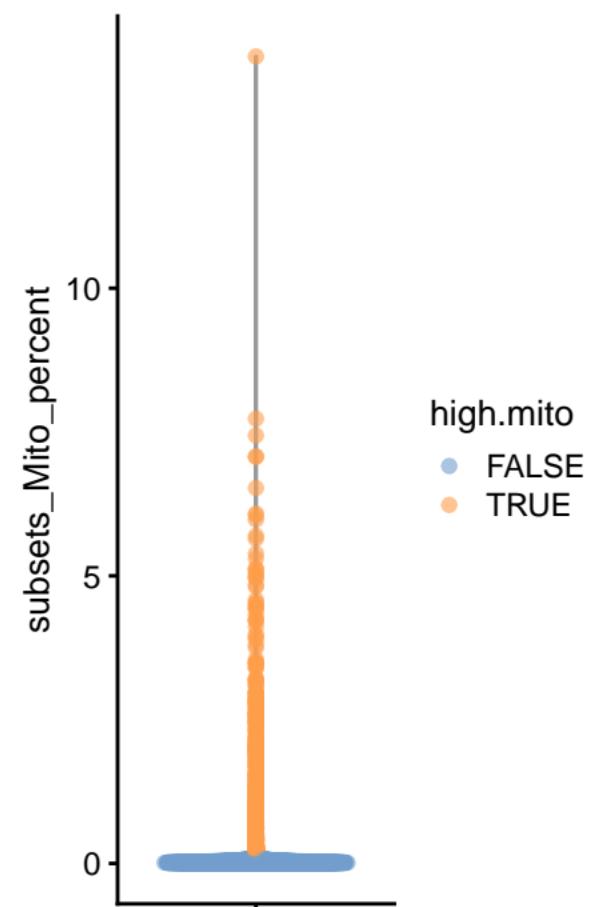
Sample: br5701.nac.neun; pre-QC nNuclei: 814; nNuclei kept: 599 (0.74%)



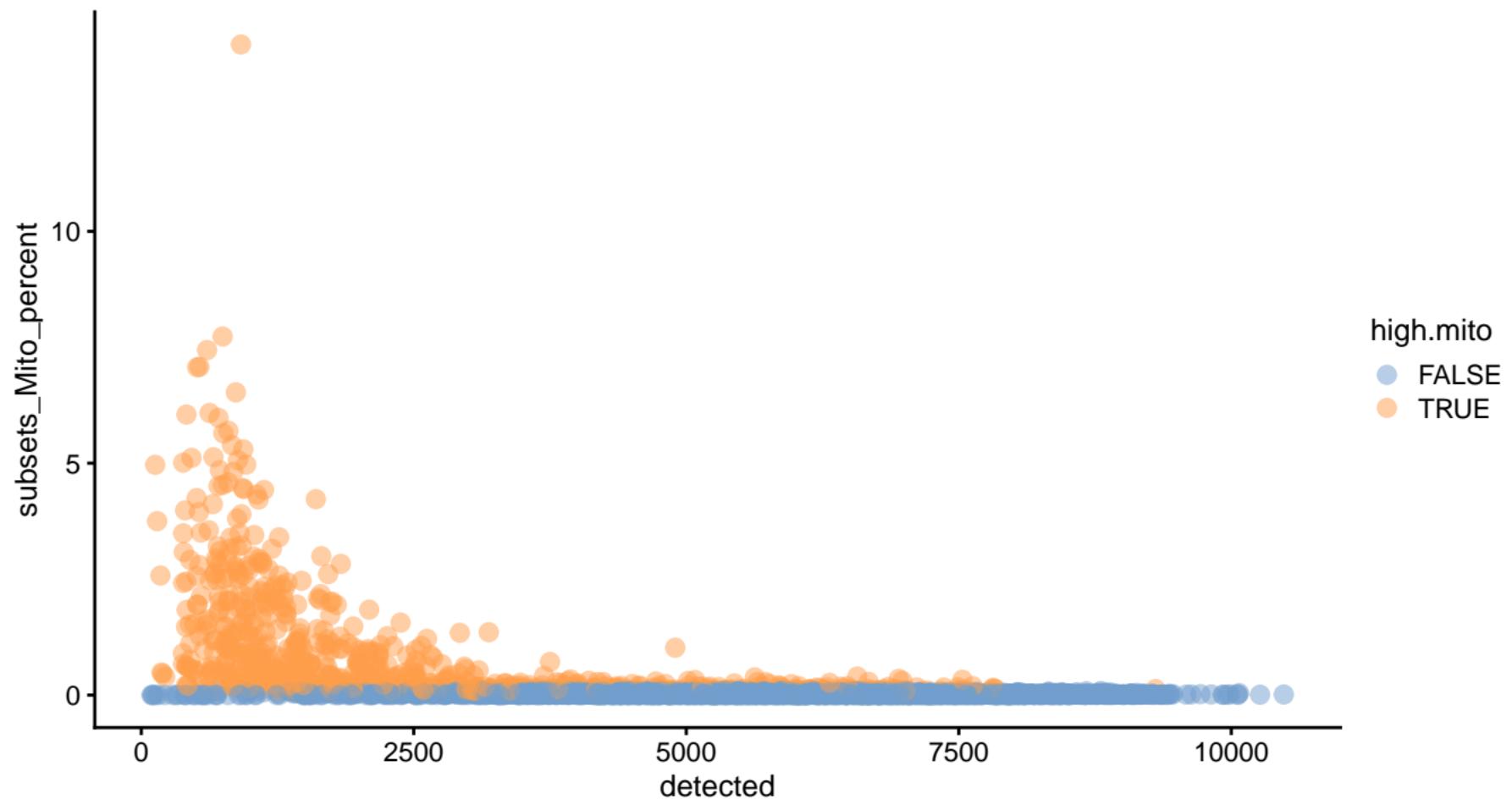
### Total count: br5701.sacc.neun    Detected features



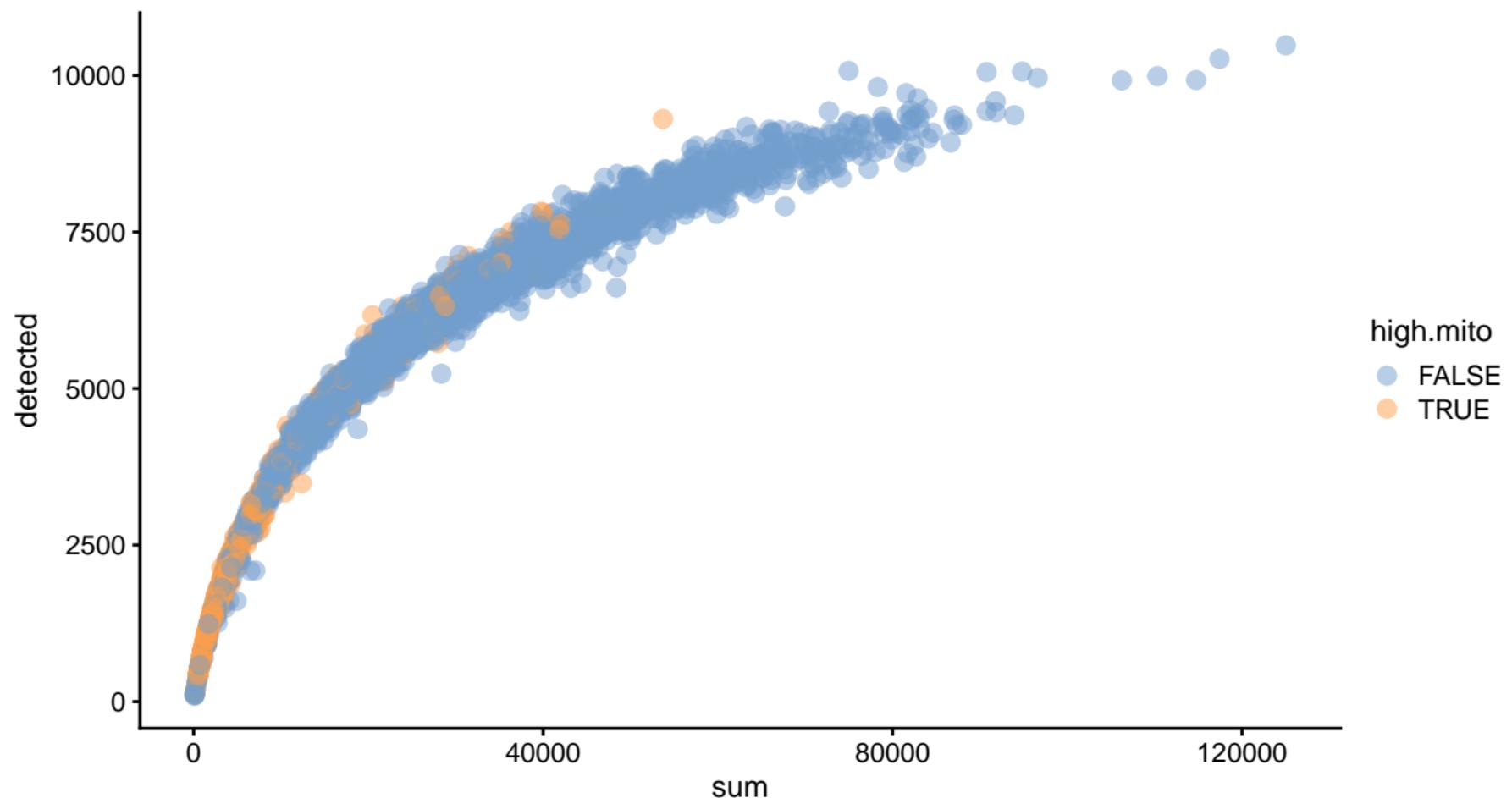
### Mito % (cutoff = 0.086)



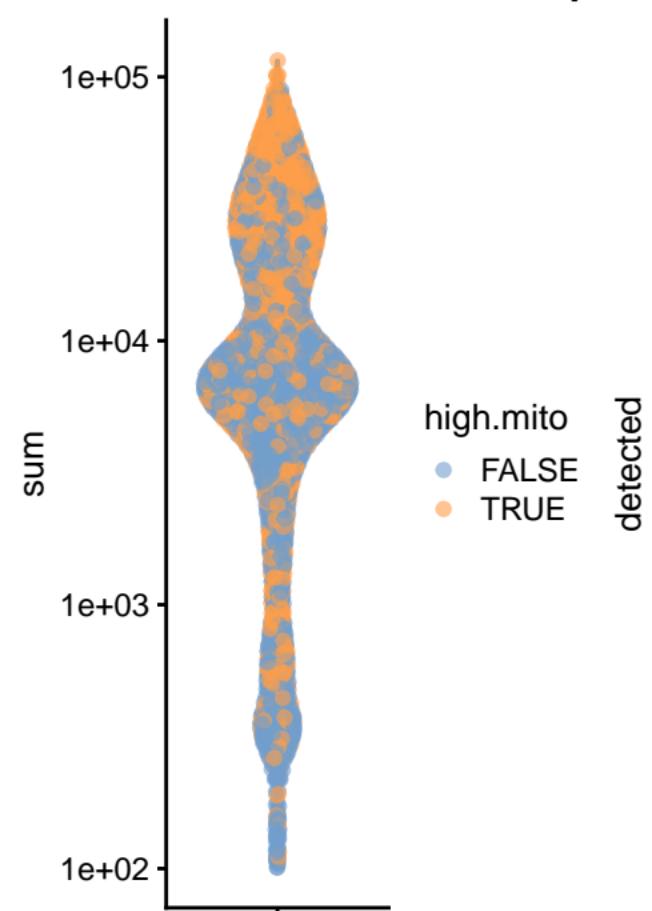
Sample: br5701.sacc.neun; pre-QC nNuclei: 4518; nNuclei kept: 3829 (0.85%)



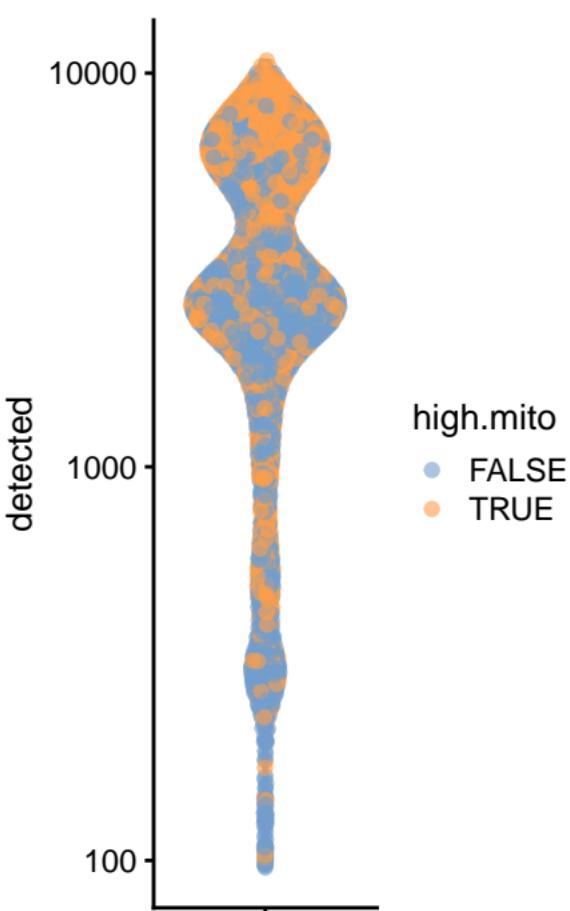
Sample: br5701.sacc.neun; pre-QC nNuclei: 4518; nNuclei kept: 3829 (0.85%)



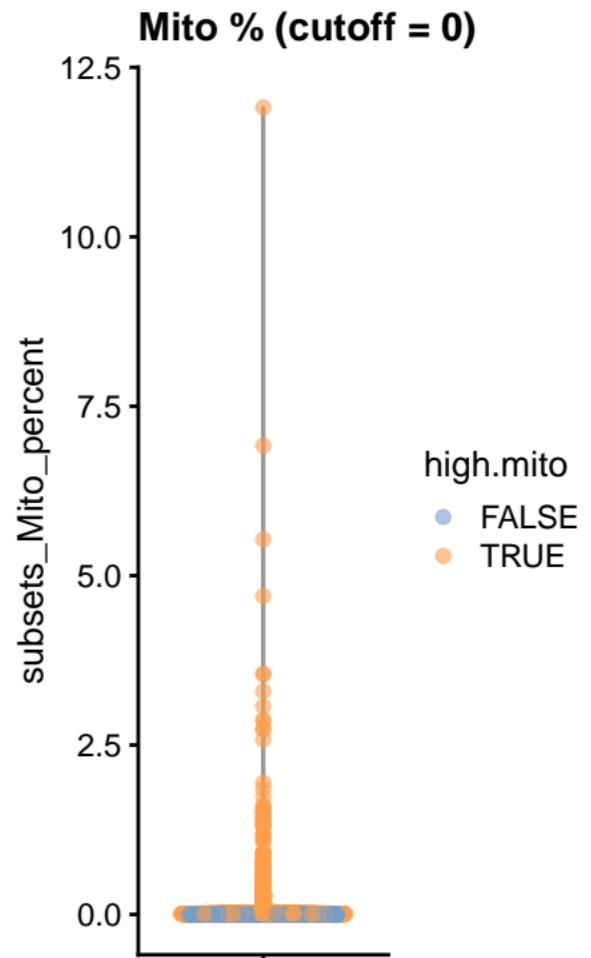
**Total count: br5207.dlpfc**



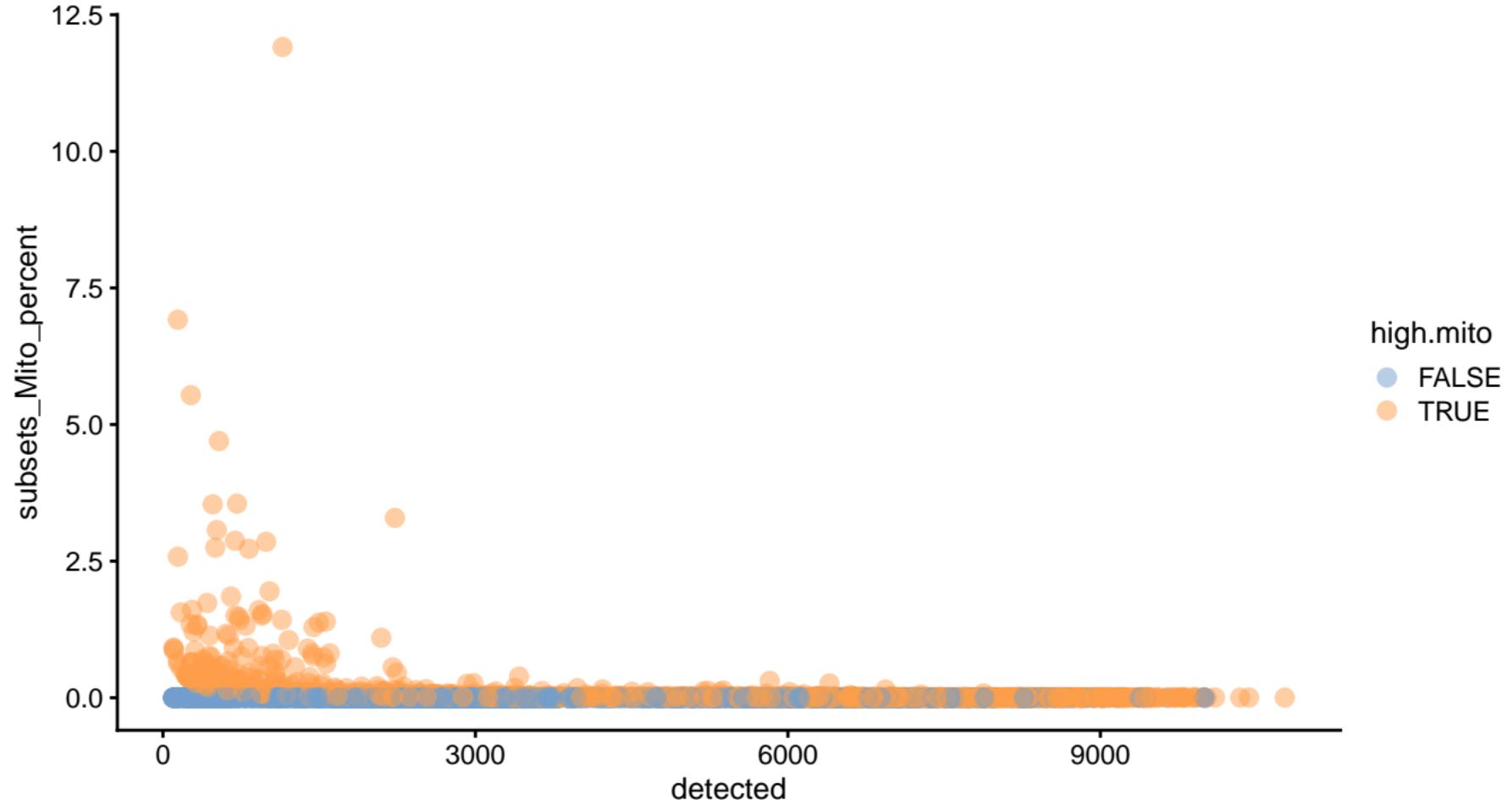
**Detected features**



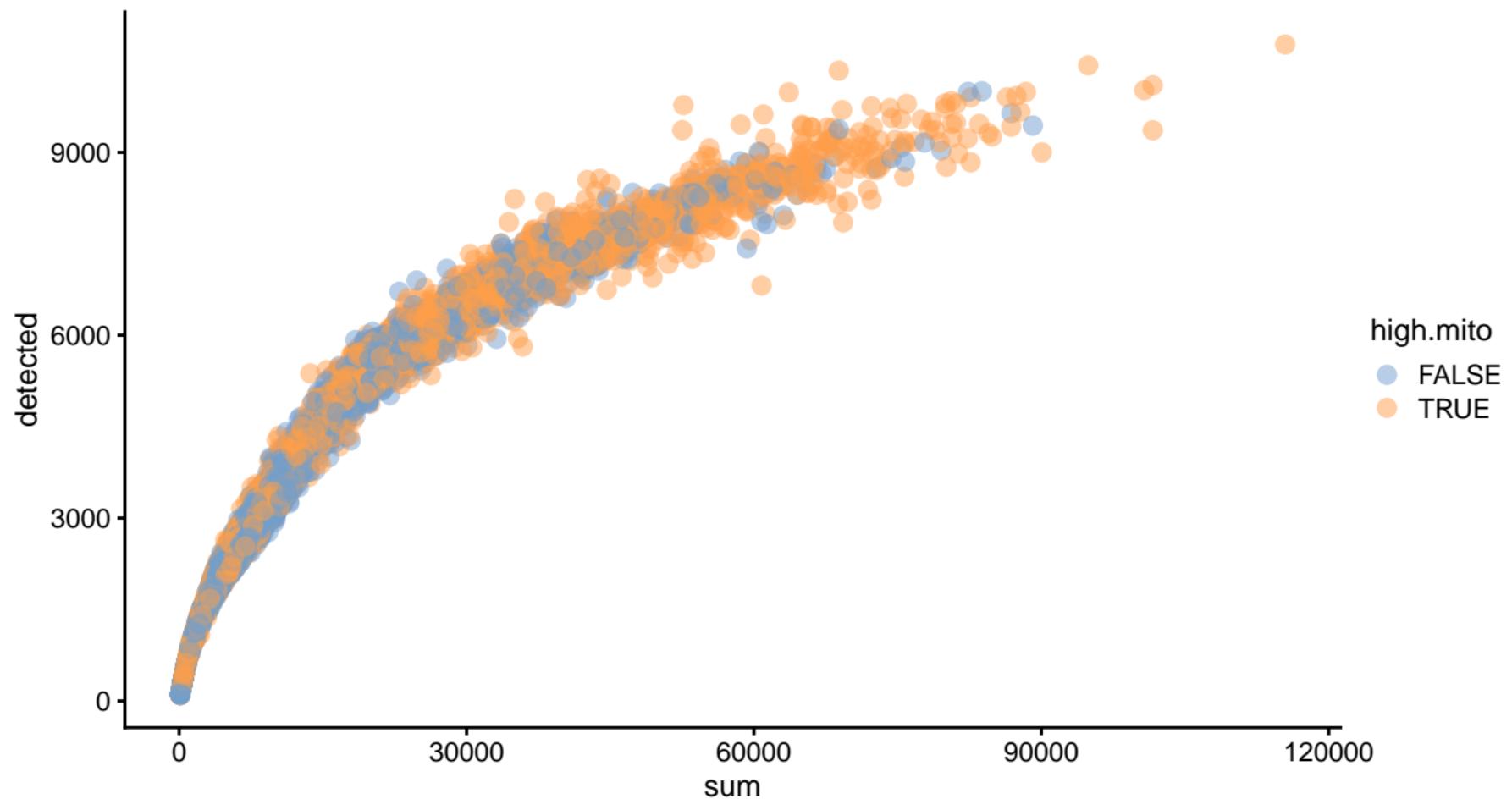
**Mito % (cutoff = 0)**



Sample: br5207.dlpfc; pre-QC nNuclei: 6453; nNuclei kept: 3628 (0.56%)



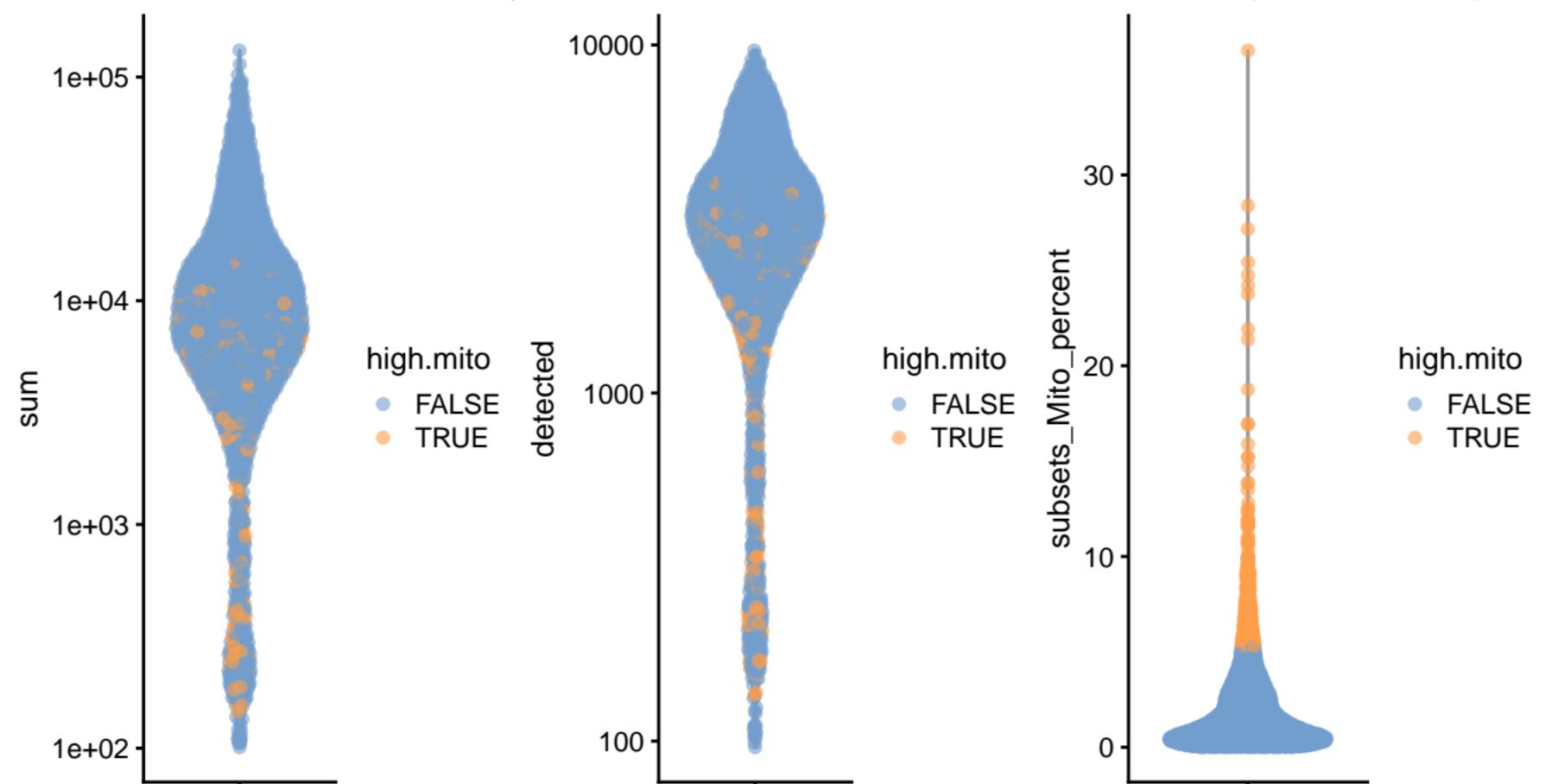
Sample: br5207.dlpfc; pre-QC nNuclei: 6453; nNuclei kept: 3628 (0.56%)



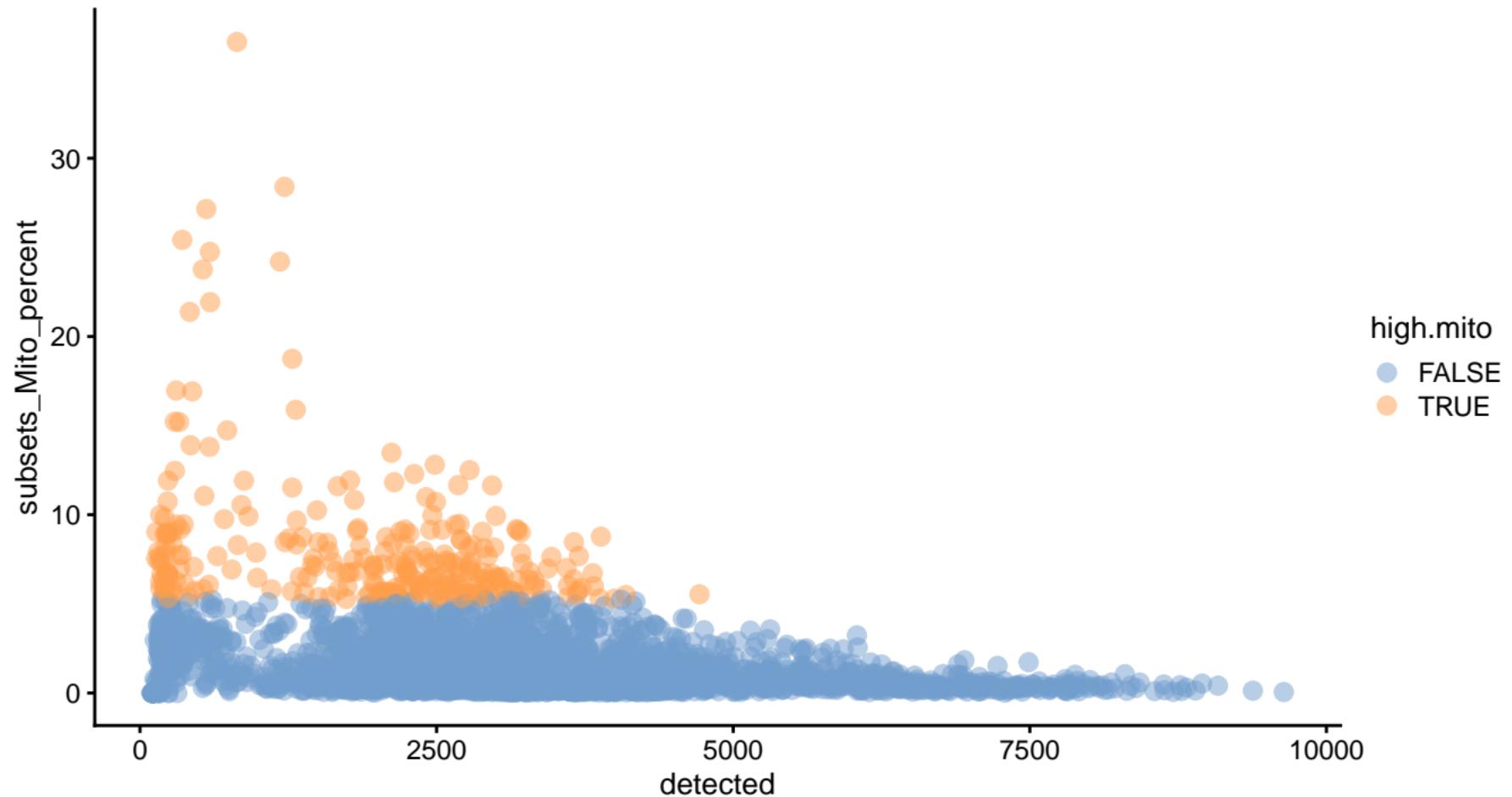
Total count: br5276.amy.neun

Detected features

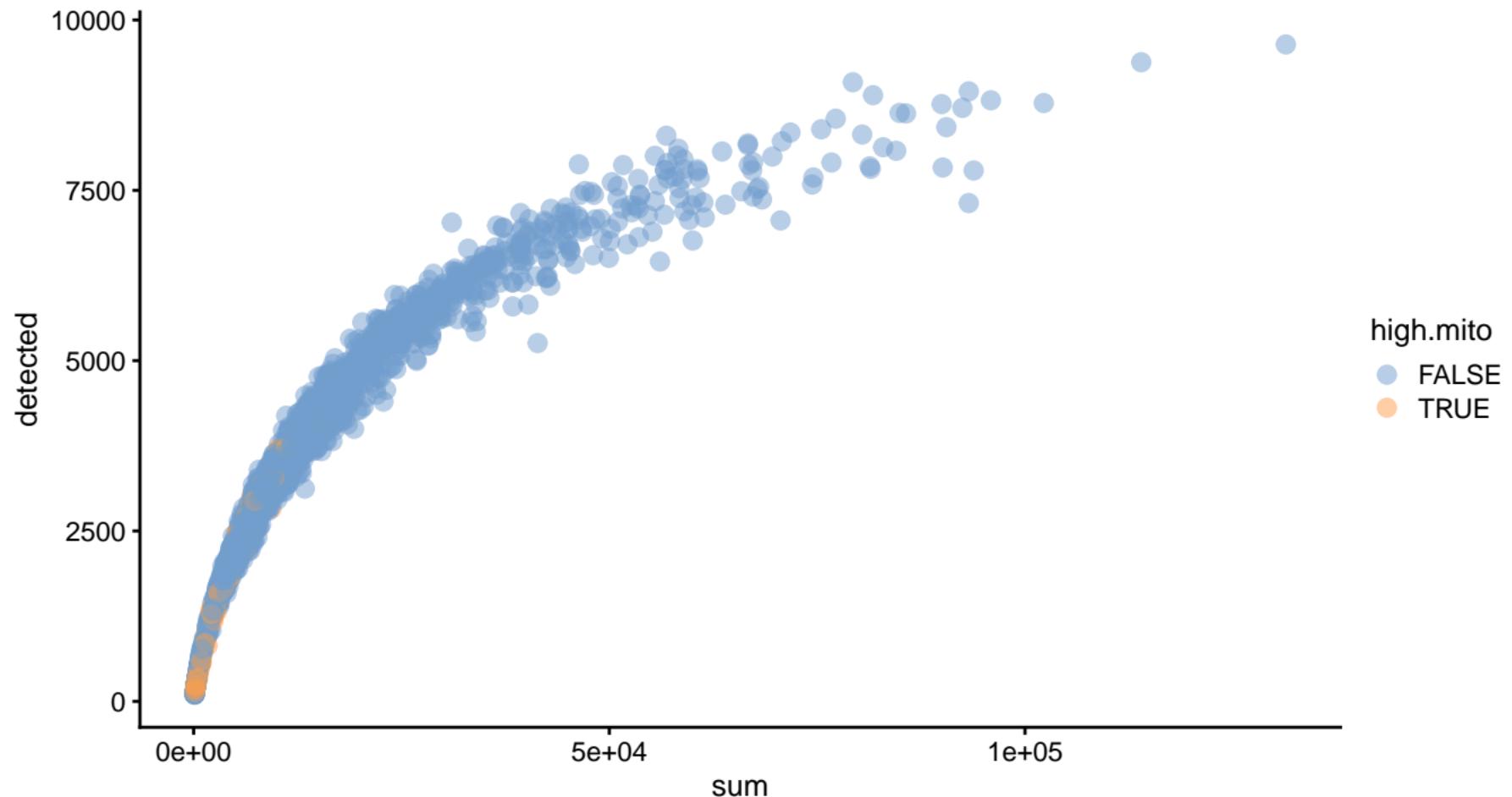
Mito % (cutoff = 5.259)

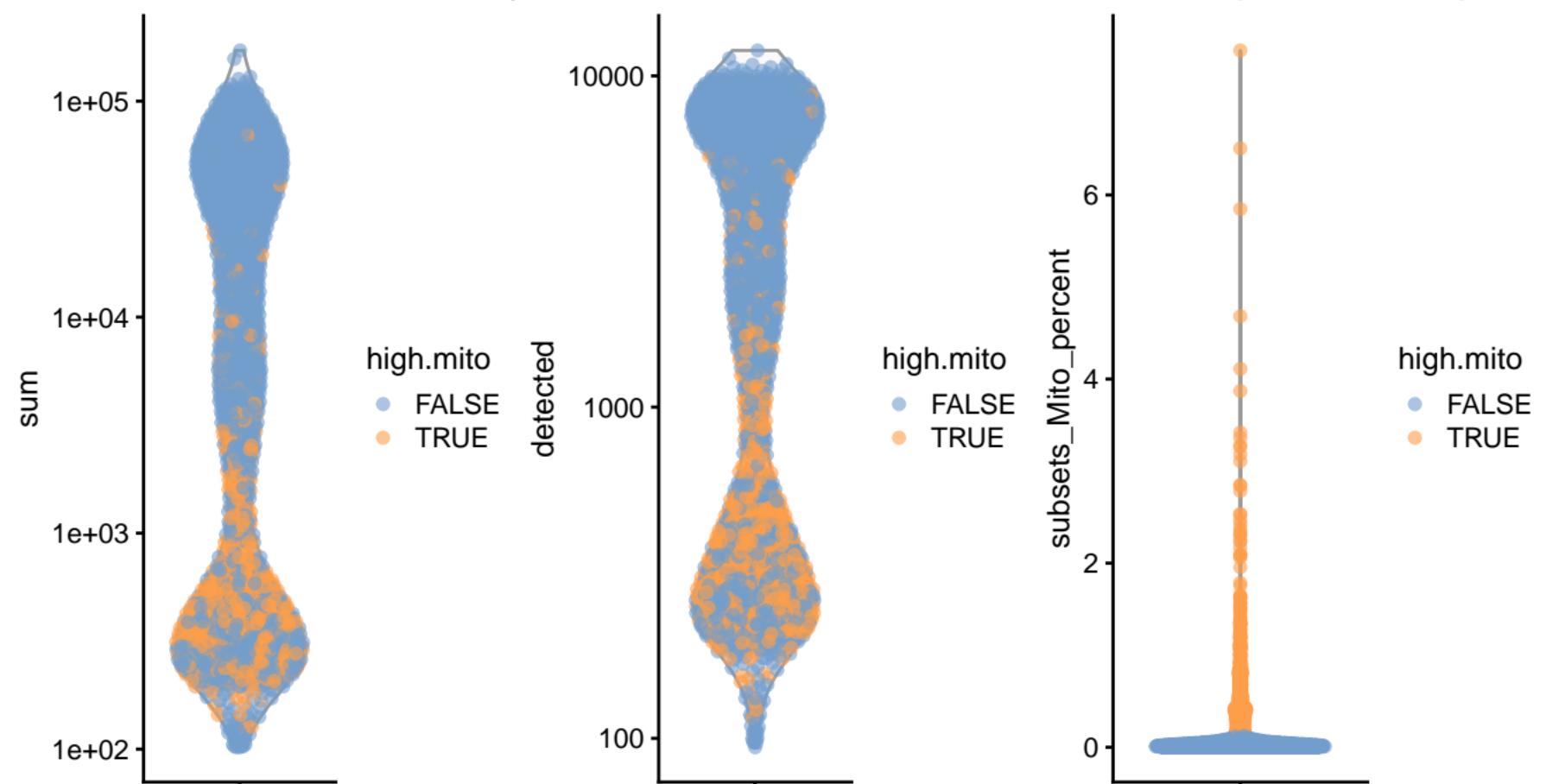


Sample: br5276.amy.neun; pre-QC nNuclei: 2738; nNuclei kept: 2469 (0.9%)

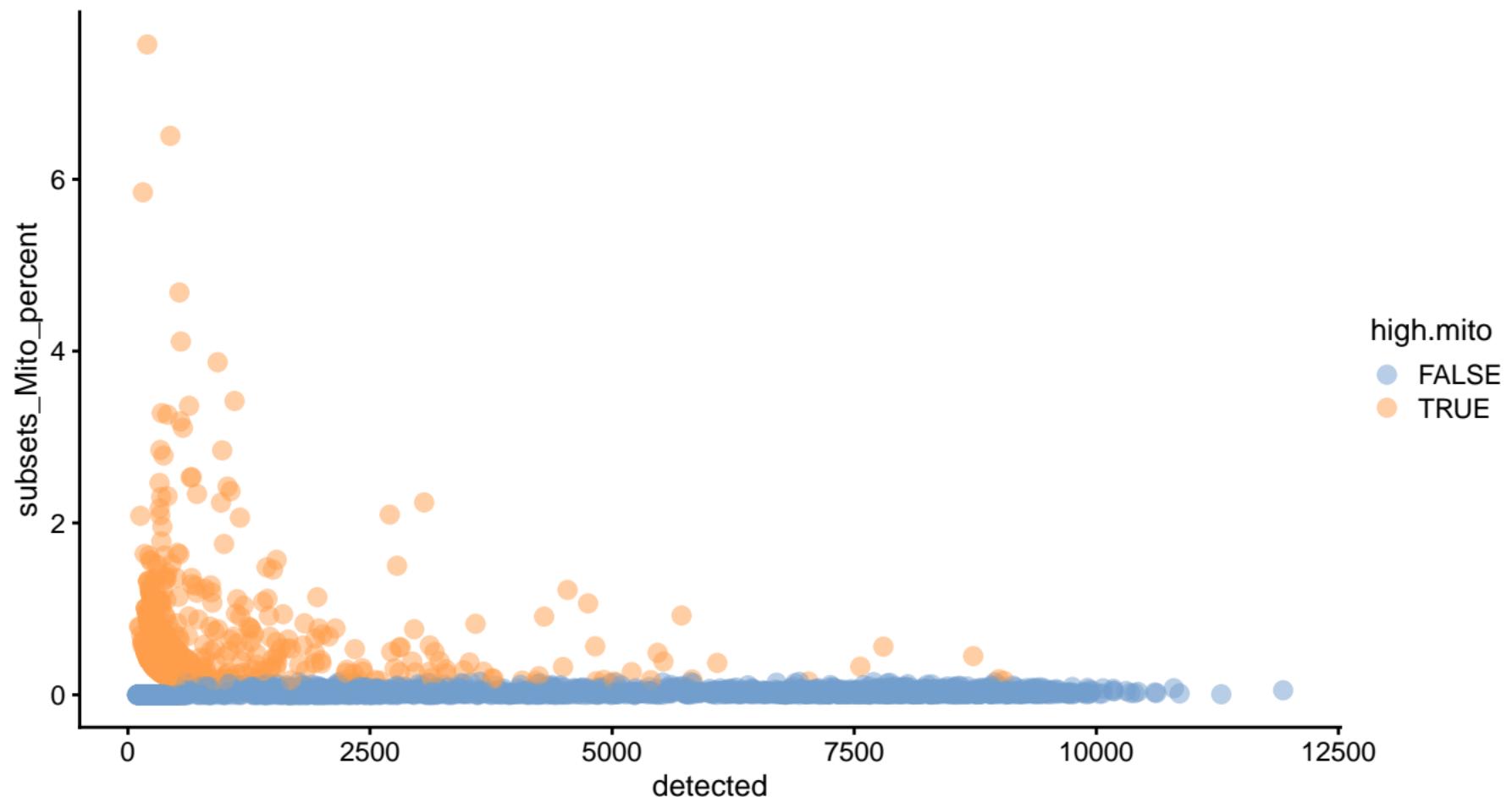


Sample: br5276.amy.neun; pre-QC nNuclei: 2738; nNuclei kept: 2469 (0.9%)

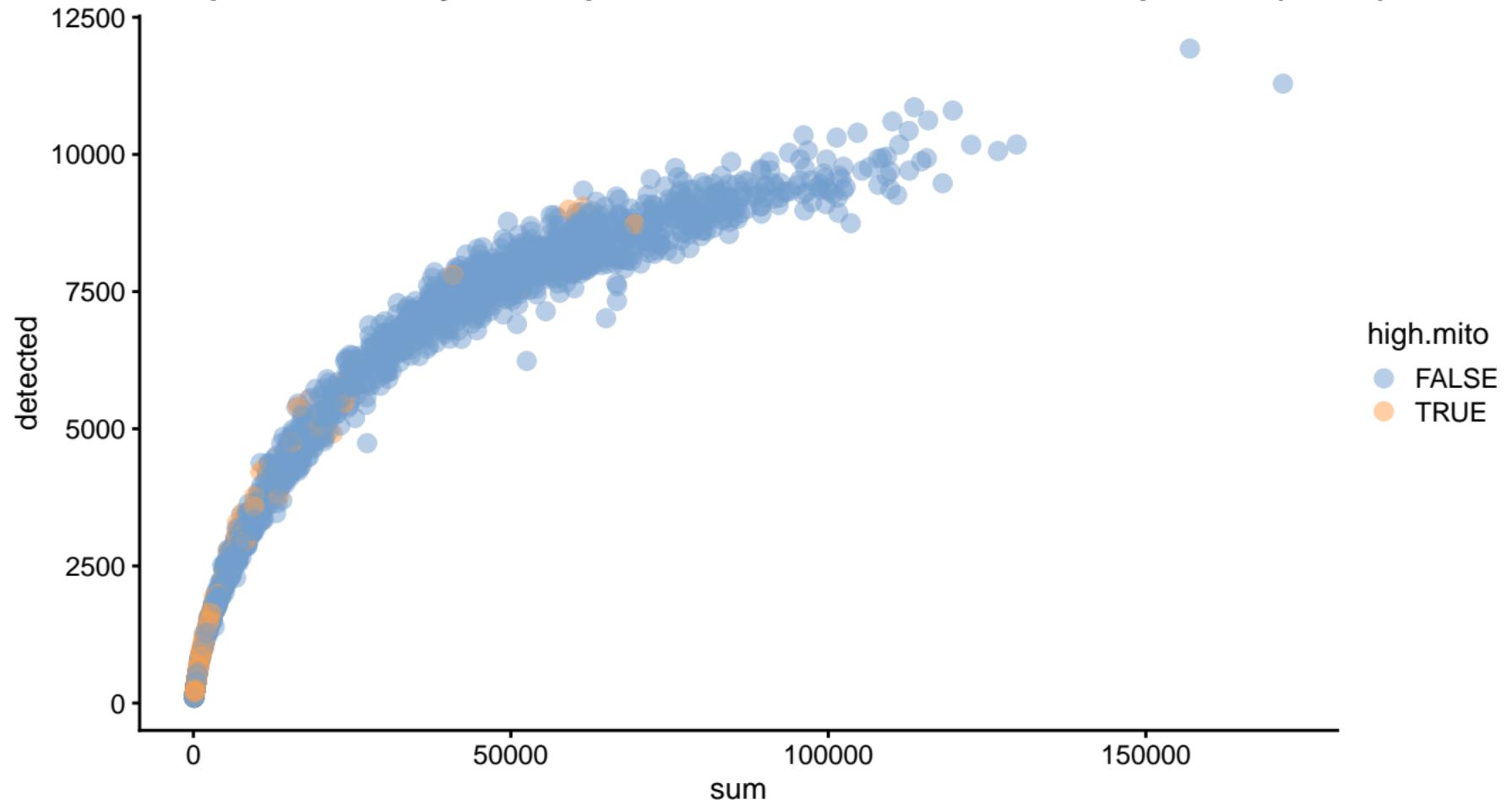


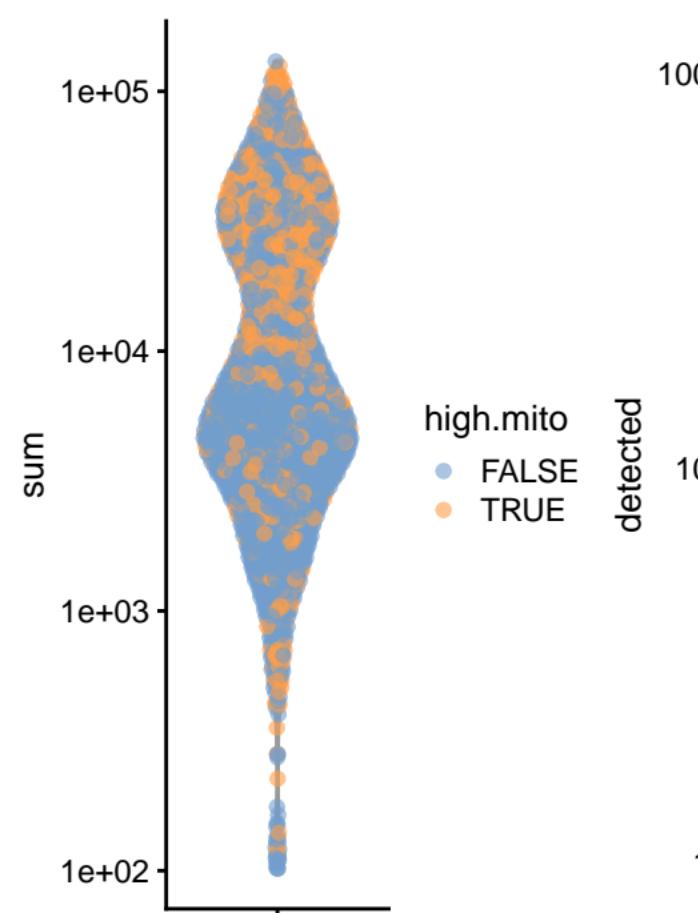
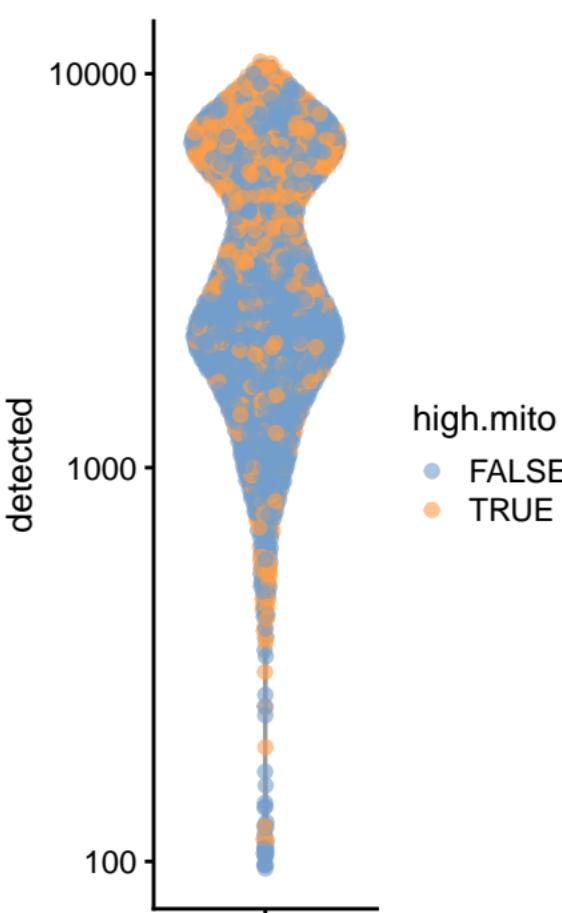
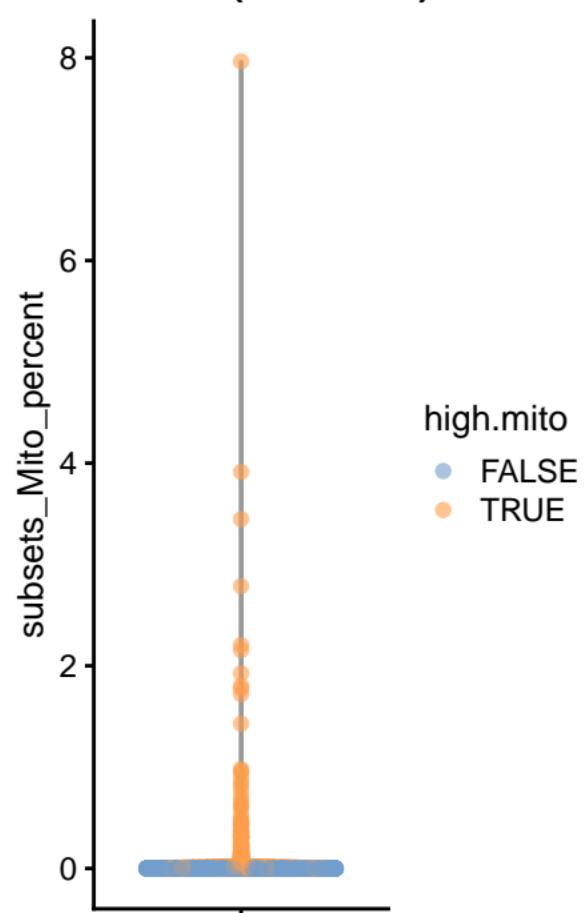
**Total count: br5400.amy.neun****Detected features****Mito % (cutoff = 0.152)**

Sample: br5400.amy.neun; pre-QC nNuclei: 3185; nNuclei kept: 2368 (0.74%)

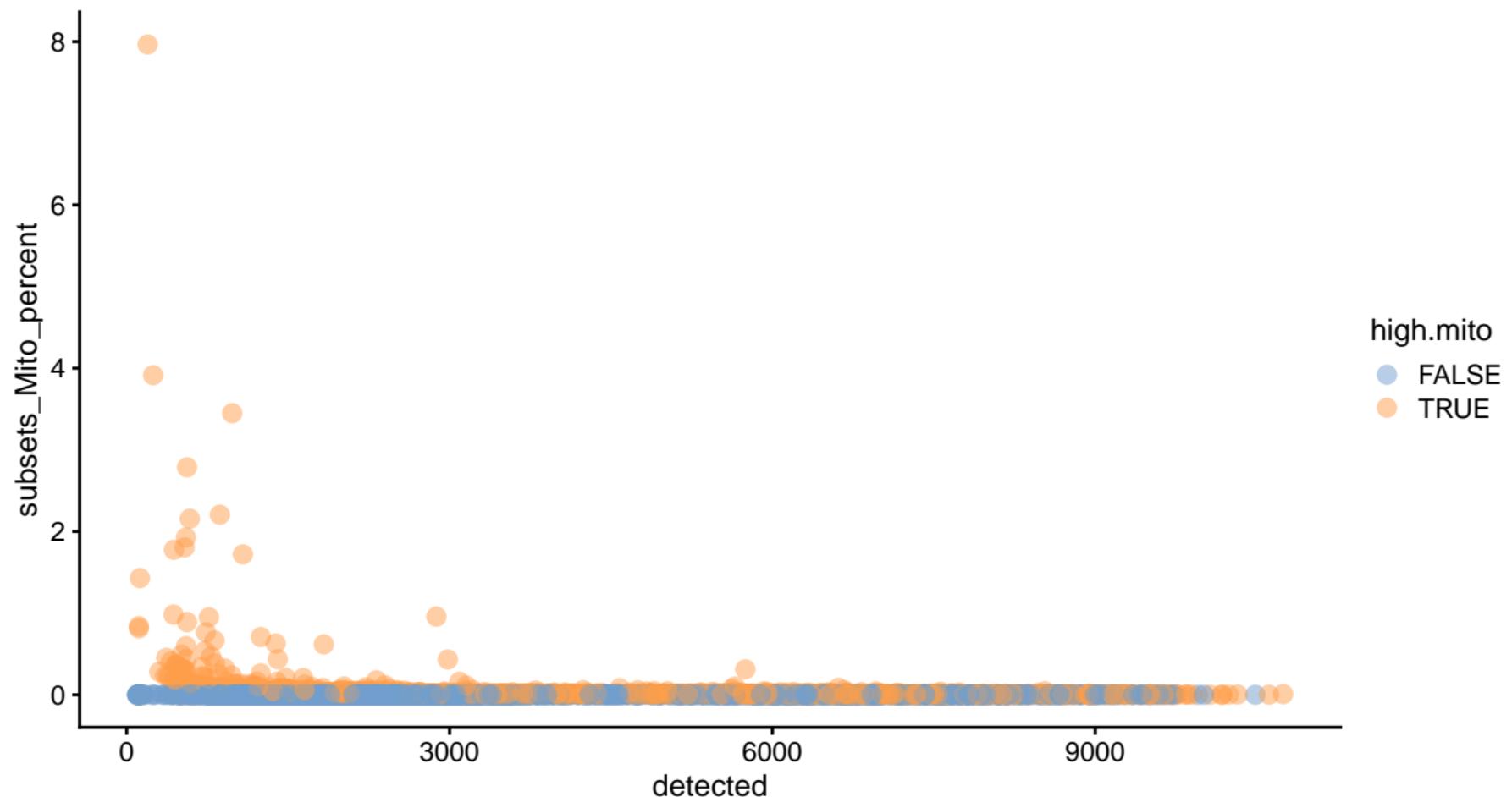


Sample: br5400.amy.neun; pre-QC nNuclei: 3185; nNuclei kept: 2368 (0.74%)

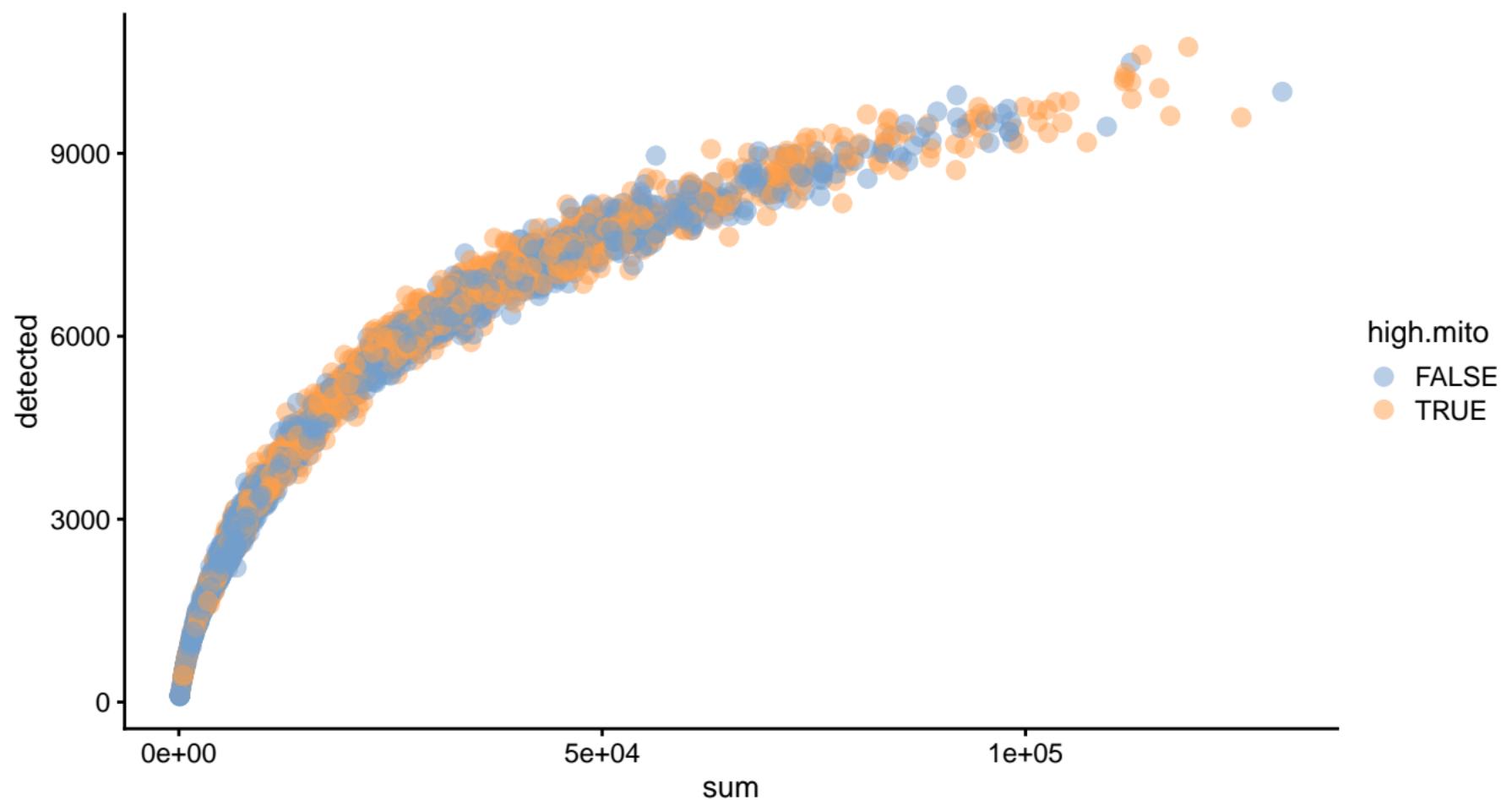


**Total count: br5400.sacc****Detected features****Mito % (cutoff = 0)**

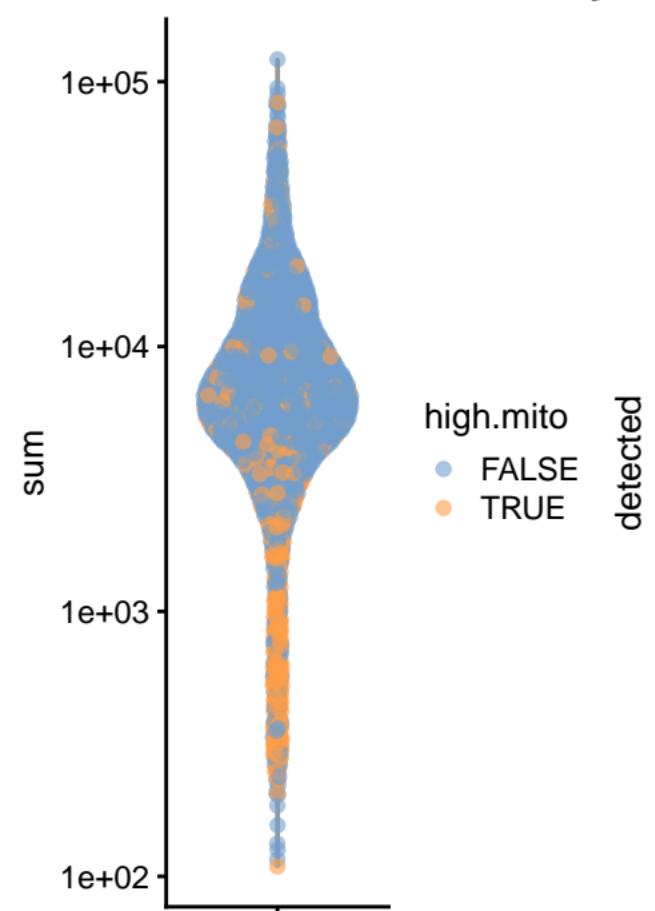
Sample: br5400.sacc; pre-QC nNuclei: 4364; nNuclei kept: 2932 (0.67%)



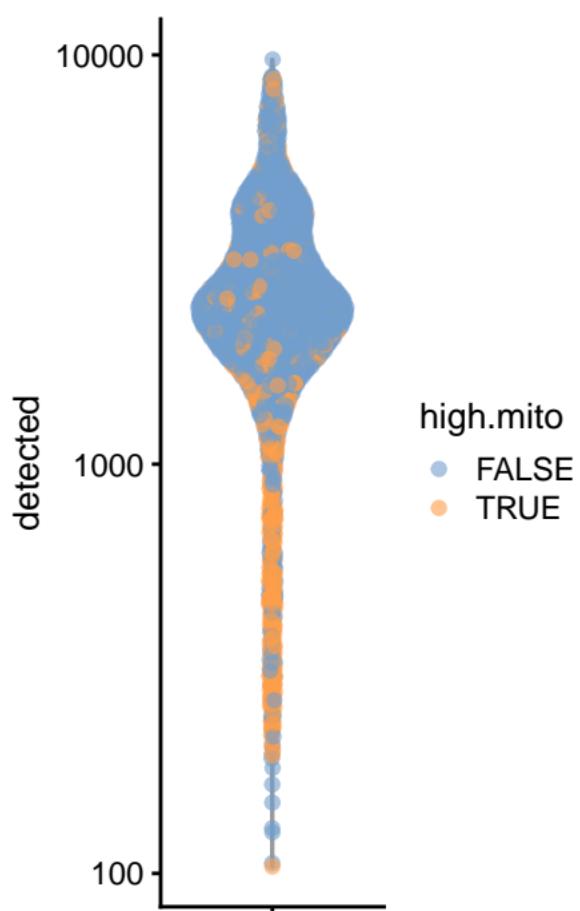
Sample: br5400.sacc; pre-QC nNuclei: 4364; nNuclei kept: 2932 (0.67%)



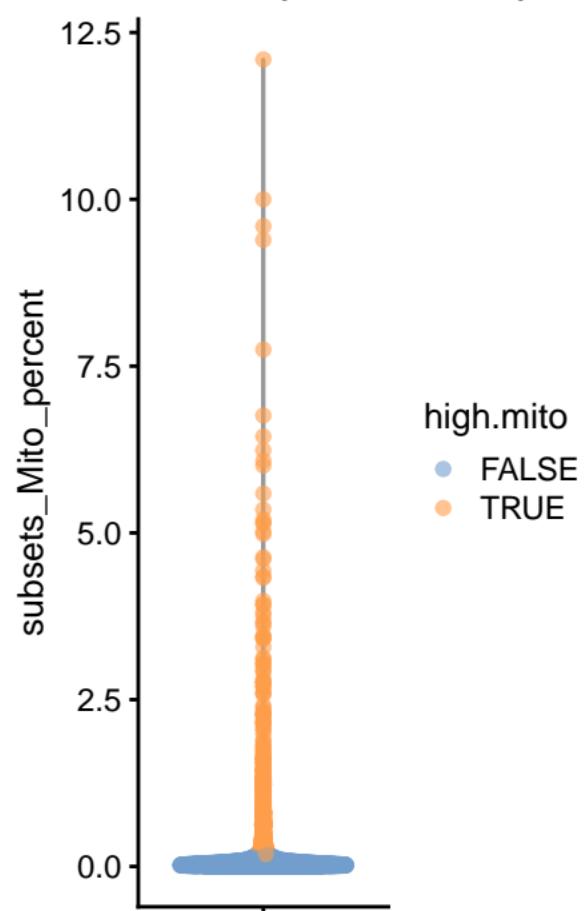
**Total count: br5701.amy**



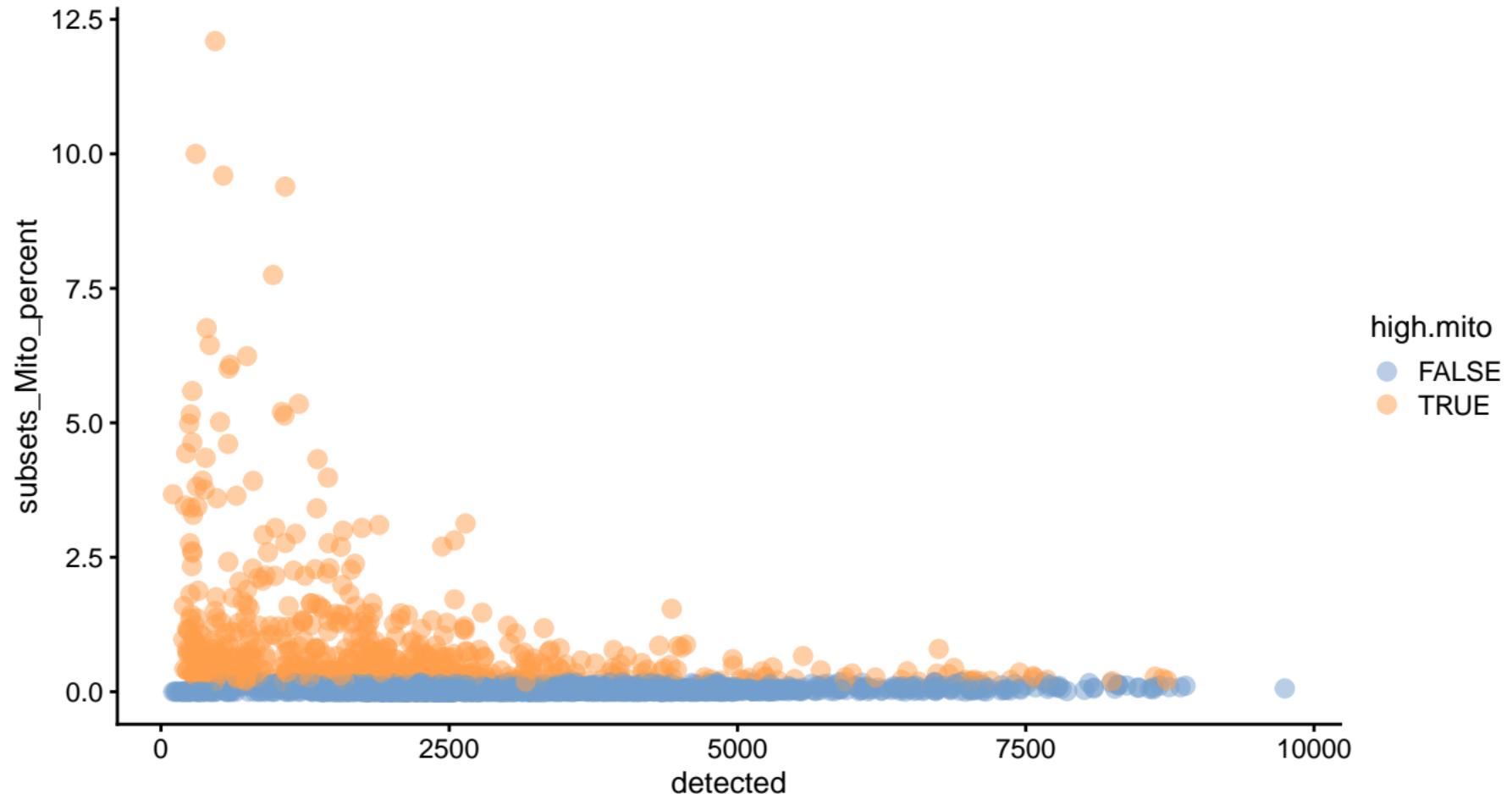
**Detected features**



**Mito % (cutoff = 0.18)**



Sample: br5701.amy; pre-QC nNuclei: 4199; nNuclei kept: 3530 (0.84%)



Sample: br5701.amy; pre-QC nNuclei: 4199; nNuclei kept: 3530 (0.84%)

