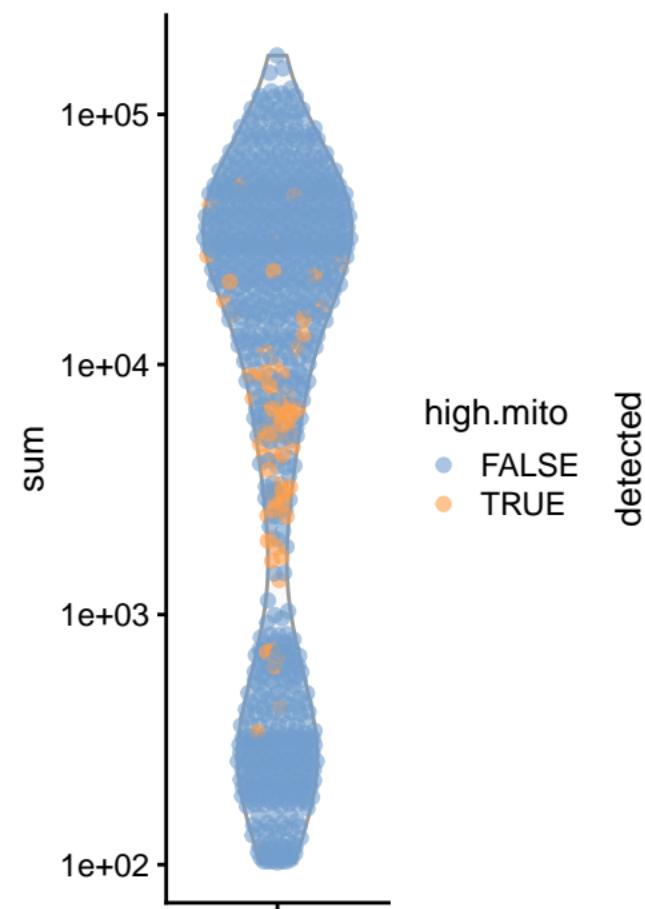
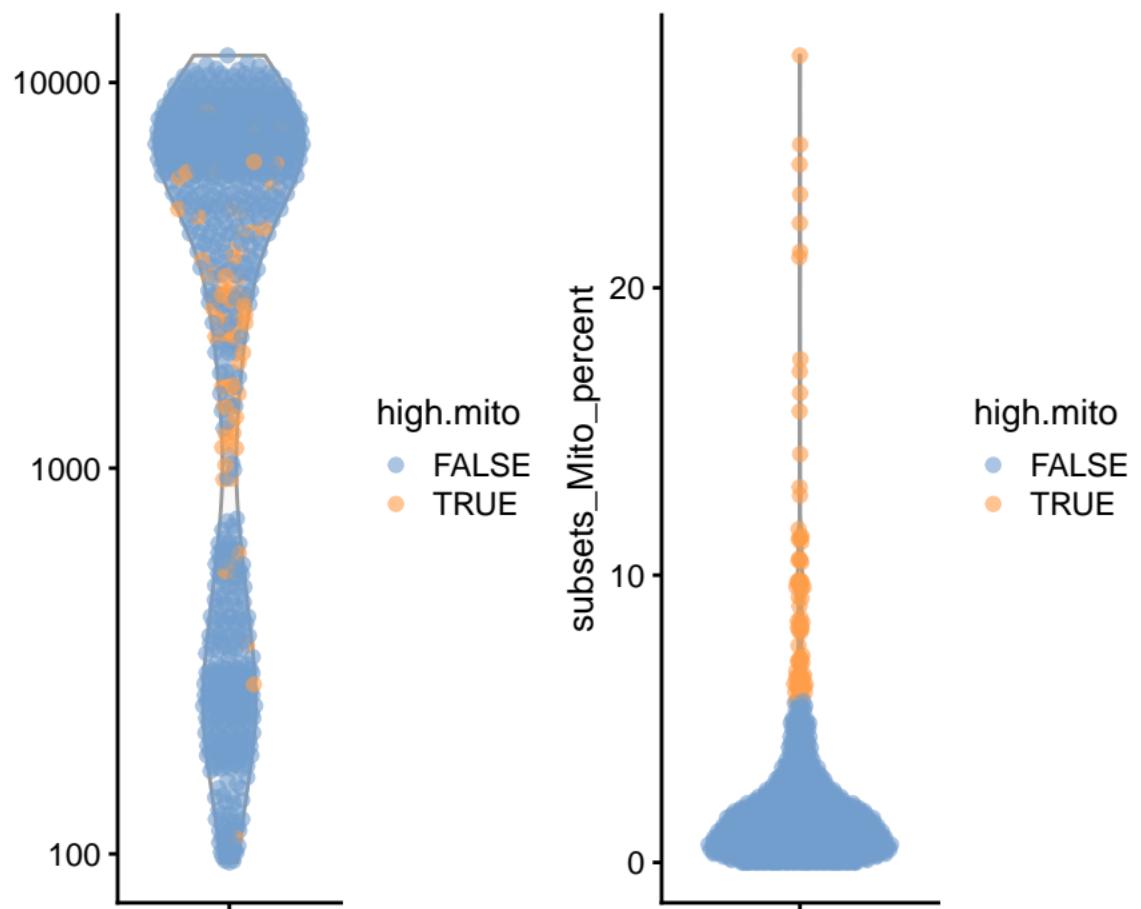


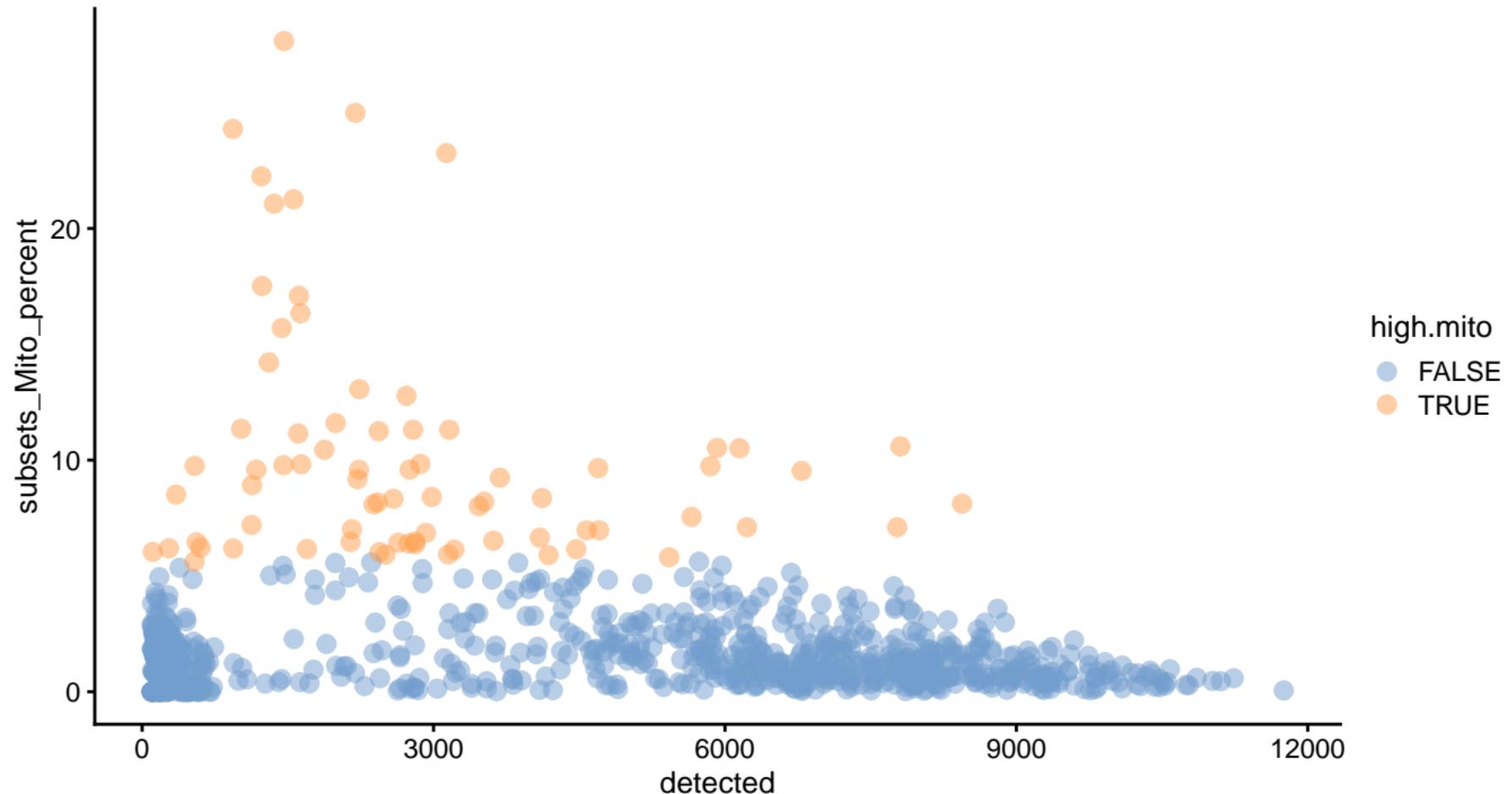
Total count: br5276.sacc.neun Detected features



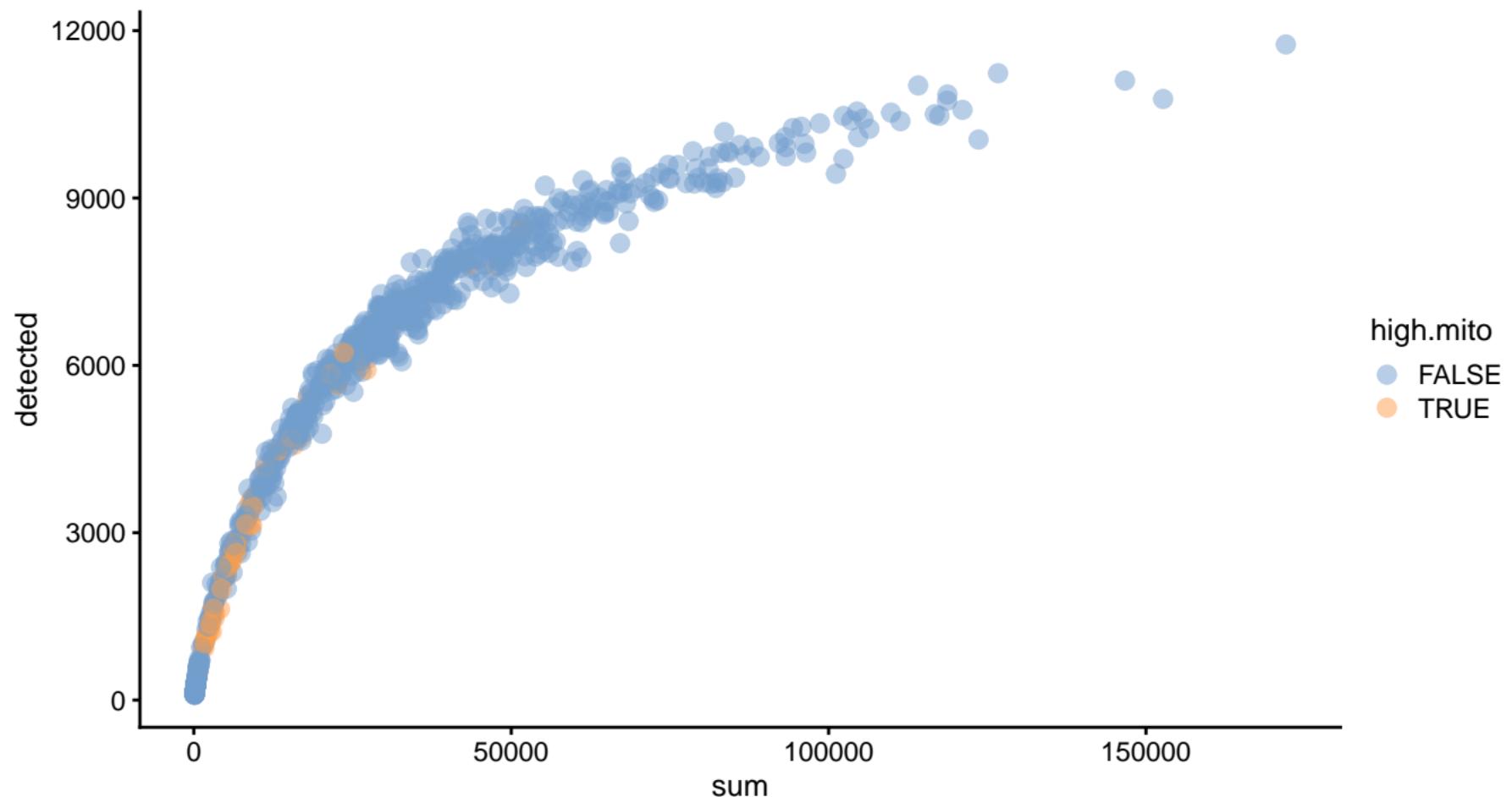
Mito % (cutoff = 5.662)

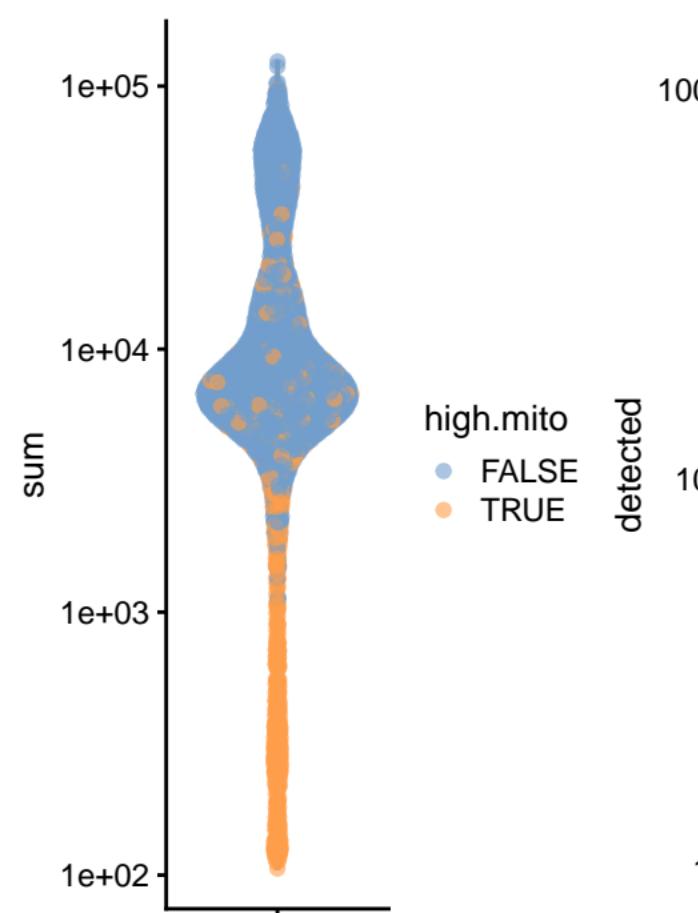
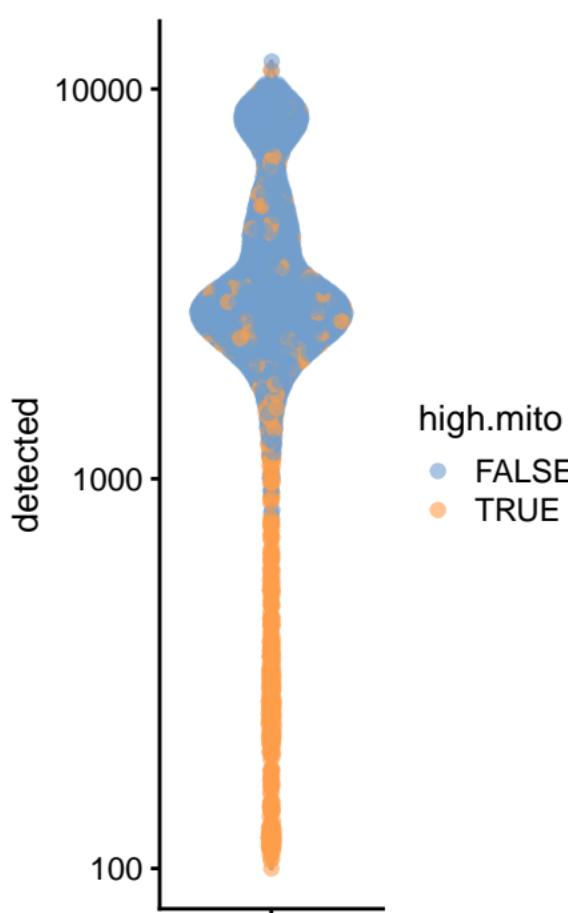
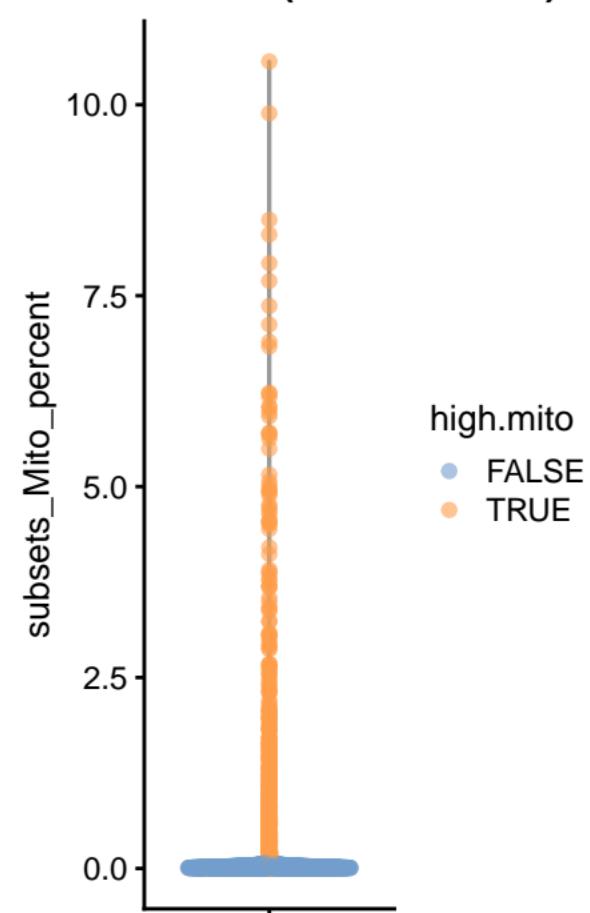


Sample: br5276.sacc.neun; pre-QC nNuclei: 926; nNuclei kept: 851 (0.92%)

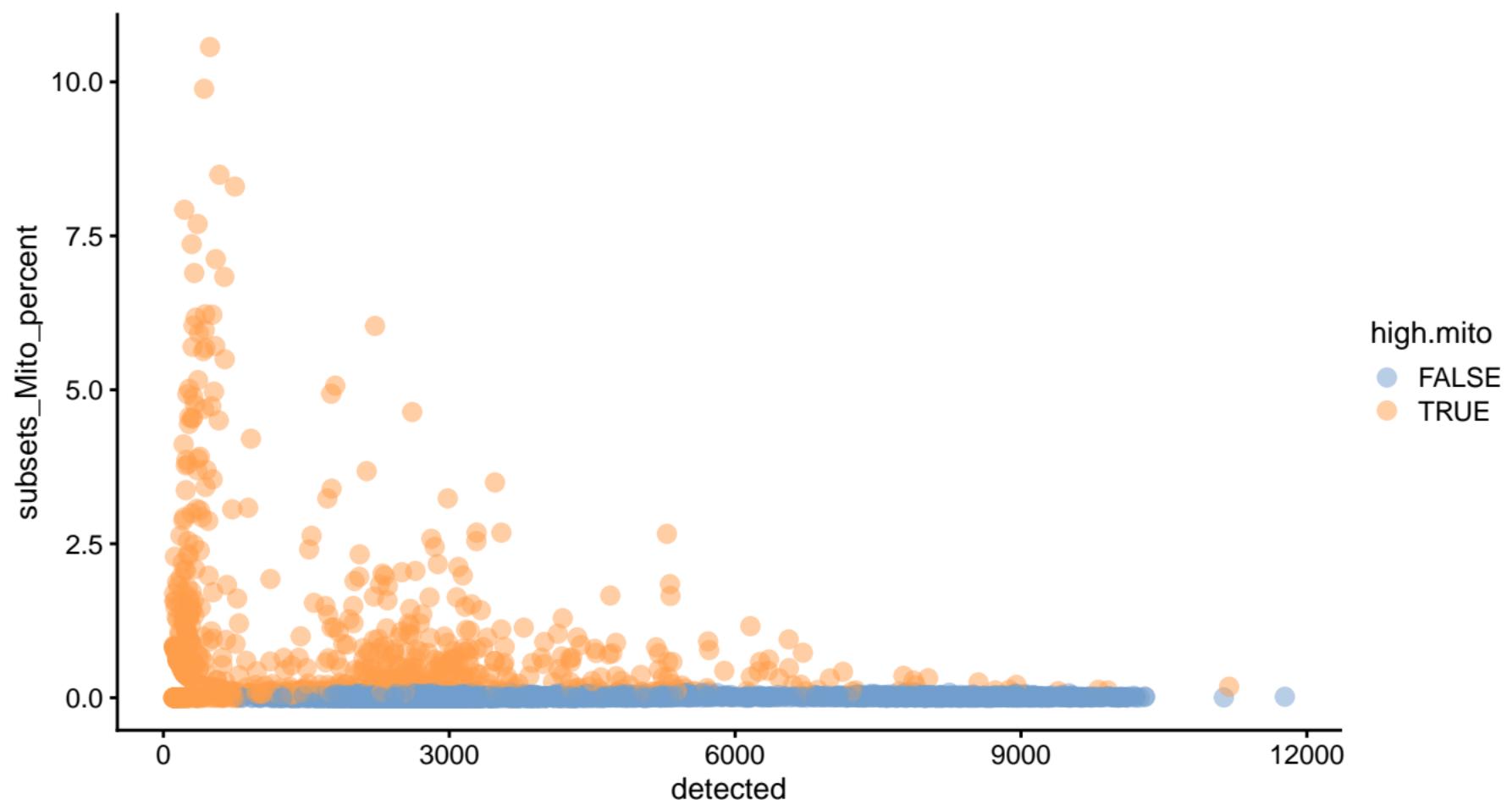


Sample: br5276.sacc.neun; pre-QC nNuclei: 926; nNuclei kept: 851 (0.92%)

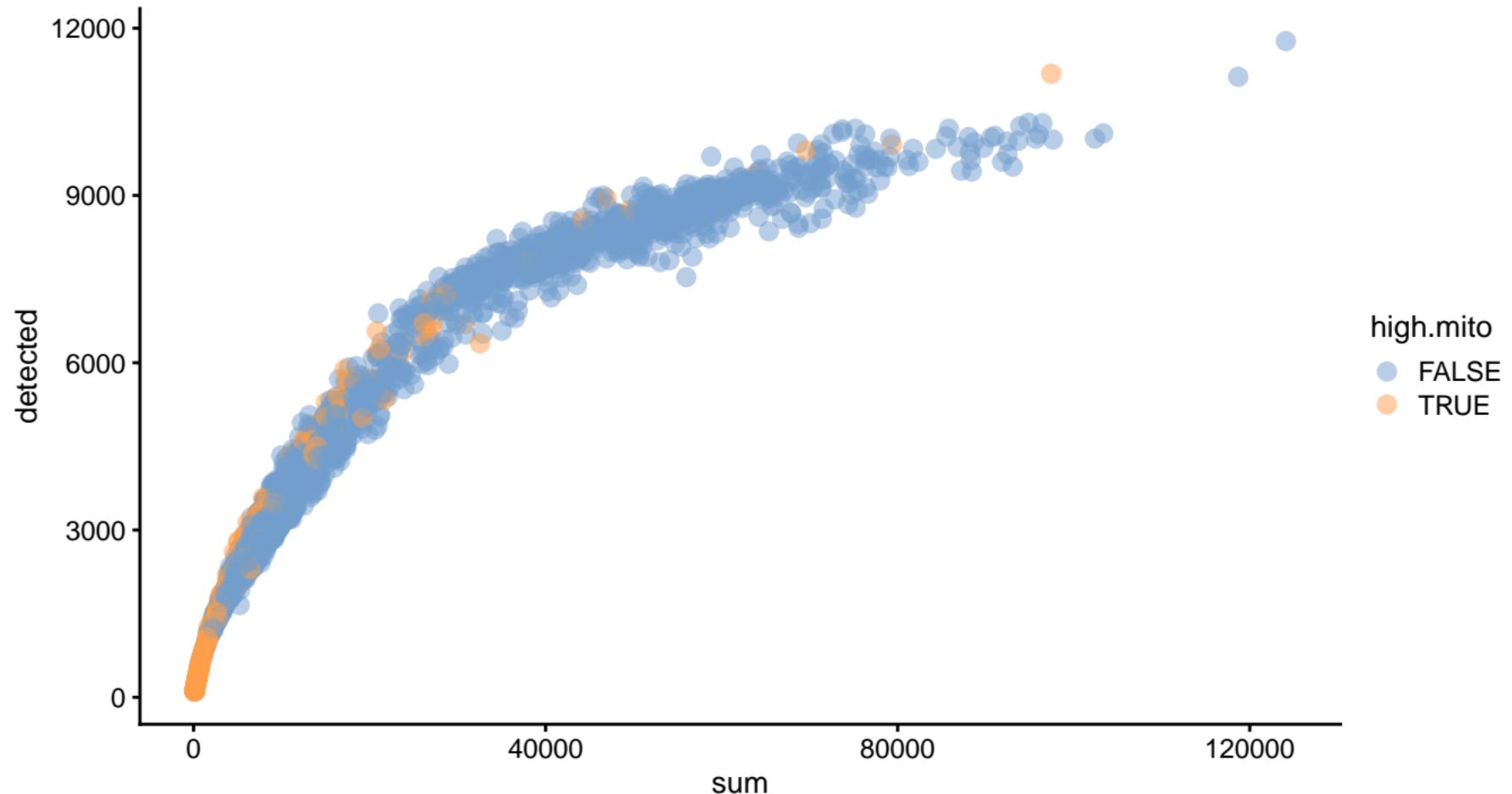


Total count: br5400.nac**Detected features****Mito % (cutoff = 0.092)**

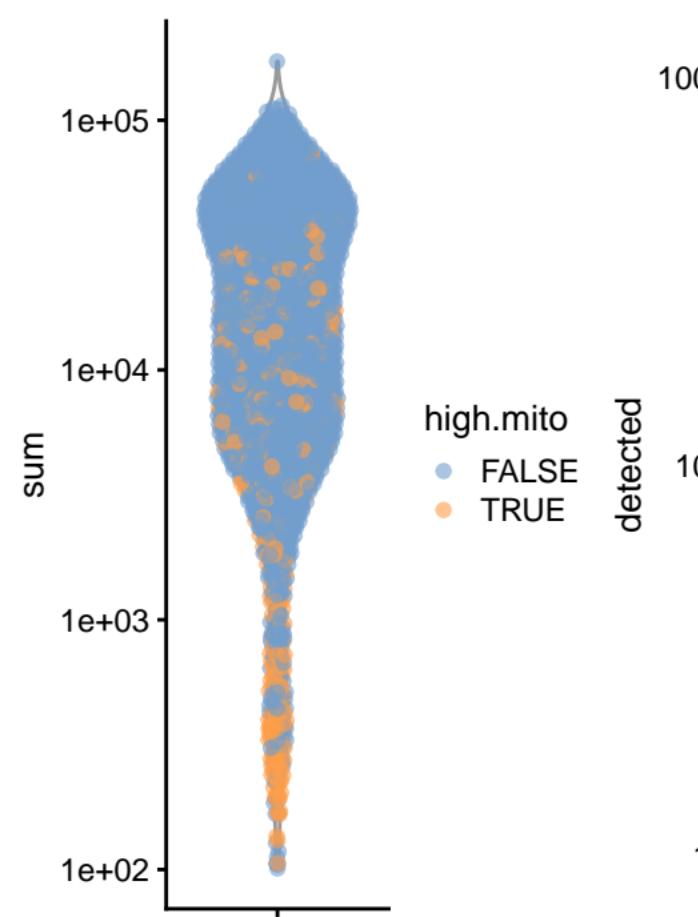
Sample: br5400.nac; pre-QC nNuclei: 4859; nNuclei kept: 4108 (0.85%)



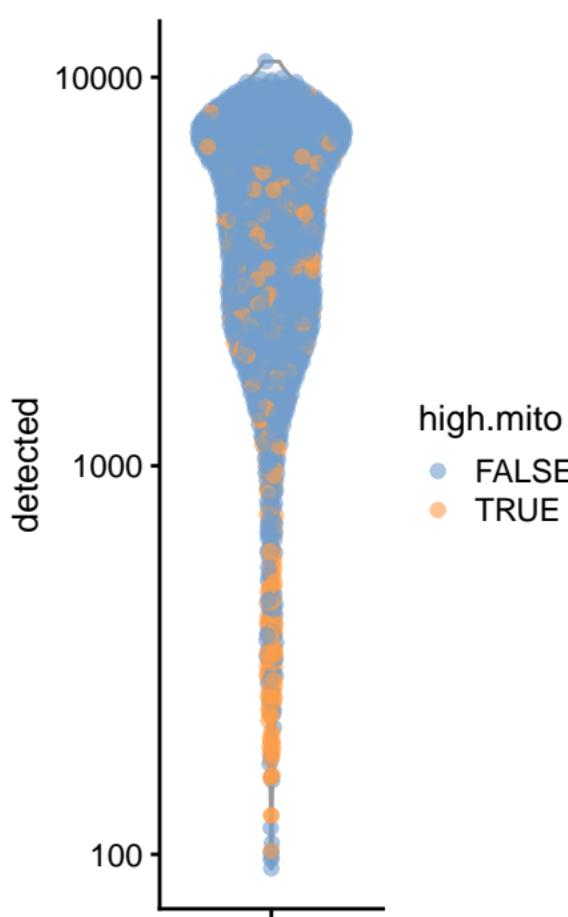
Sample: br5400.nac; pre-QC nNuclei: 4859; nNuclei kept: 4108 (0.85%)



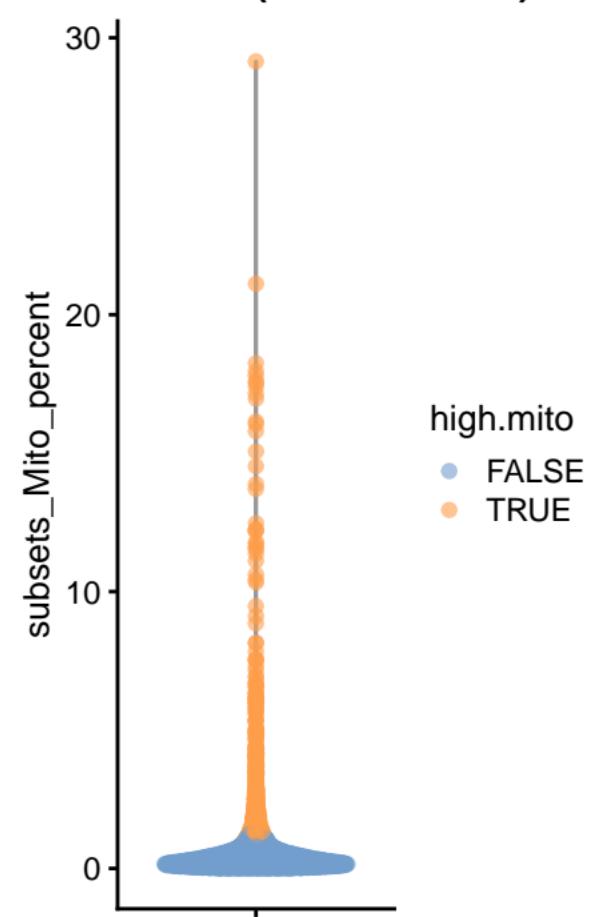
Total count: br5276.nac



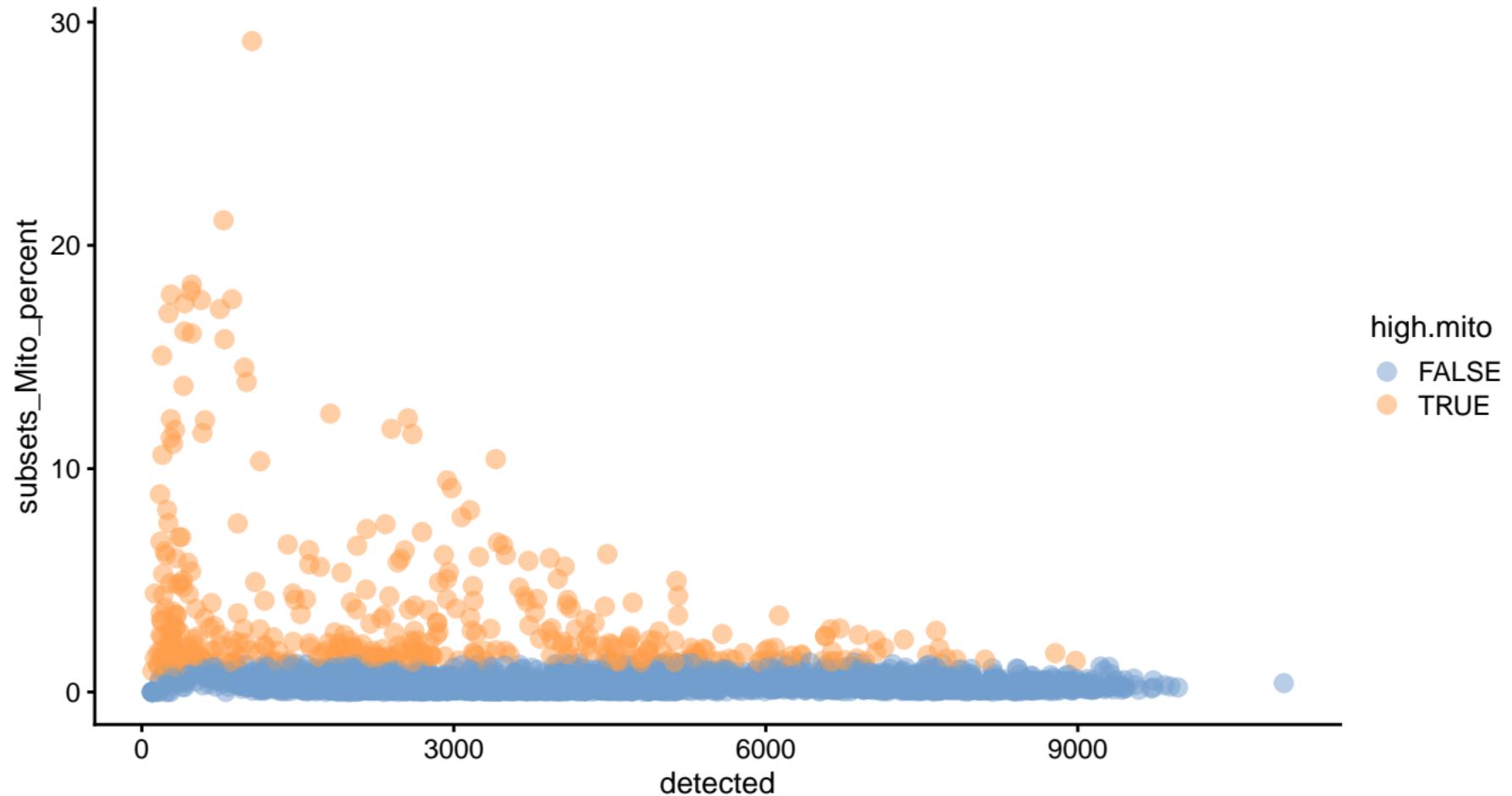
Detected features



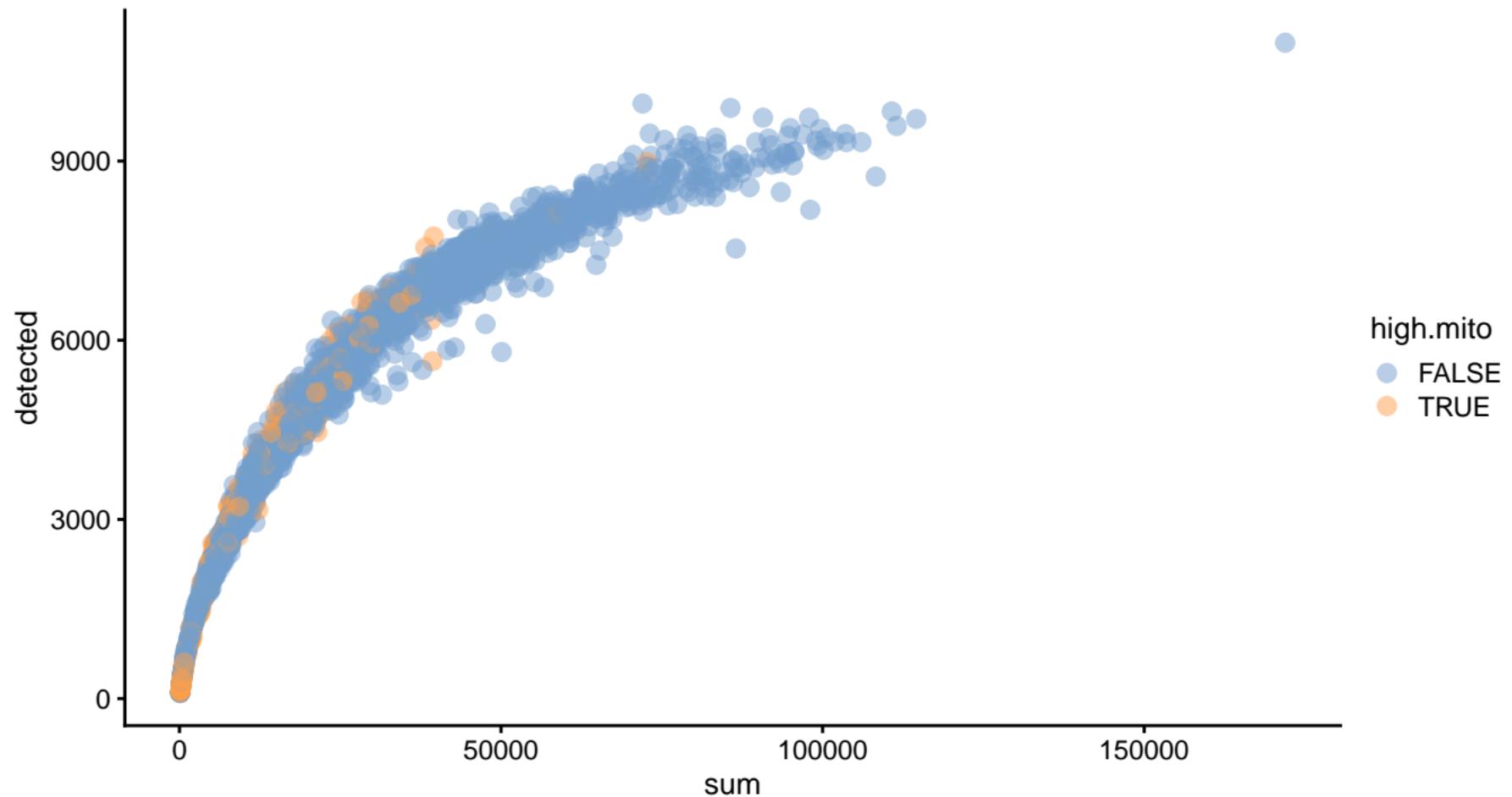
Mito % (cutoff = 1.338)

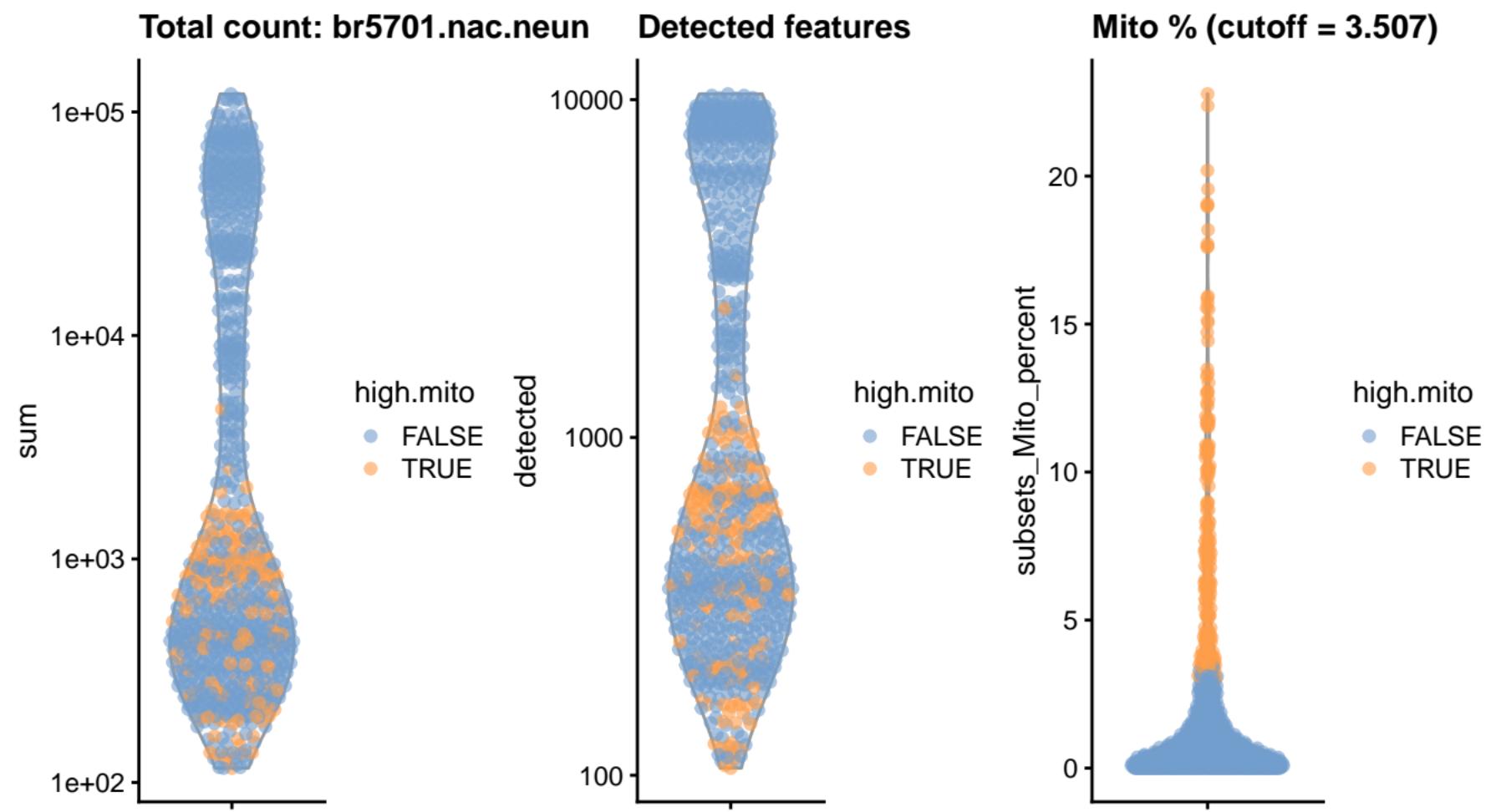


Sample: br5276.nac; pre-QC nNuclei: 3012; nNuclei kept: 2626 (0.87%)

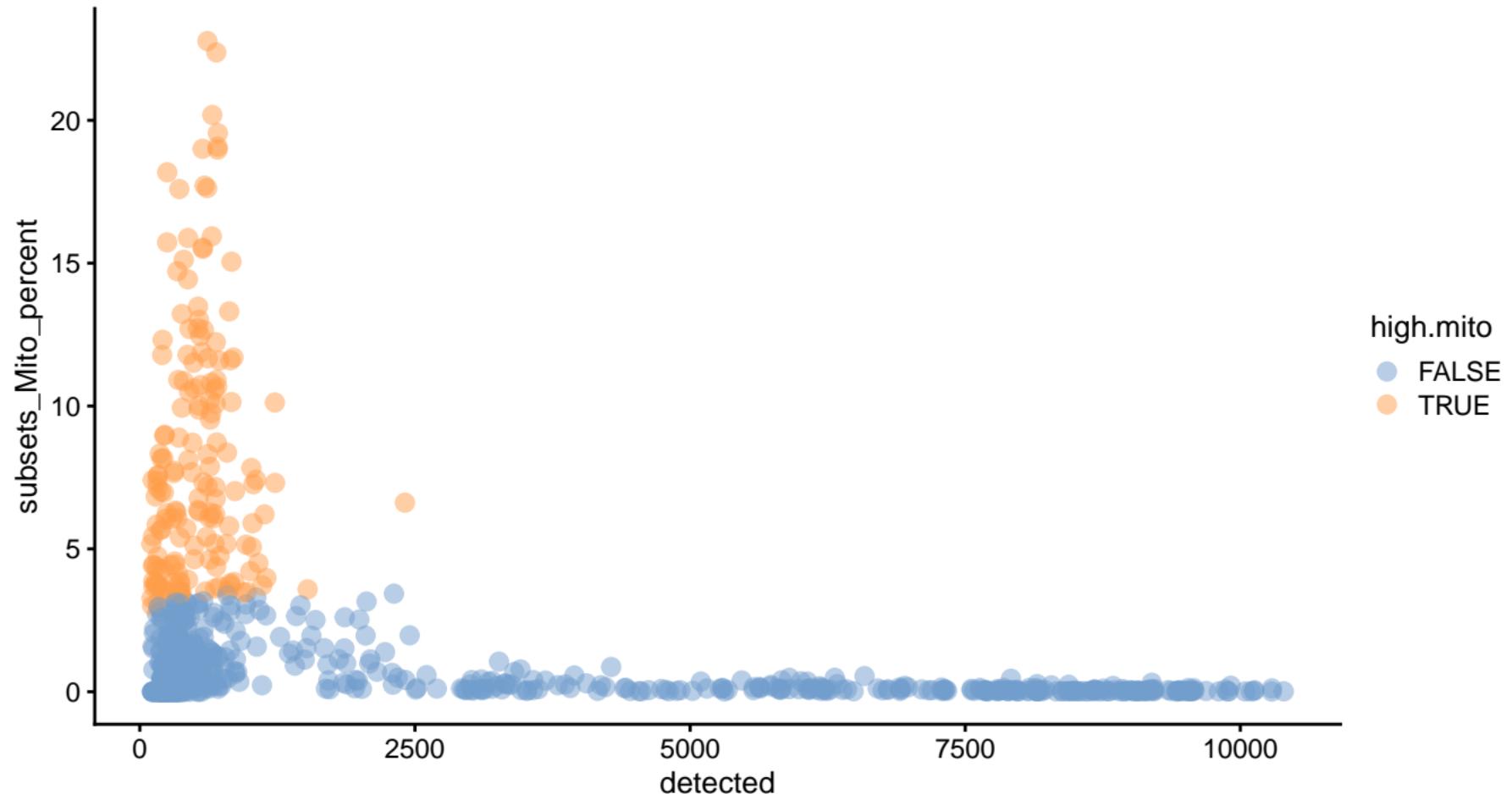


Sample: br5276.nac; pre-QC nNuclei: 3012; nNuclei kept: 2626 (0.87%)

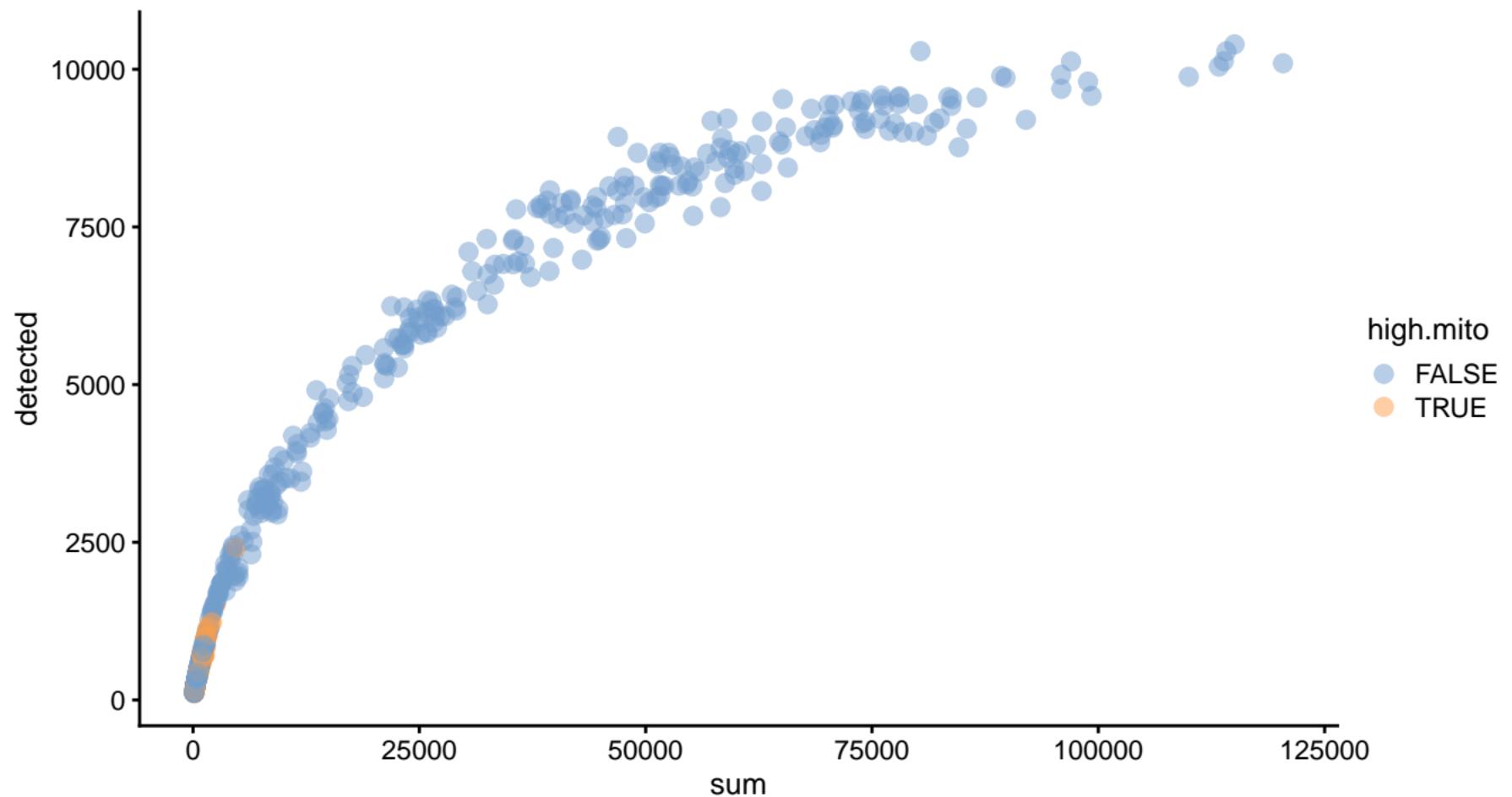


Total count: br5701.nac.neun**Detected features****Mito % (cutoff = 3.507)**

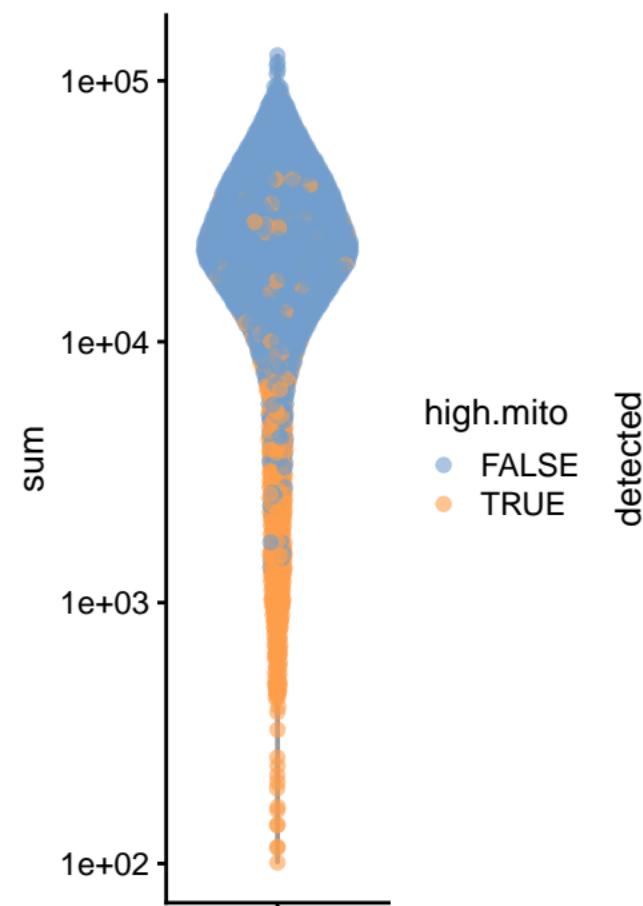
Sample: br5701.nac.neun; pre-QC nNuclei: 814; nNuclei kept: 647 (0.79%)



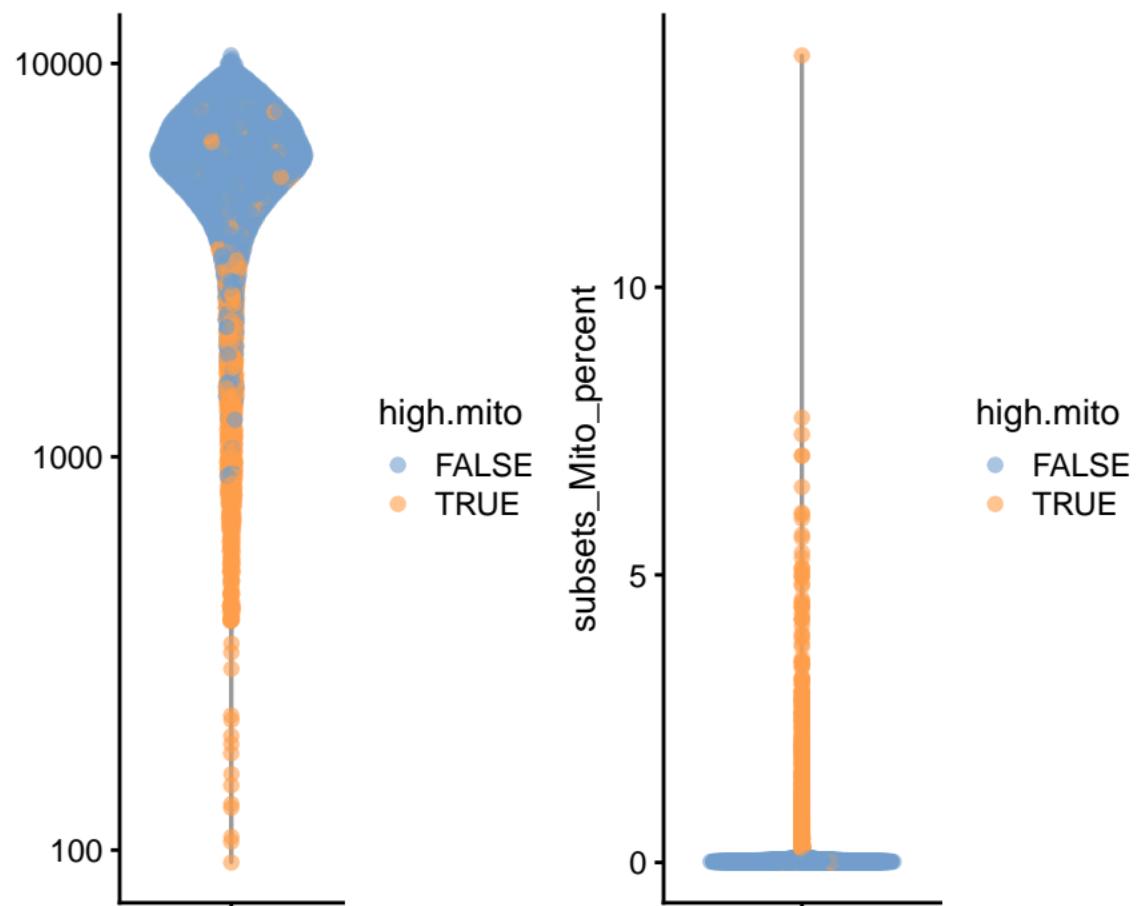
Sample: br5701.nac.neun; pre-QC nNuclei: 814; nNuclei kept: 647 (0.79%)



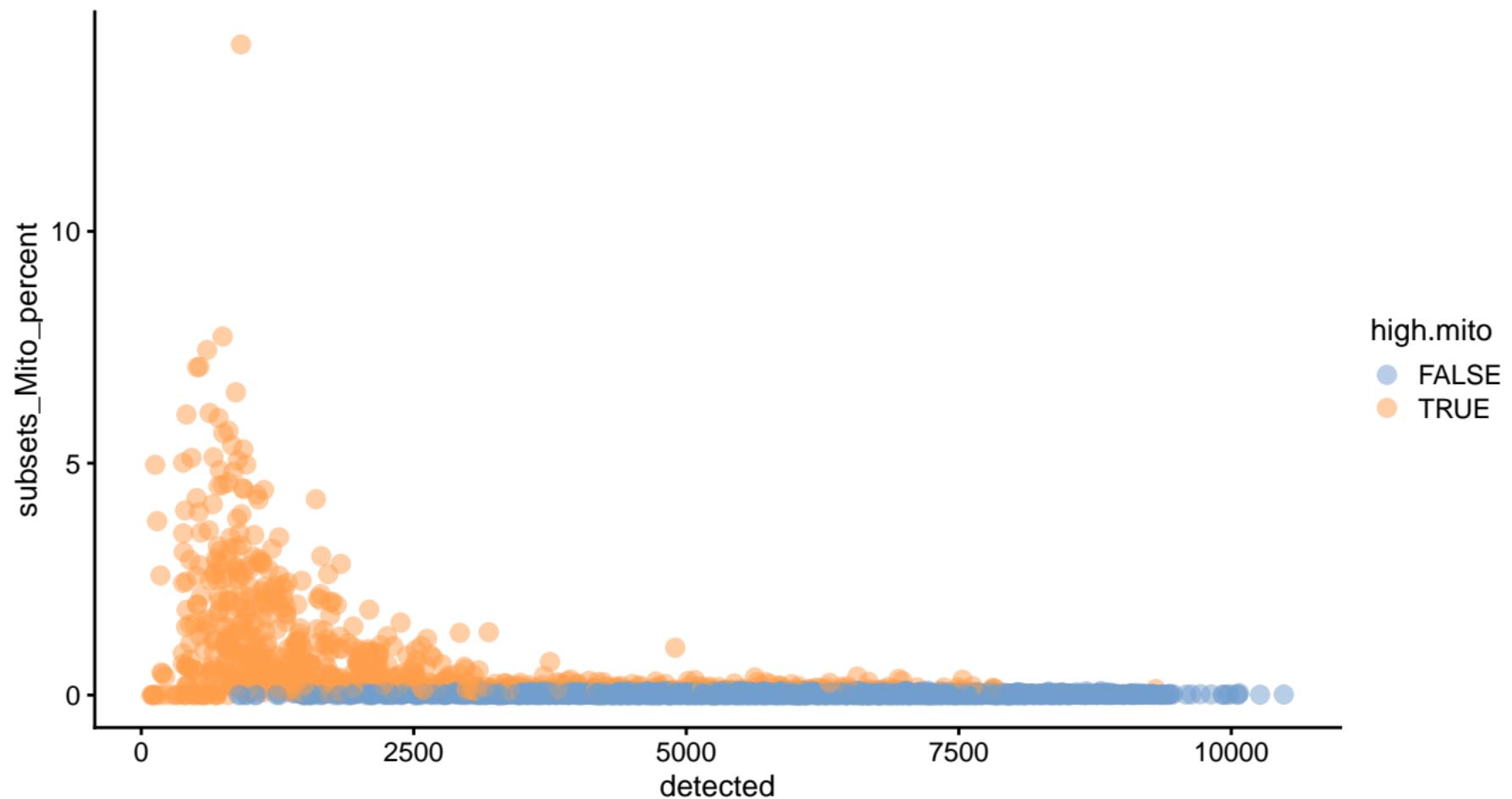
Total count: br5701.sacc.neun Detected features



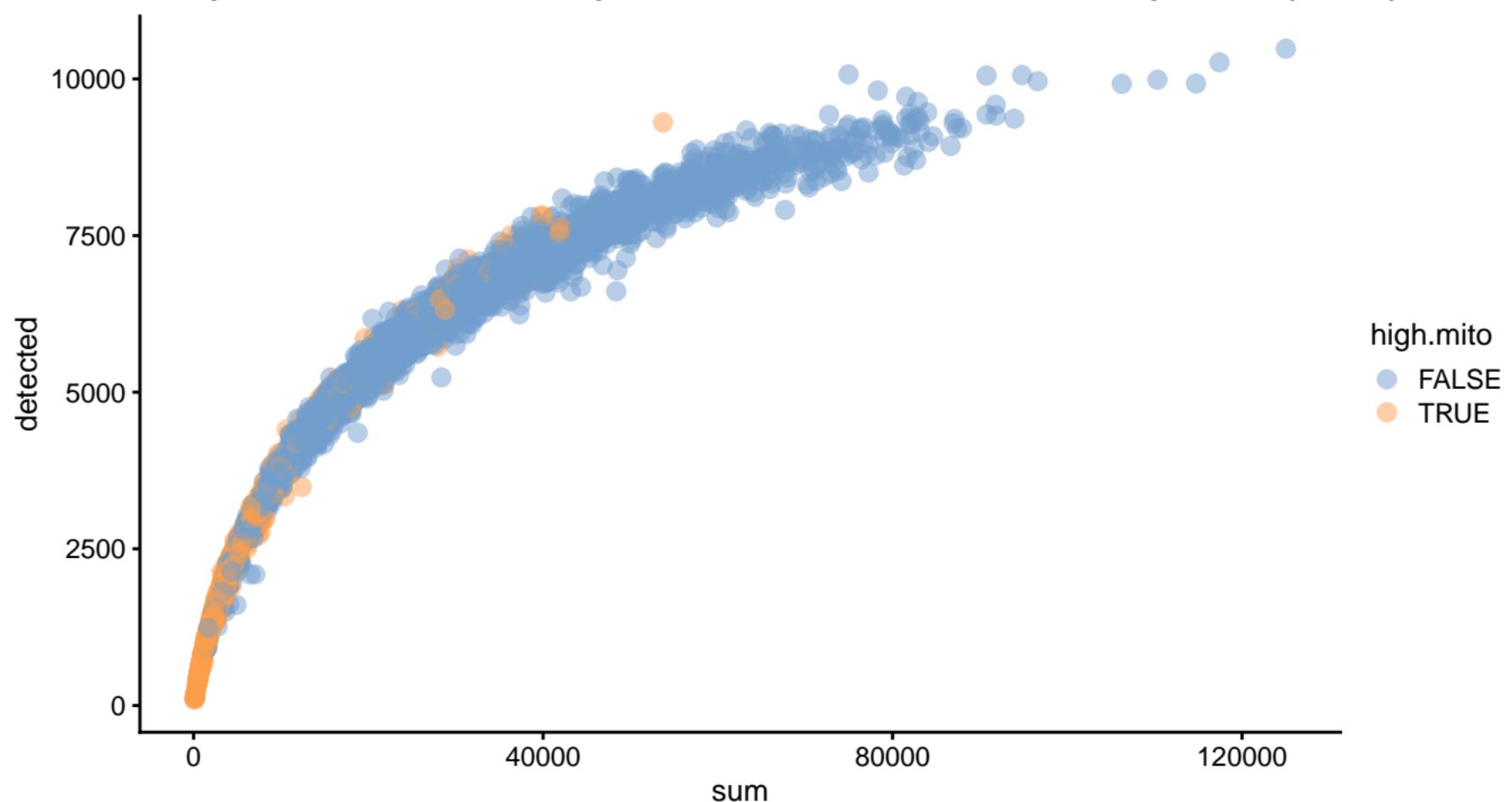
Mito % (cutoff = 0.098)



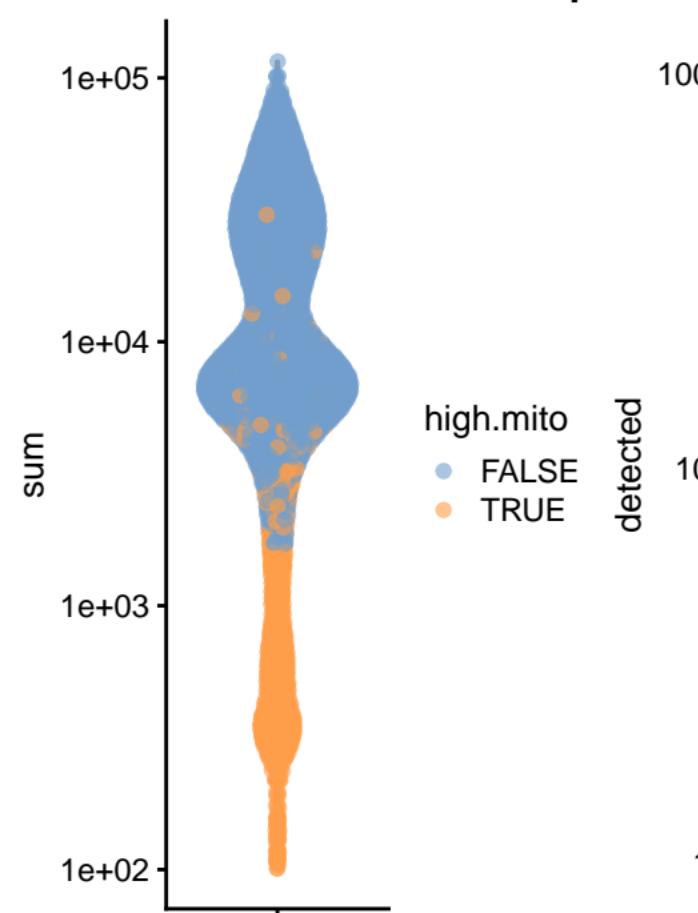
Sample: br5701.sacc.neun; pre-QC nNuclei: 4518; nNuclei kept: 3805 (0.84%)



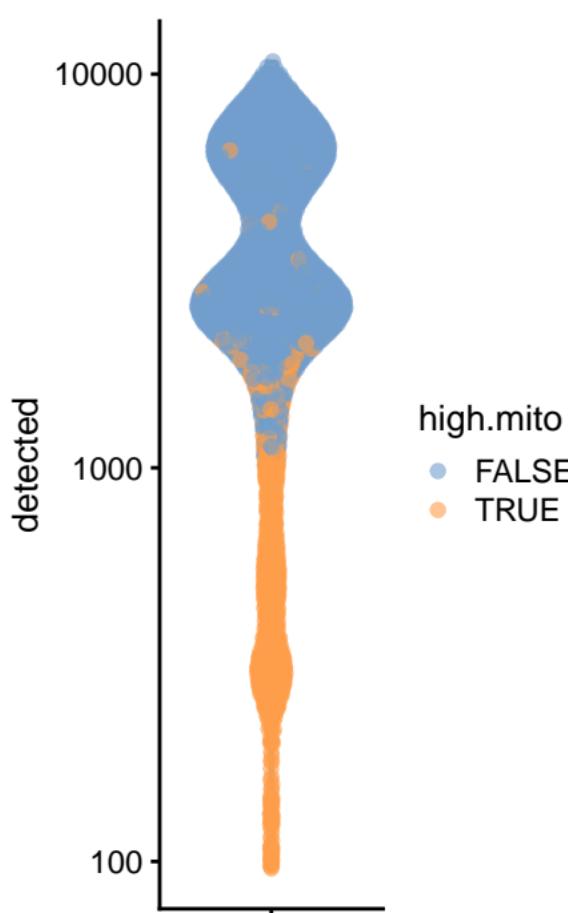
Sample: br5701.sacc.neun; pre-QC nNuclei: 4518; nNuclei kept: 3805 (0.84%)



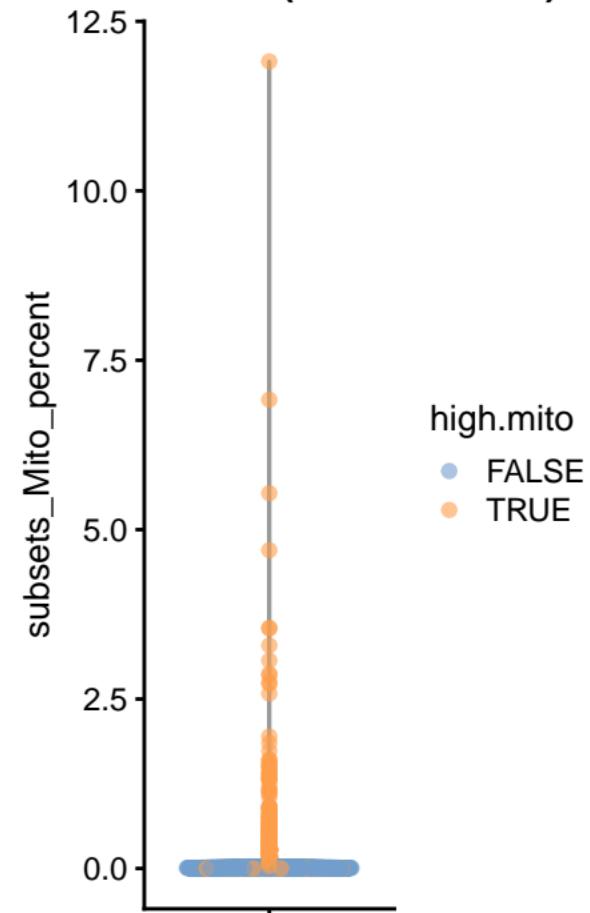
Total count: br5207.dlpfc



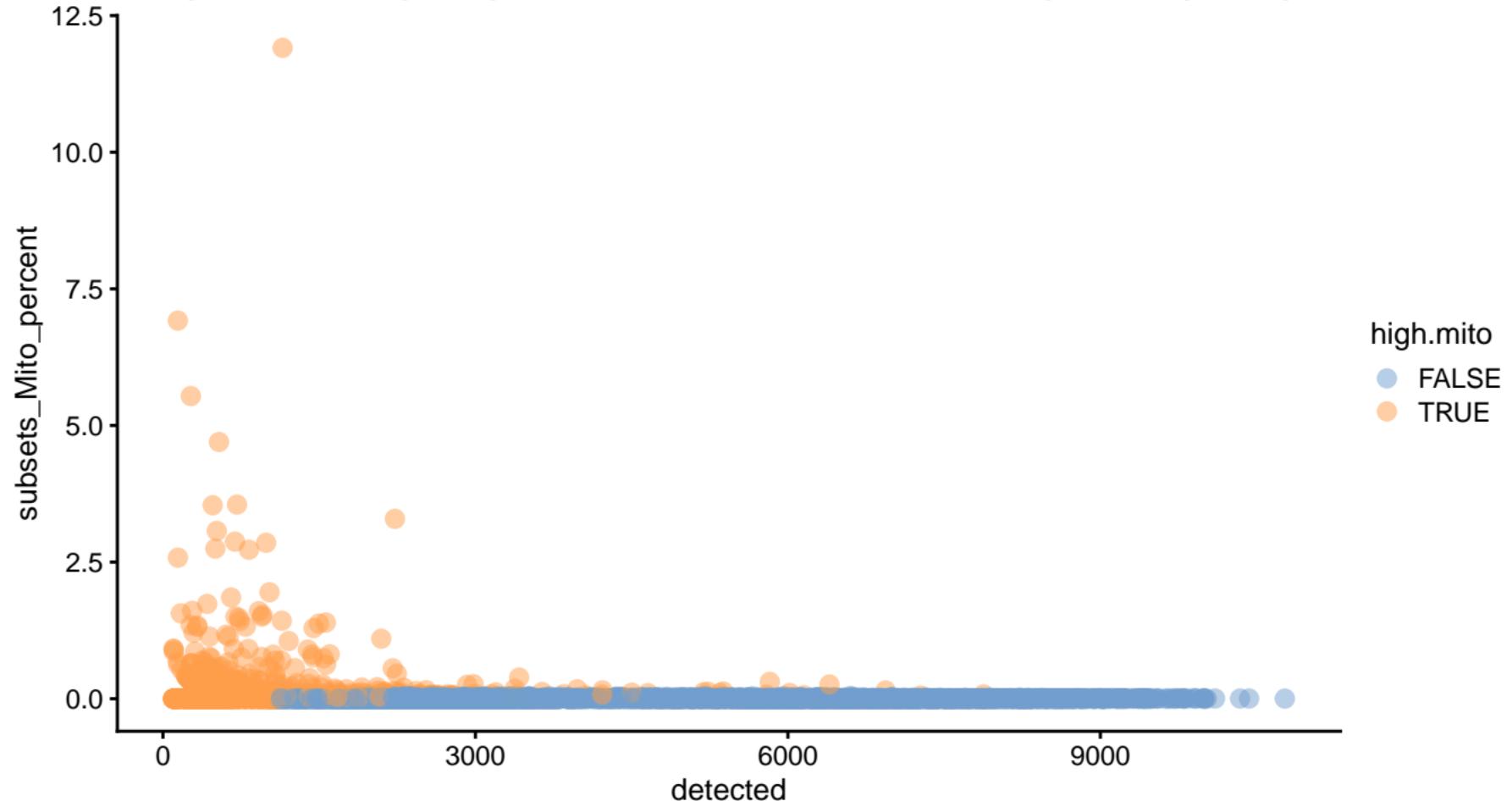
Detected features



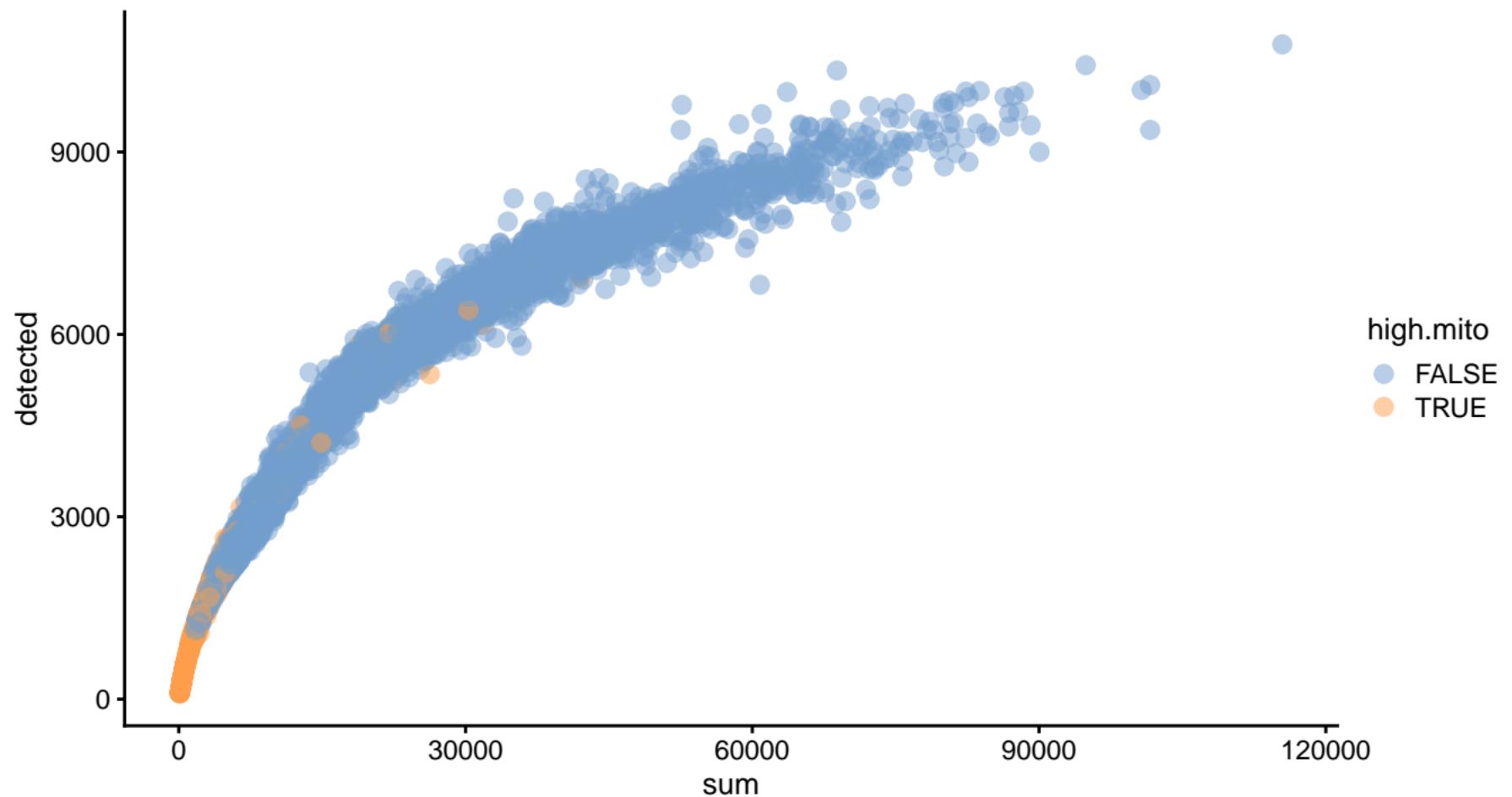
Mito % (cutoff = 0.059)



Sample: br5207.dlpfc; pre-QC nNuclei: 6453; nNuclei kept: 5294 (0.82%)



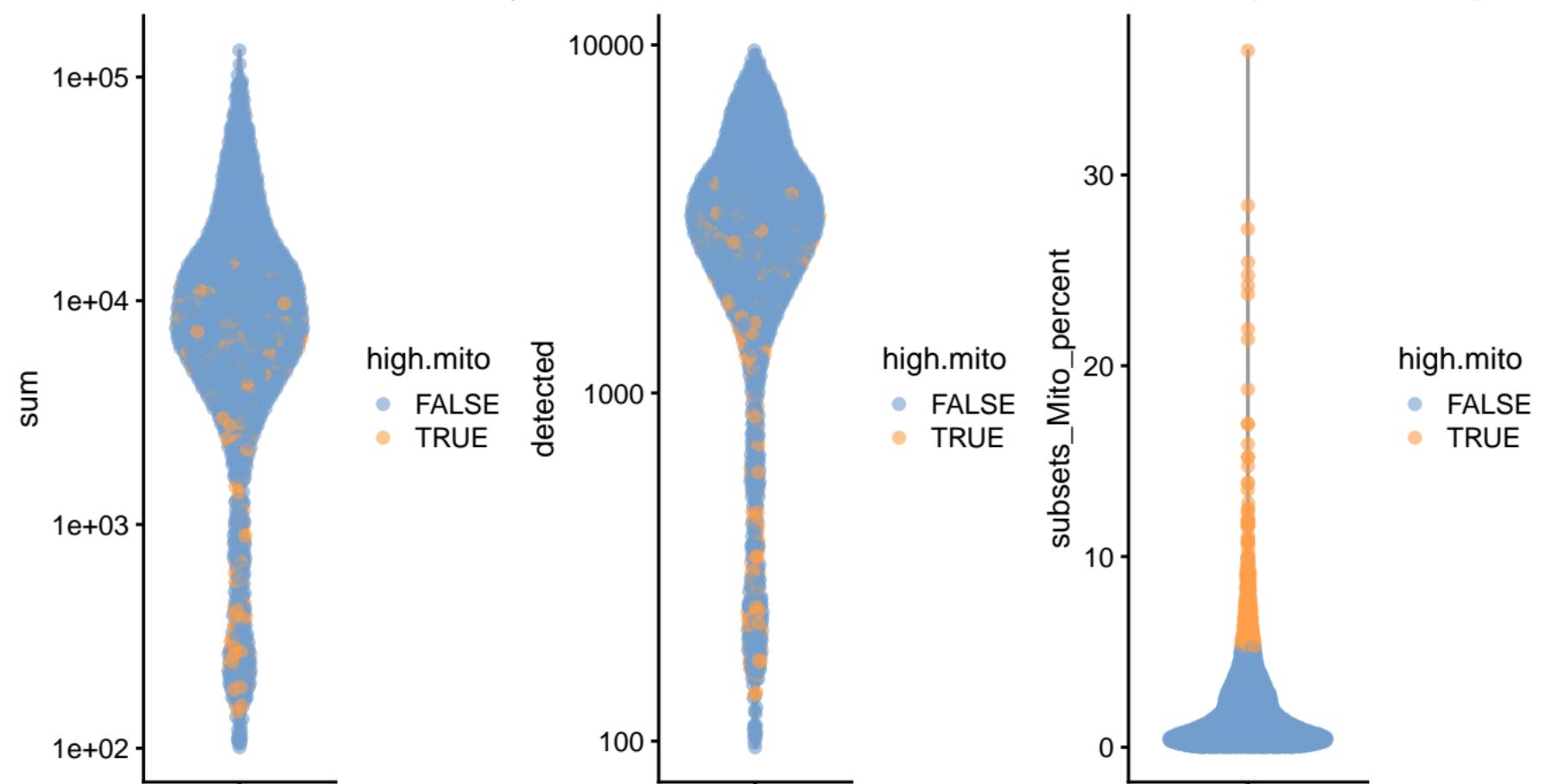
Sample: br5207.dlpfc; pre-QC nNuclei: 6453; nNuclei kept: 5294 (0.82%)



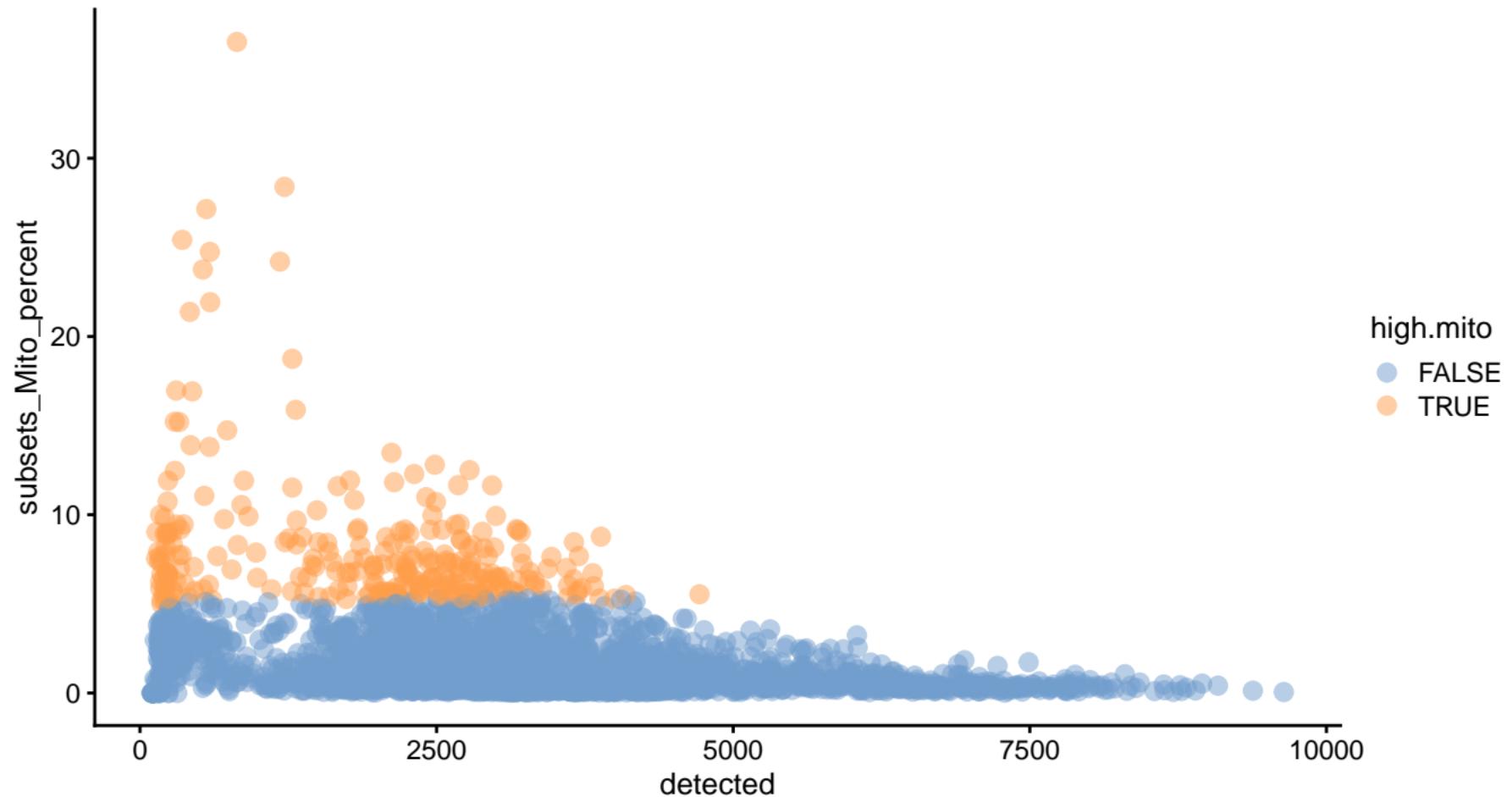
Total count: br5276.amy.neun

Detected features

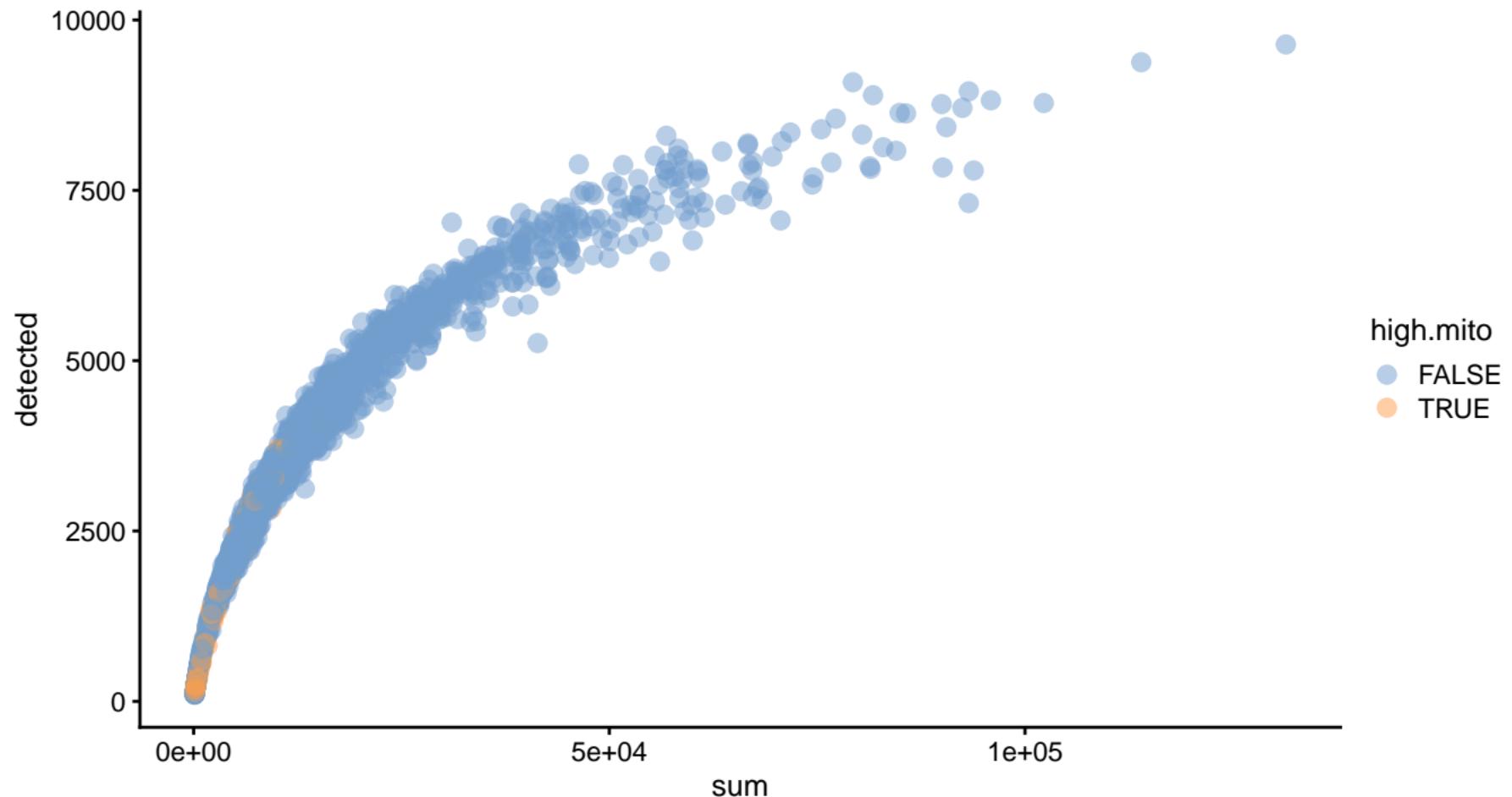
Mito % (cutoff = 5.295)

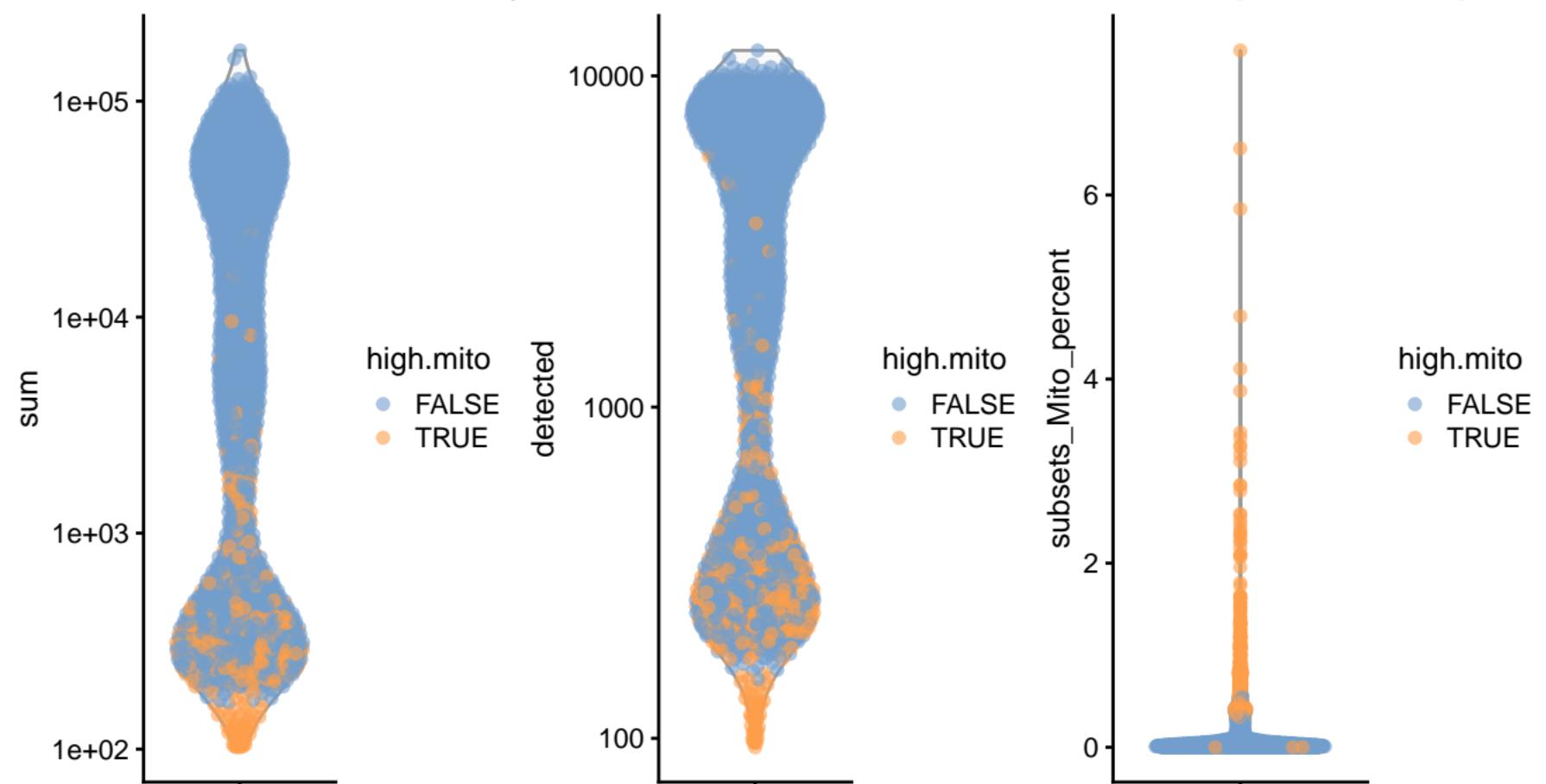


Sample: br5276.amy.neun; pre-QC nNuclei: 2738; nNuclei kept: 2465 (0.9%)

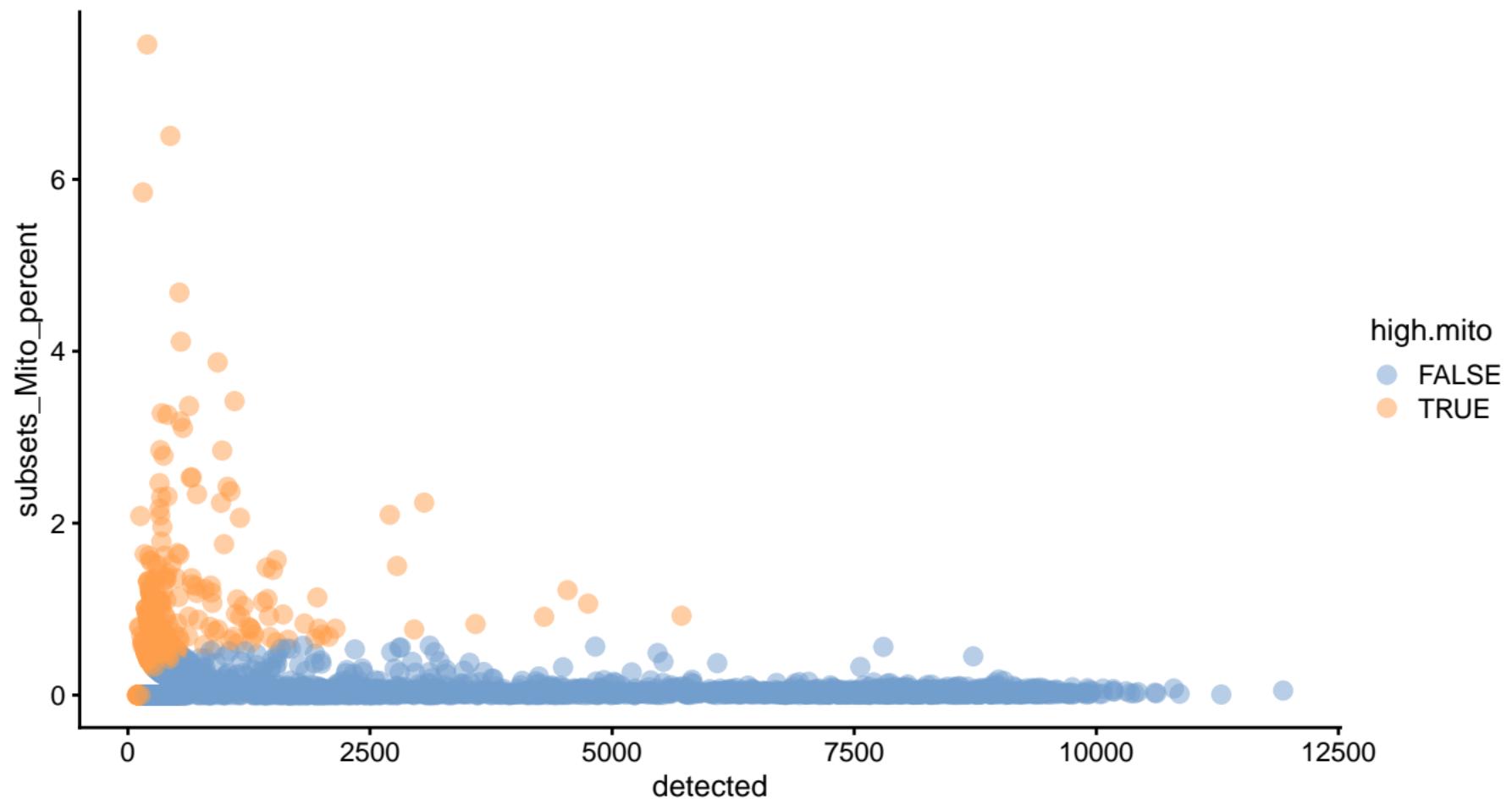


Sample: br5276.amy.neun; pre-QC nNuclei: 2738; nNuclei kept: 2465 (0.9%)

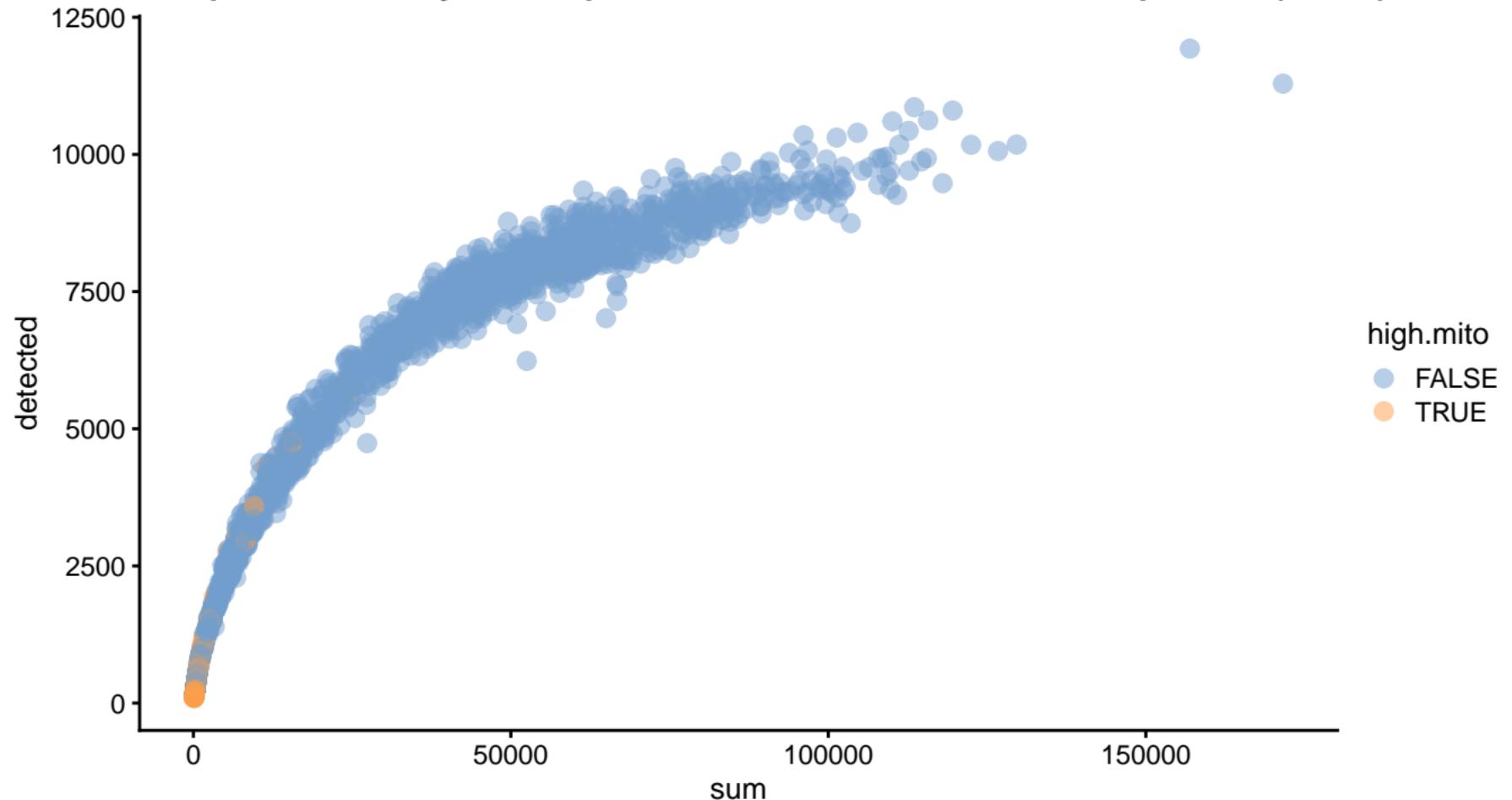


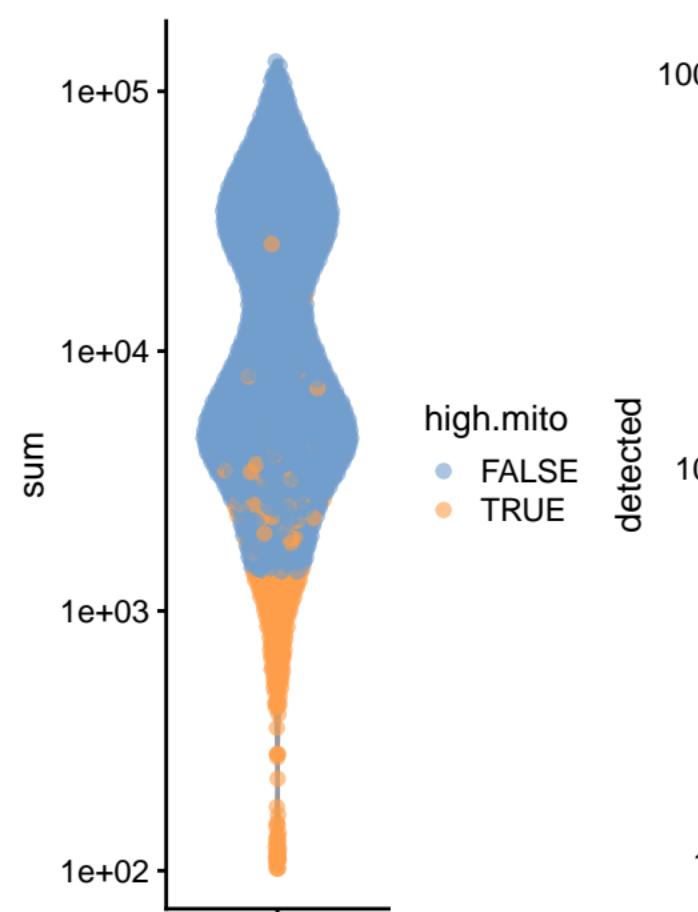
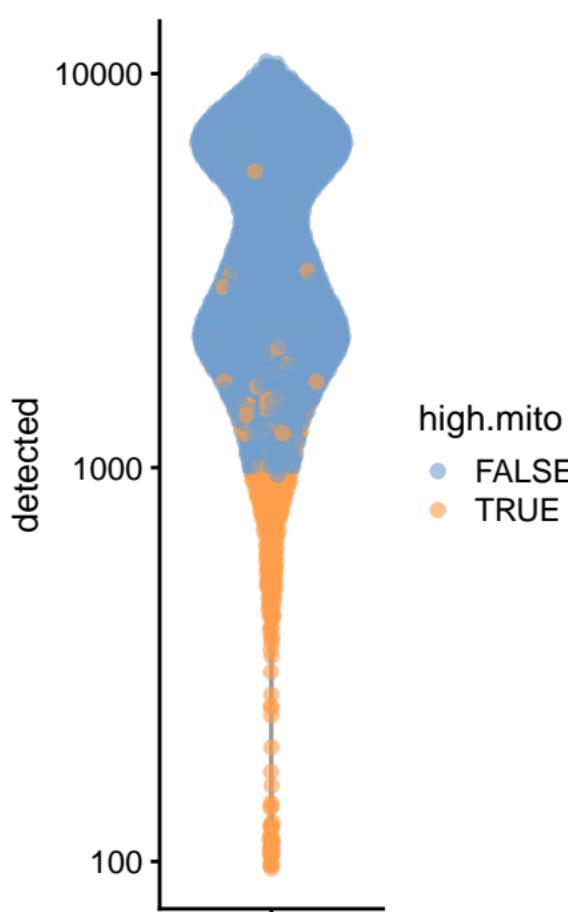
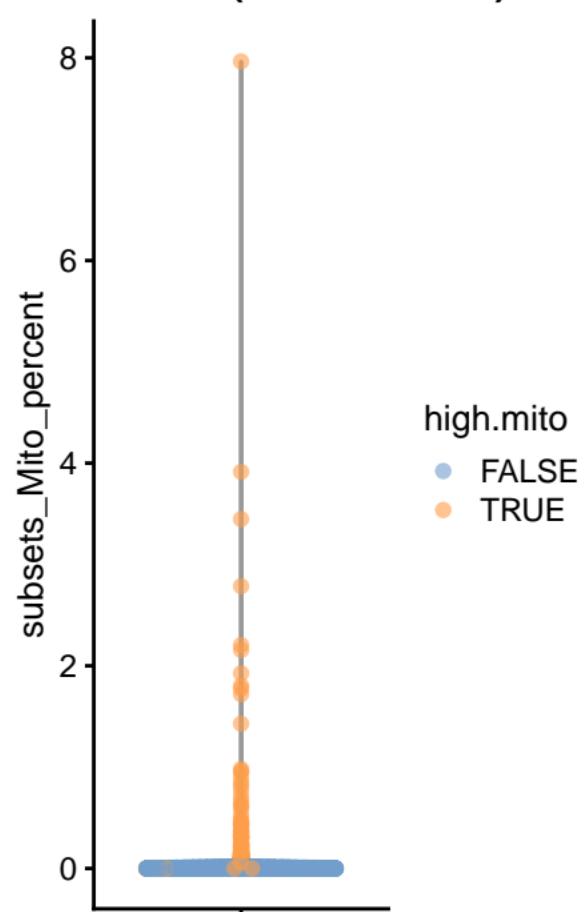
Total count: br5400.amy.neun**Detected features****Mito % (cutoff = 0.619)**

Sample: br5400.amy.neun; pre-QC nNuclei: 3185; nNuclei kept: 2635 (0.83%)

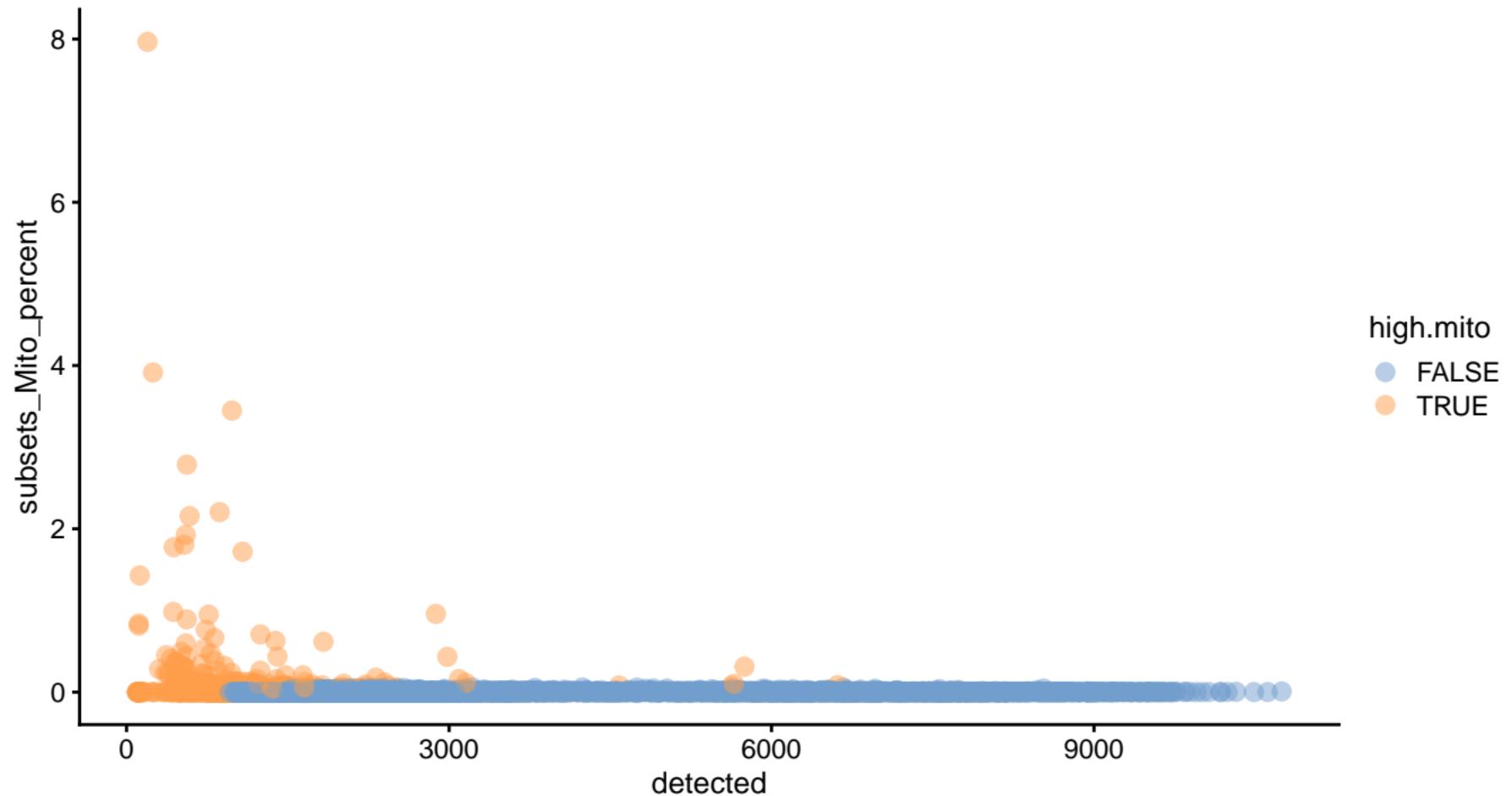


Sample: br5400.amy.neun; pre-QC nNuclei: 3185; nNuclei kept: 2635 (0.83%)

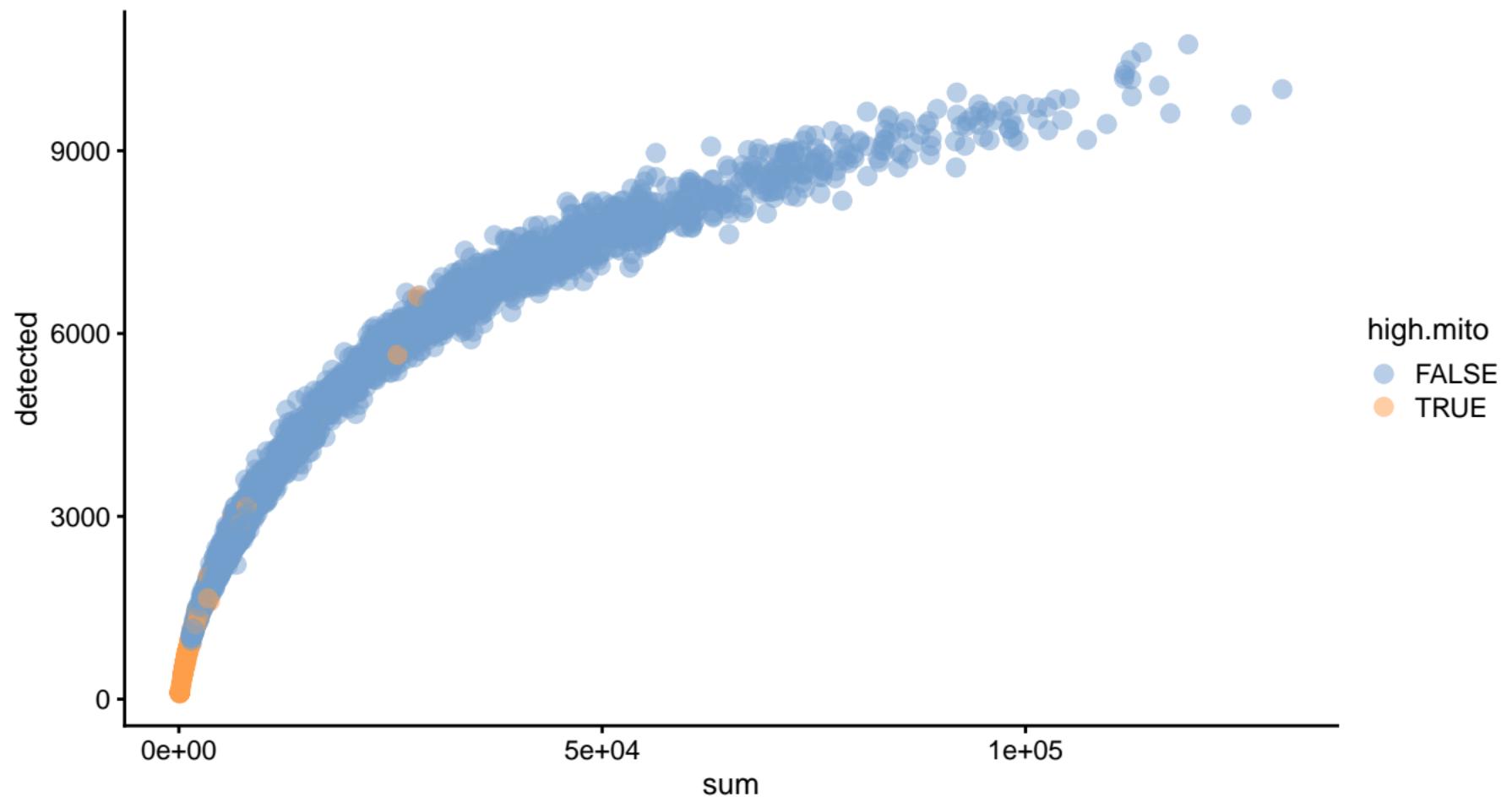


Total count: br5400.sacc**Detected features****Mito % (cutoff = 0.071)**

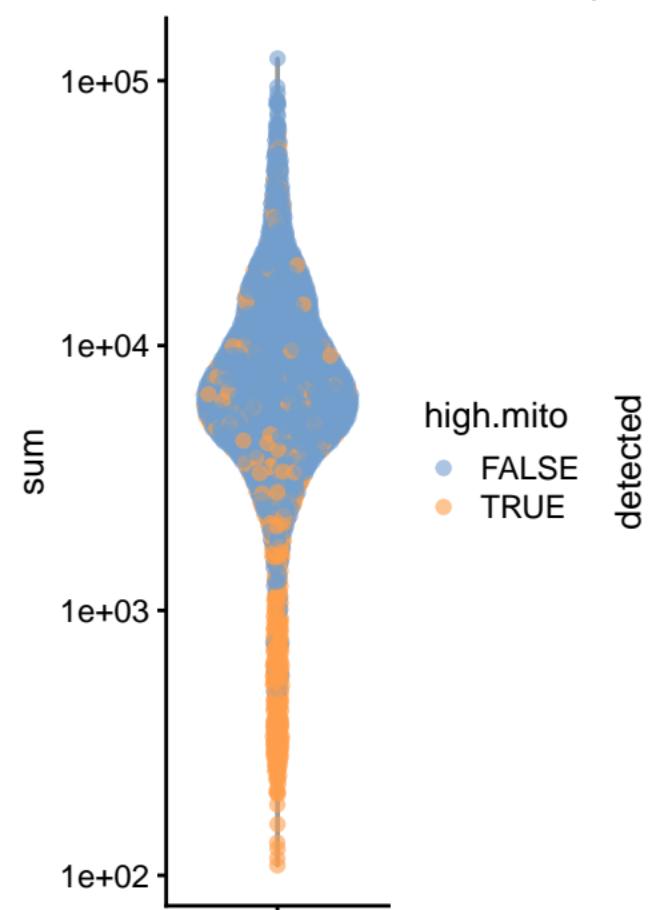
Sample: br5400.sacc; pre-QC nNuclei: 4364; nNuclei kept: 3959 (0.91%)



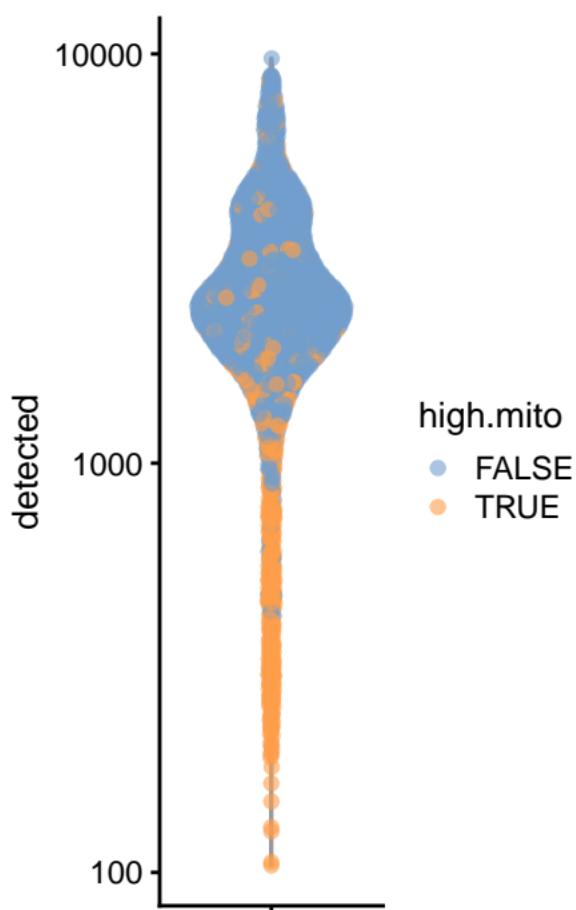
Sample: br5400.sacc; pre-QC nNuclei: 4364; nNuclei kept: 3959 (0.91%)



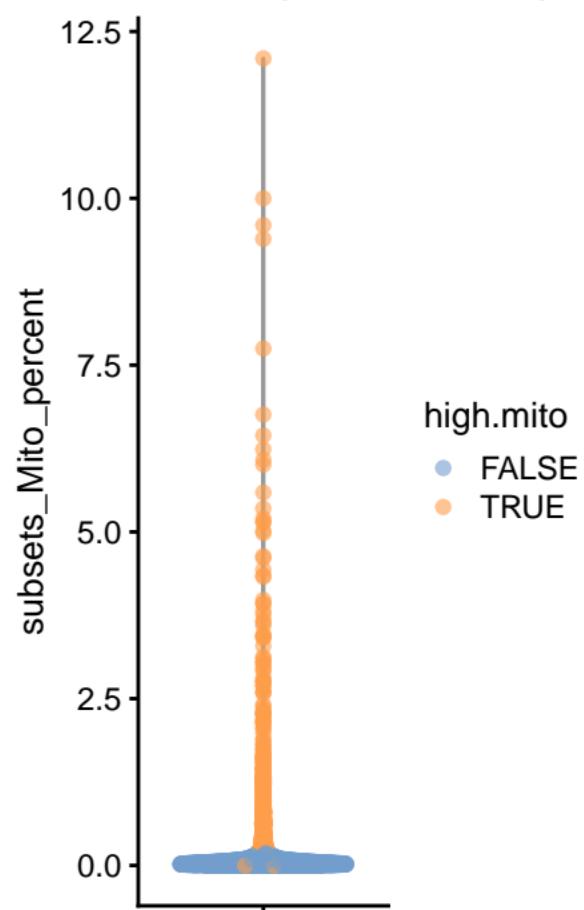
Total count: br5701.amy



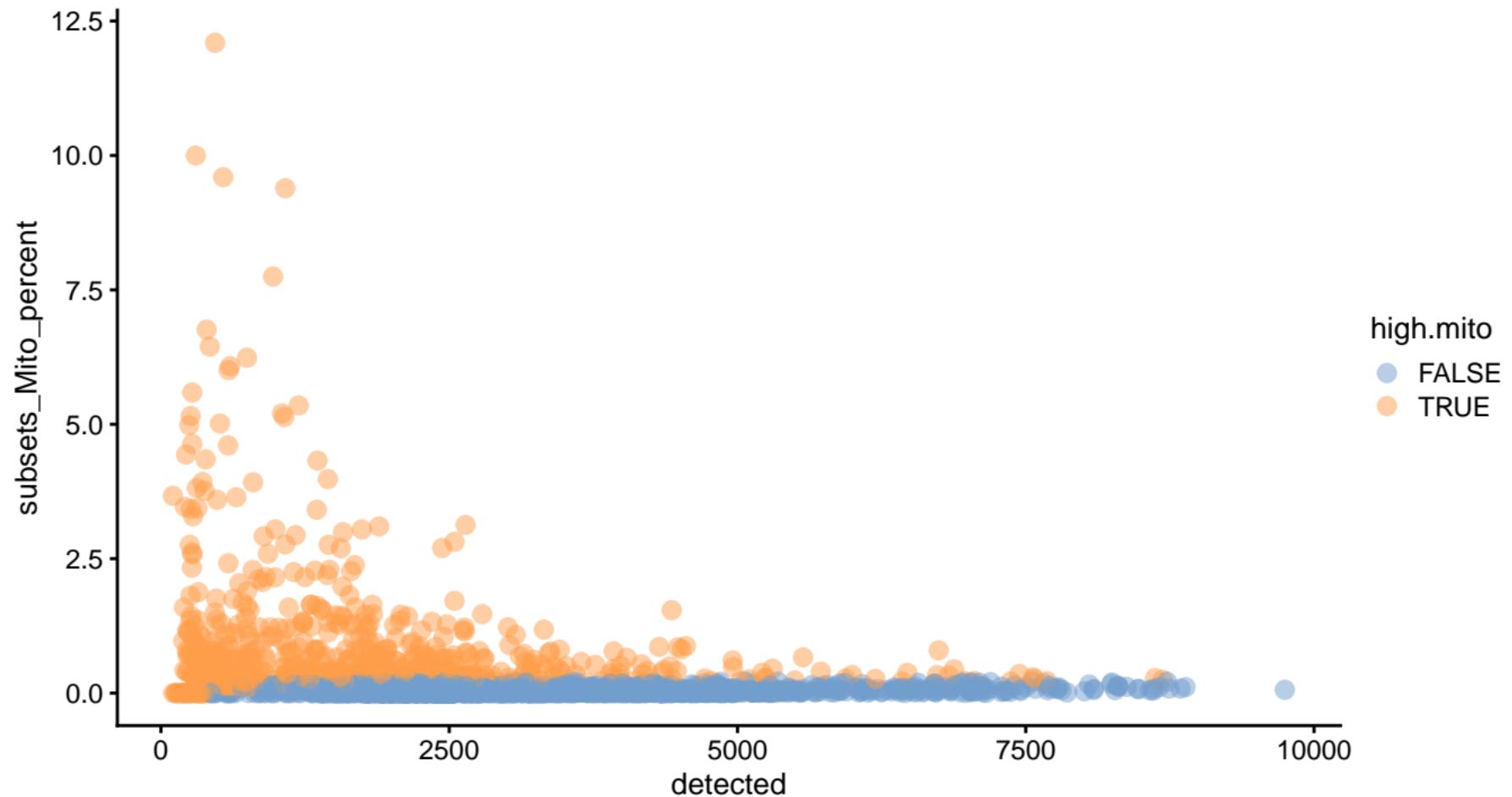
Detected features



Mito % (cutoff = 0.218)



Sample: br5701.amy; pre-QC nNuclei: 4199; nNuclei kept: 3524 (0.84%)



Sample: br5701.amy; pre-QC nNuclei: 4199; nNuclei kept: 3524 (0.84%)

