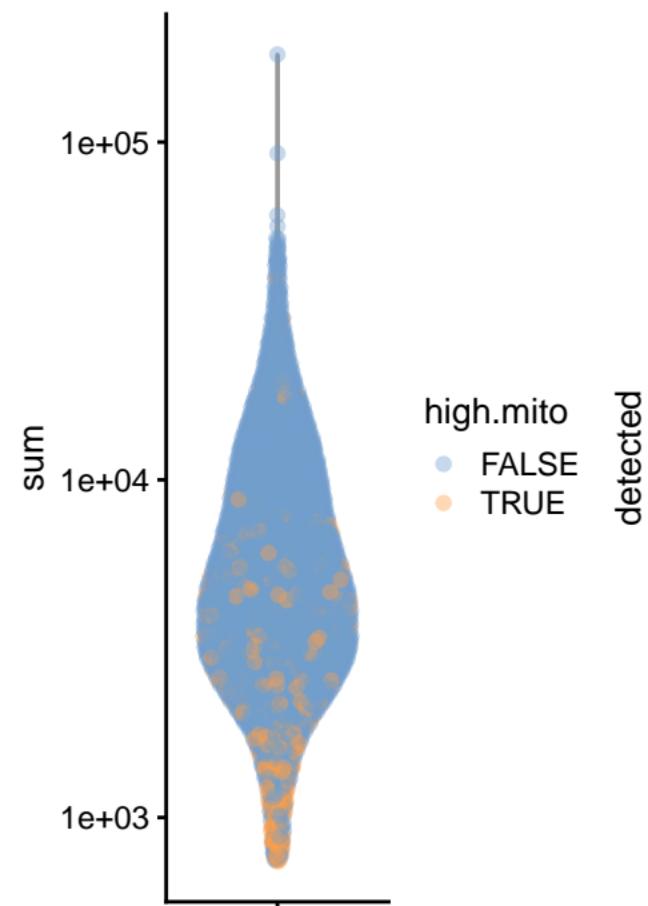
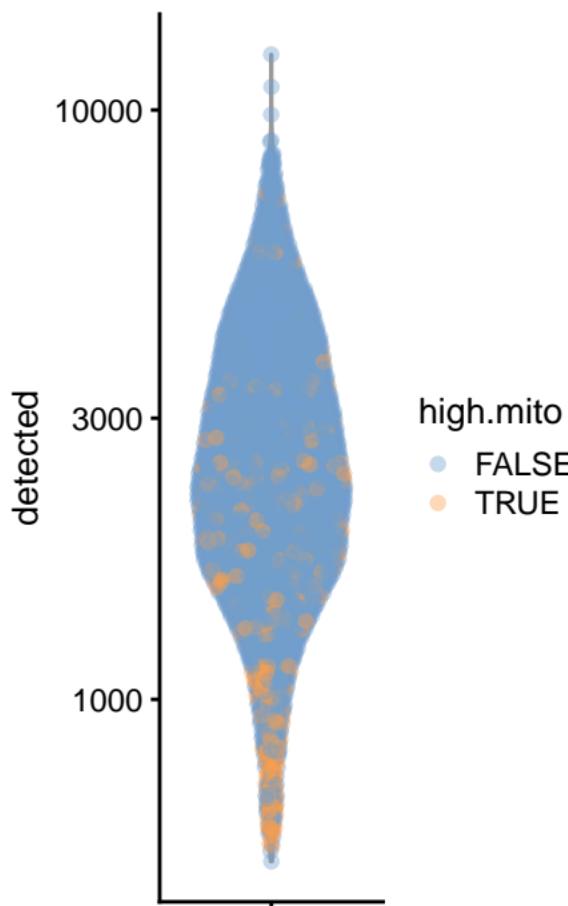


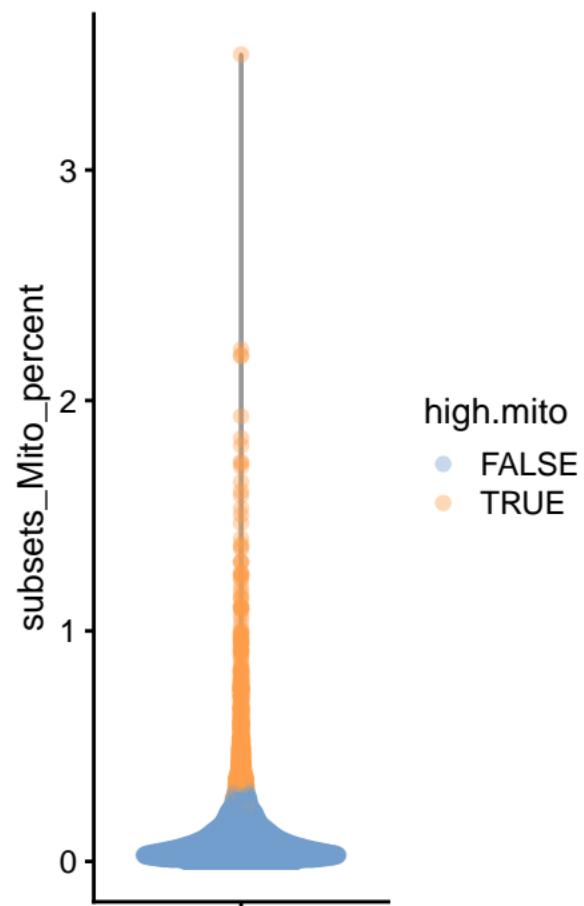
Total count: 1M



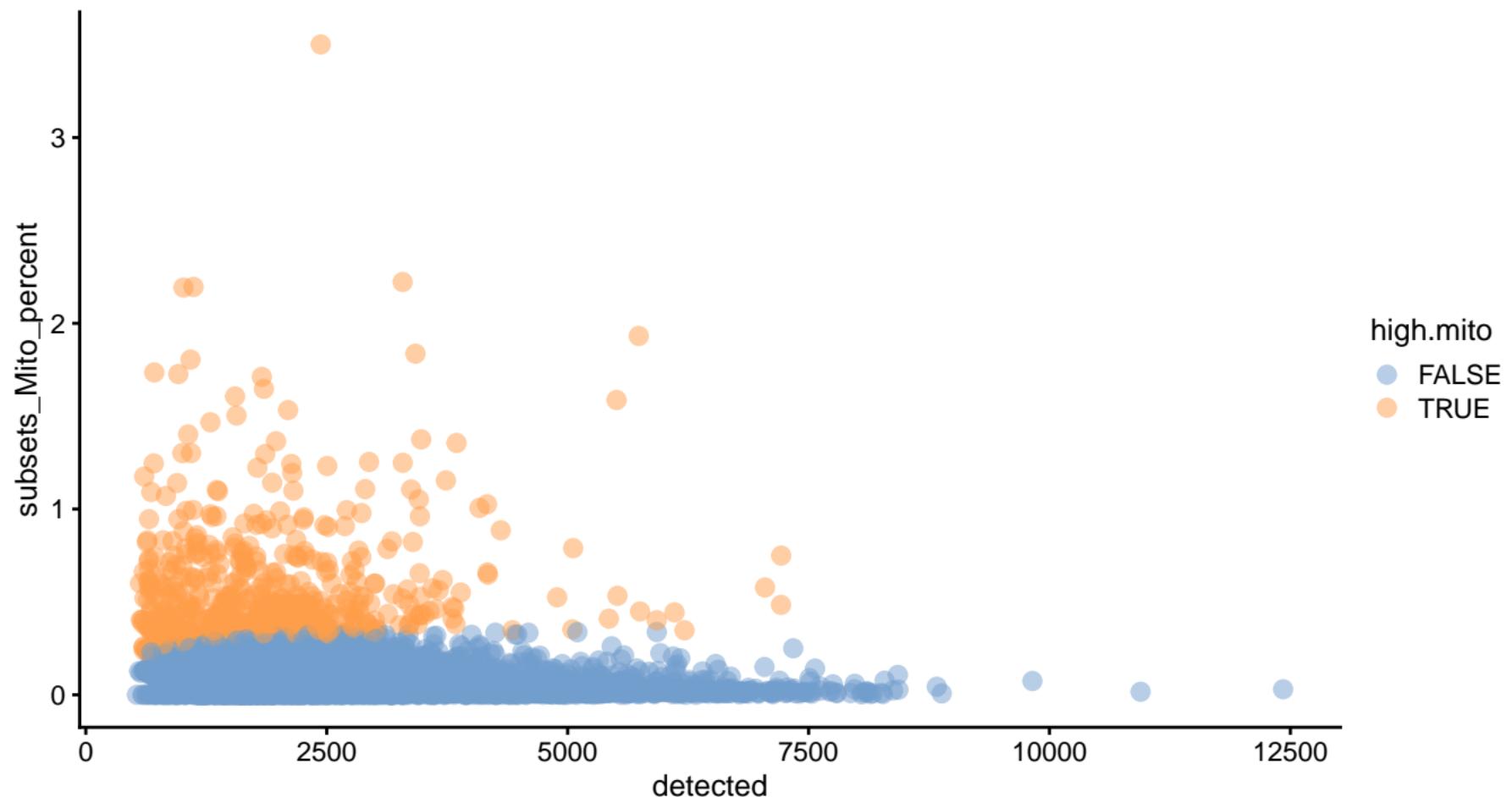
Detected features



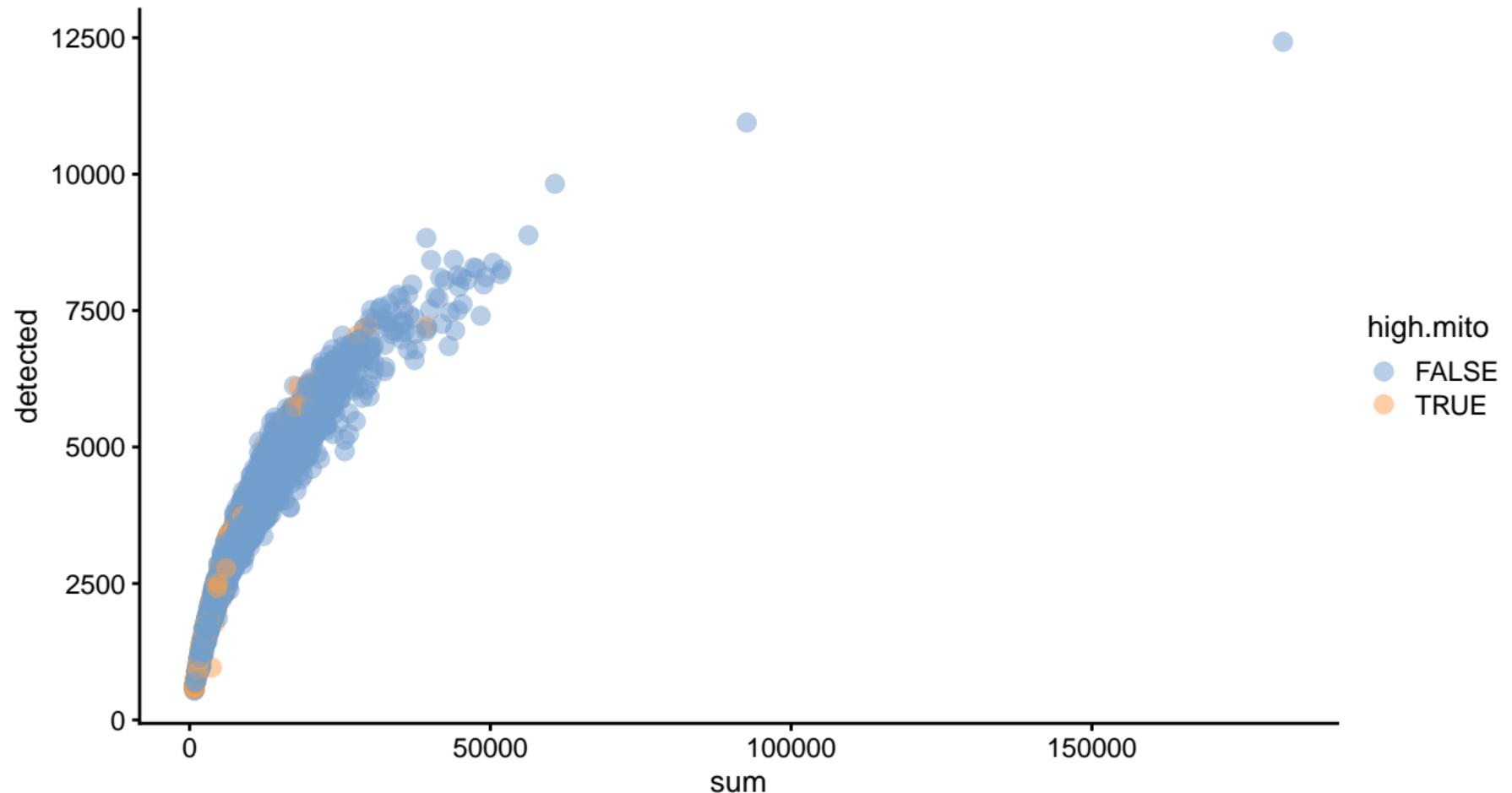
Mito % (cutoff = 0.348)

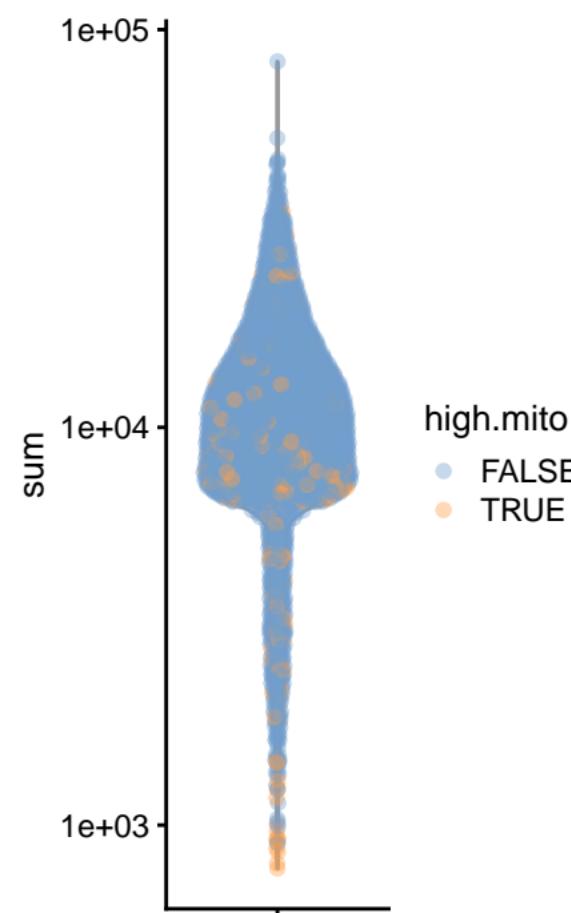
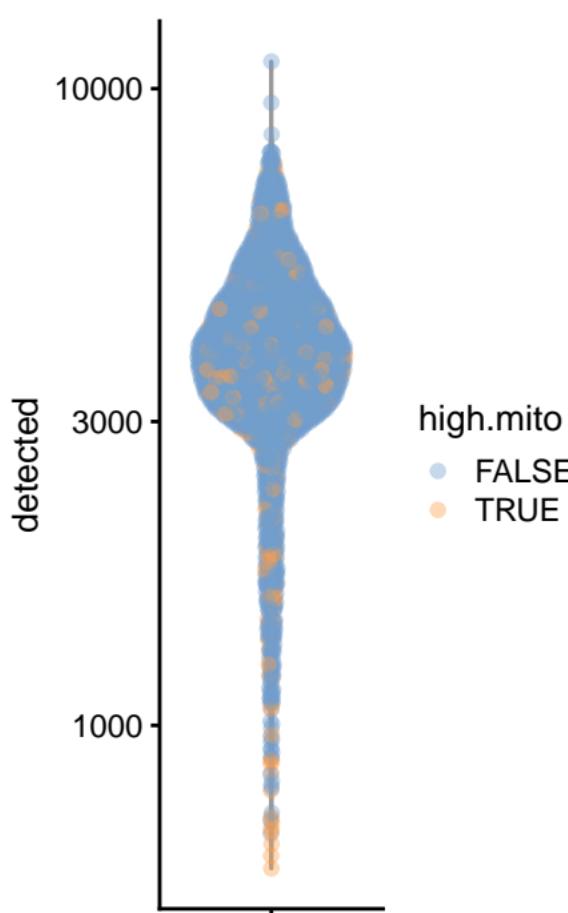
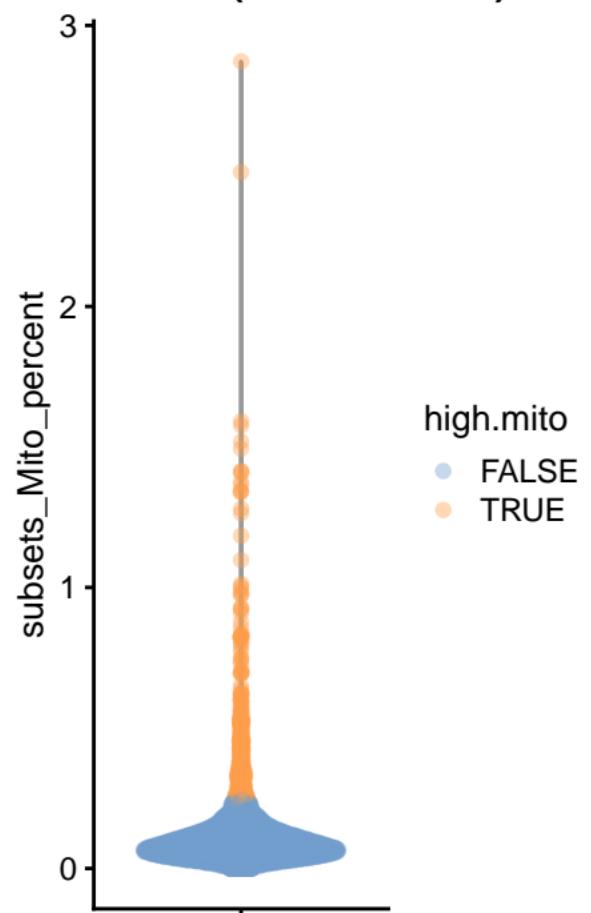


Sample: 1M; pre-QC nNuclei: 6728; nNuclei kept: 6084 (90.43%)

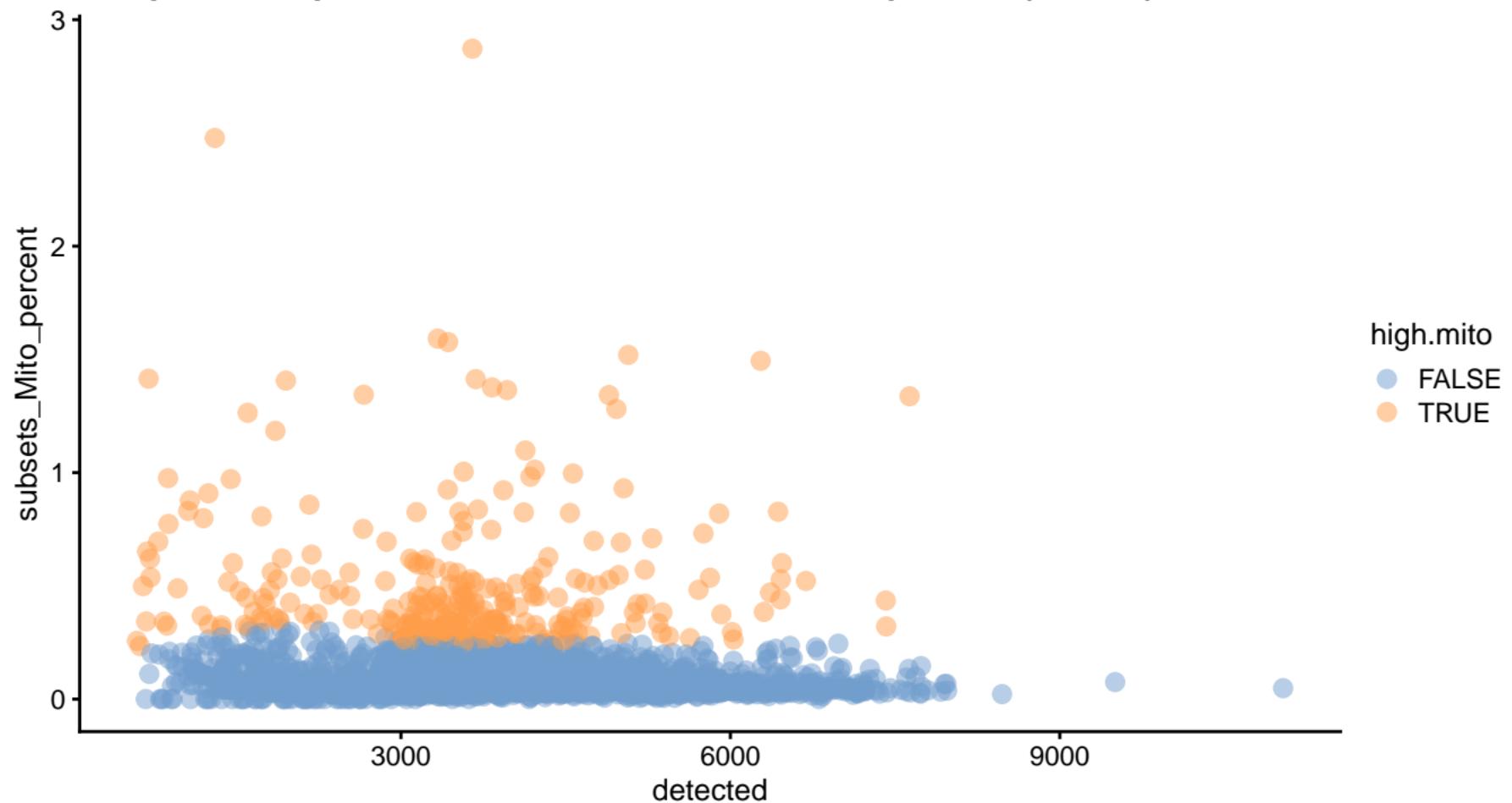


Sample: 1M; pre-QC nNuclei: 6728; nNuclei kept: 6084 (90.43%)

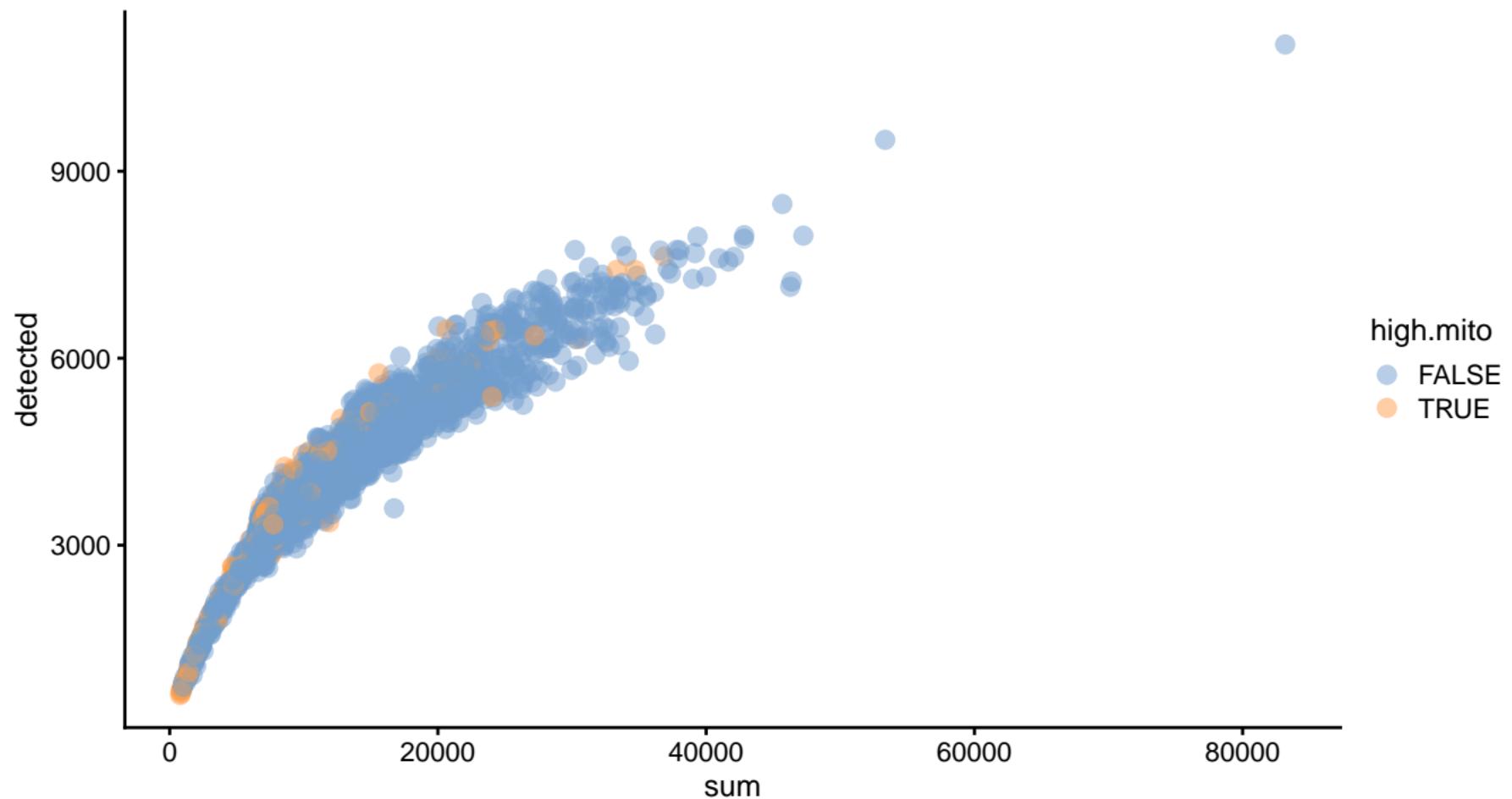


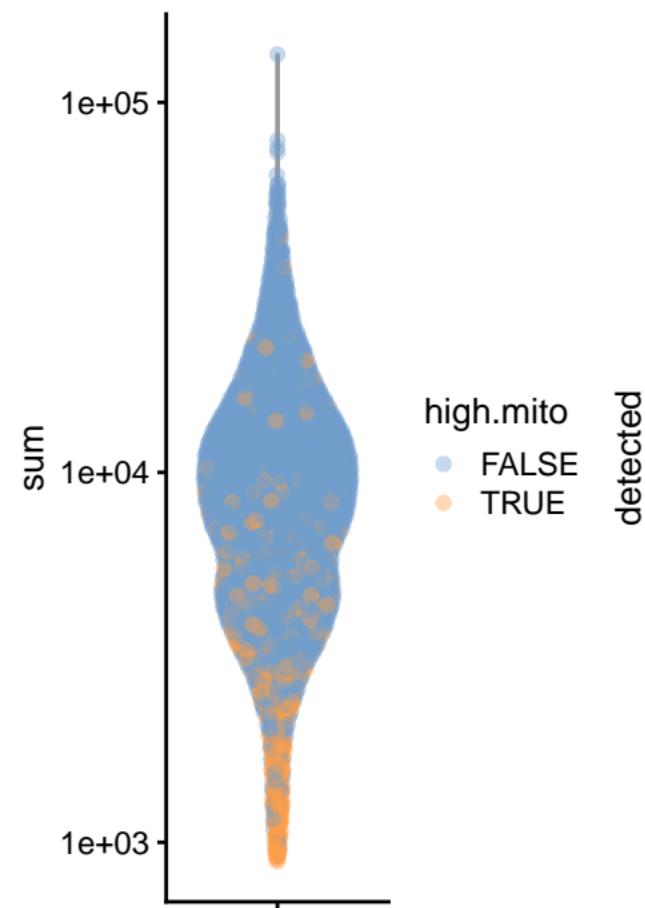
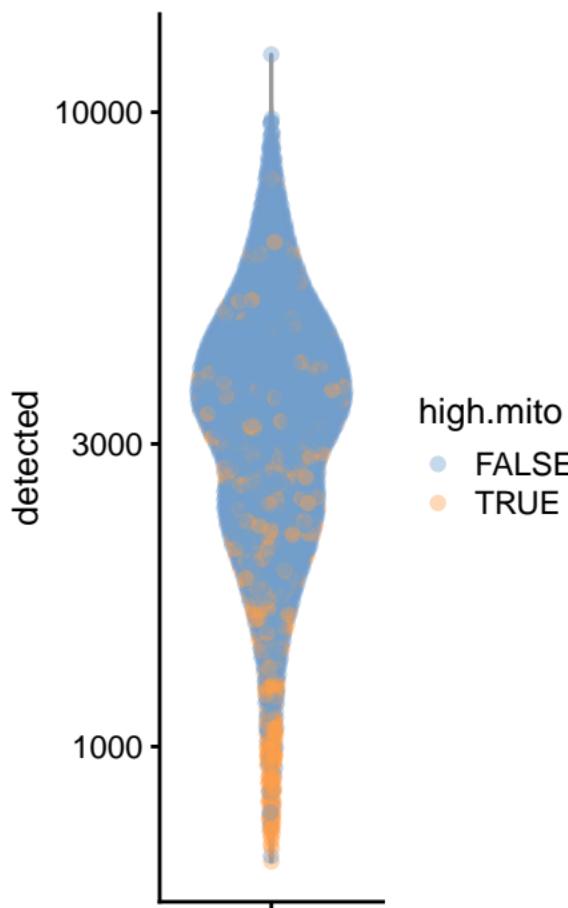
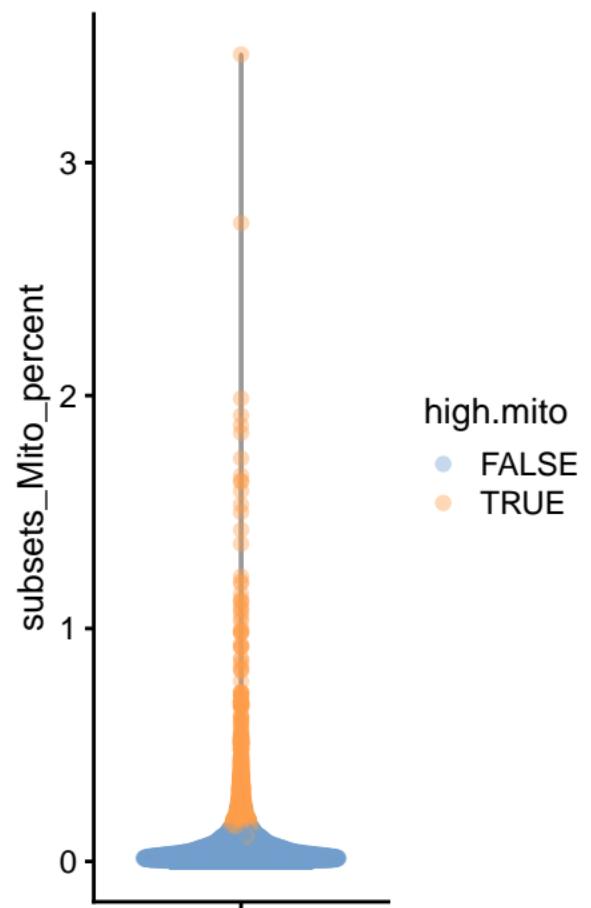
Total count: 2F**Detected features****Mito % (cutoff = 0.255)**

Sample: 2F; pre-QC nNuclei: 3982; nNuclei kept: 3693 (92.74%)

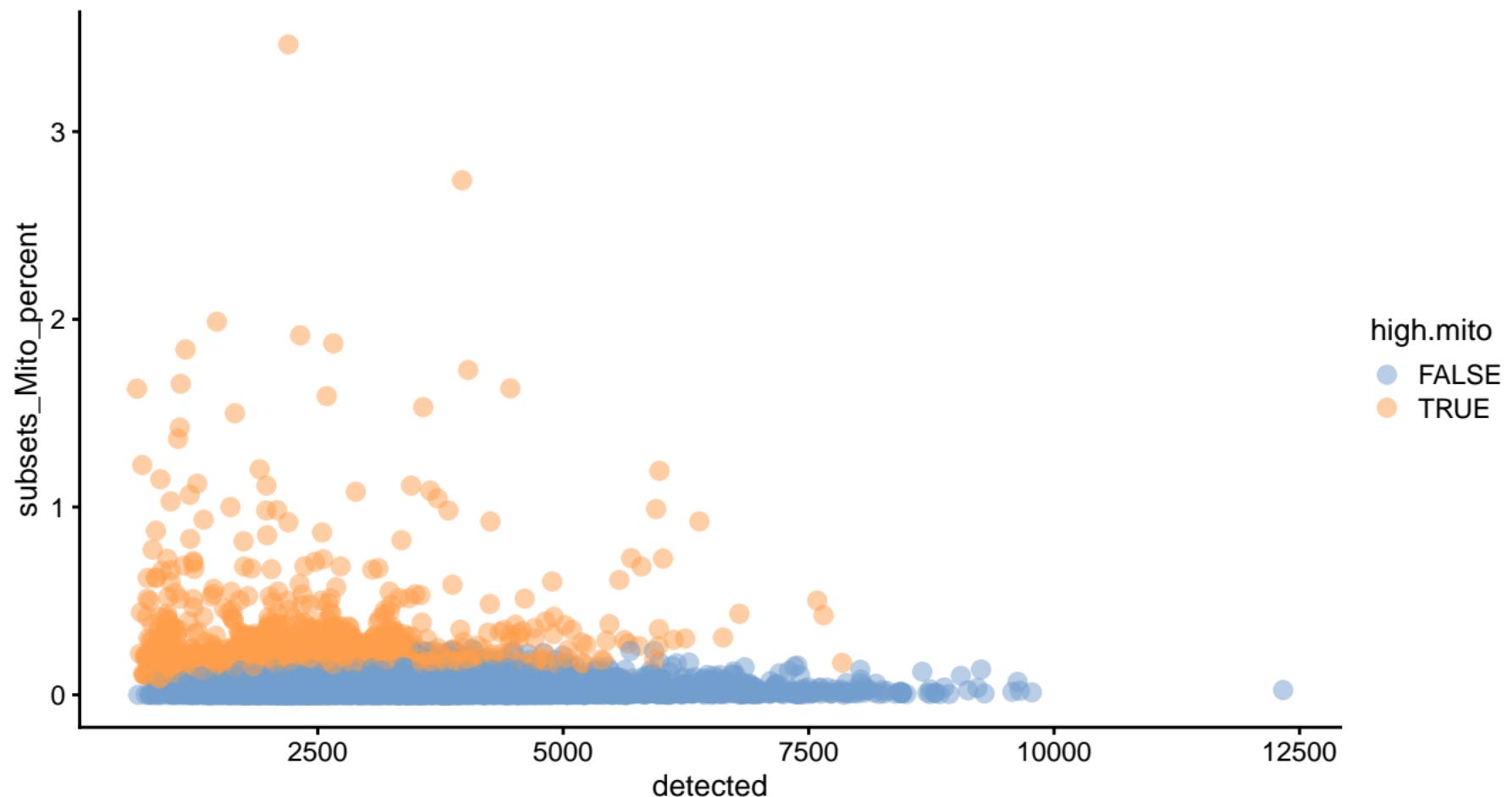


Sample: 2F; pre-QC nNuclei: 3982; nNuclei kept: 3693 (92.74%)

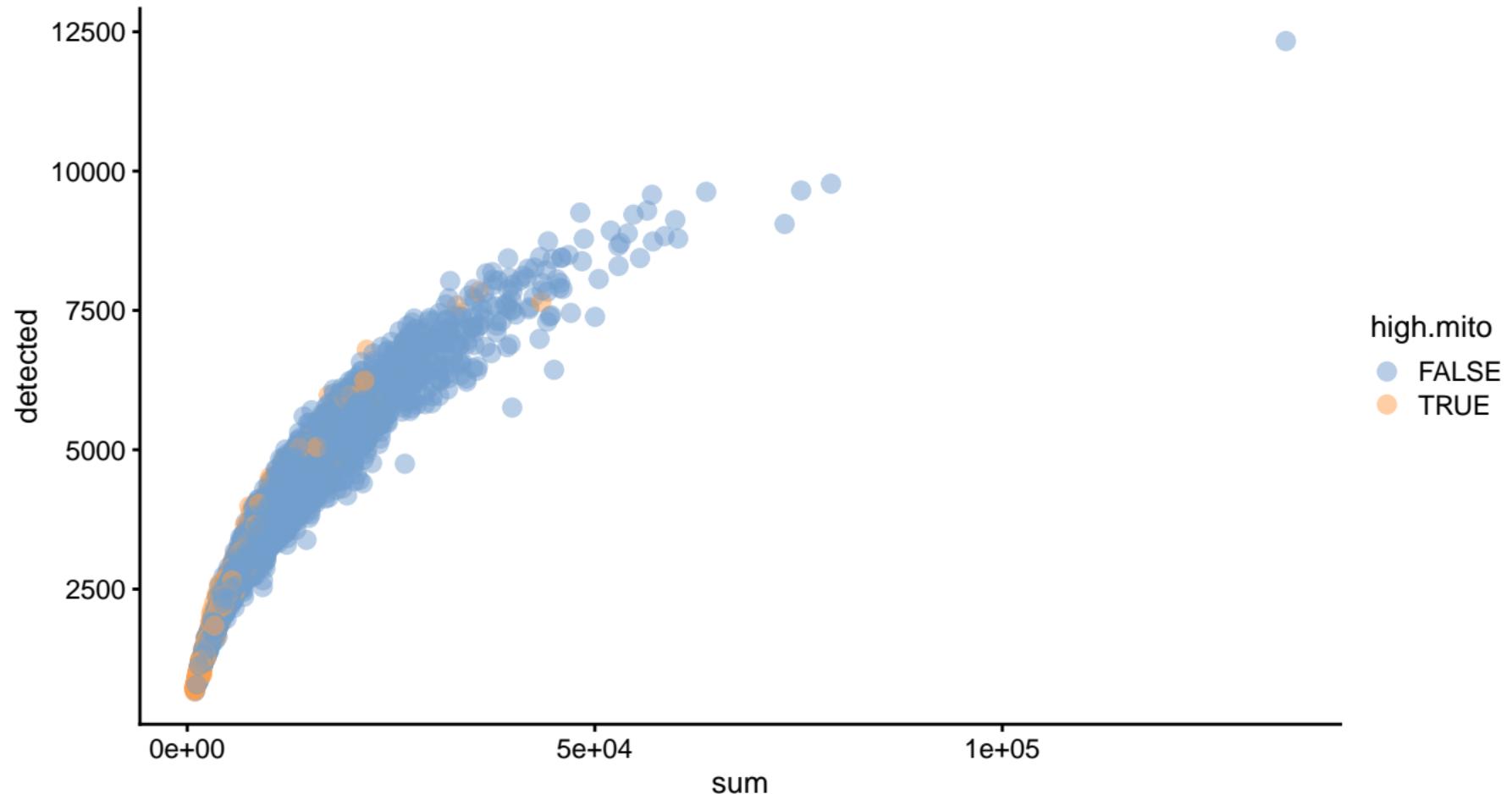


Total count: 3M**Detected features****Mito % (cutoff = 0.164)**

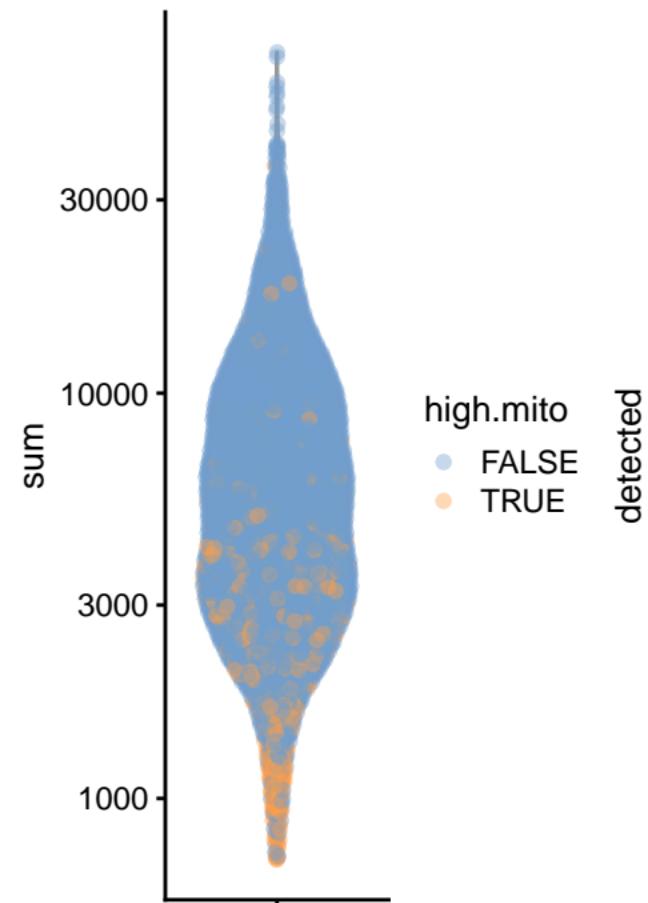
Sample: 3M; pre-QC nNuclei: 6961; nNuclei kept: 6068 (87.17%)



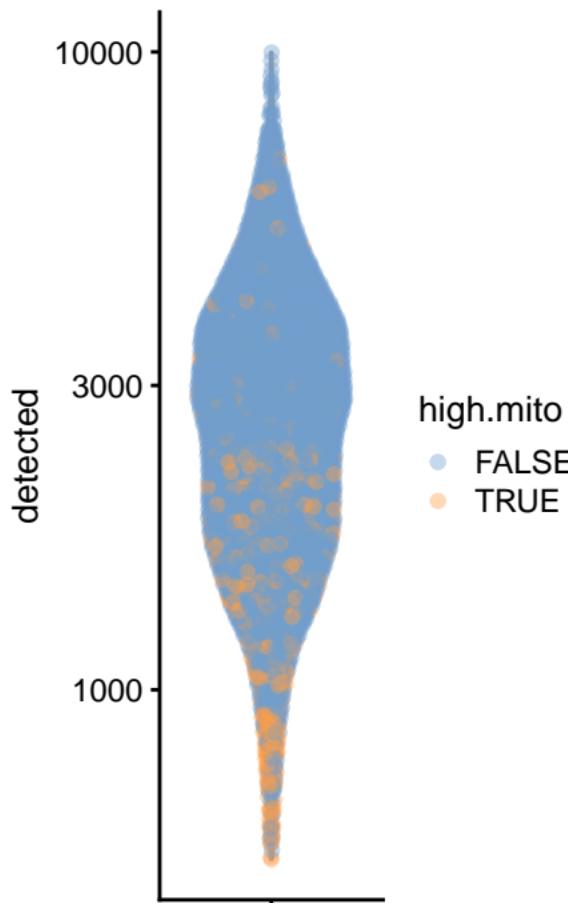
Sample: 3M; pre-QC nNuclei: 6961; nNuclei kept: 6068 (87.17%)



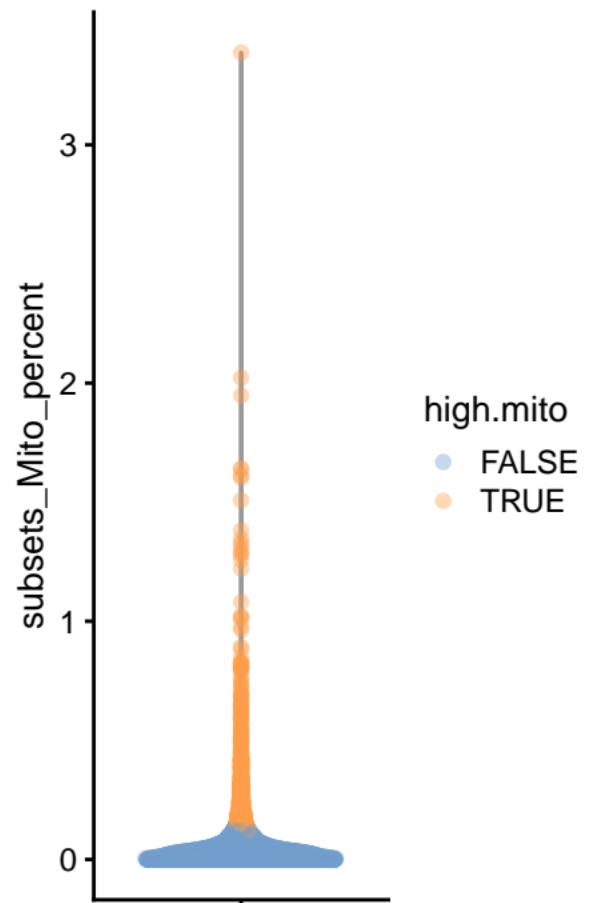
Total count: 4F



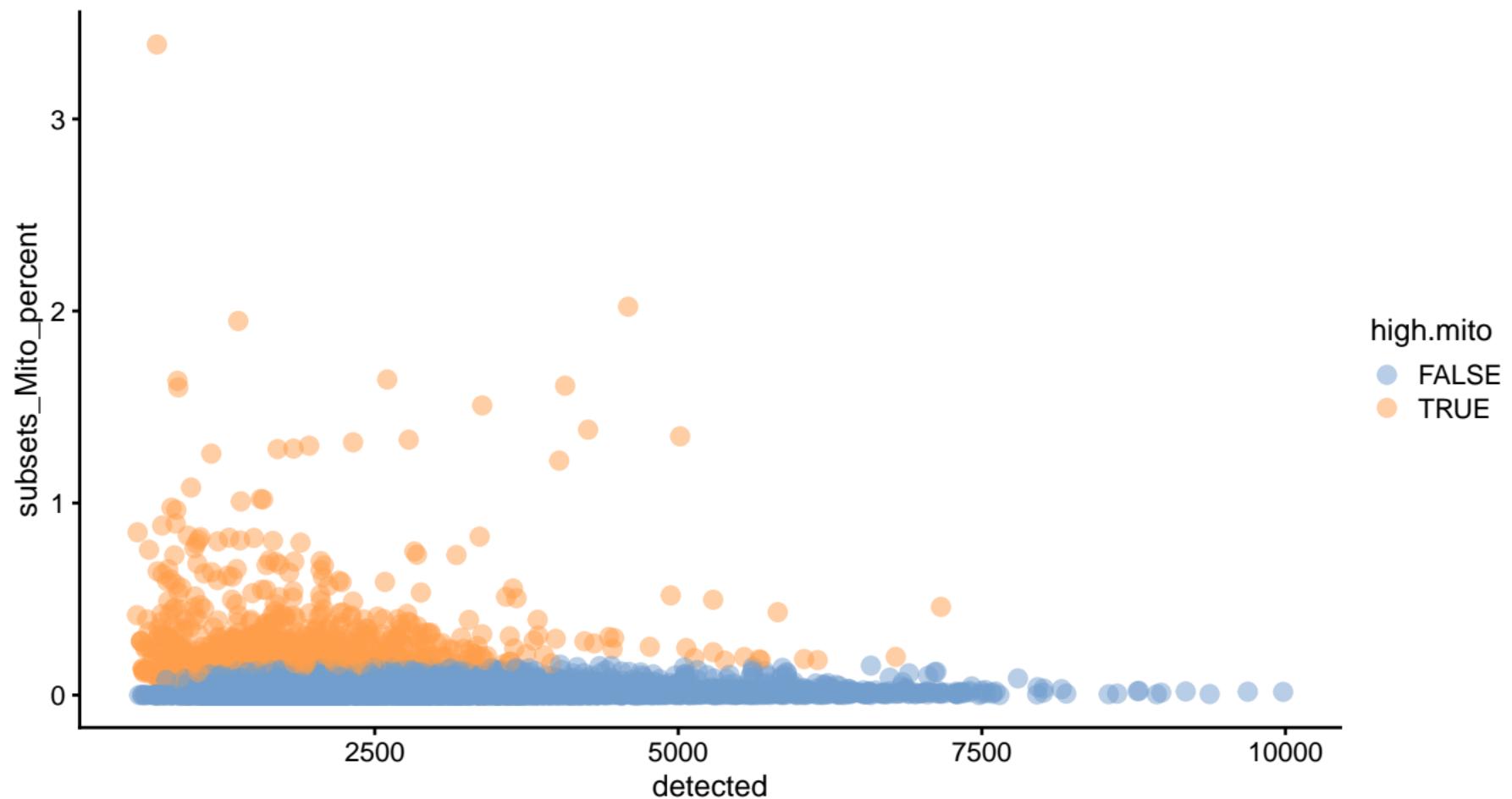
Detected features



Mito % (cutoff = 0.169)



Sample: 4F; pre-QC nNuclei: 7856; nNuclei kept: 7015 (89.29%)



Sample: 4F; pre-QC nNuclei: 7856; nNuclei kept: 7015 (89.29%)

