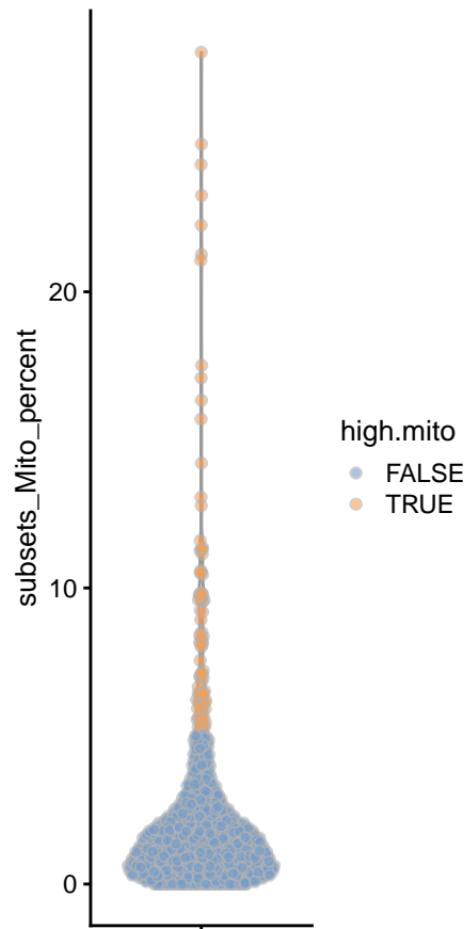
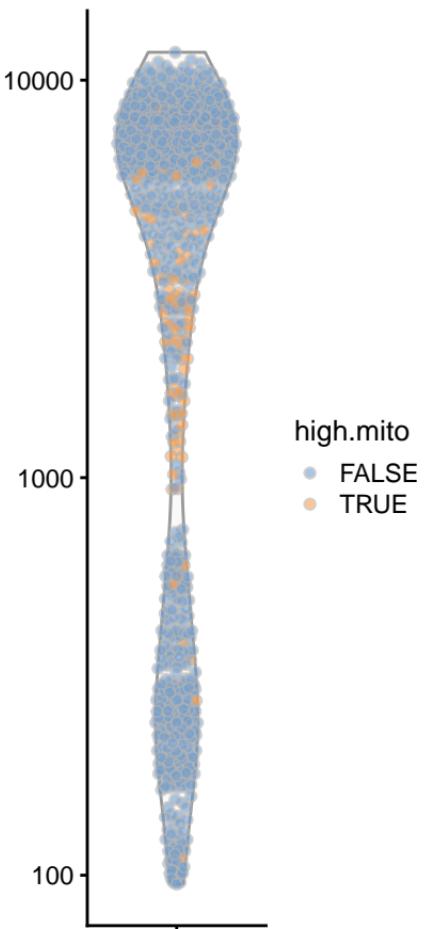
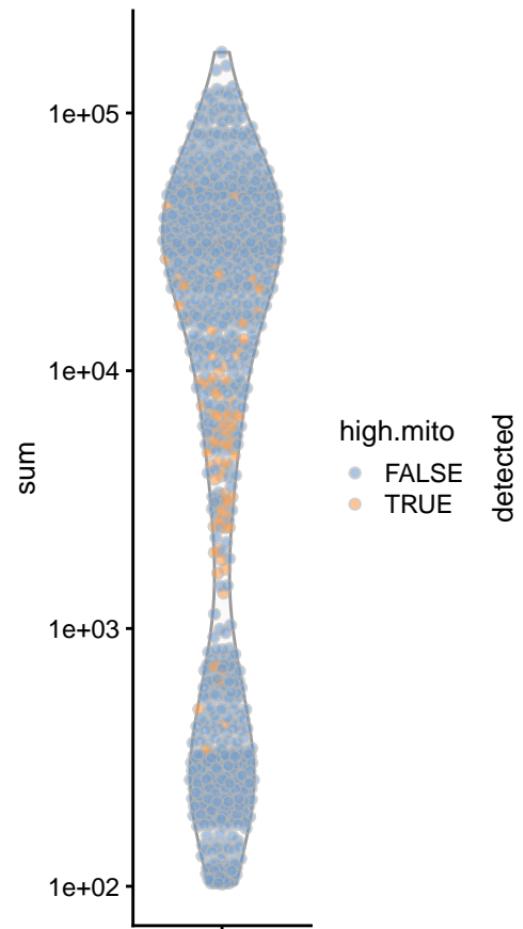


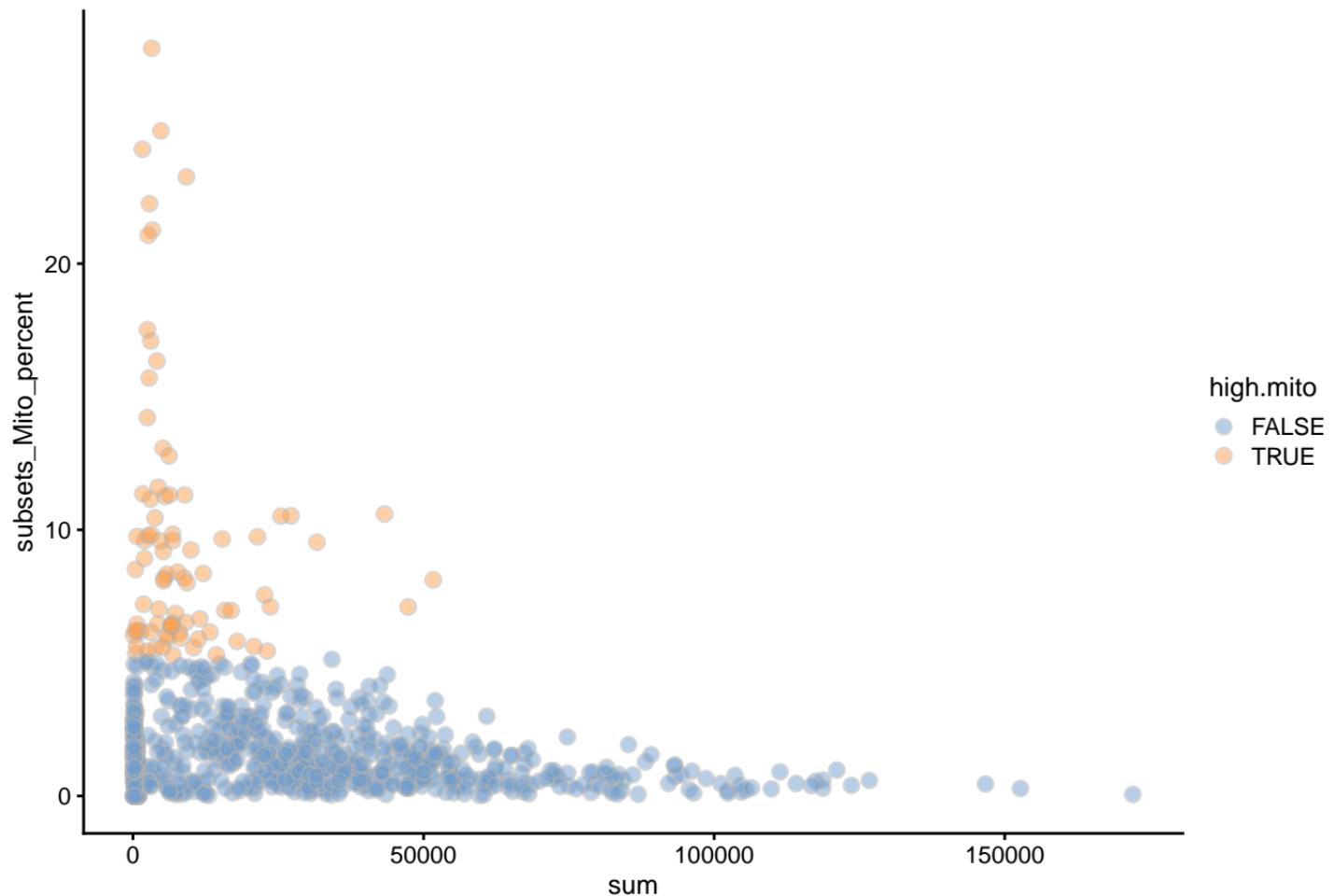
Total count: br5276.sacc.neun

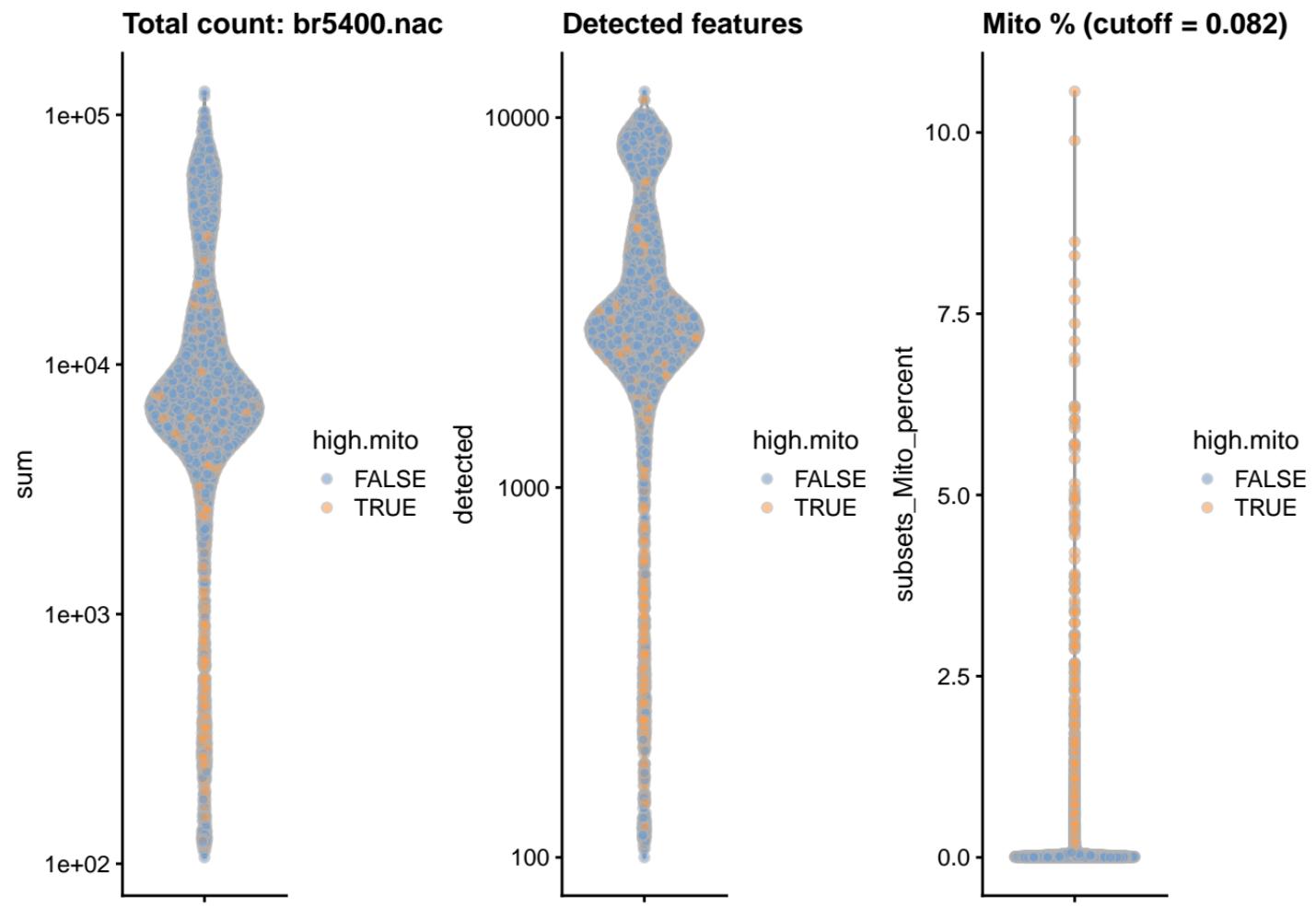
Detected features

Mito % (cutoff = 5.191)

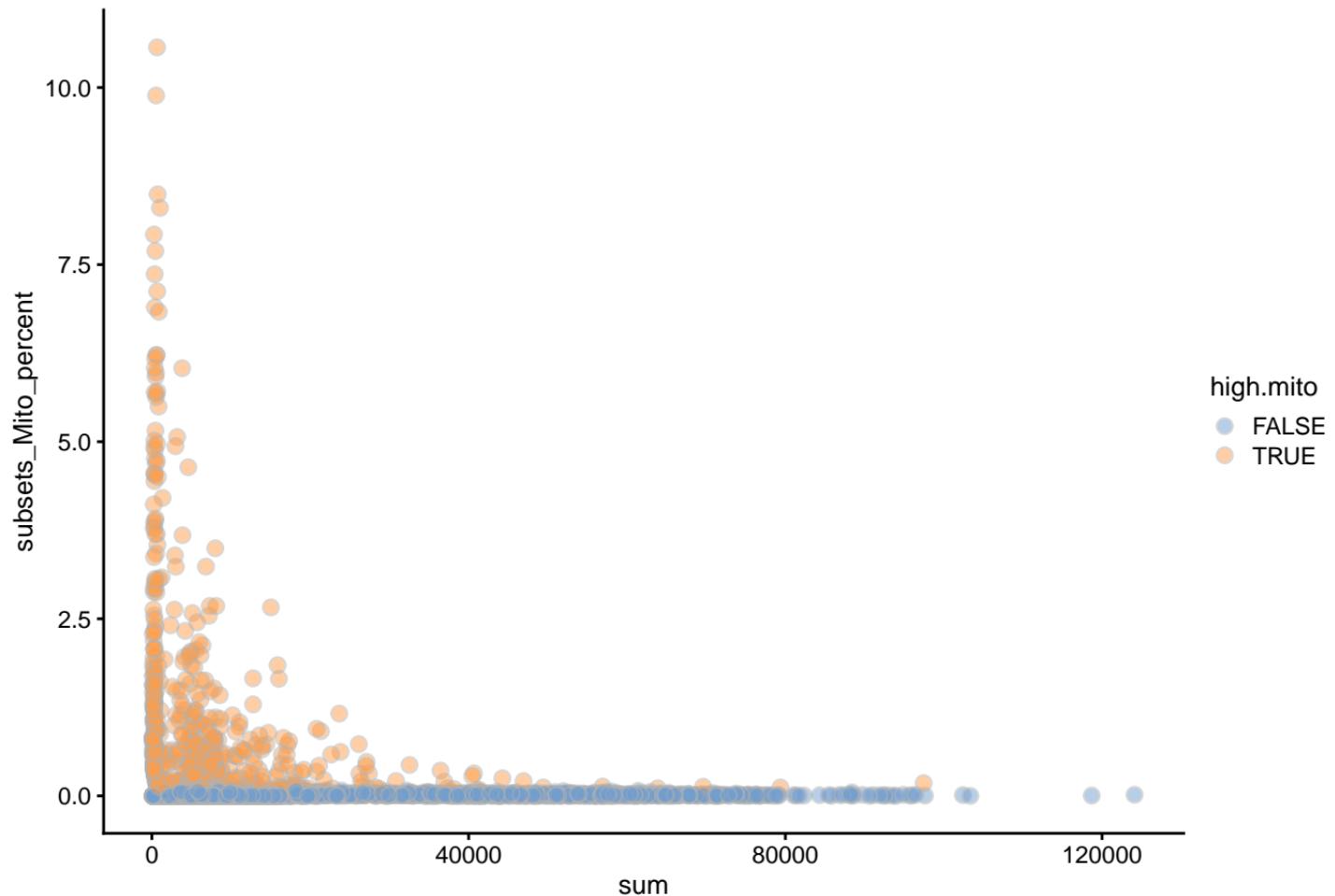


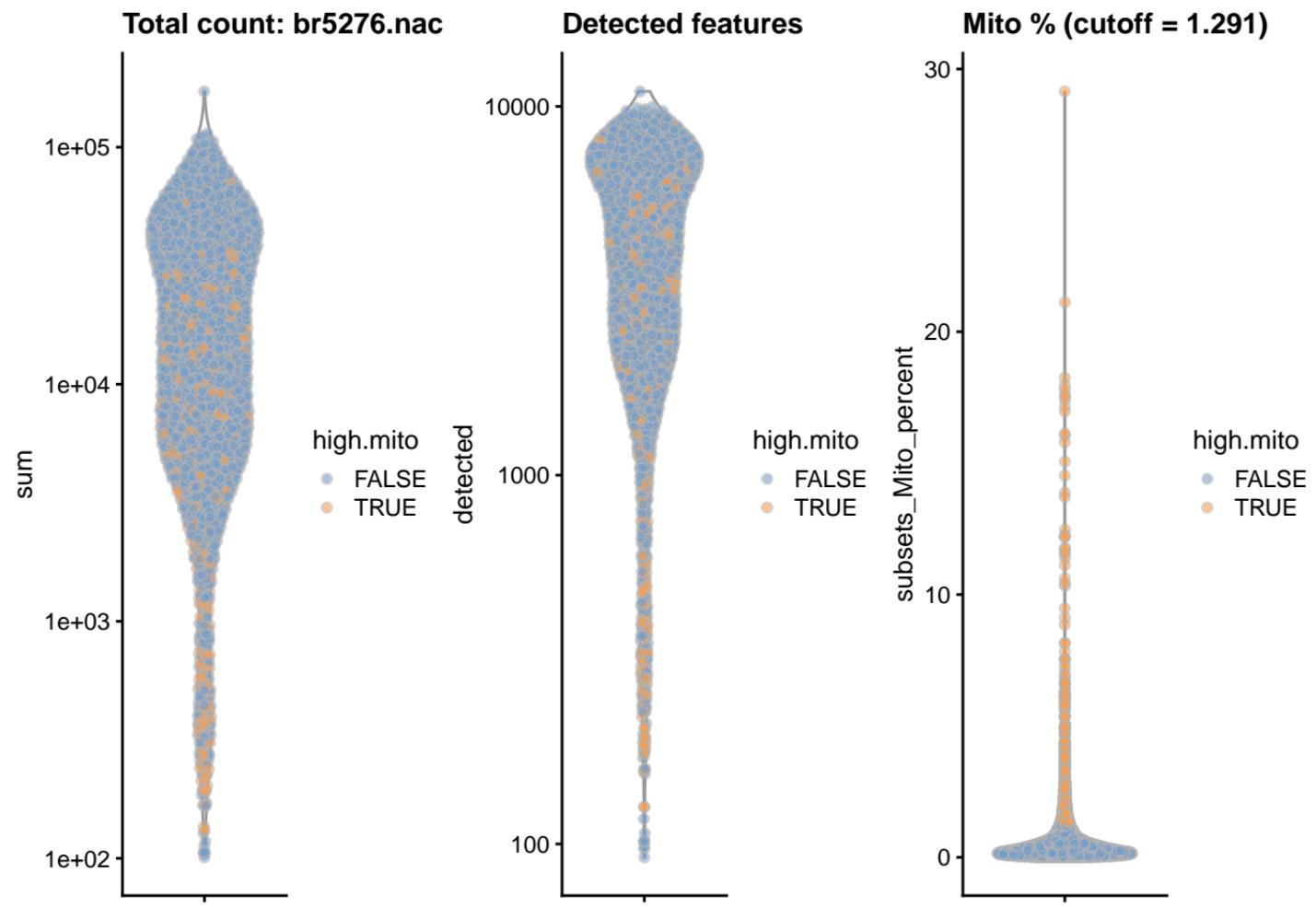
Sample: br5276.sacc.neun; nNuclei (pre-QC): 926; Mito % (cutoff = 5.191)



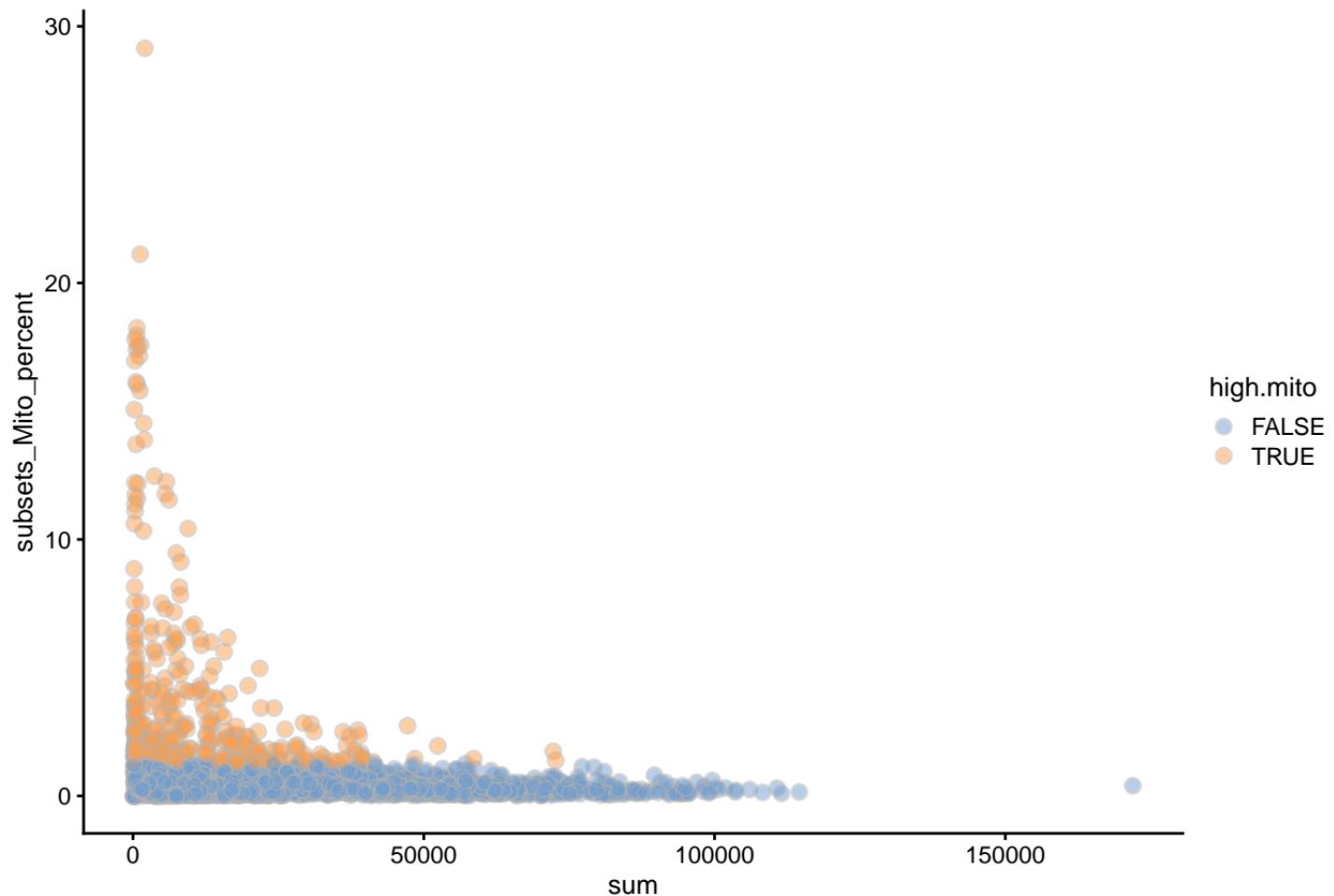


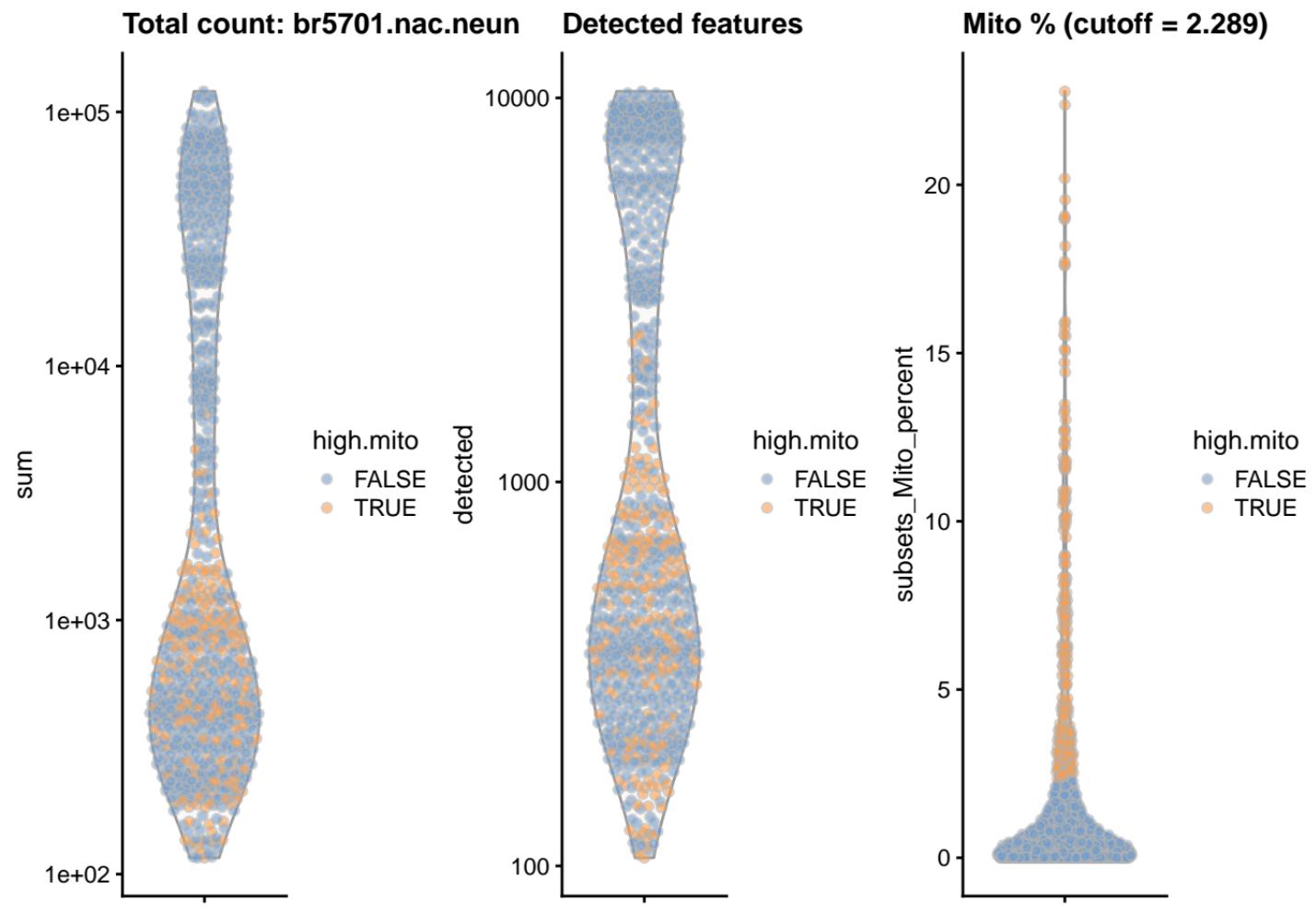
Sample: br5400.nac; nNuclei (pre-QC): 4859; Mito % (cutoff = 0.082)



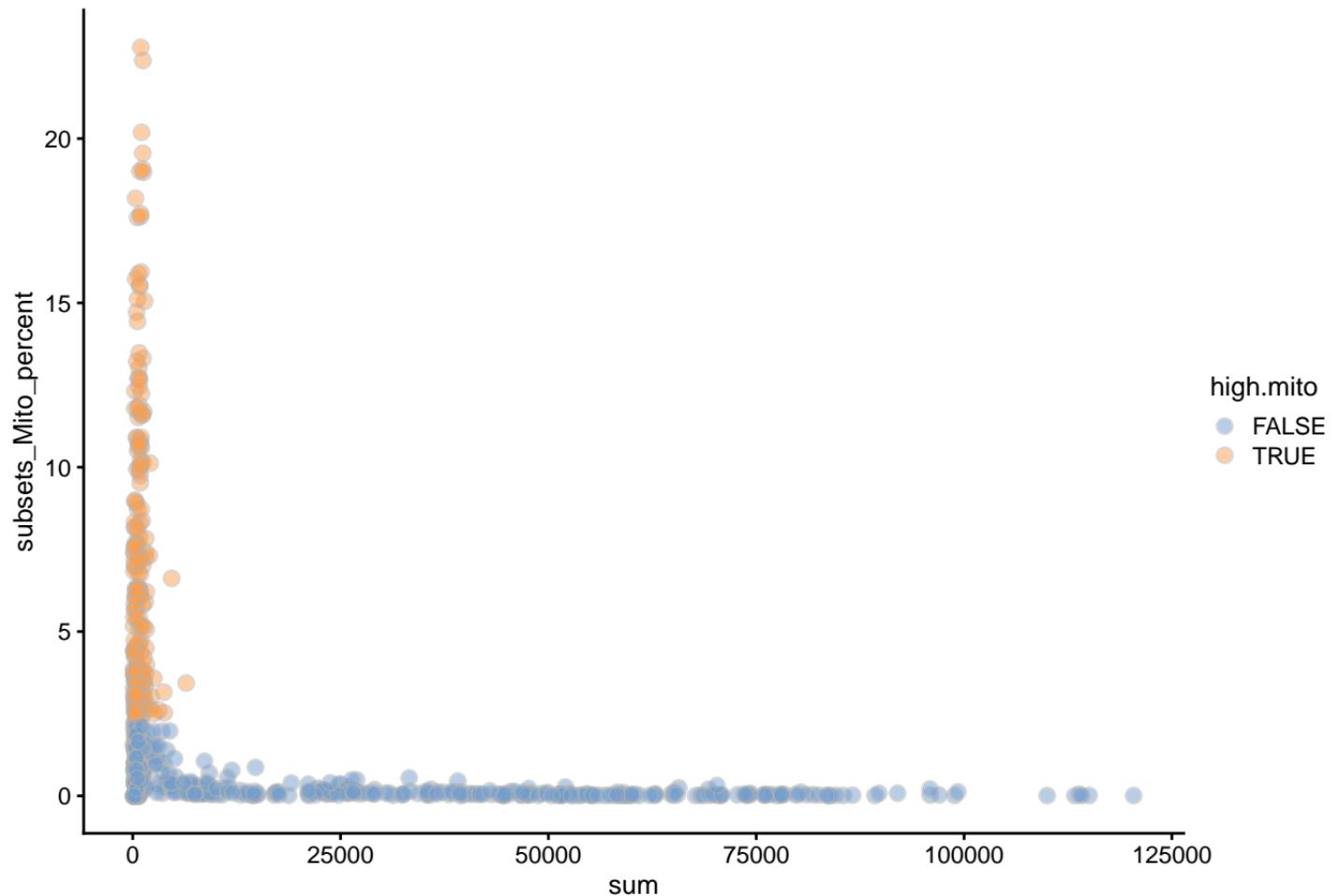


Sample: br5276.nac; nNuclei (pre-QC): 3012; Mito % (cutoff = 1.291)





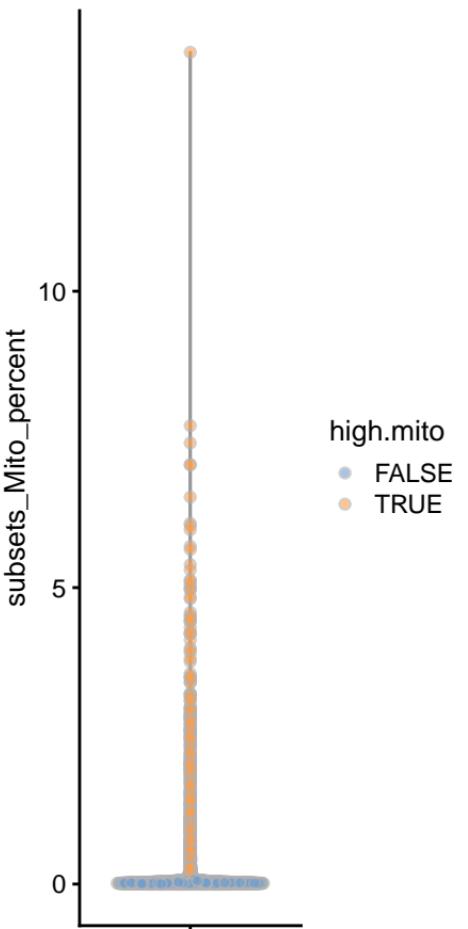
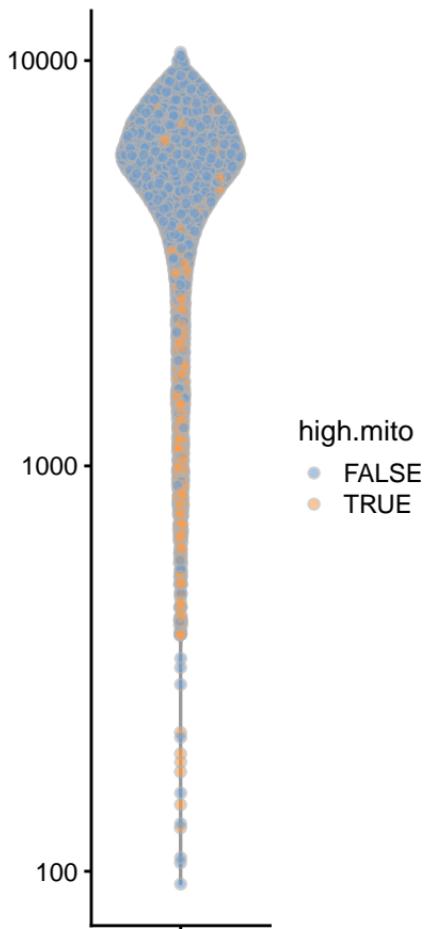
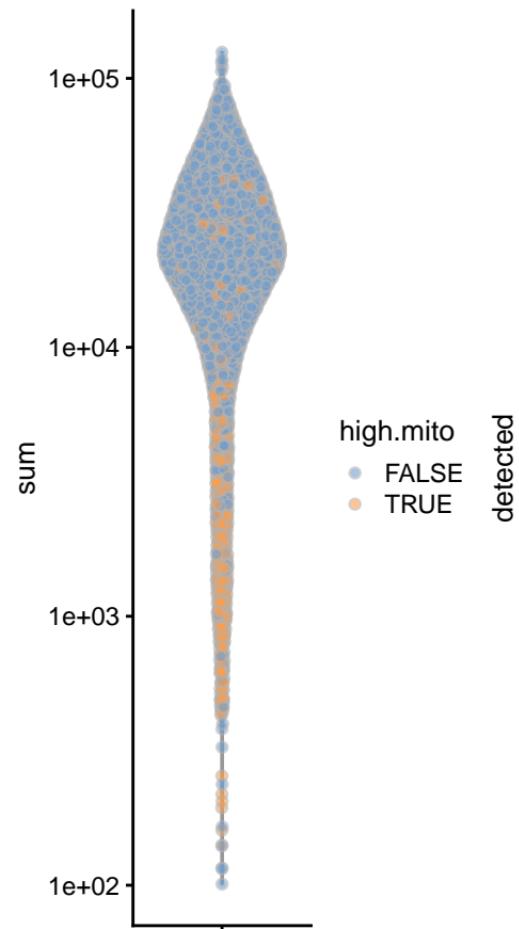
Sample: br5701.nac.neun; nNuclei (pre-QC): 814; Mito % (cutoff = 2.289)



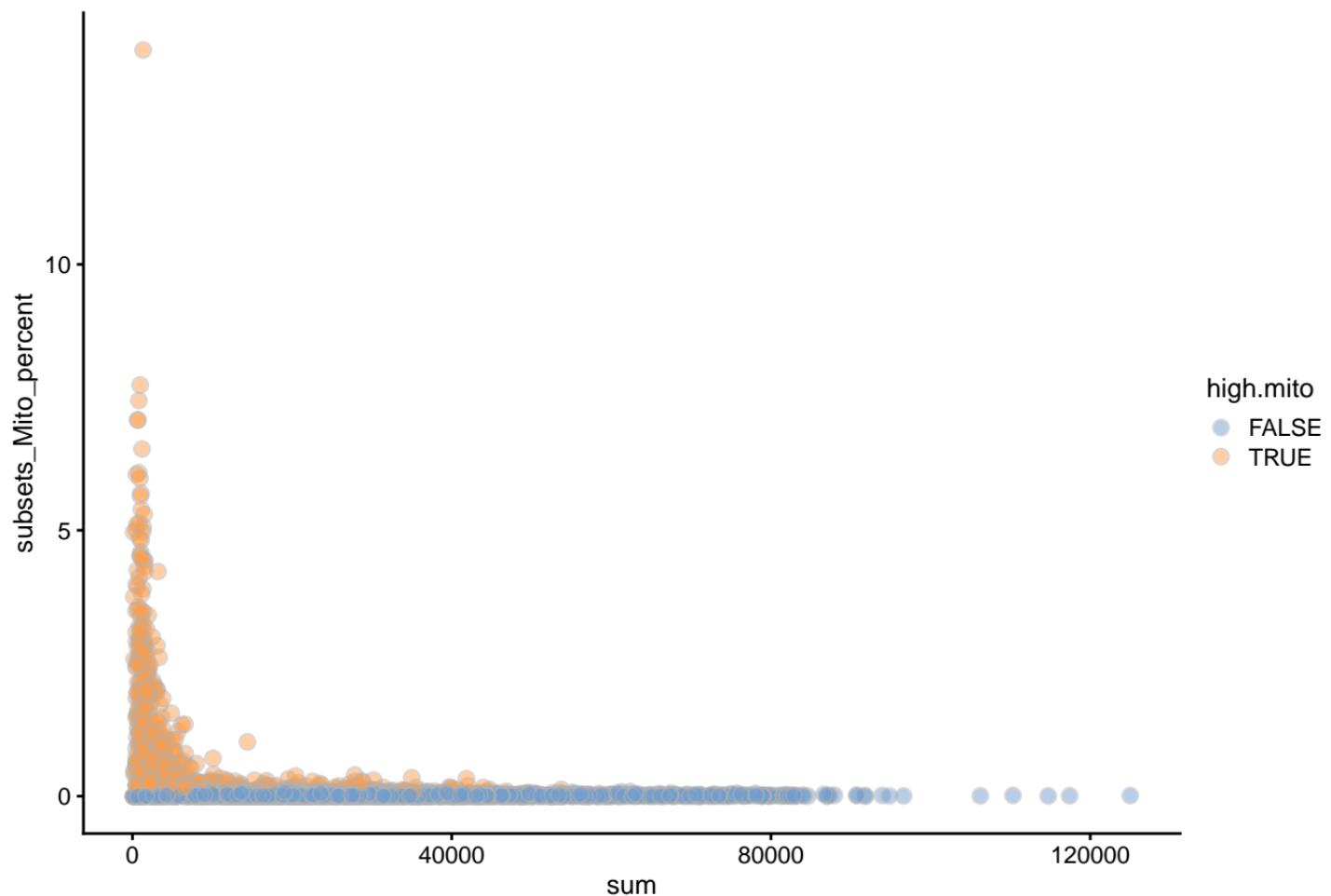
Total count: br5701.sacc.neun

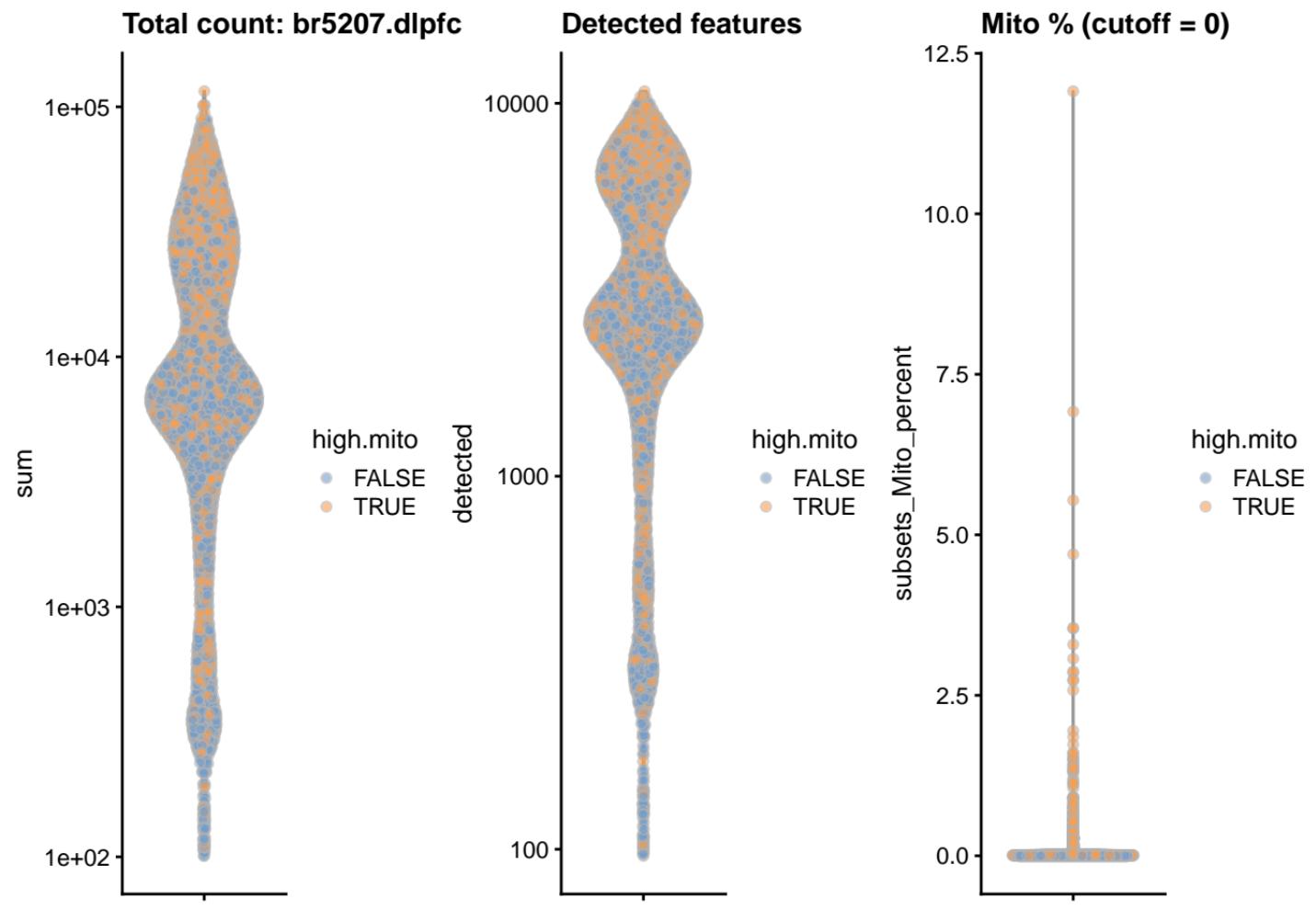
Detected features

Mito % (cutoff = 0.086)

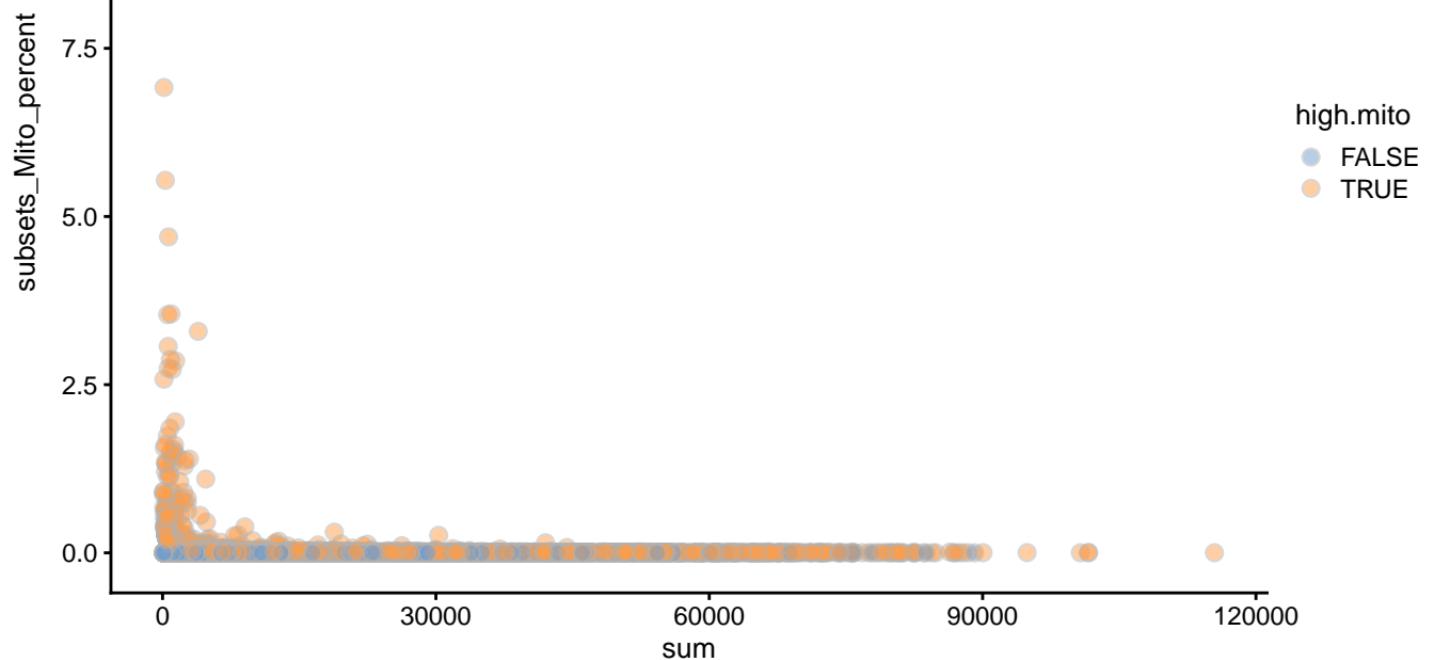


Sample: br5701.sacc.neun; nNuclei (pre-QC): 4518; Mito % (cutoff = 0.086)





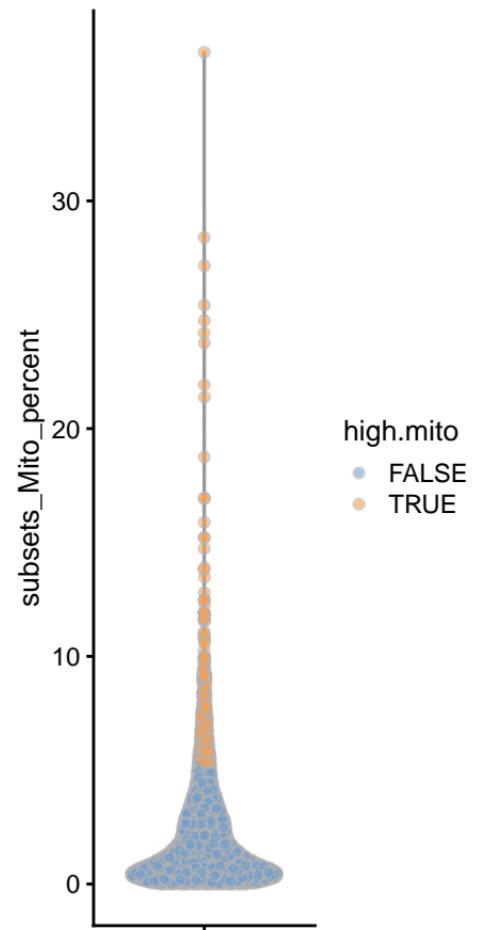
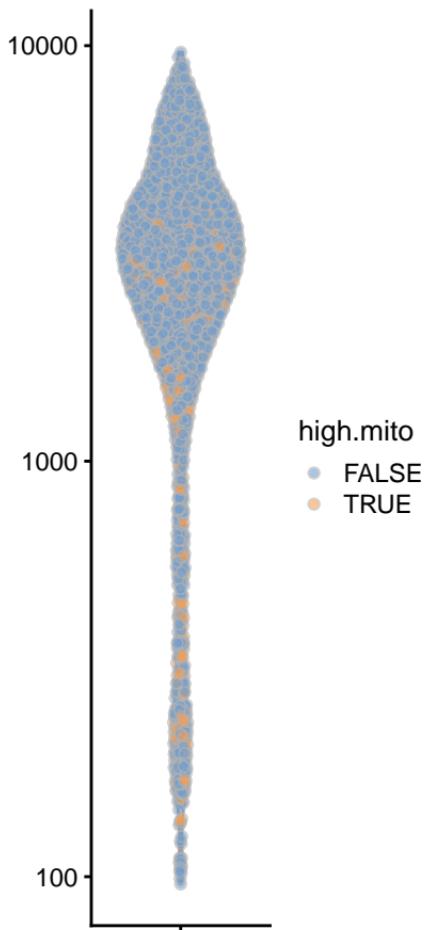
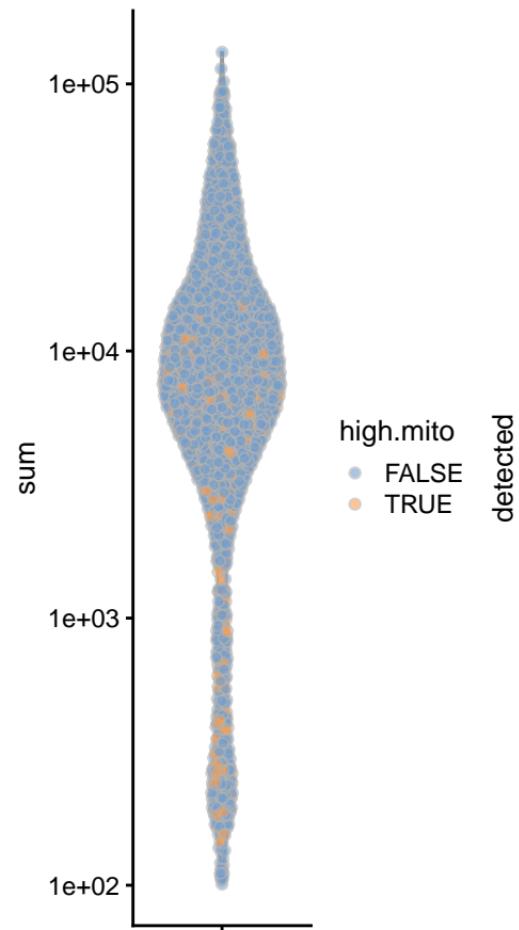
Sample: br5207.dlpfc; nNuclei (pre-QC): 6453; Mito % (cutoff = 0)



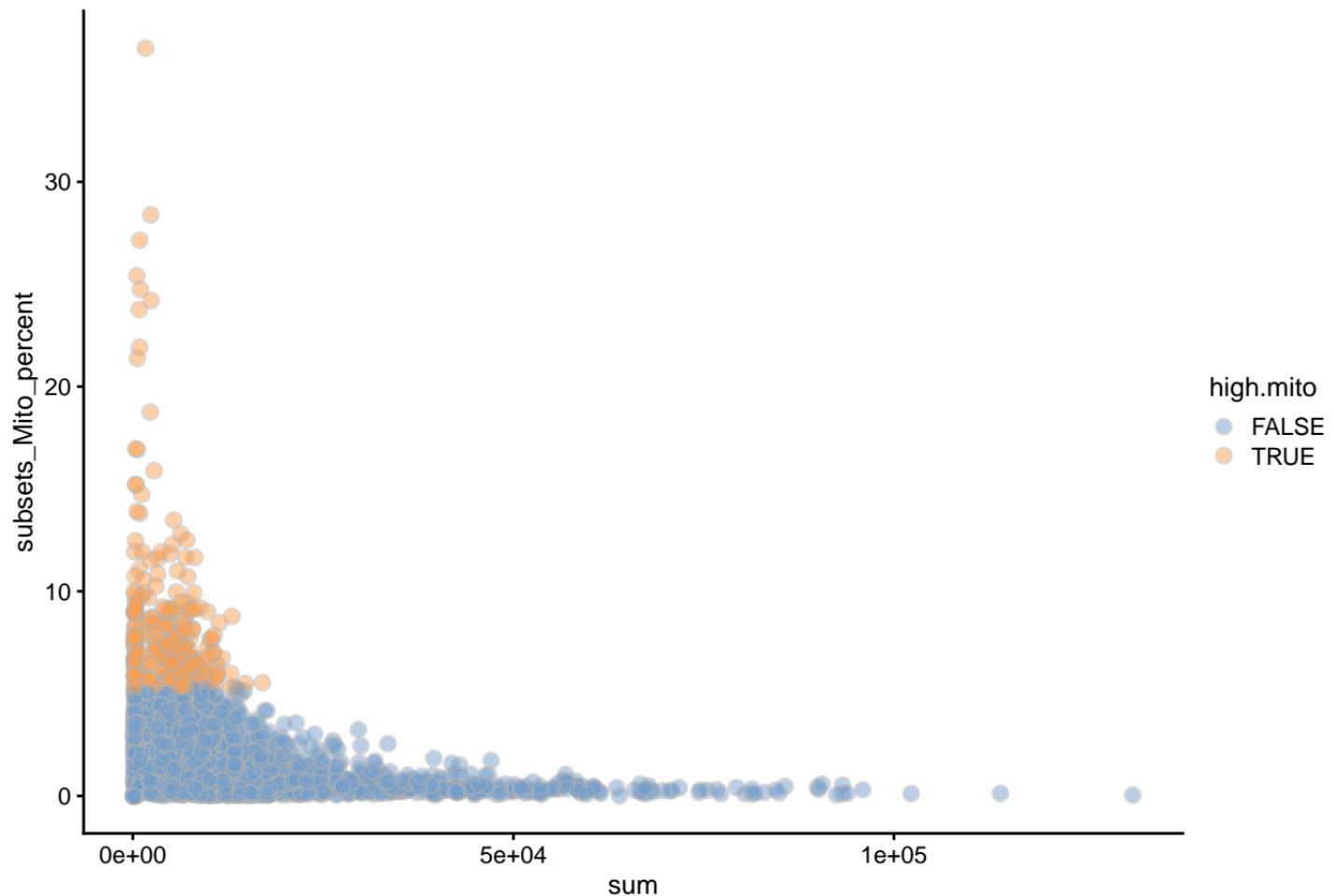
Total count: br5276.amy.neun

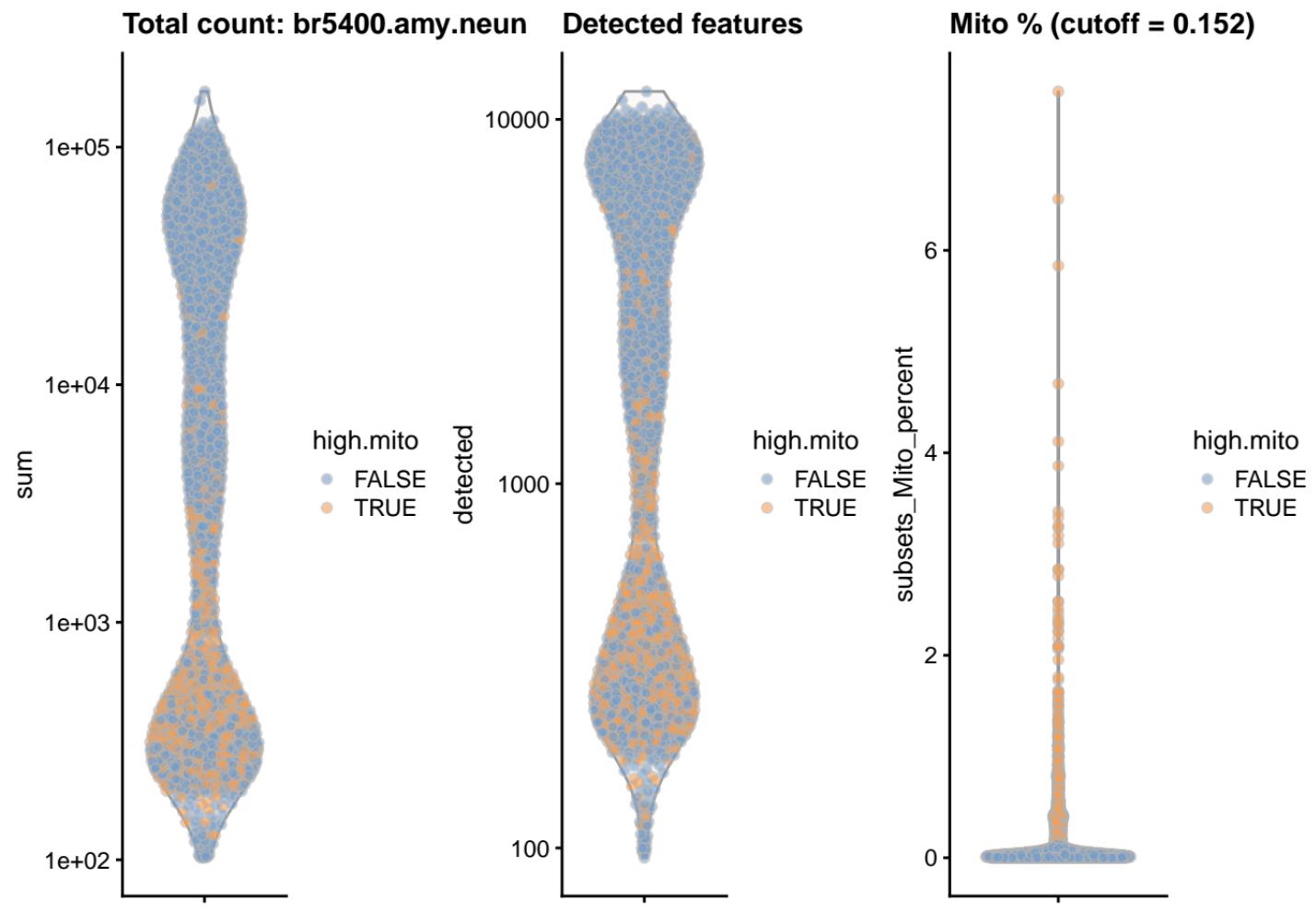
Detected features

Mito % (cutoff = 5.259)

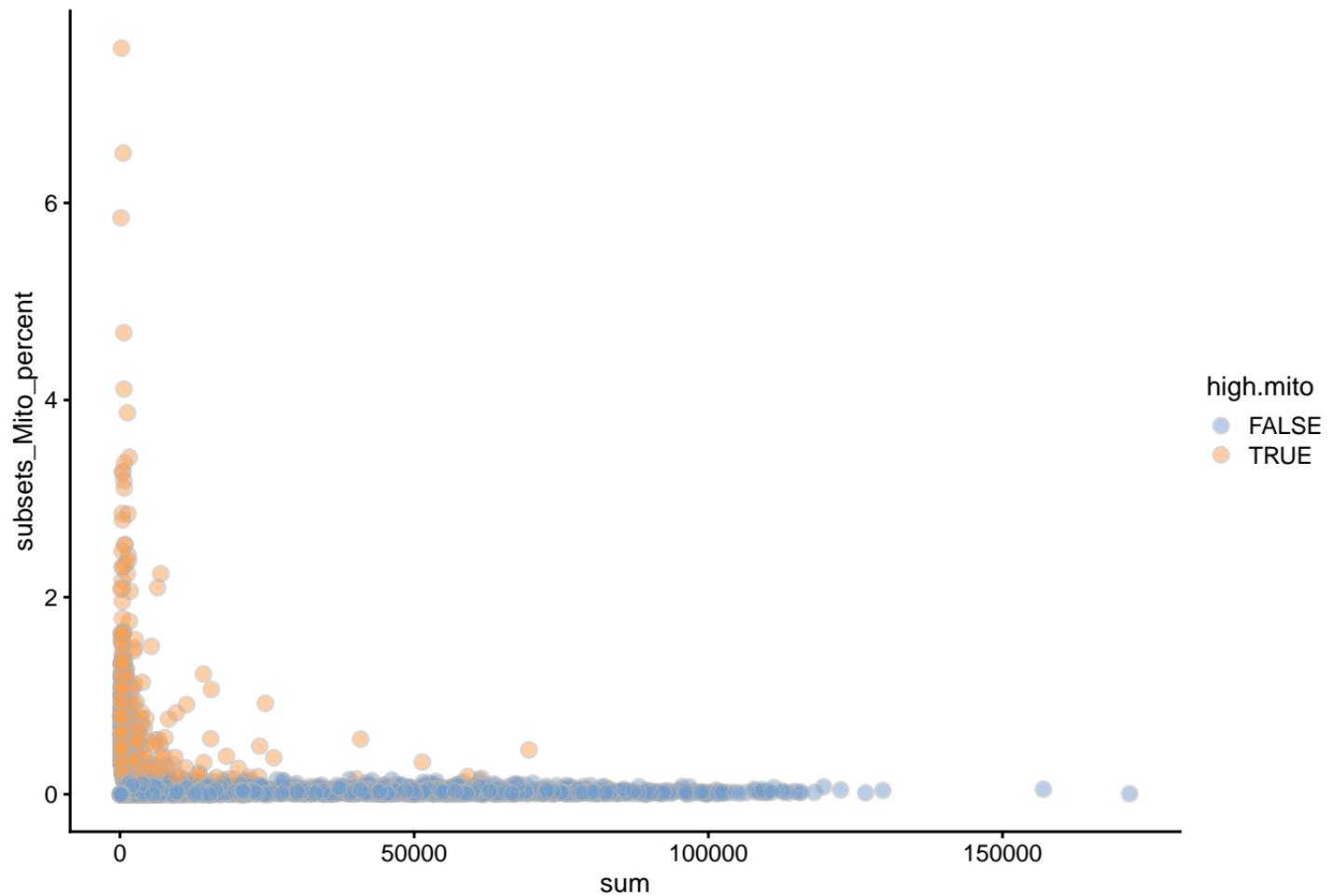


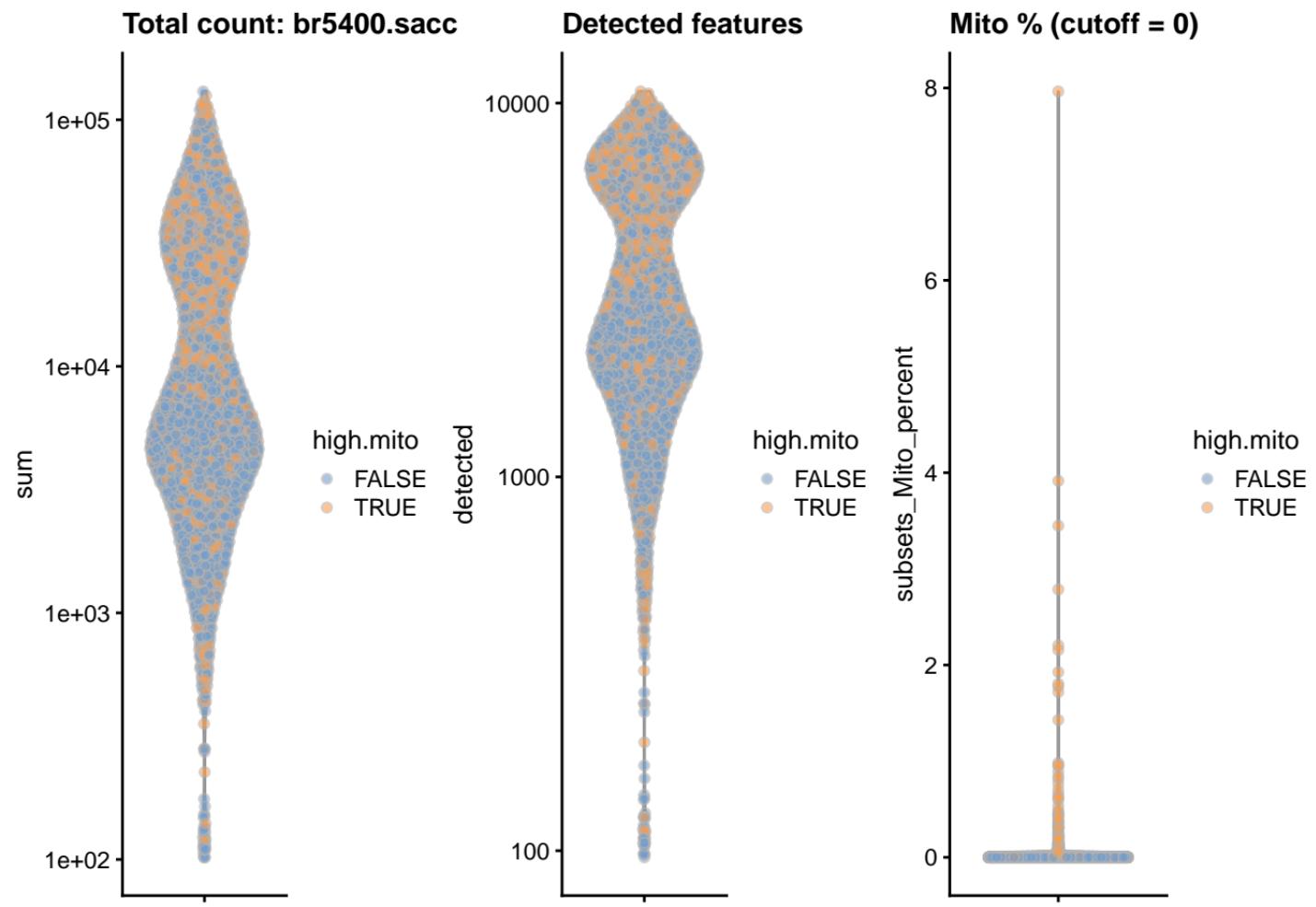
Sample: br5276.amy.neun; nNuclei (pre-QC): 2737; Mito % (cutoff = 5.259)



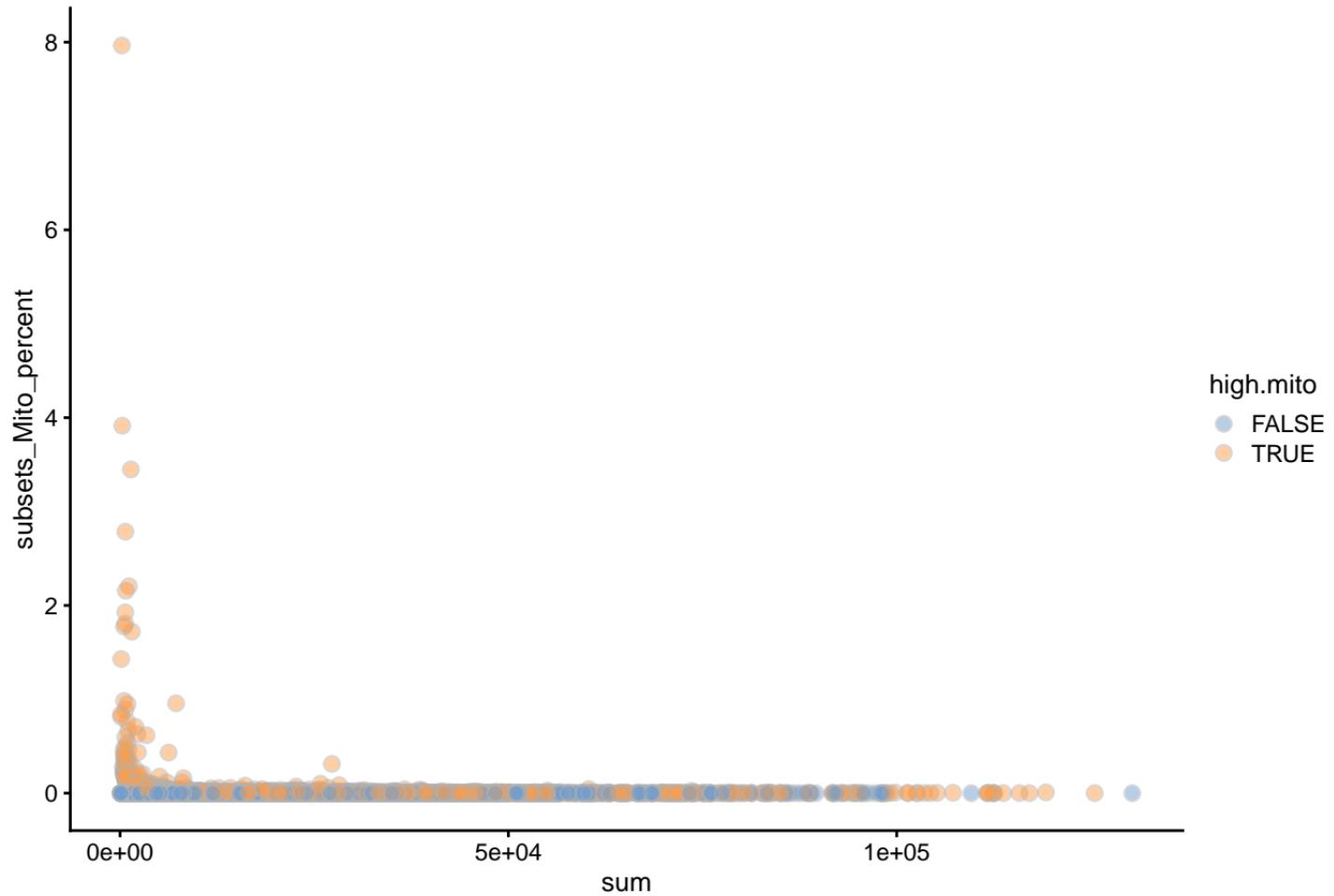


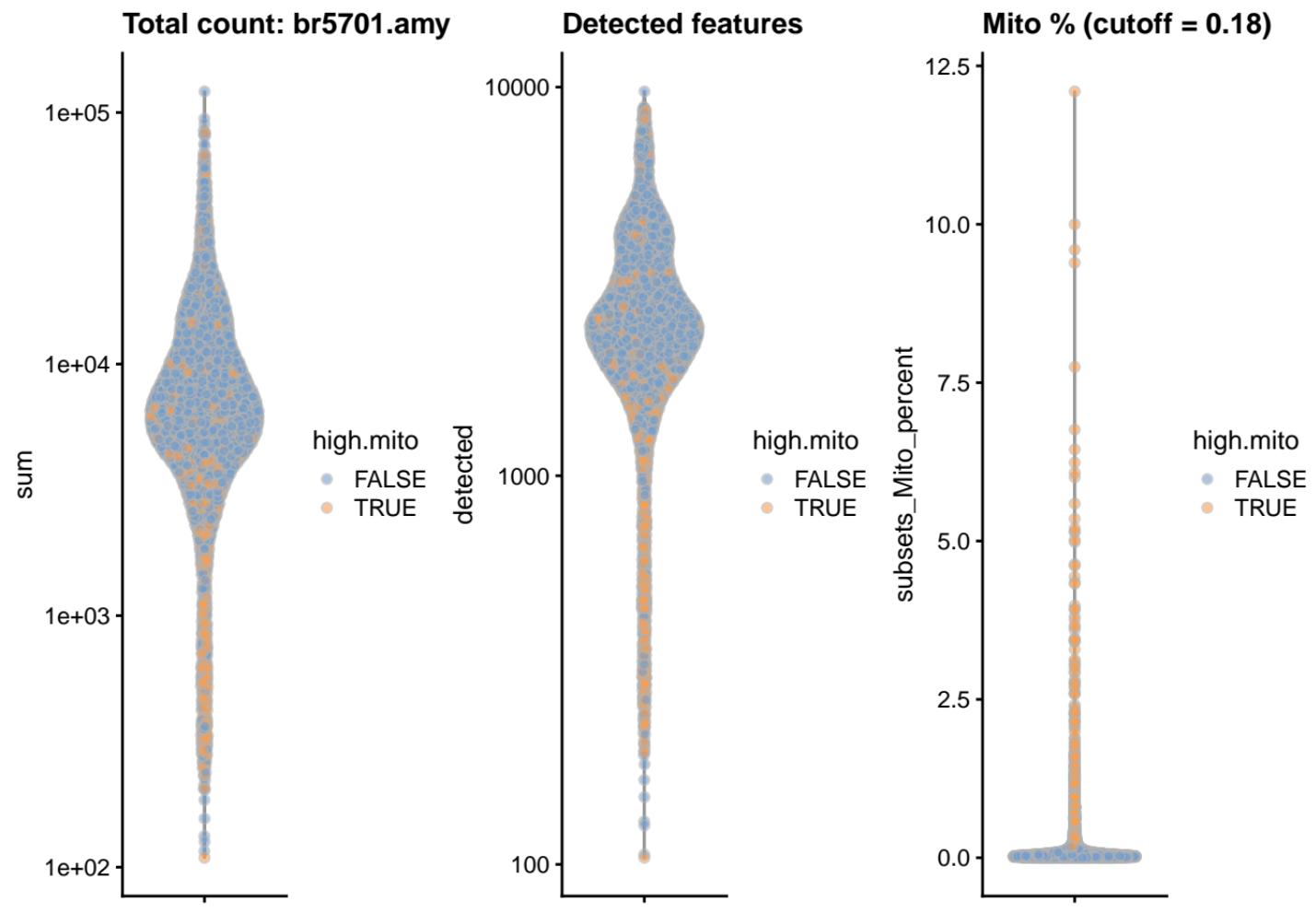
Sample: br5400.amy.neun; nNuclei (pre-QC): 3185; Mito % (cutoff = 0.152)





Sample: br5400.sacc; nNuclei (pre-QC): 4364; Mito % (cutoff = 0)





Sample: br5701.amy; nNuclei (pre-QC): 4195; Mito % (cutoff = 0.18)

