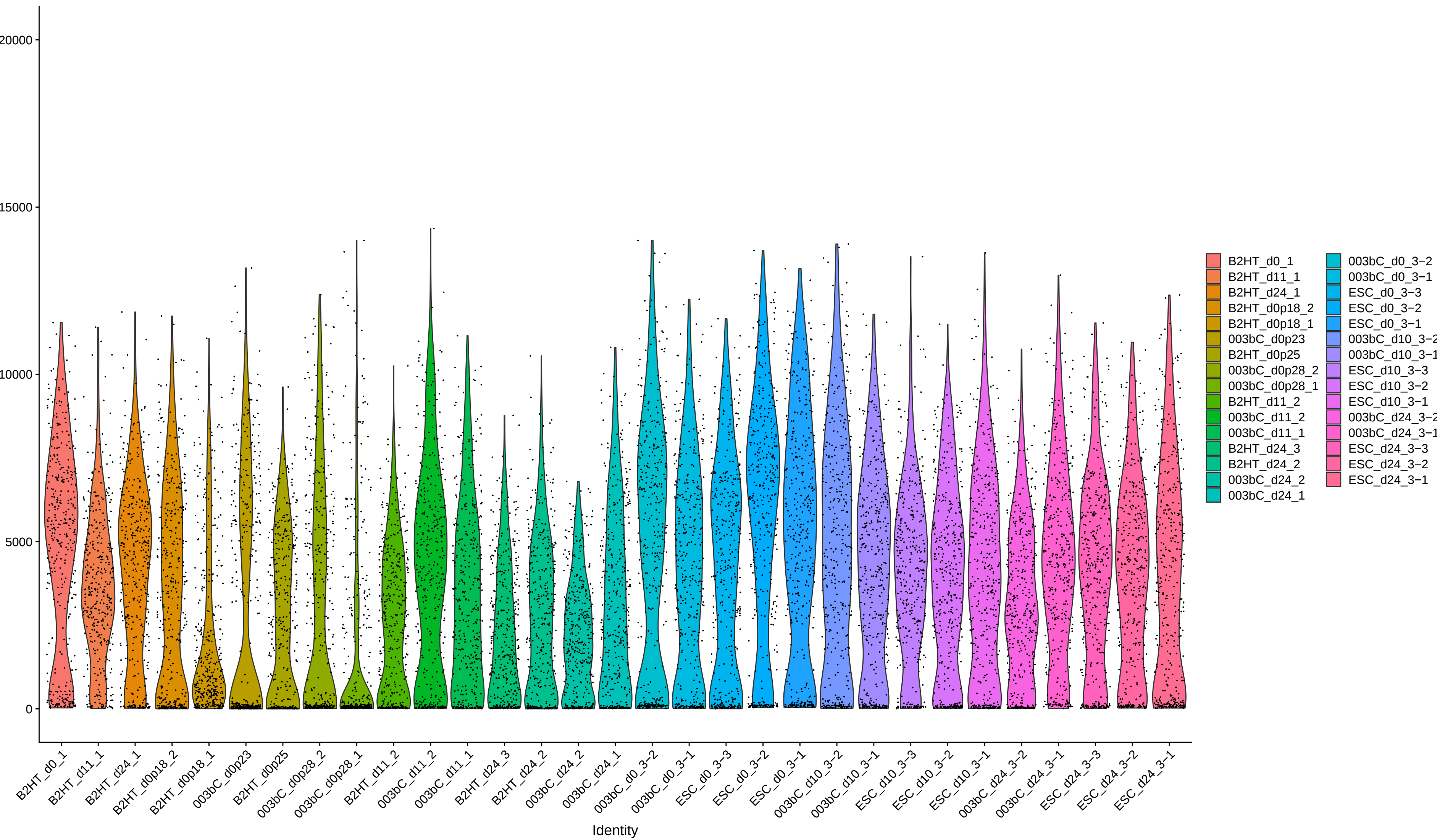
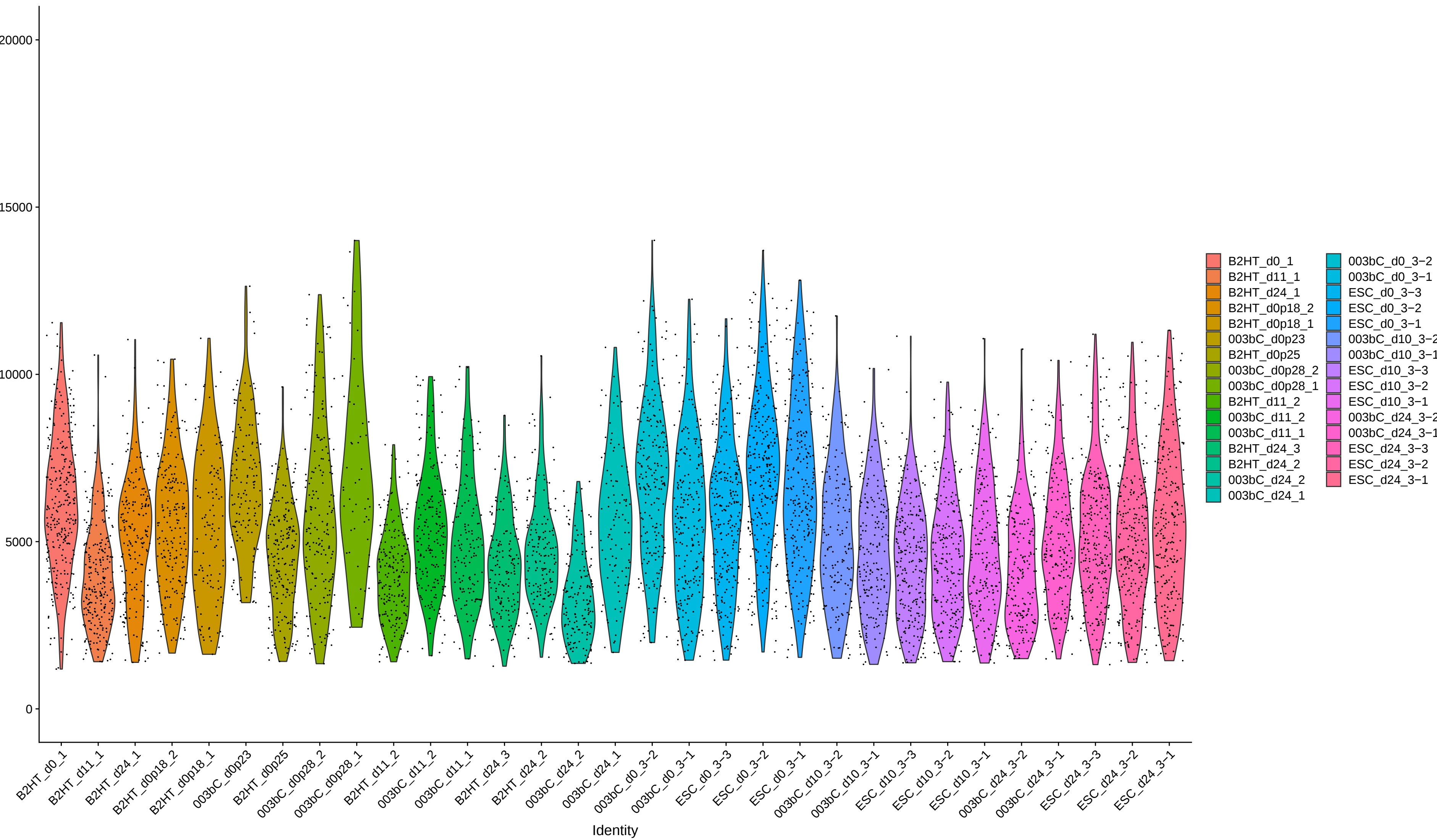


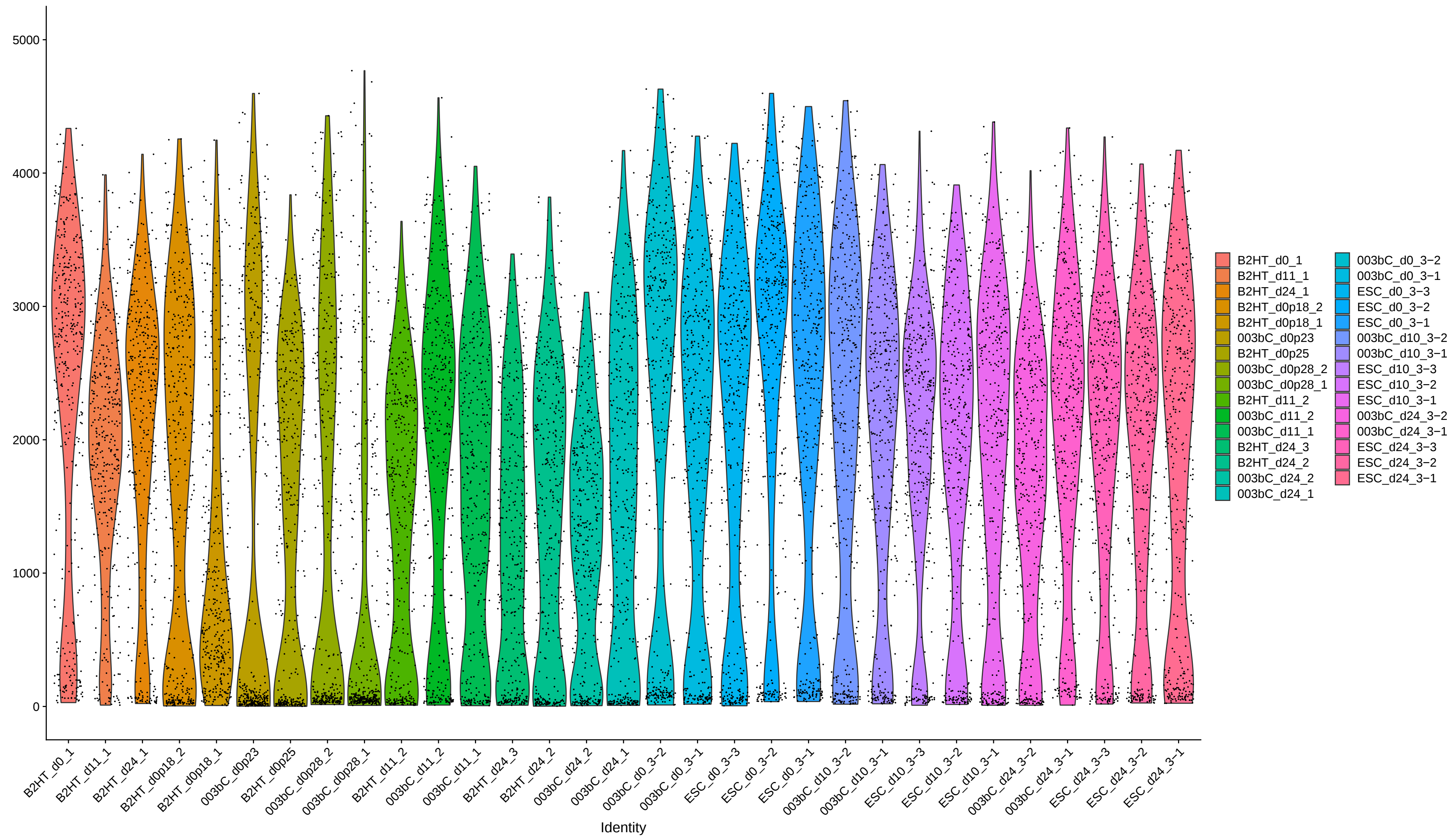
total UMI counts all cells



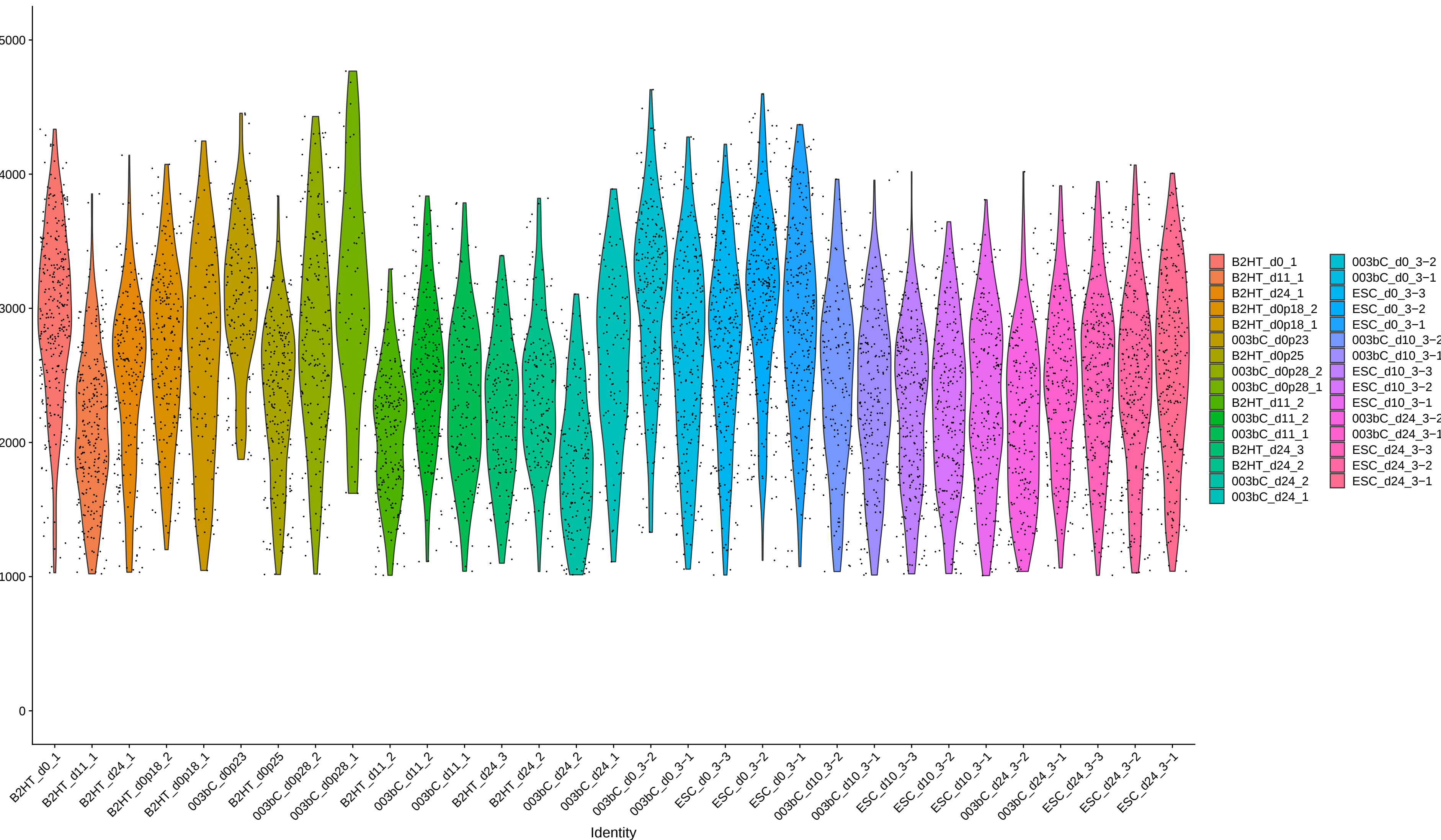
total UMI counts filtered cells



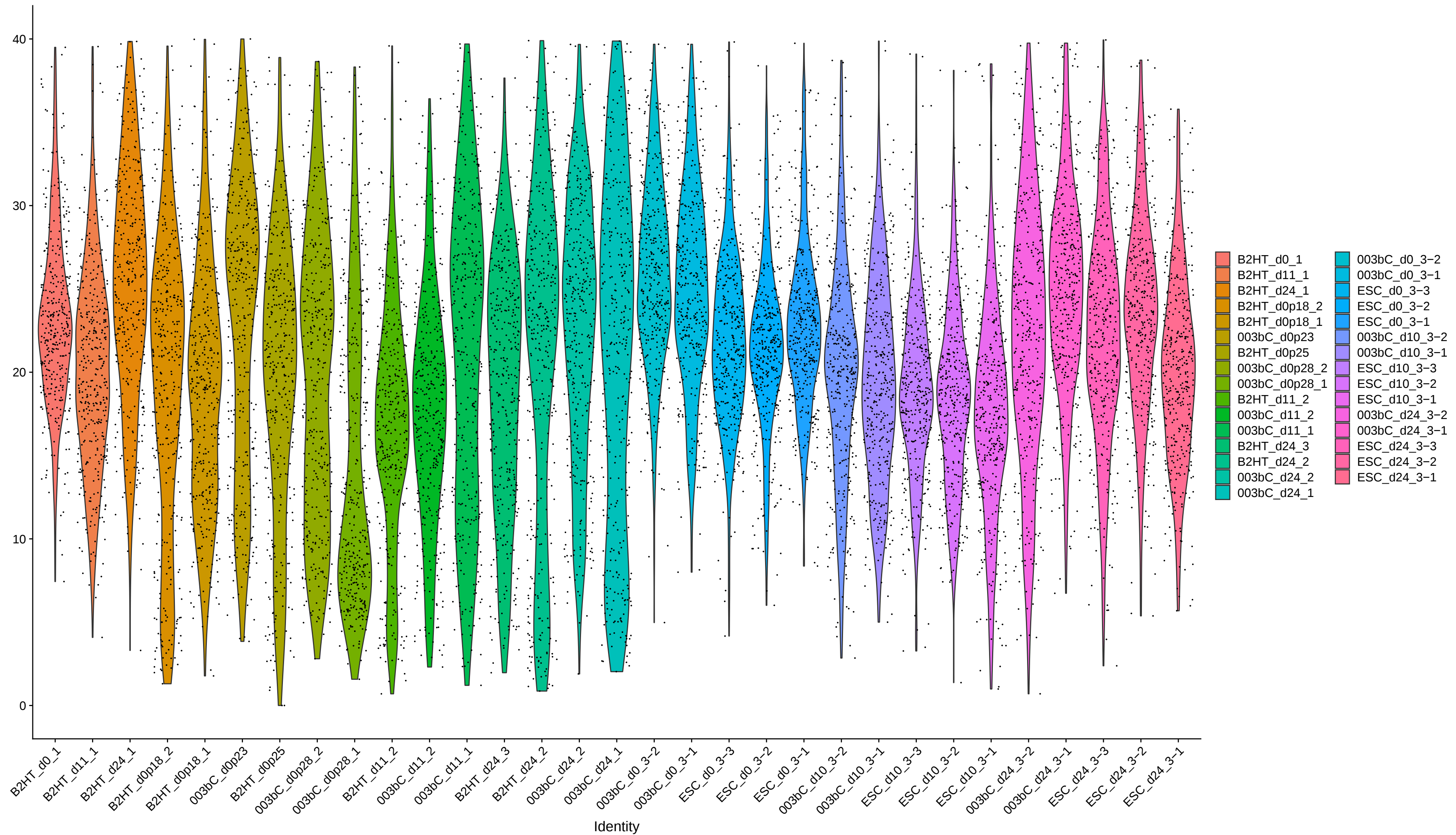
genes measured all cells



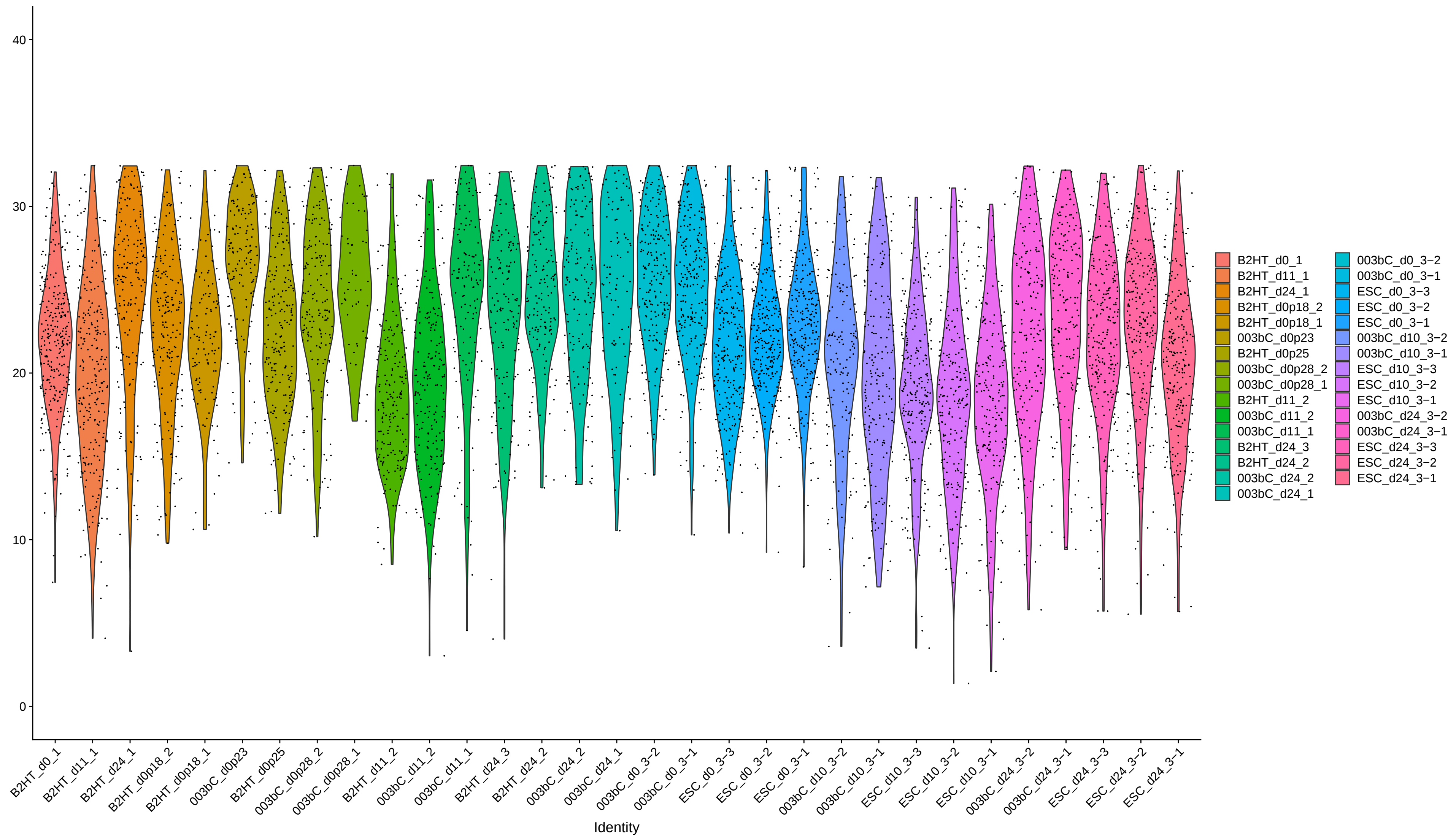
genes measured filtered cells



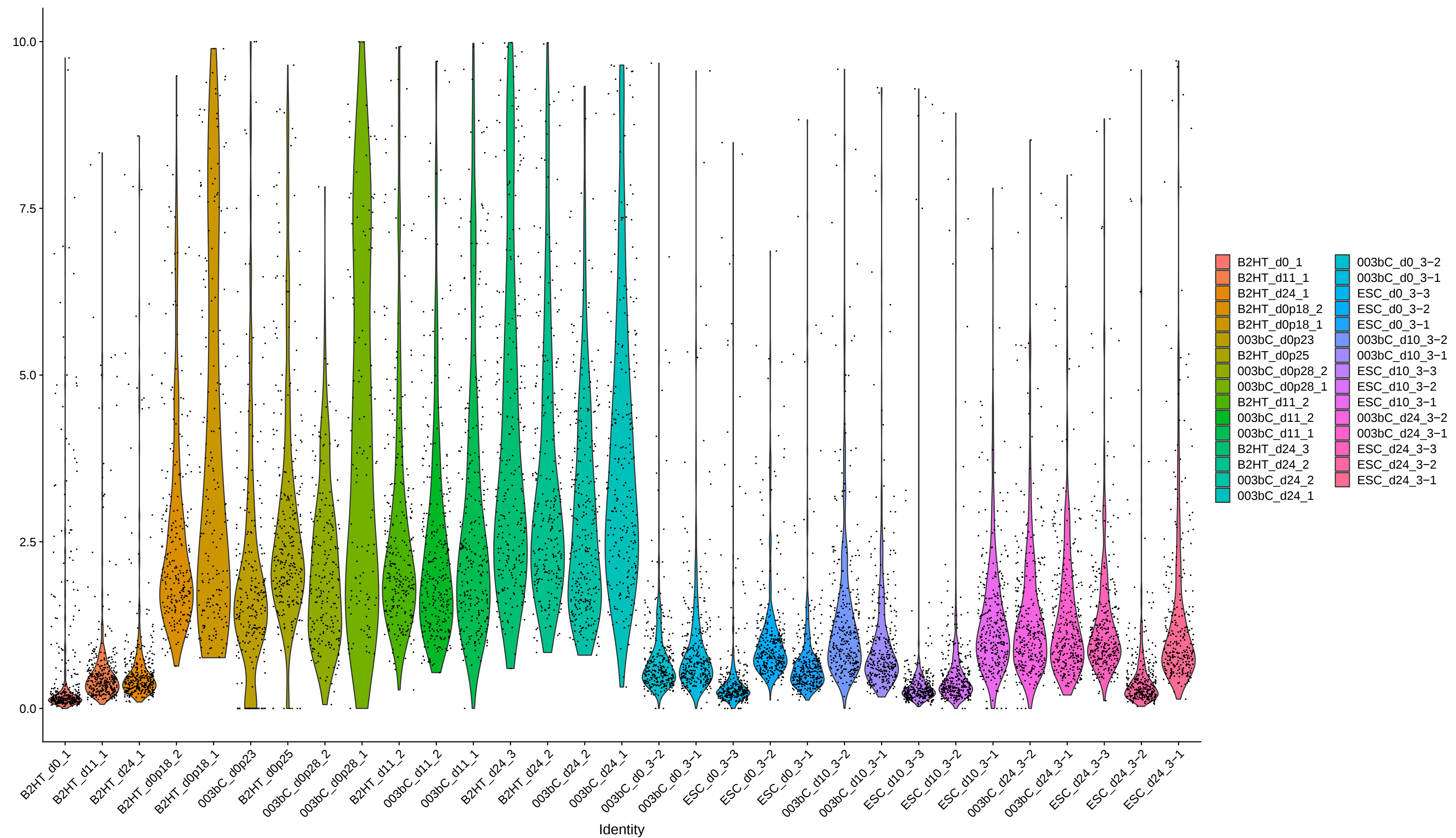
% mitochondrial reads all cells



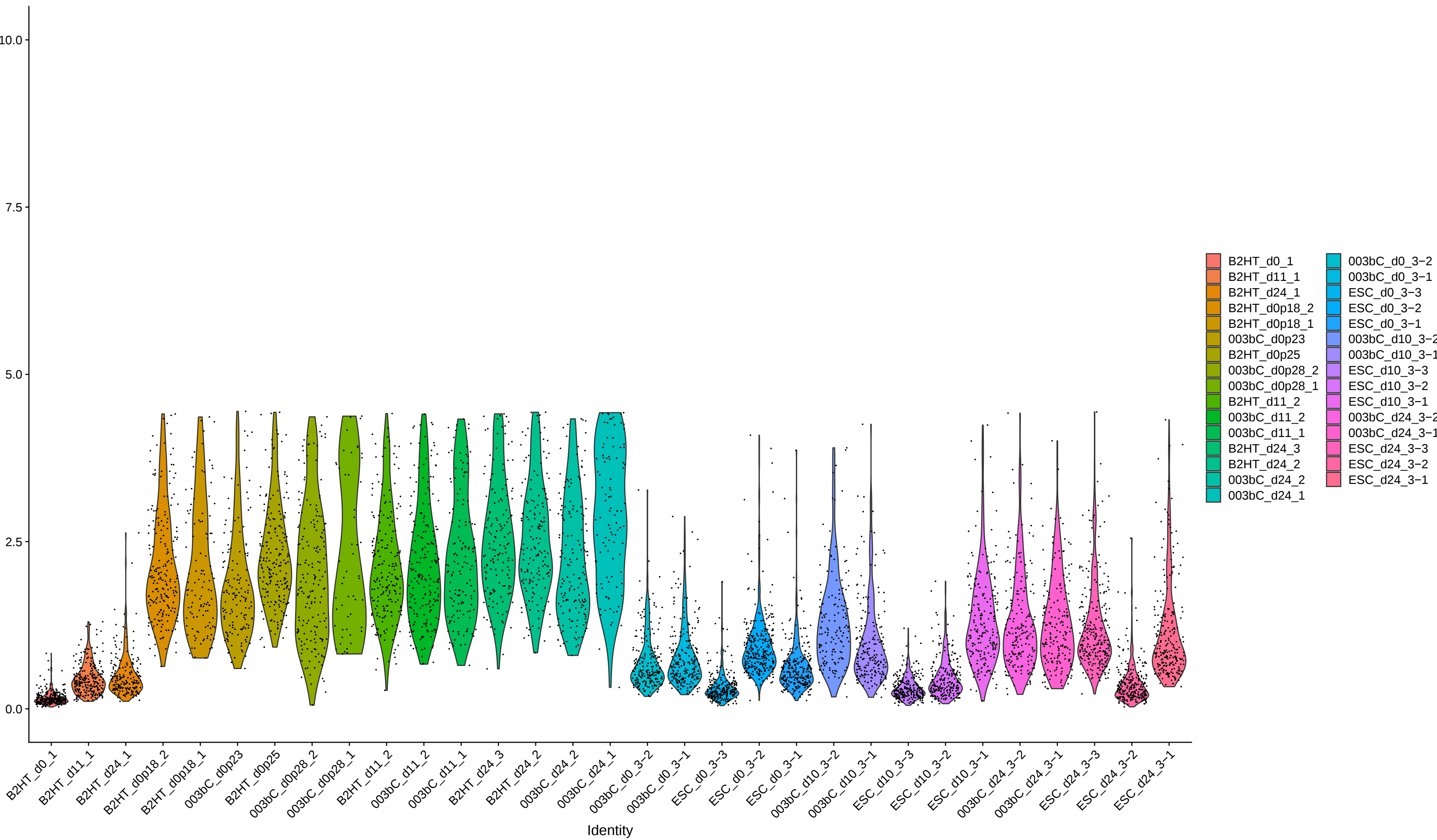
% mitochondrial reads filtered cells



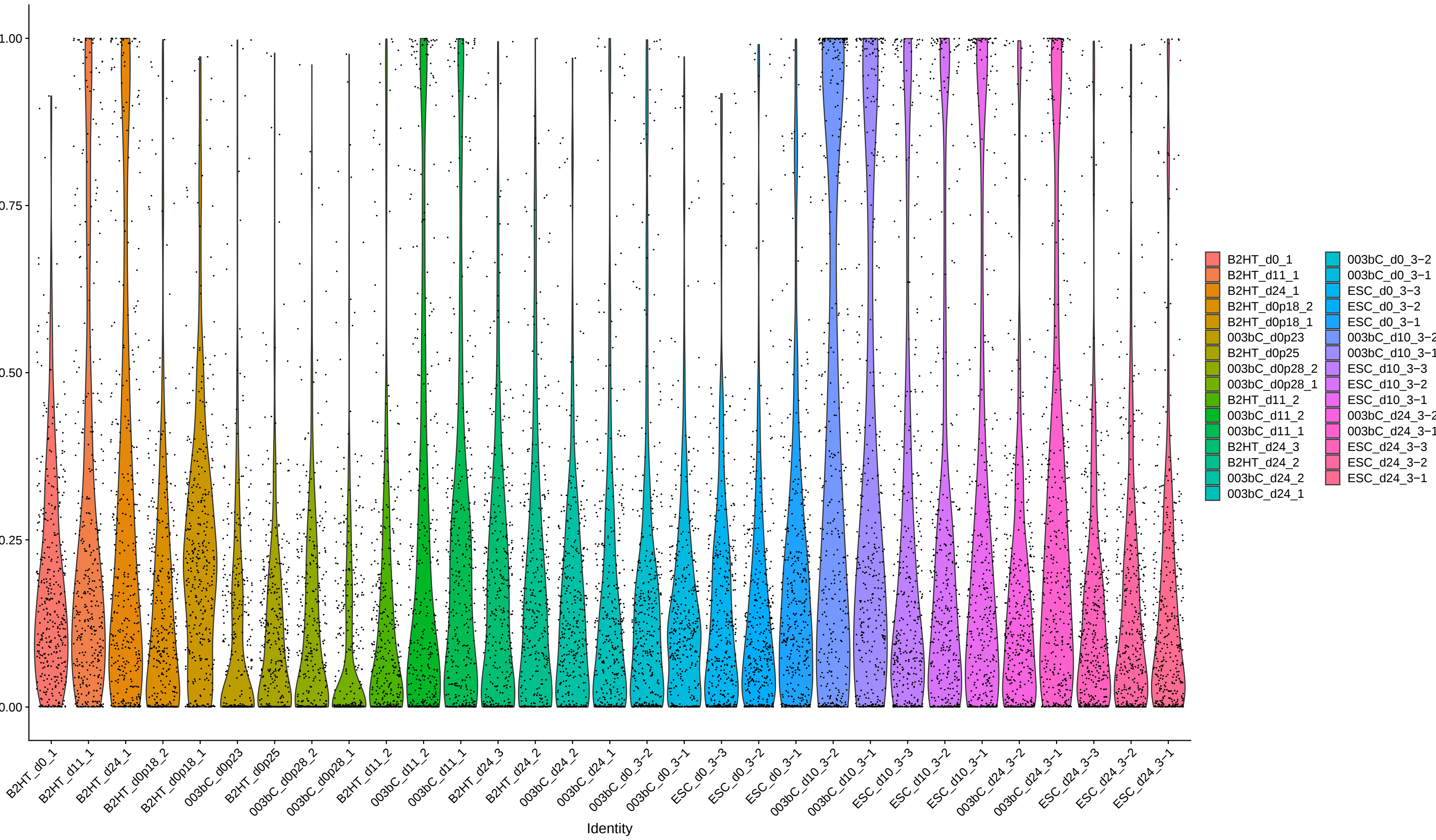
% ERCC reads all cells



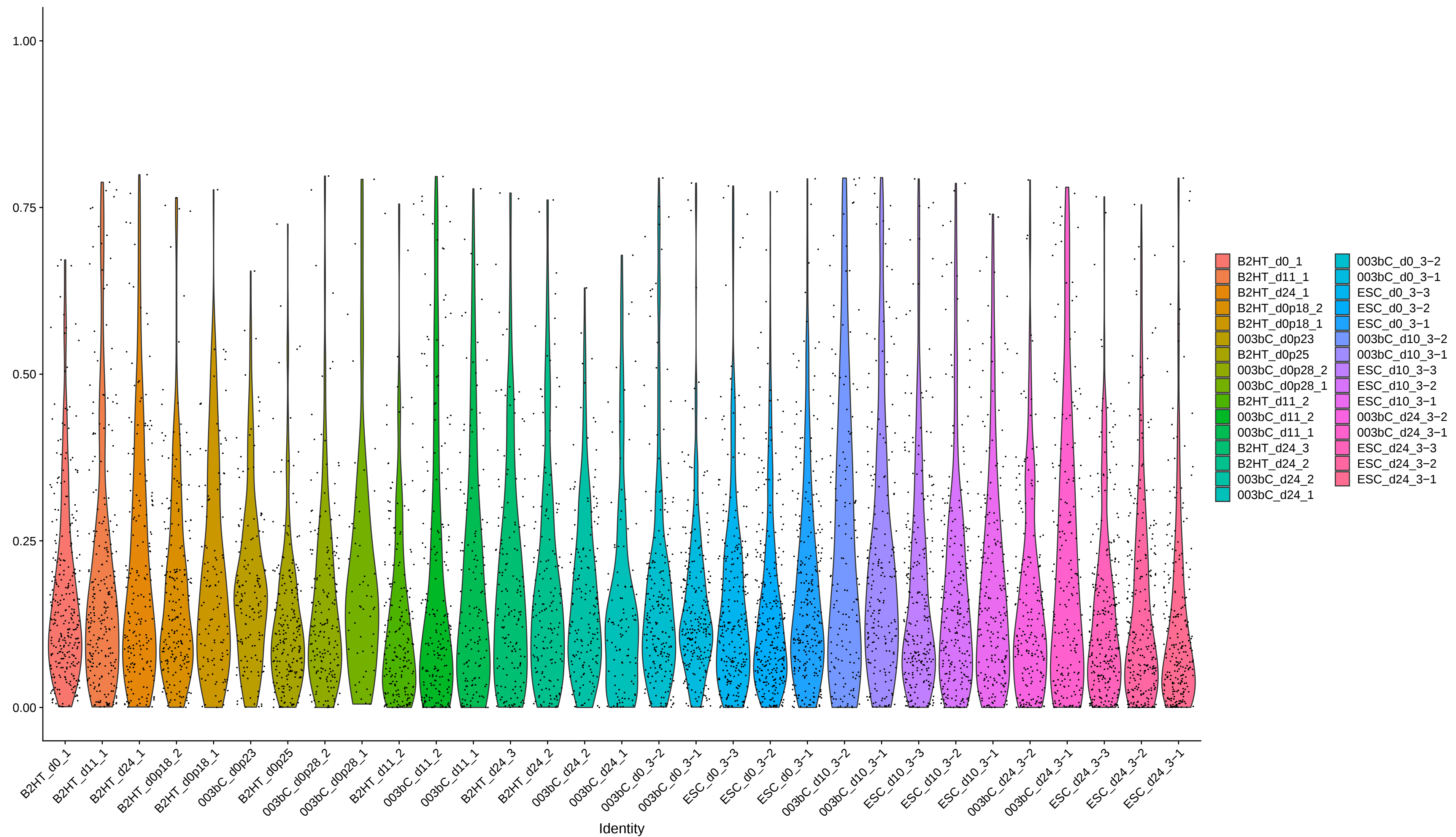
% ERCC reads filtered cells



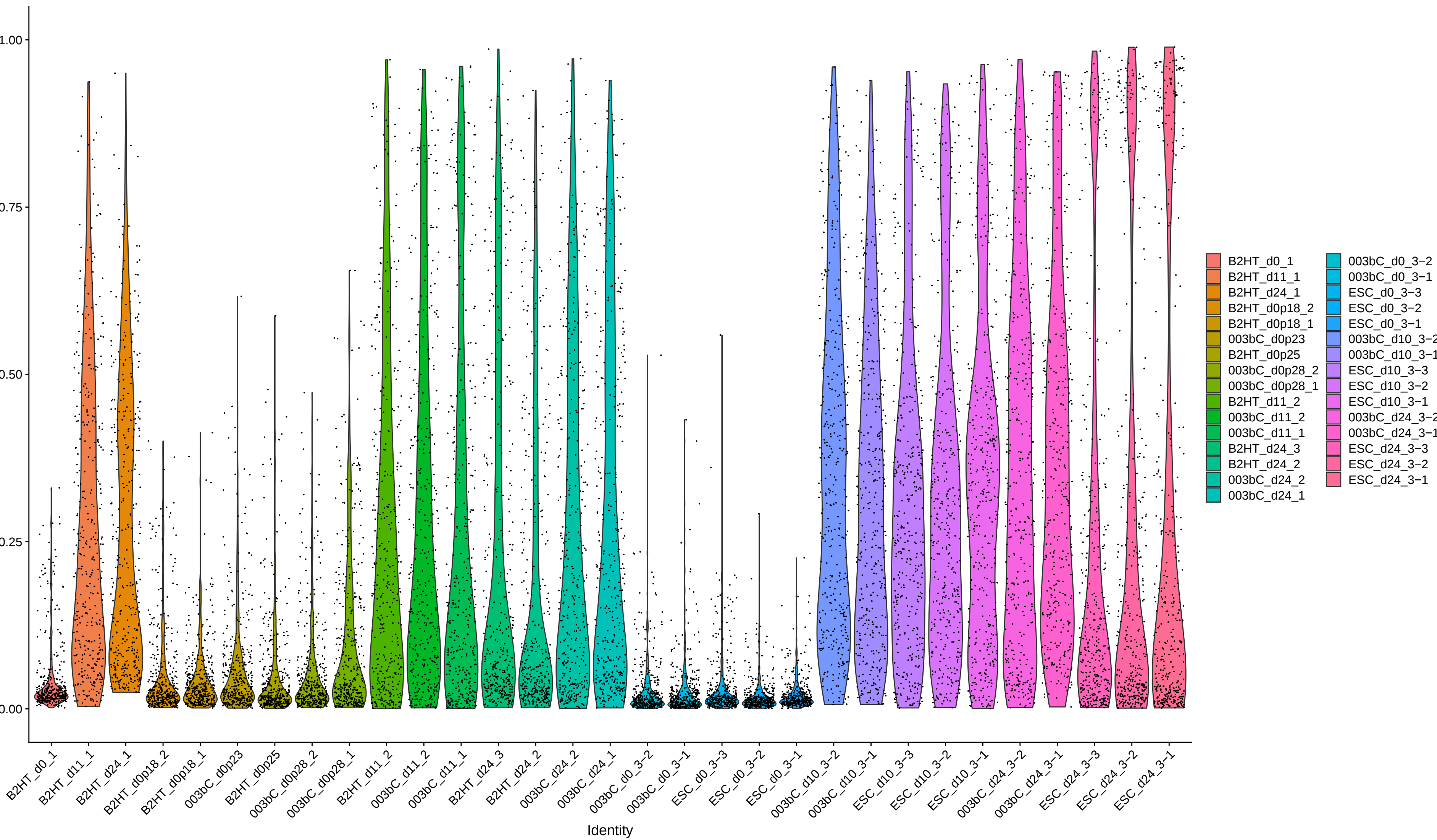
scDbIFinder_doublet_score



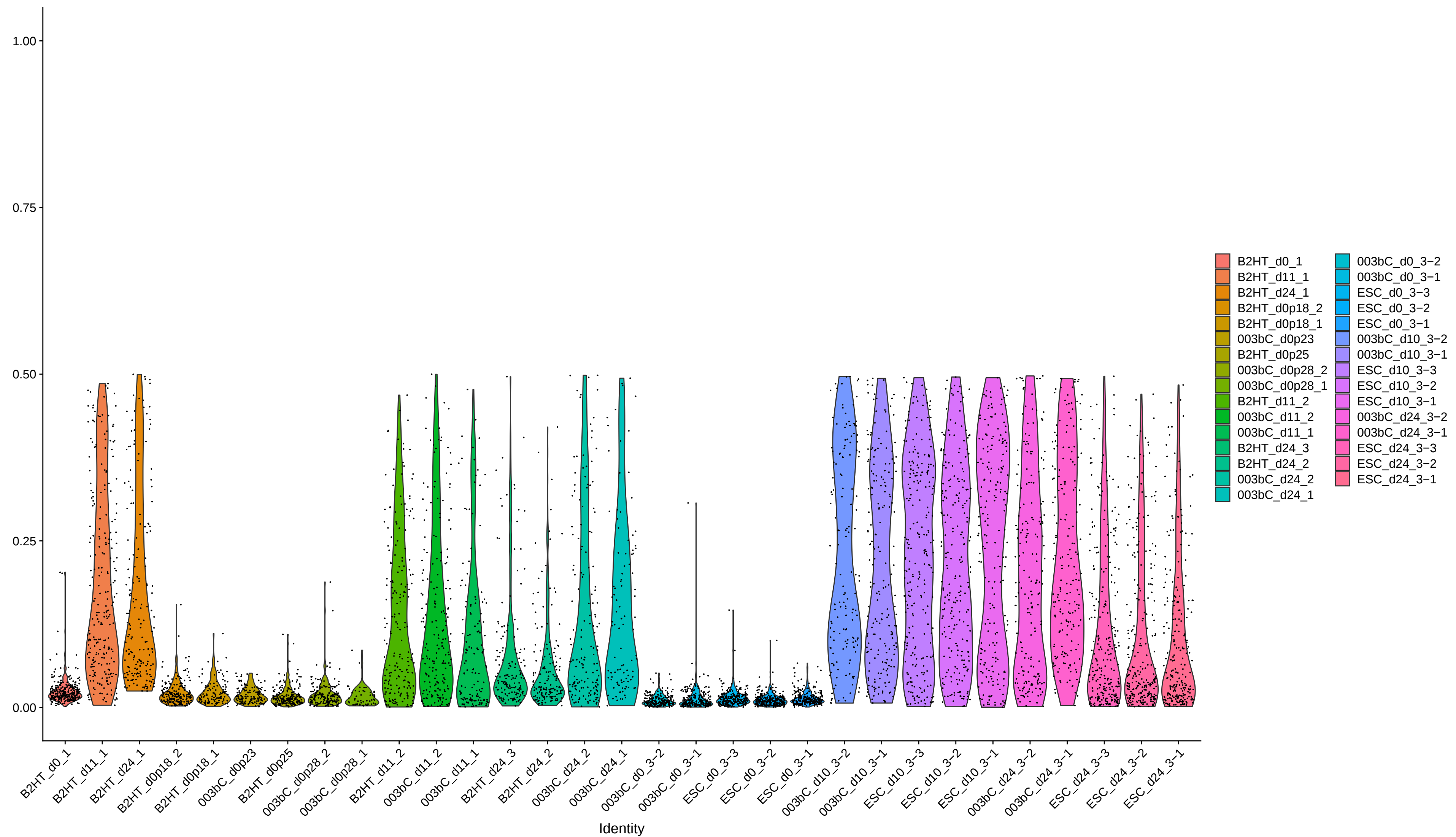
scDbIFinder_doublet_score_filtered



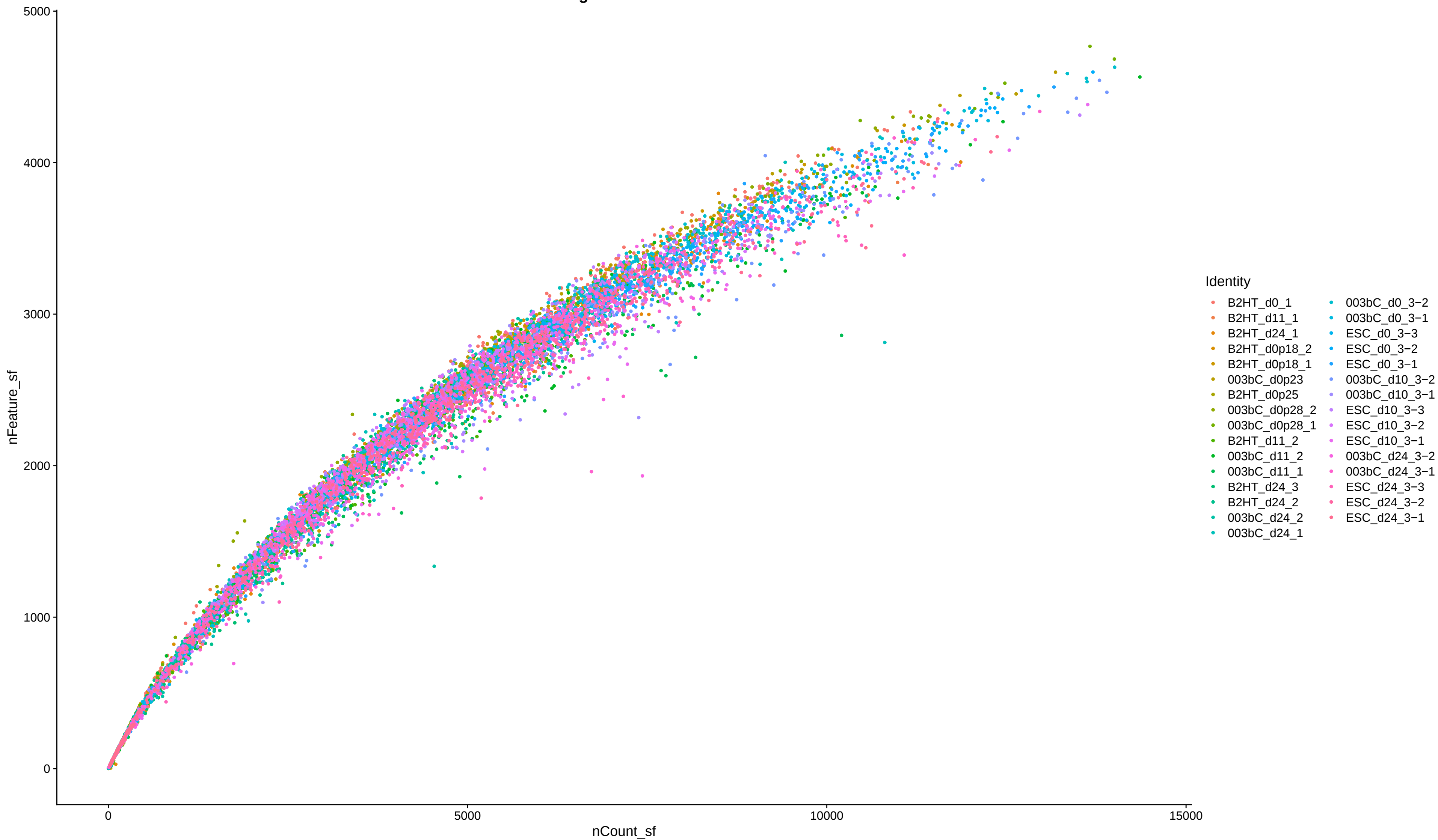
decontX_contamination



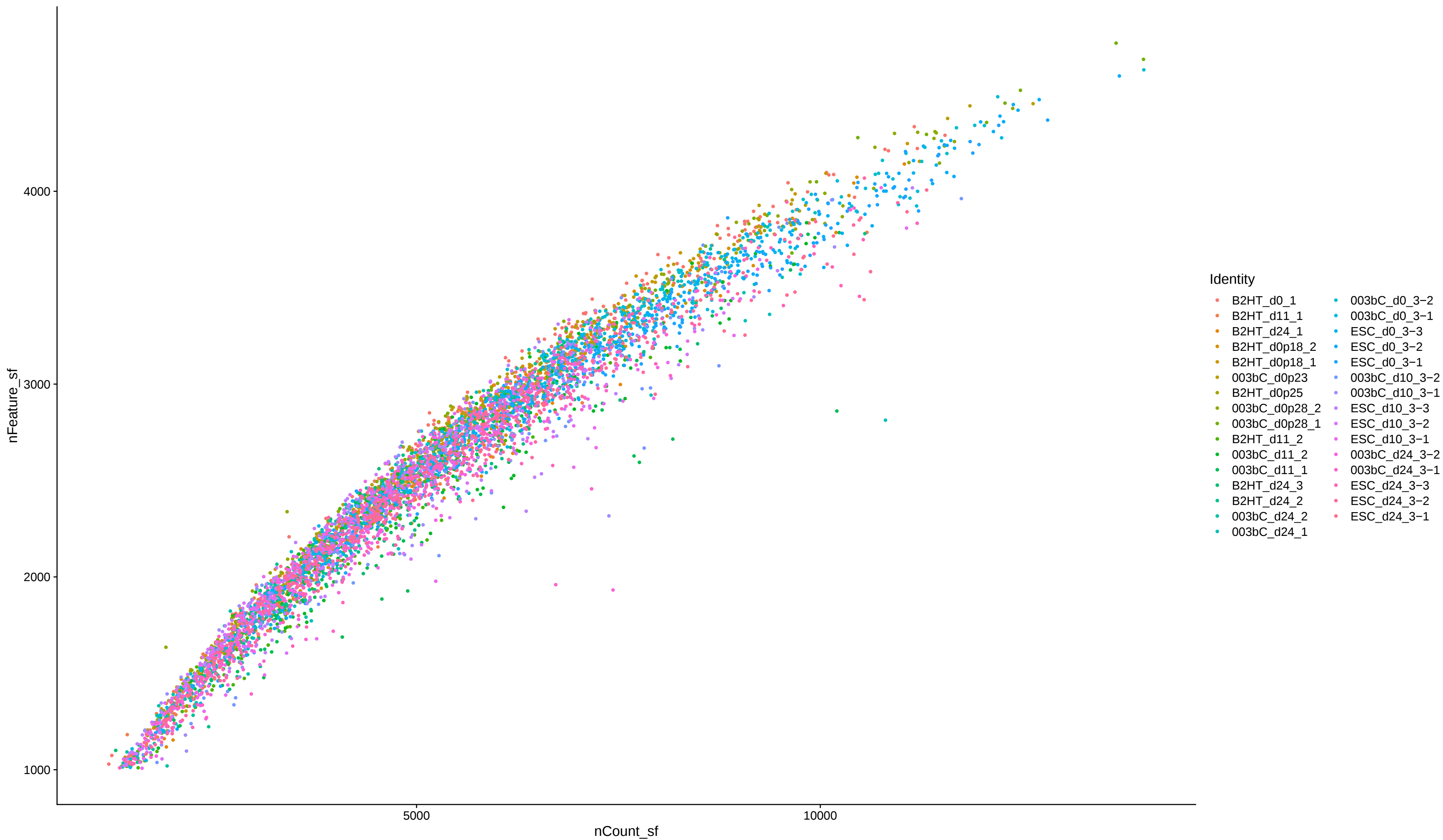
decontX_contamination_filtered



% counts vs genes measured all cells



% counts vs genes measured filtered cells



	descriptive_name	cells after QC
1	003bC_d0	825
2	003bC_d11	690
3	003bC_d24	649
4	B2HT_d0	817
5	B2HT_d11	453
6	B2HT_d24	502
7	ESC_d0	805
8	ESC_d11	690
9	ESC_d24	770