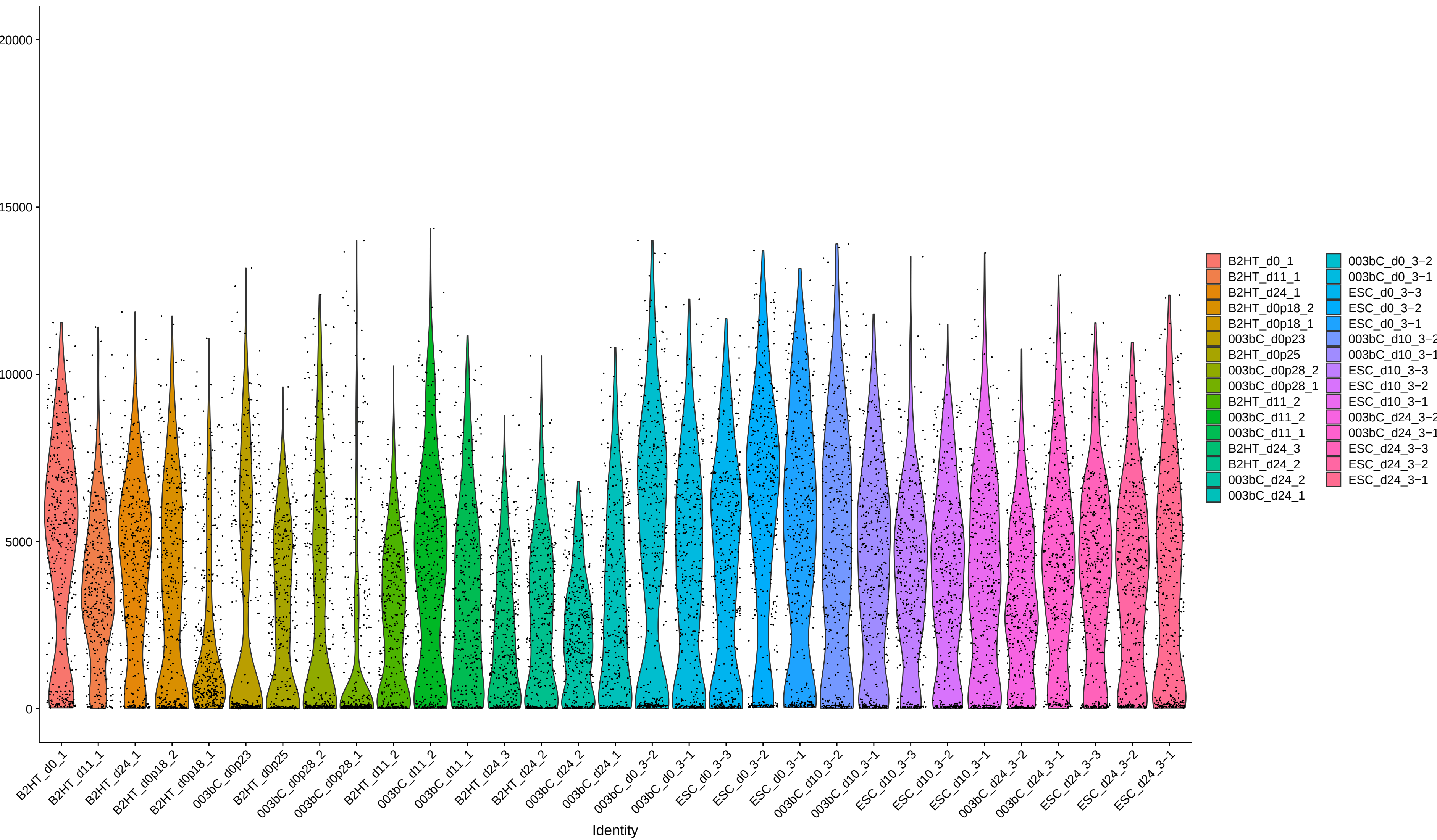
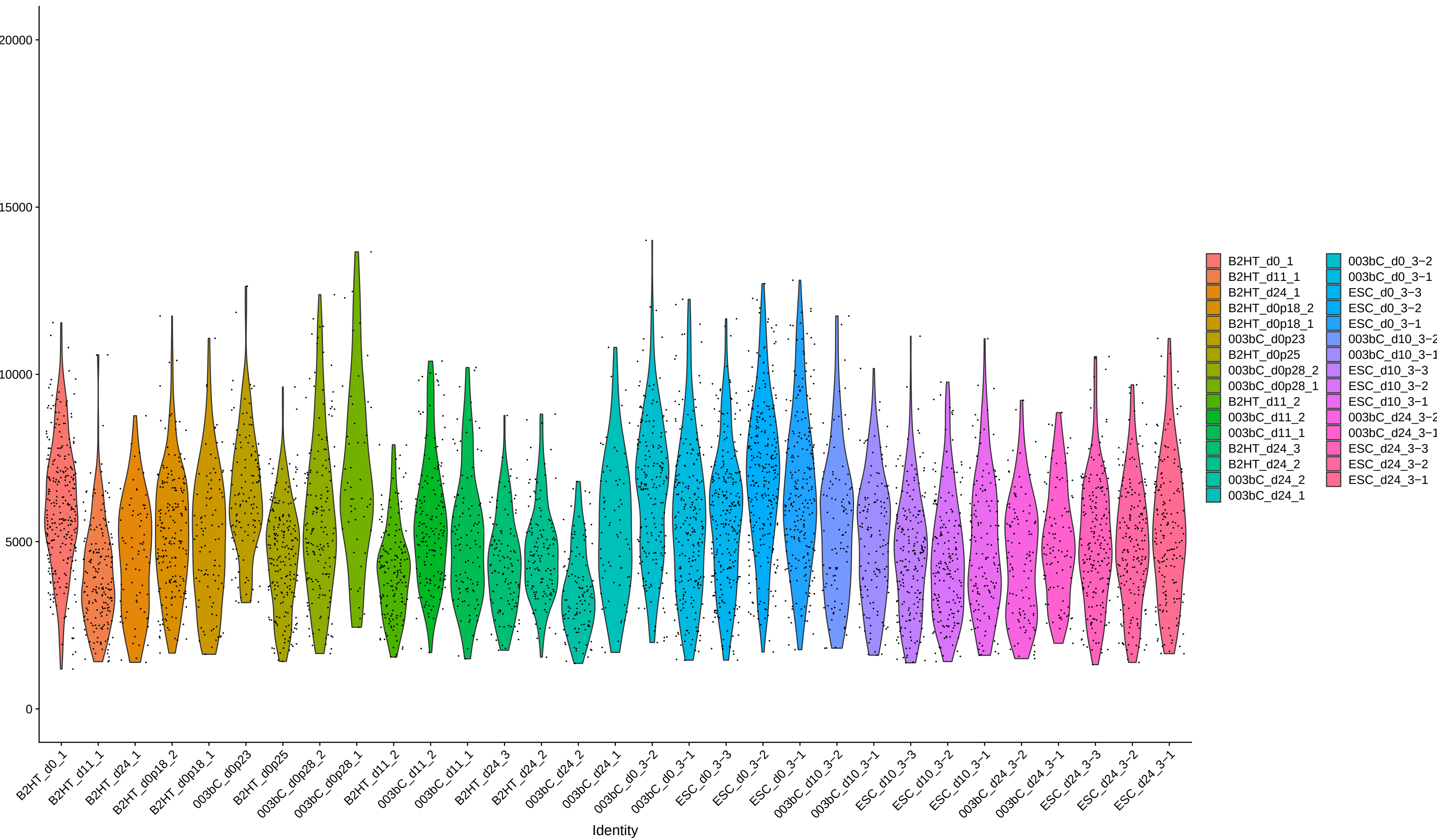


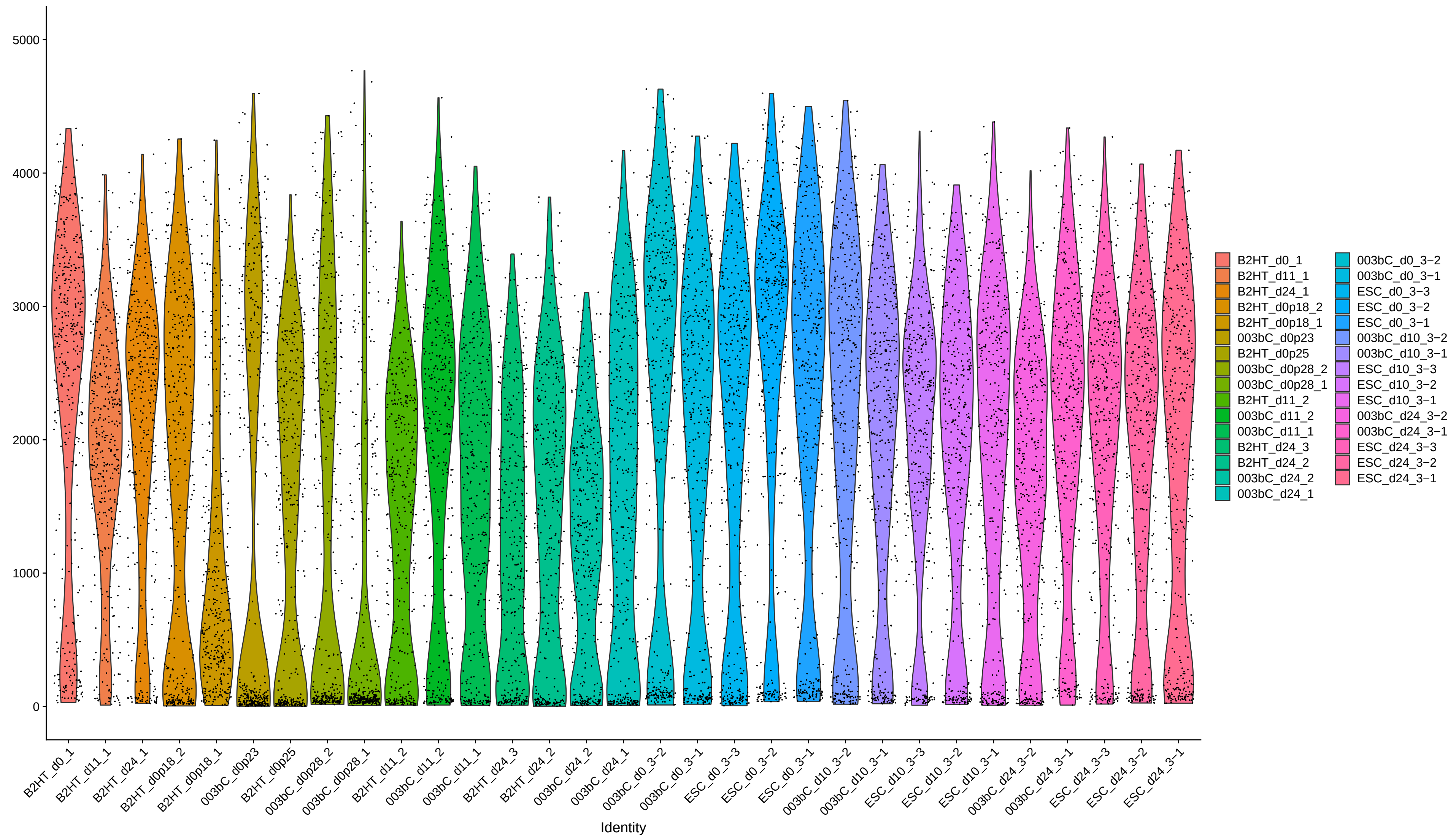
total UMI counts all cells



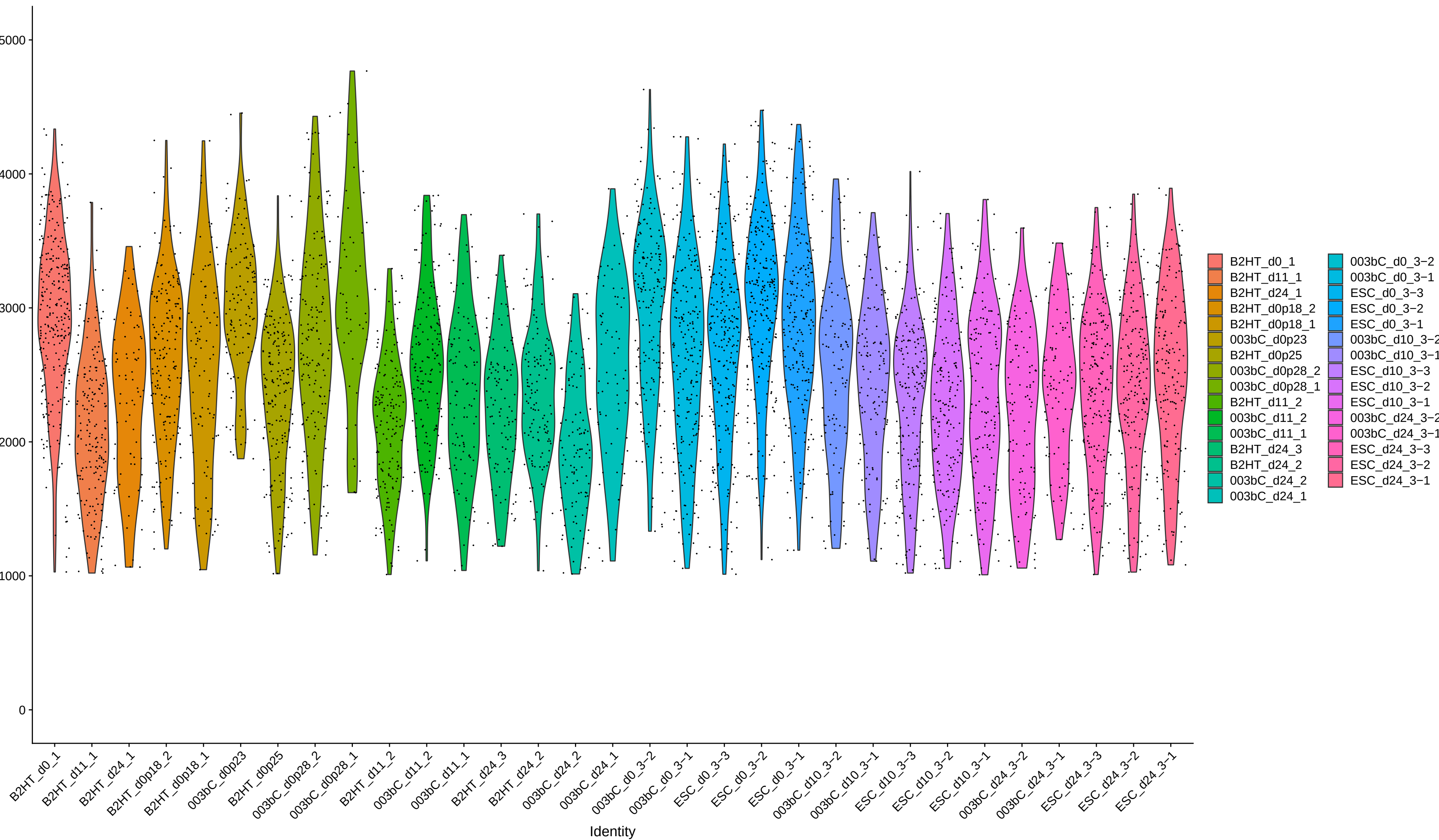
total UMI counts filtered cells



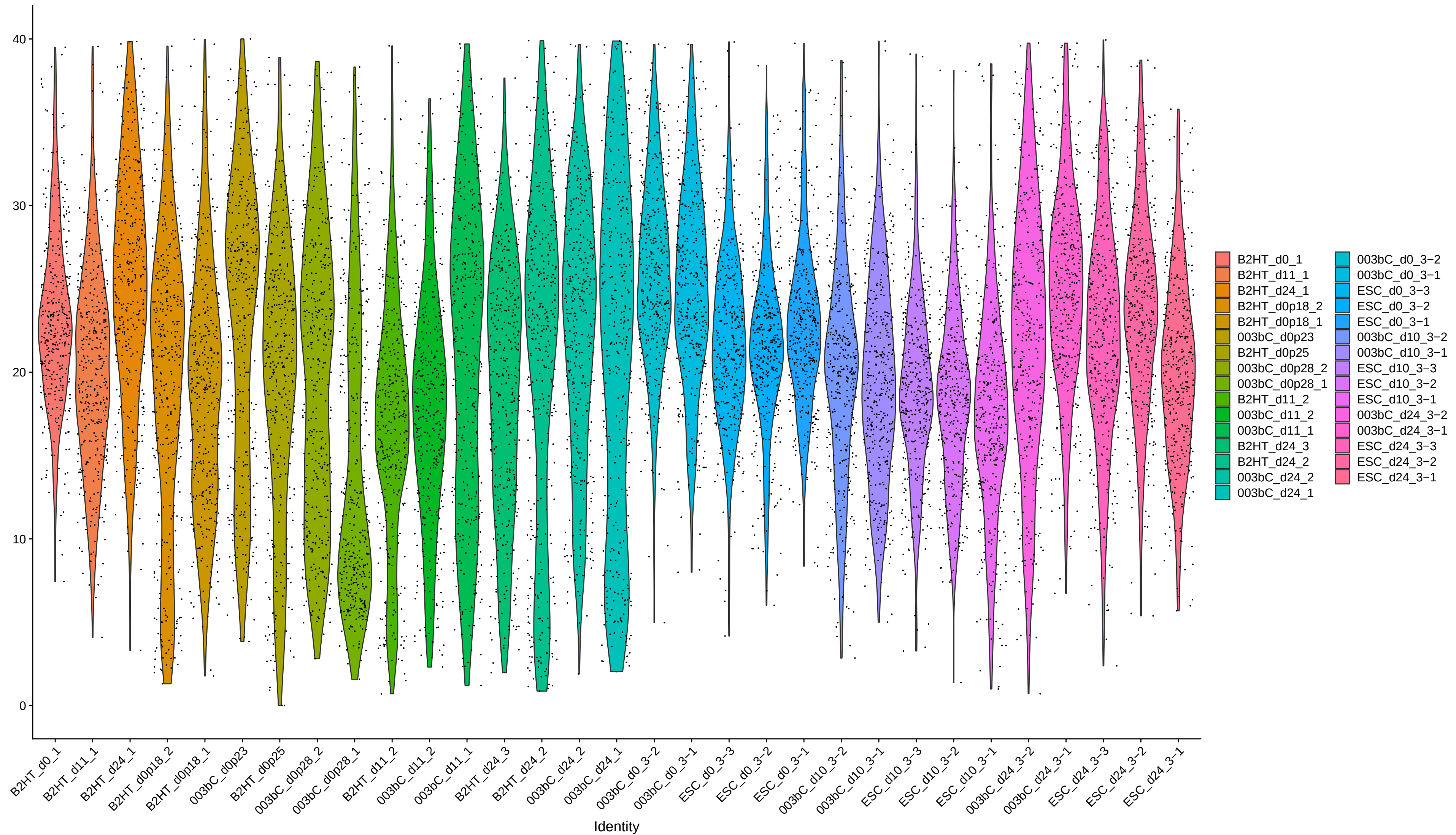
genes measured all cells



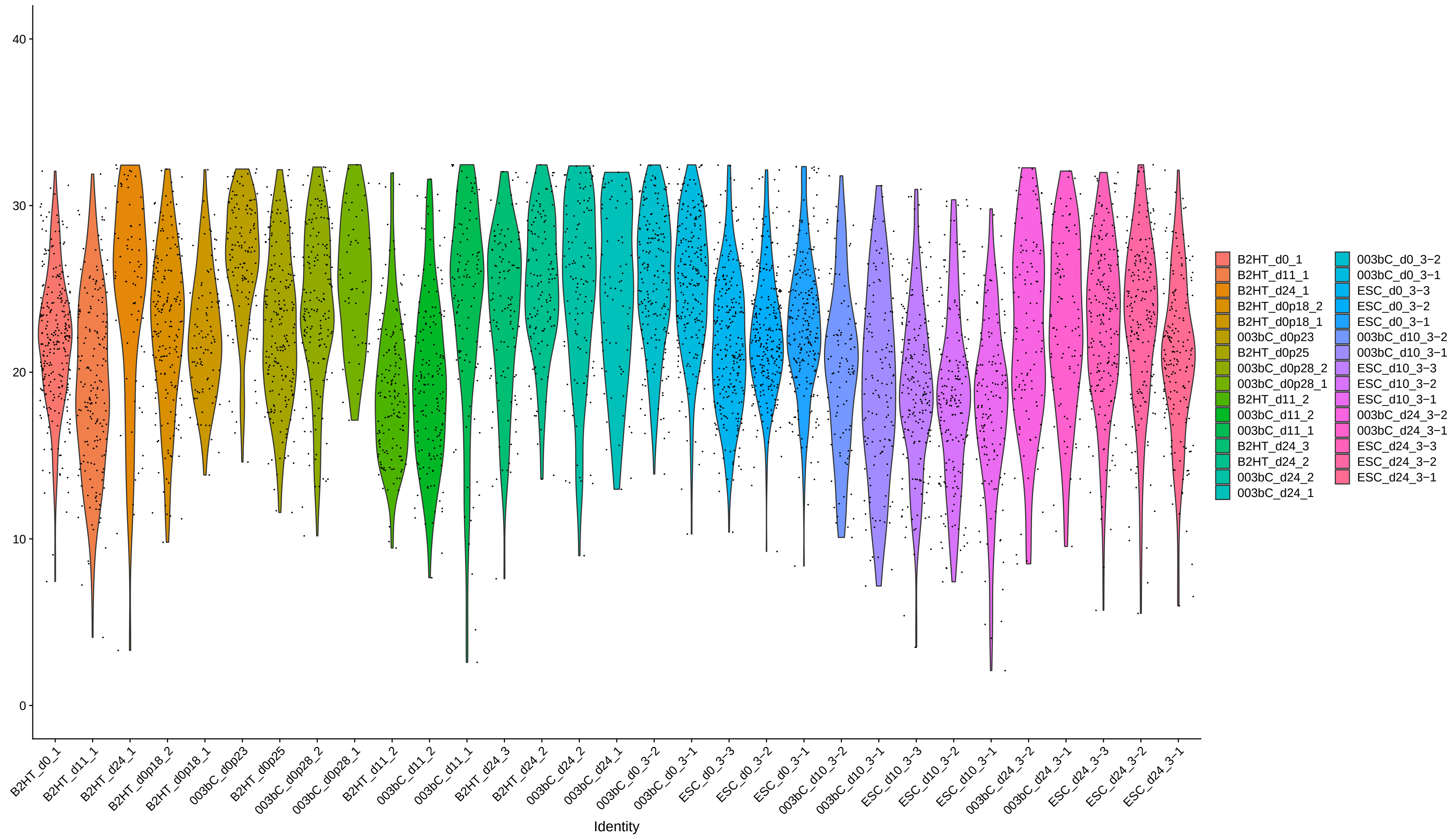
genes measured filtered cells



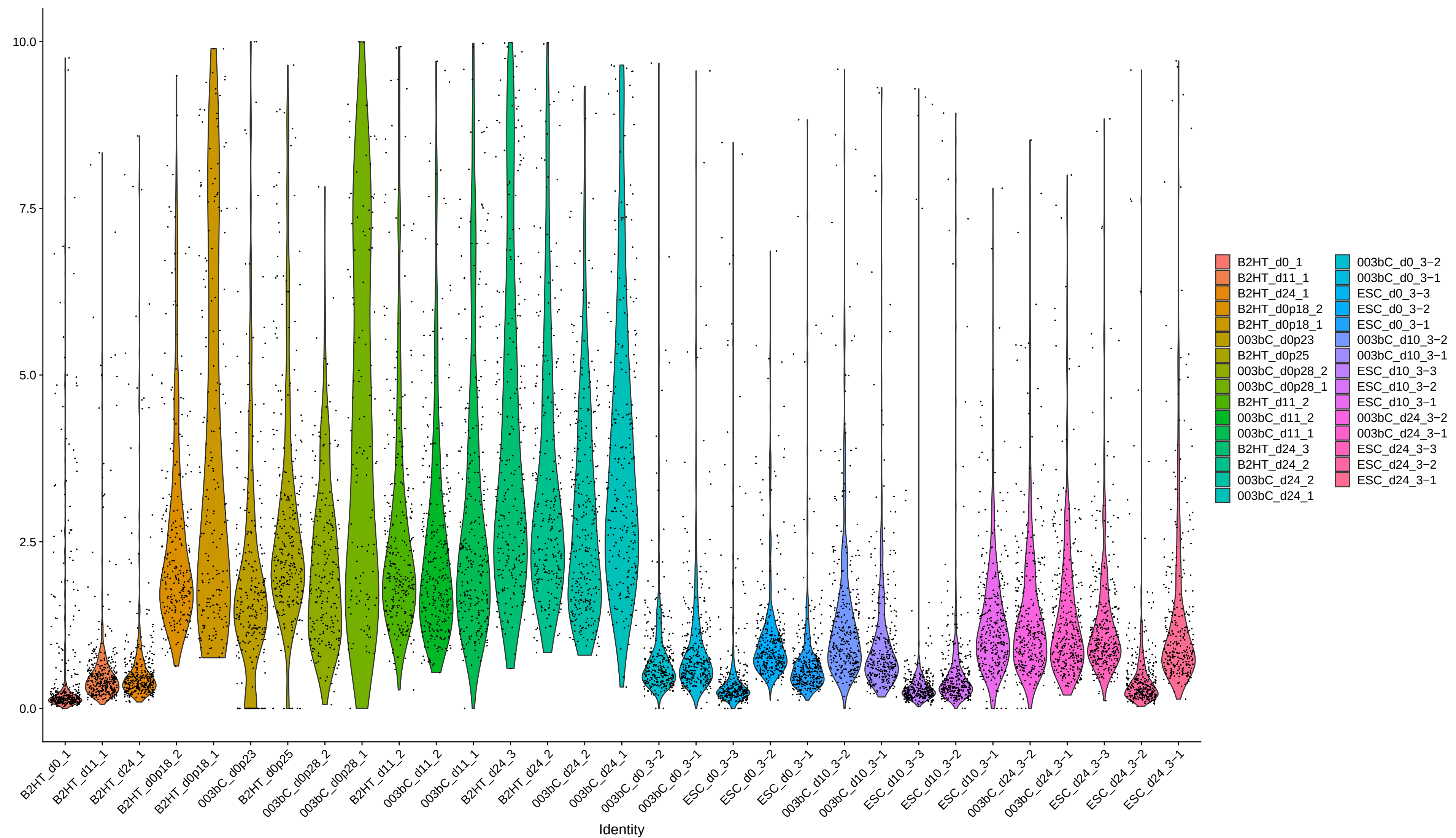
% mitochondrial reads all cells



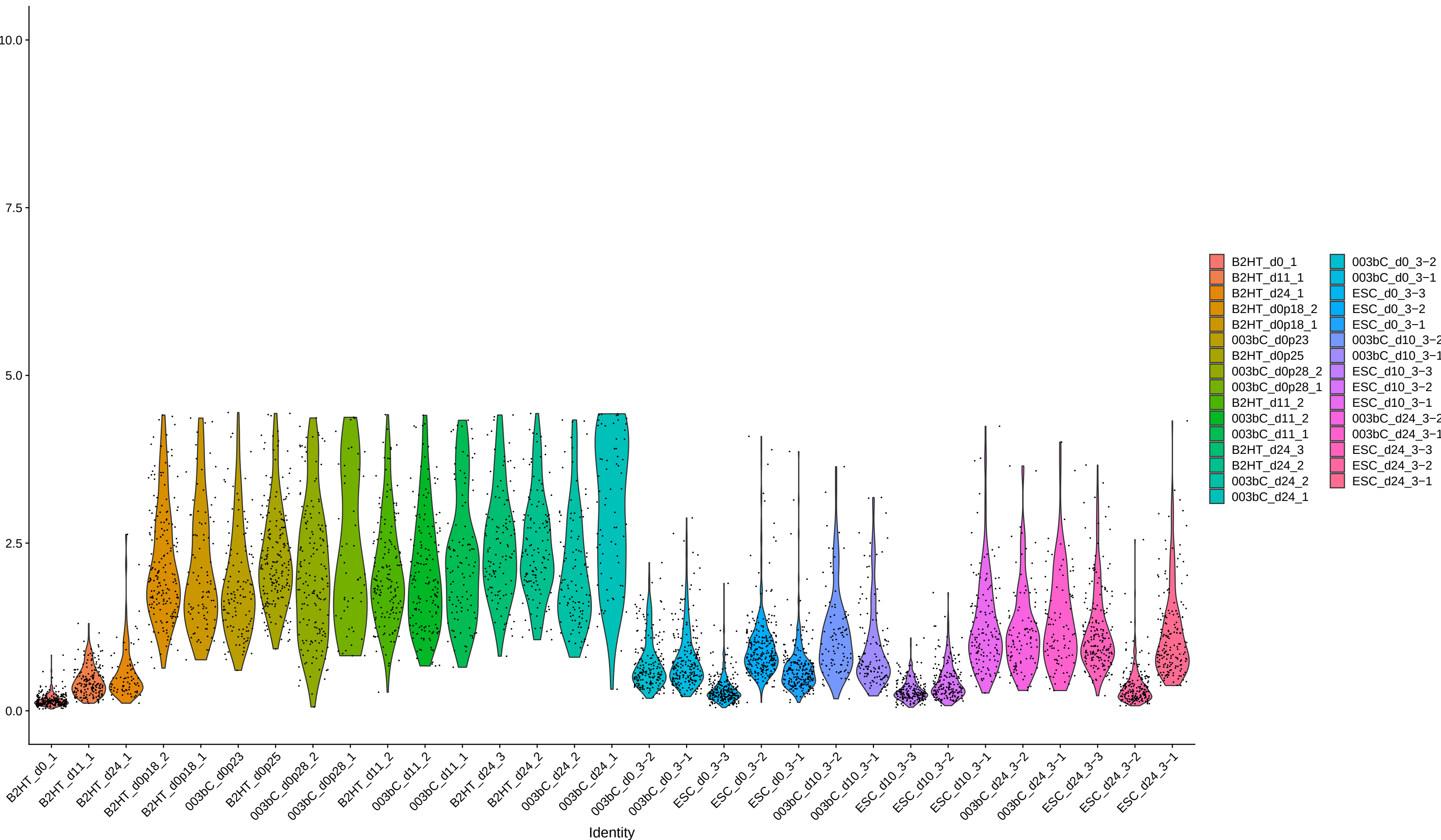
% mitochondrial reads filtered cells



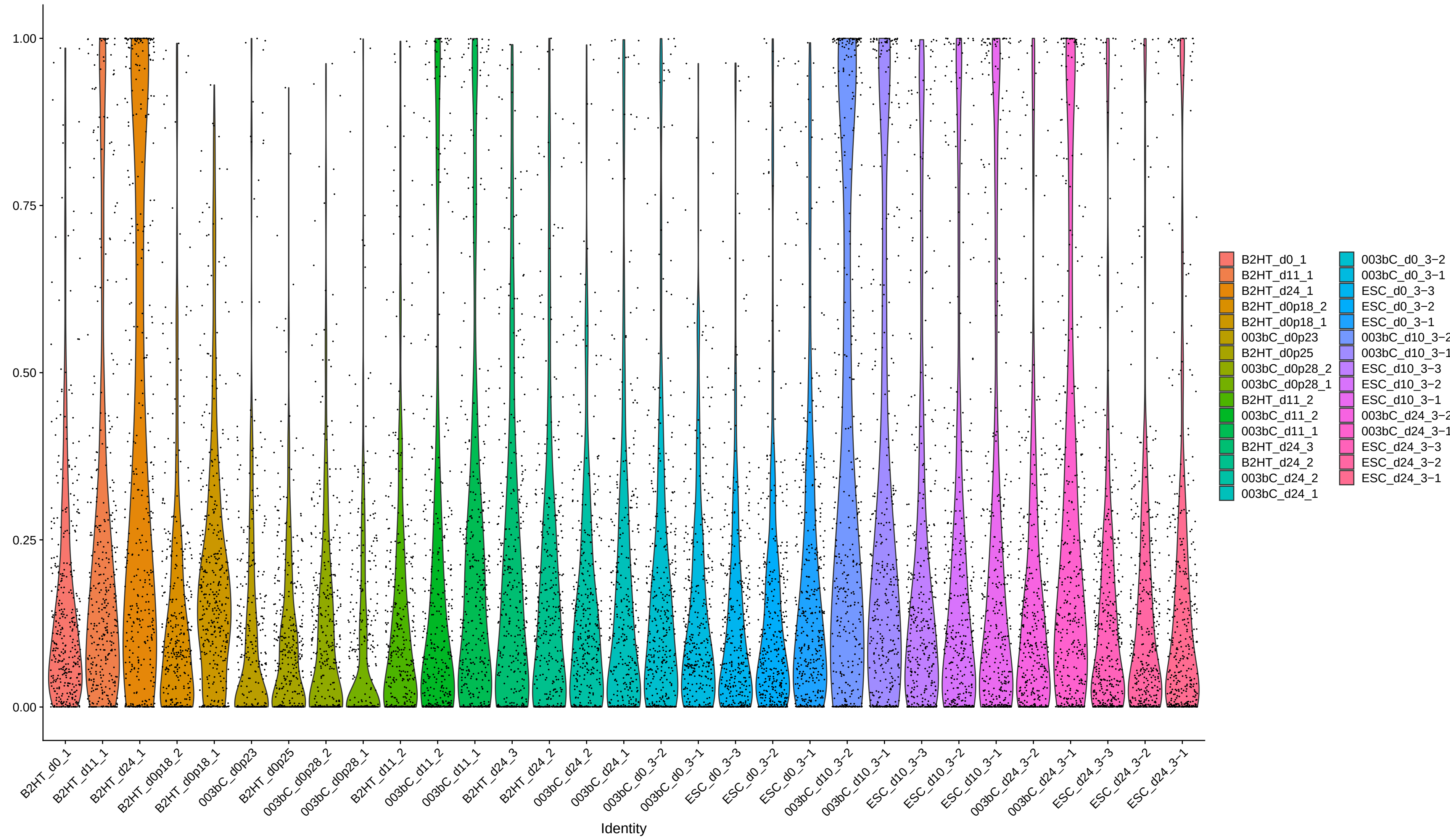
% ERCC reads all cells



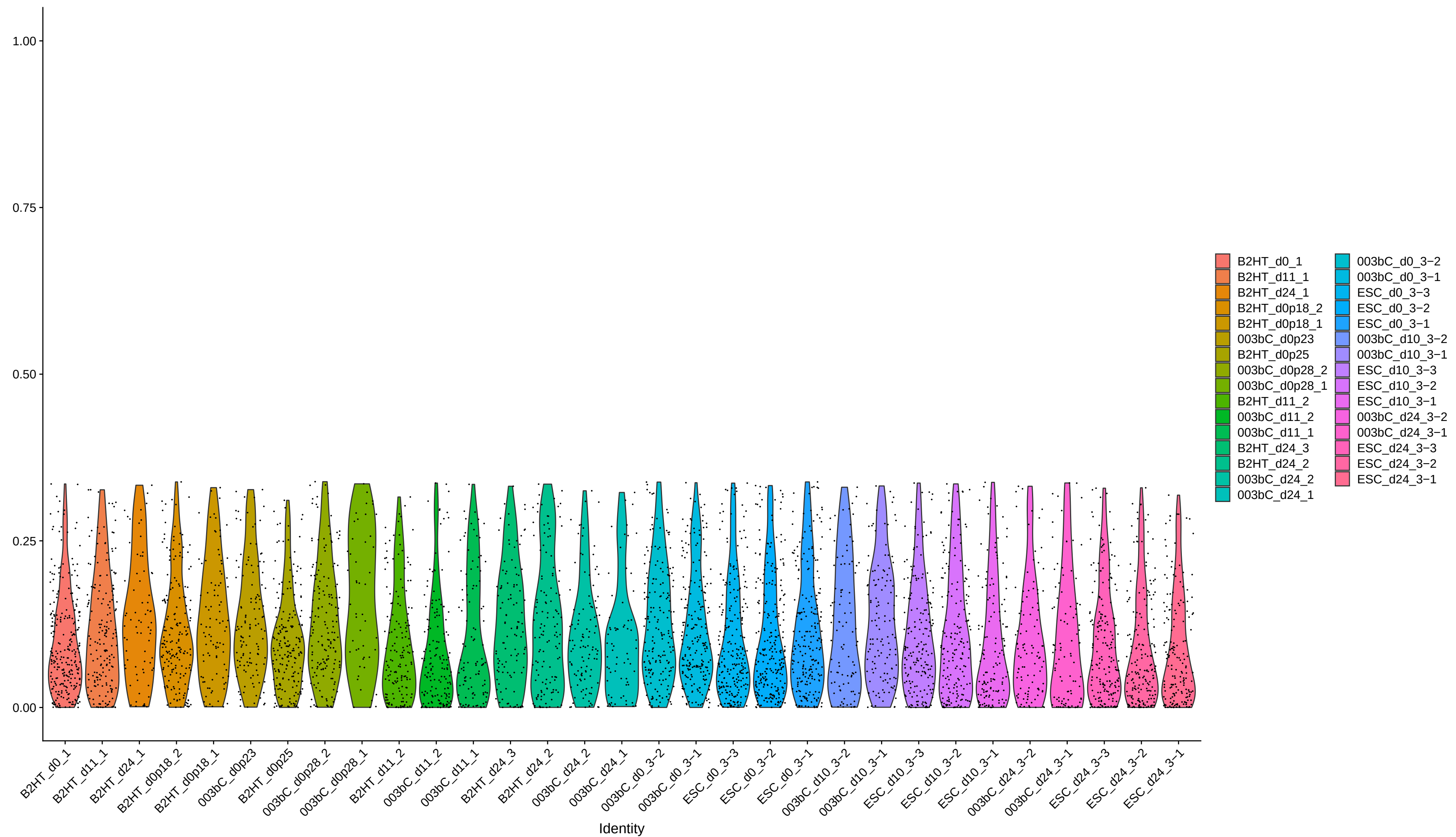
% ERCC reads filtered cells



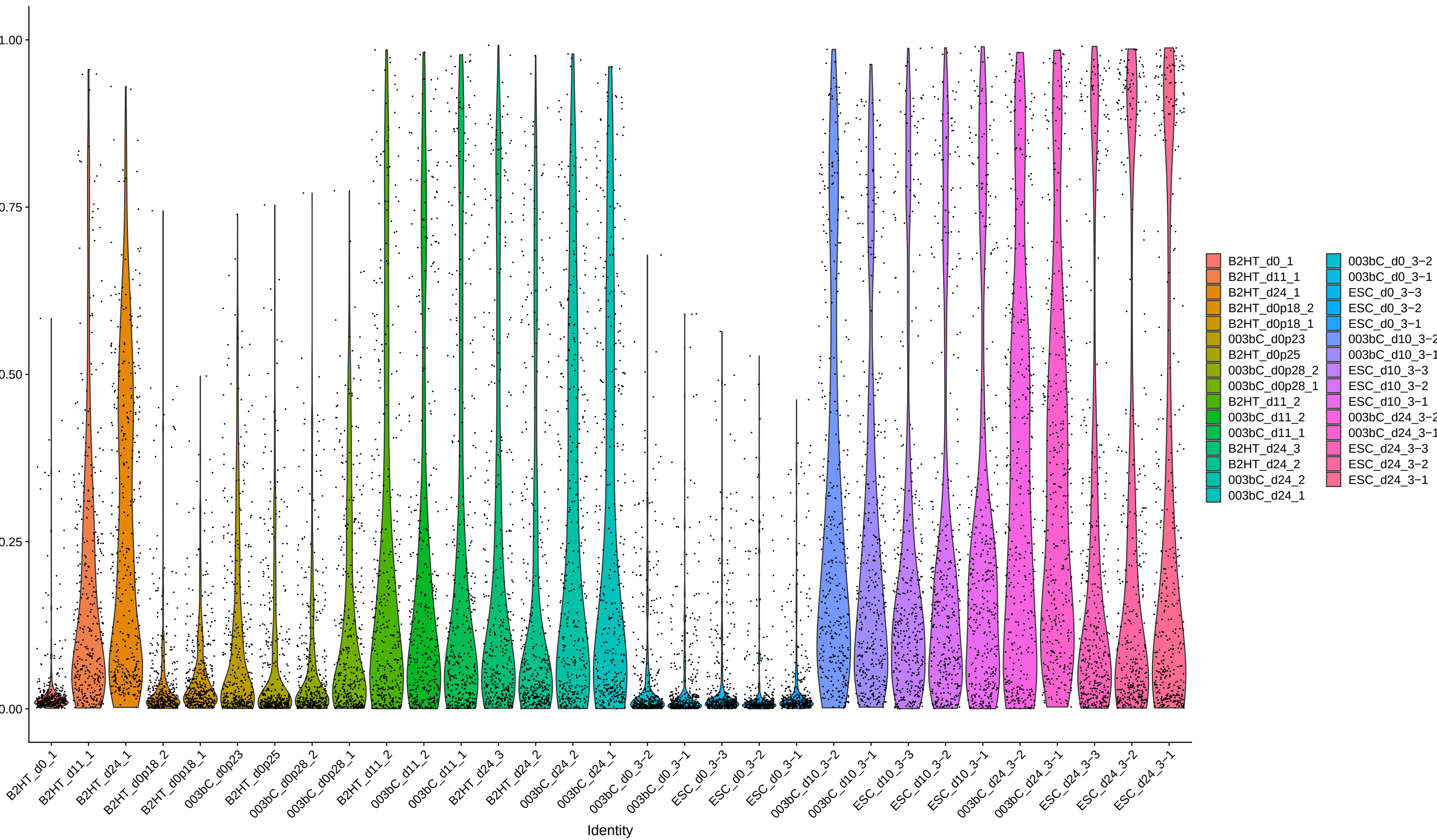
scDbfFinder_doublet_score



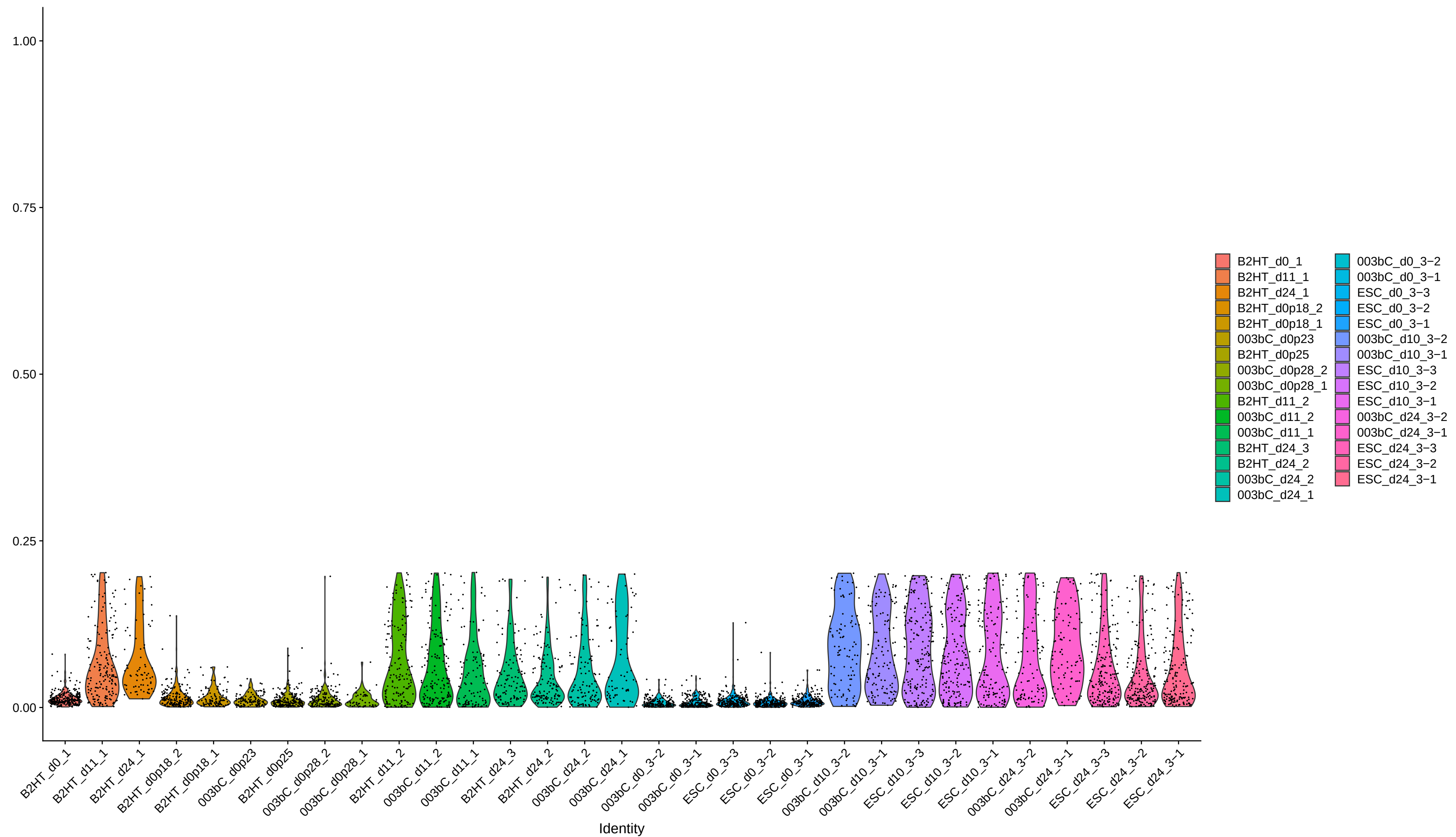
scDbIFinder_doublet_score_filtered



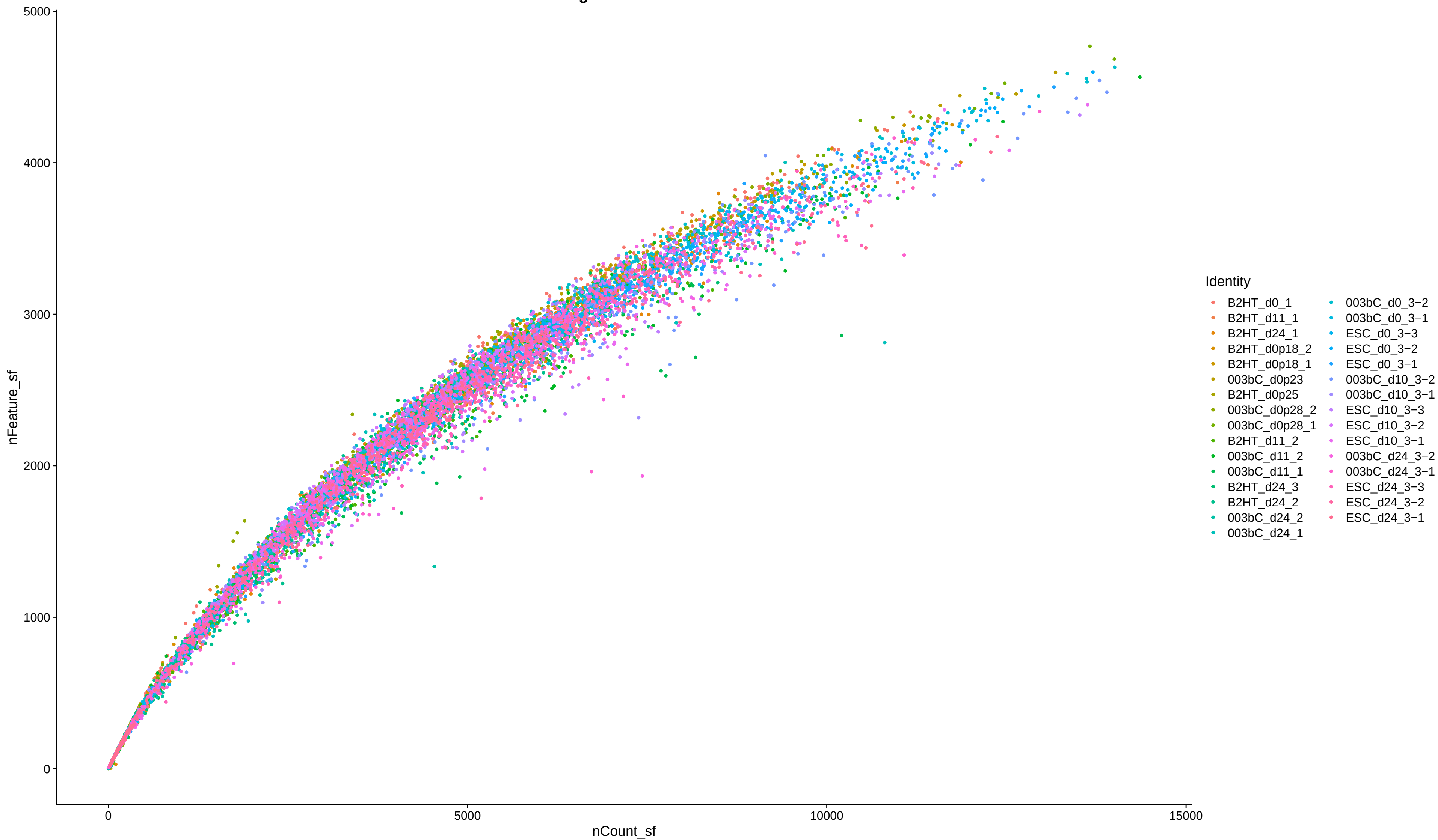
decontX_contamination



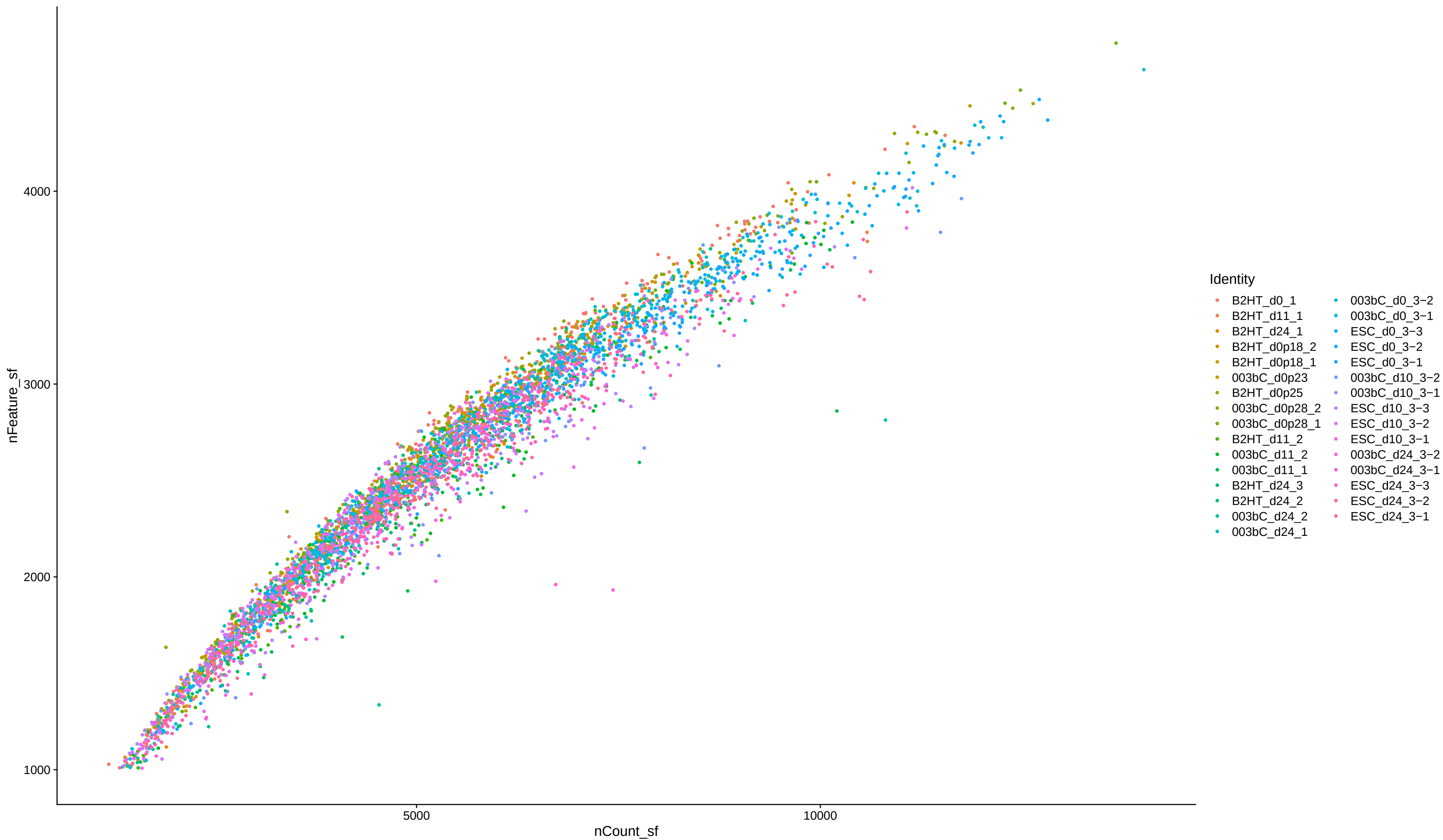
decontX_contamination_filtered



% counts vs genes measured all cells



% counts vs genes measured filtered cells



	descriptive_name	cells after QC
1	003bC_d0	735
2	003bC_d11	483
3	003bC_d24	375
4	B2HT_d0	737
5	B2HT_d11	338
6	B2HT_d24	309
7	ESC_d0	710
8	ESC_d11	500
9	ESC_d24	550