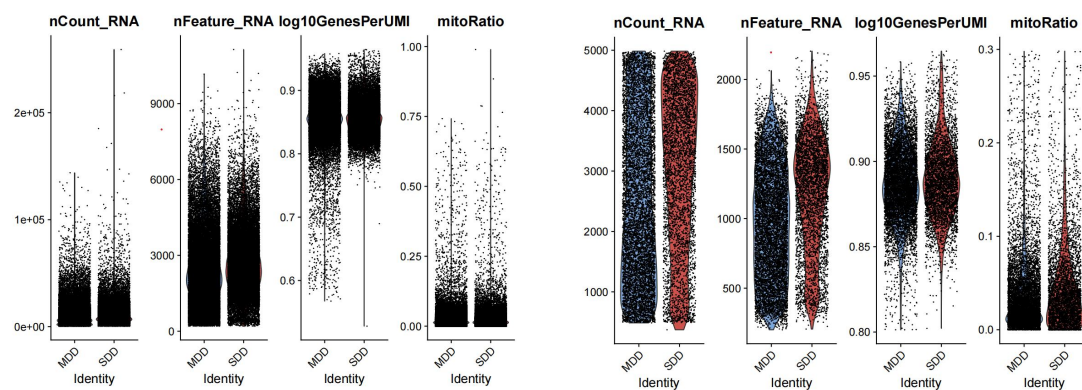


Summary of single-cell RNA-sequencing quality control and doublet removal per sample.

sample	Raw cells	After quality control cells	After doublet removal cells	Doublet rate
GSM7831813	5421	874	860	0.0160183066361556
GSM7831814	9654	5120	4871	0.048632813
GSM7831815	7474	1384	1325	0.0426300578034682
GSM7831816	7350	1156	1134	0.0190311418685121
GSM7831817	5517	307	297	0.0325732899022801
GSM7831818	11544	3024	2899	0.0413359788359788
GSM7831819	7837	1205	1157	0.0398340248962656

Quality control metrics of single-cell data before and after filtering



Batch correction effect evaluation before and after Harmony integration

