

									type	
	0.94	0.94	0.99	0.73	1.00	0.00	0.97	0.96	0.68	Seurat v4 WNN
	0.77	0.94	0.99	0.56	0.31	0.81	0.59	0.90	0.57	scMDC
	0.78	0.95	0.82	0.52	0.42	0.82	0.62	0.85	0.60	scMVP
	0.71	0.93	NA	0.46	0.41	0.94	0.12	0.82	0.48	scAI
	0.72	0.76	NA	0.38	0.28	0.83	NA	0.74	0.49	DCCA
	0.54	0.75	0.86	0.67	0.30	0.93	0.10	0.72	0.50	Multigrade
	0.68	0.73	NA	0.62	0.41	NA	NA	0.70	0.51	scMVAE
	0.50	0.67	0.76	0.30	0.33	0.84	0.20	0.64	0.42	MOFA+
	0.63	0.90	0.00	0.40	0.21	0.85	0.54	0.51	0.50	uniPort
	0.47	0.69	0.66	0.25	0.27	0.54	0.40	0.61	0.37	cobolt
	0.49	0.65	0.76	0.28	0.18	0.34	0.40	0.63	0.30	MultiVI
	0.17	0.00	NA	0.64	0.50	0.85	0.60	0.08	0.65	scDEC
Batch removal score										
Bio conservation score										
10X mouse brain Batch removal										
10X PBMC Batch removal										
HSPC Multiome p10 Batch removal										
BMMC Multiome p10 Batch removal										
SHARE-seq skin Bio conservation										
BMMC Multiome s3d10 Bio conservation										
BMMC Multiome p10 Bio conservation										

