												type	1	ty	
0.87	0.92	0.70	0.49	0.99	0.89	NA	0.80	0.98	0.95	0.83	0.85	bindSC	0.	8	Batch removal Bio Conservation
0.83	0.99	0.75	0.48	0.61	0.49	NA	NA	0.74	1.00	0.85	0.66	Seurat v4 WNN			Ranking
0.79	0.97	0.37	0.57	0.84	0.58	0.72	0.77	0.68	0.87	0.71	0.72	MaxFuse	0.	6	
0.91	0.94	0.49	0.24	0.64	0.80	NA	0.81	0.47	0.60	0.78	0.59	scMDC	0.	4	
0.83	0.88	0.60	0.42	0.73	0.03	0.69	0.58	0.79	0.70	0.77	0.56	SCALEX	0.	2	
0.74	0.99	0.84	0.24	0.90	0.21	0.49	0.16	0.55	0.68	0.86	0.46	totalVI	0		
0.75	0.82	0.85	0.13	0.54	0.45	0.42	0.50	0.63	0.63	0.81	0.47	MOFA+			
0.79	0.84	0.74	0.15	0.62	0.77	NA	NA	0.30	0.58	0.79	0.48	Multigrate			
0.59	0.33	0.68	0.60	0.49	0.51	0.18	0.00	0.73	0.46	0.53	0.42	sciPENN			
0.00	0.17	0.33	0.68	0.00	0.00	NA	NA	0.00	0.00	0.17	0.14	CiteFuse			
BMMC CITE p10 Bio conservation	BMMC CITE s2d1 Bio conservation	Lymph node spatial Bio conservation	BMMC CITE p10 Batch removal	HSPC CITE p10 Batch removal	Lymph node spatial Batch removal	COVID19 Batch removal	WBC Batch removal	NSCLC Batch removal	Kidney Batch removal	Bio conservation score	Batch removal score				