Raw distance based on 80599 peaks, and Normalized UMAP distances from 27 parameter choices

Raw distances with	ou UDistances between non-replicate clones
	Mean dist between non-replicate clones
	Distances between replicate clones
***************************************	Distance between 5a/5b sister clones
• (CD CONTROL COM	
	,
• CD1• CD30 CD30 CD30 CD30 CD30 CD30 CD30 CD30	
	0 (00 010 0 0
CONTROL 0 0 000 000 000 000 000 000 000 000 0	0 000 000 0

UMAP parameters

Min dist.	# Nghbors	
0.5 0.25 0.1 0.25 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.25 0.1 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5	10 10 10 55 55 33 30 10 10 10 55 53 30 10 10 10 10 10 10 10 55 55 53 33 30 10 10 10 10 10 10 10 10 10 10 10 10 10	20 20 20 20 20 20 20 20 20 20 20 20 20 2
0.1	10	20
0.25	5	20
0.1	5	20
0.25	3	20
0.1	10	10
0.25	10	10
0.5	5	10
0.25 0.1	5 5	10 10
0.5	3	10
0.25	3	10
0.5 0.25	10 10	5 5
0.1	10	5
0.5 0.25	5	5
0.1	5	5
0.25	3	5 5
0.1	3	5