

Table S2. Impact of the co-localization parameter on the detection.

inter_gene_max_space (D)	N_components detected (known Cas proteins)	N_clusters detected	% large clusters > 13 genes	Maximal cluster size (N_total_genes)
4	10615	1638	6%	36
5	10663	1628	7%	36
6	10683	1613	8%	43