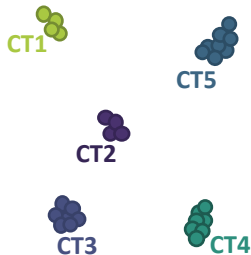


single-cell data



bed_file

		summarized activity					
#chr	start	end	CT1	CT2	CT3	CT4	CT5
chr1	1000	1200	12	42	5	6	9
chr1	2000	2200	25	2	50	20	6

-n 4



-n 4,6



-n 4-6



-n 4+

