

Same rearrangement

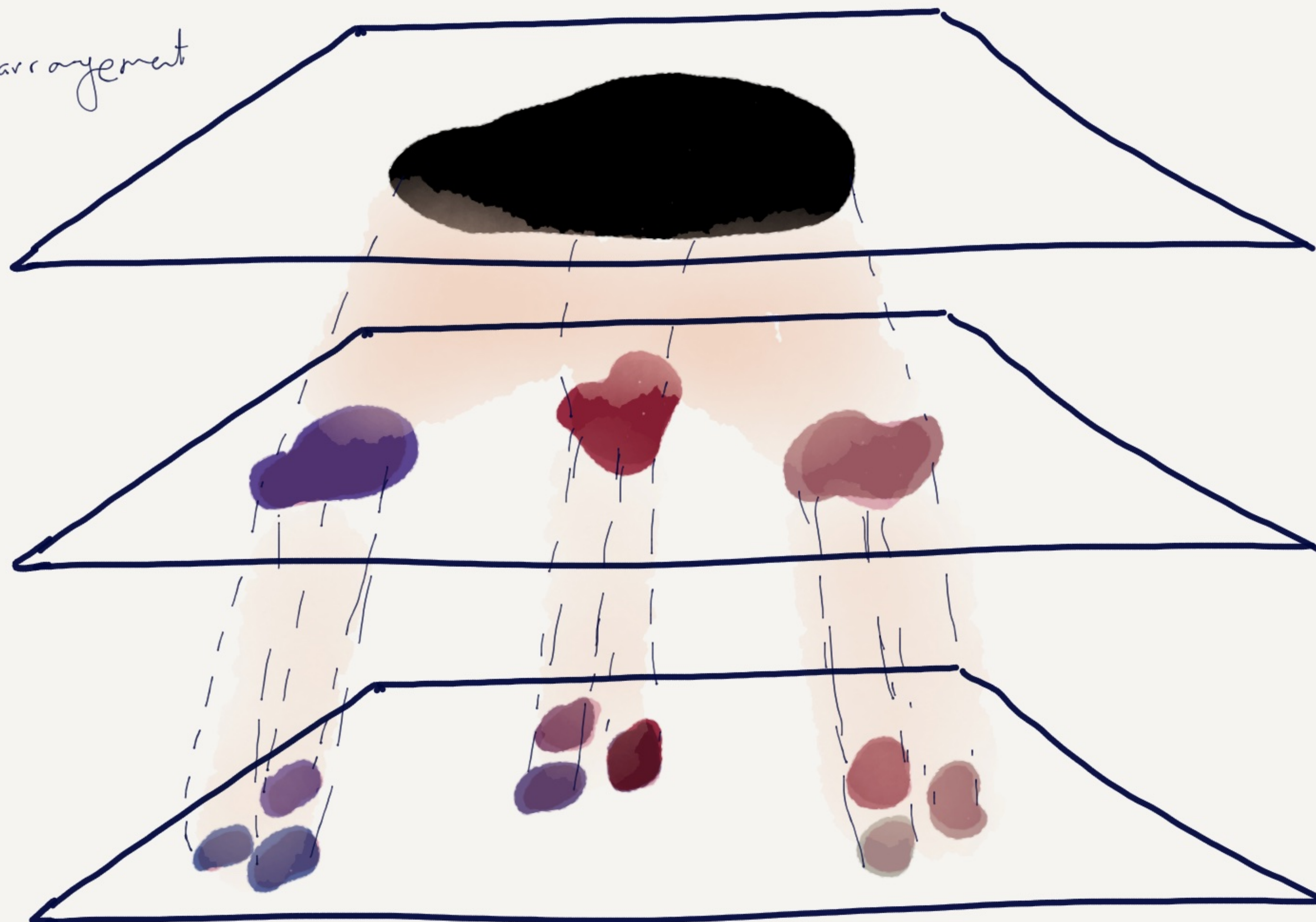
clone

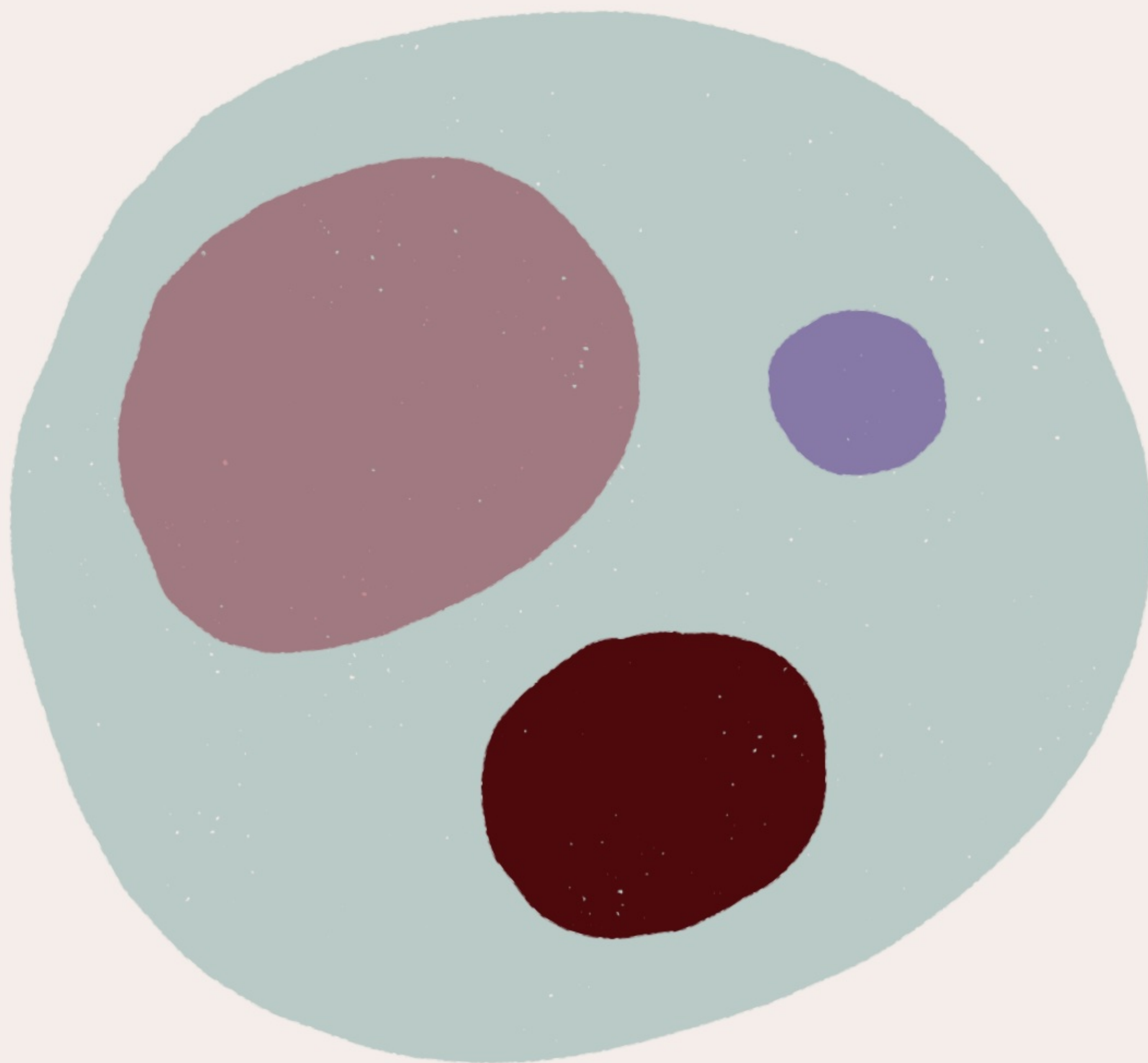
Same V
Same J
Same CDR3

clonotype

same seq

?

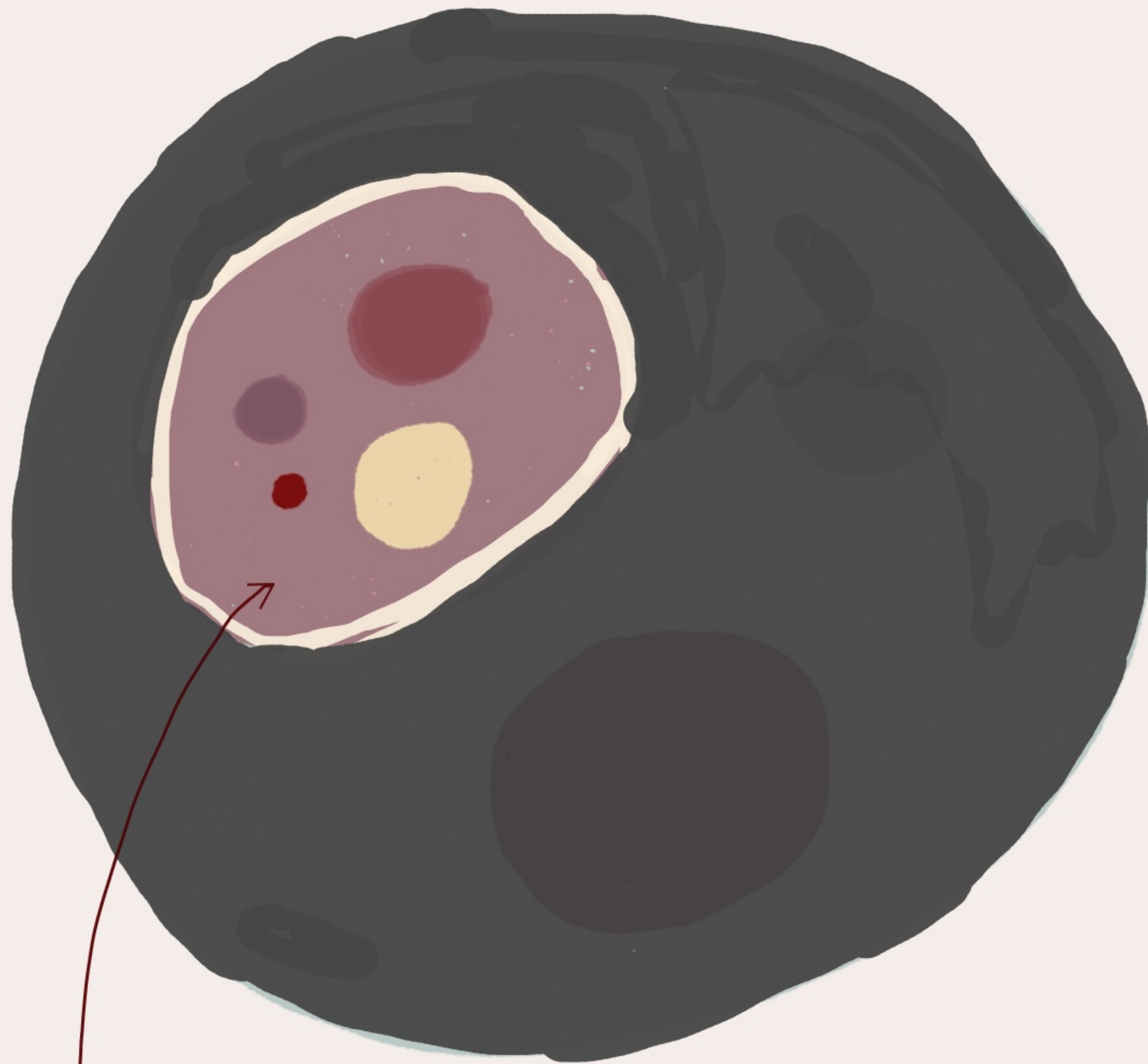




Level one

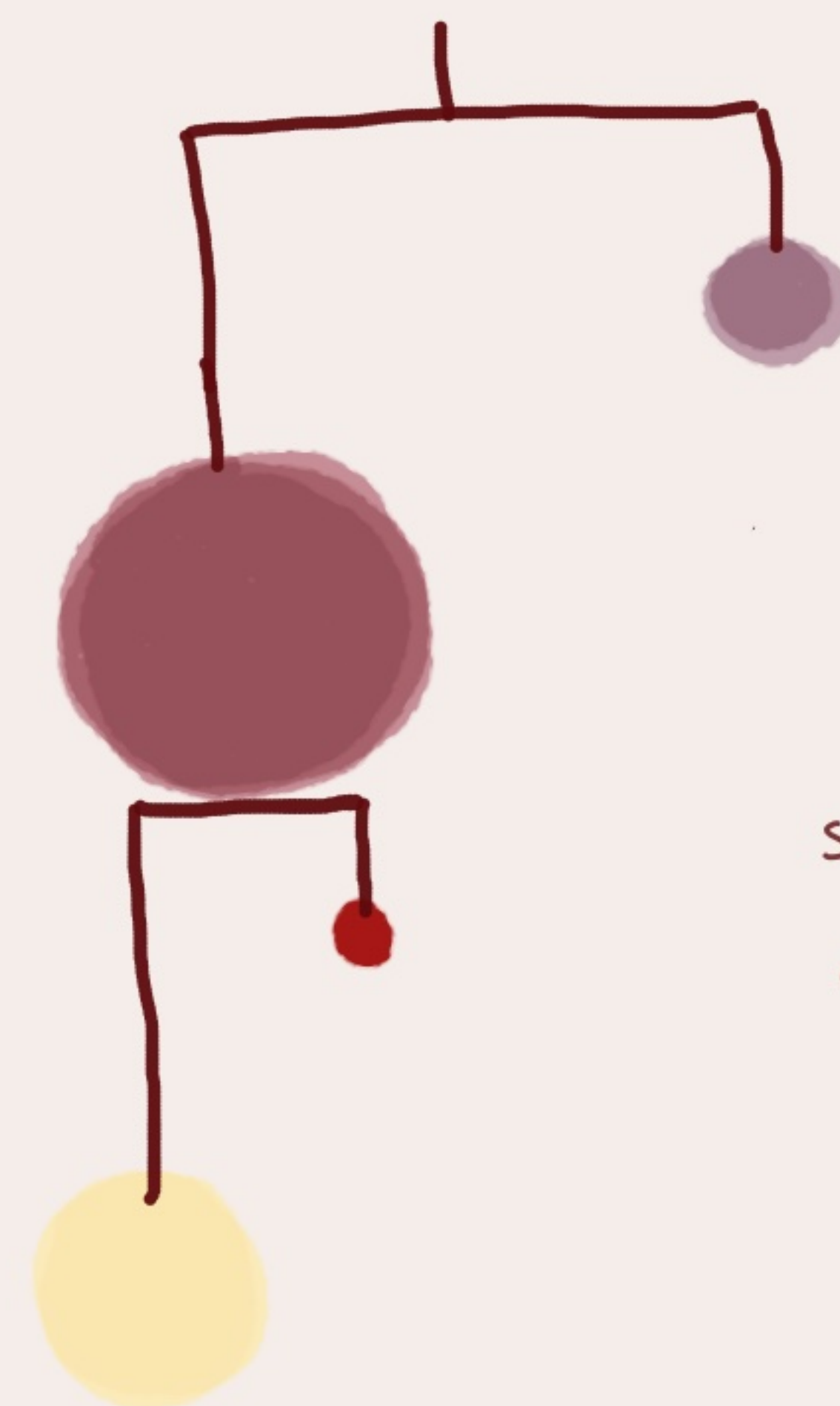
- associated with abundance plot



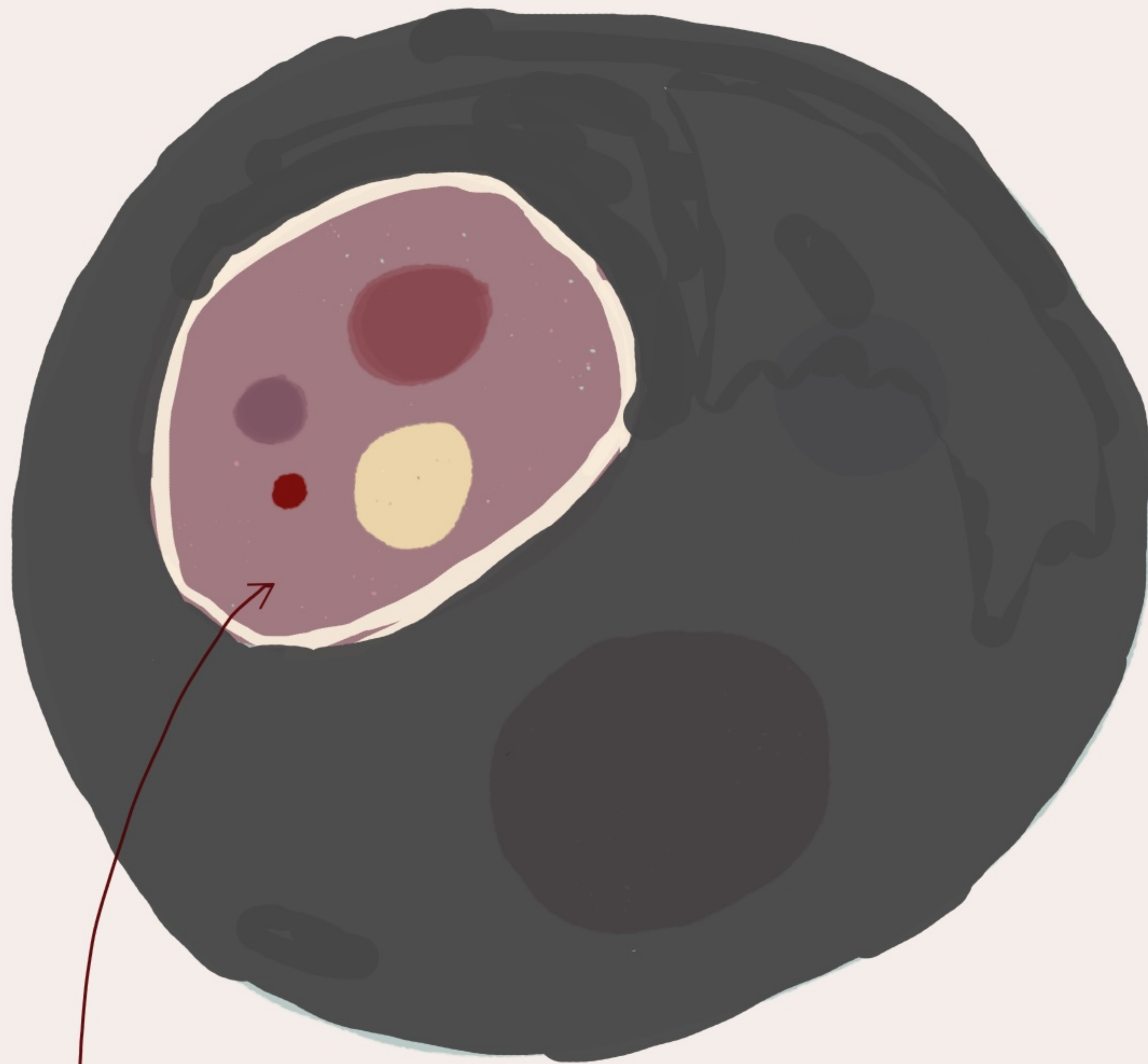


if click on it

Level one

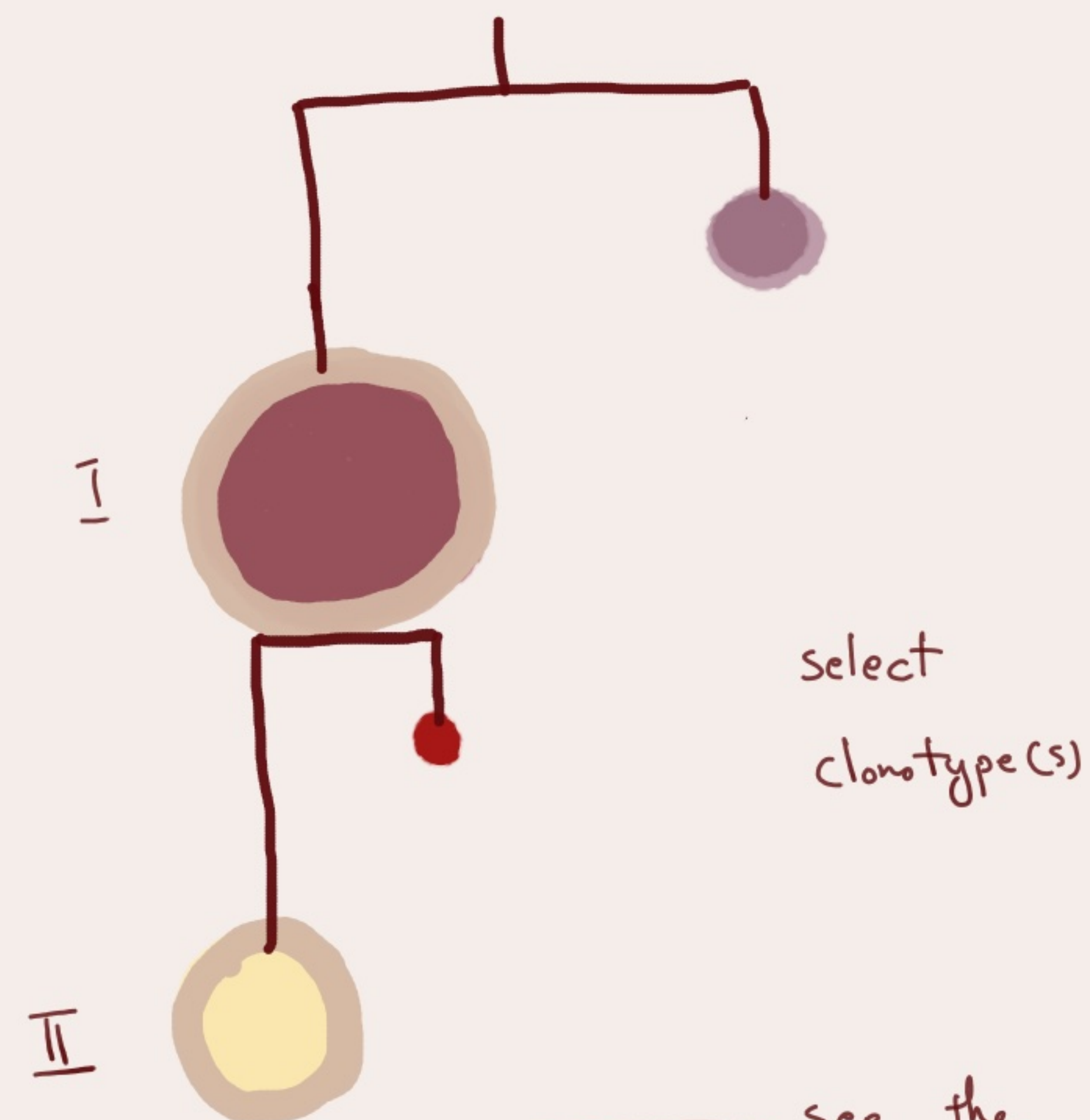


select
clonotype(s)



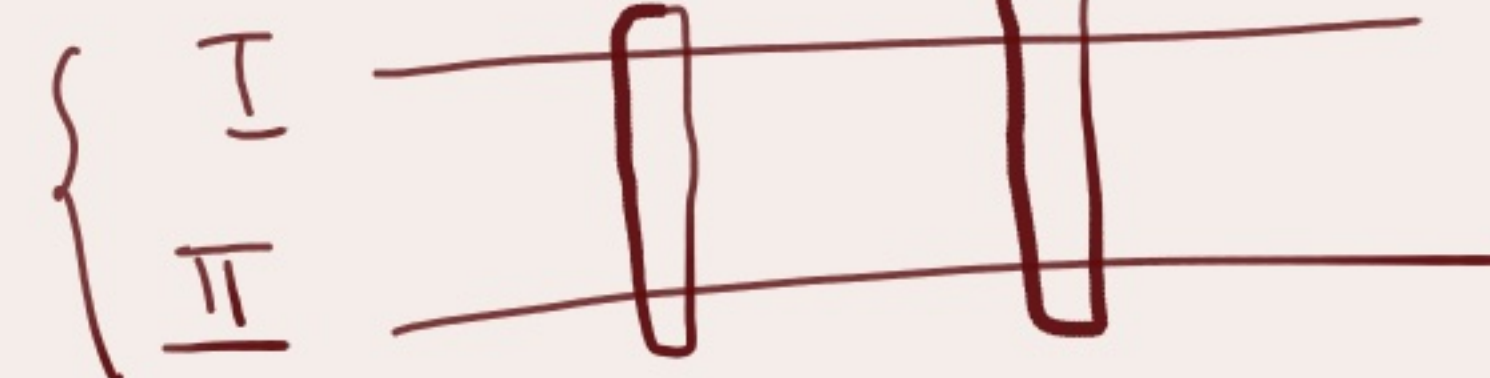
if click on it

Level two



select
clonotype(s)

sequence
alignment



see the
differences