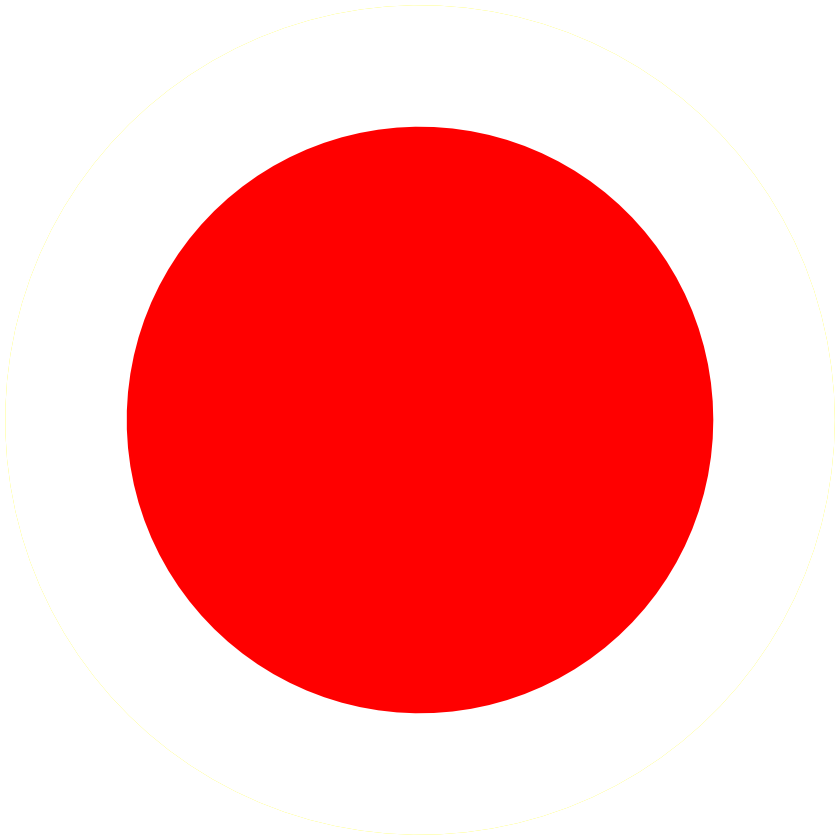


- cloud (1 , genomes<=4)
- shell (0)
- soft core (2 , genomes>=7)
- core (2 , genomes=8)



total gene clusters = 3   taxa = 8