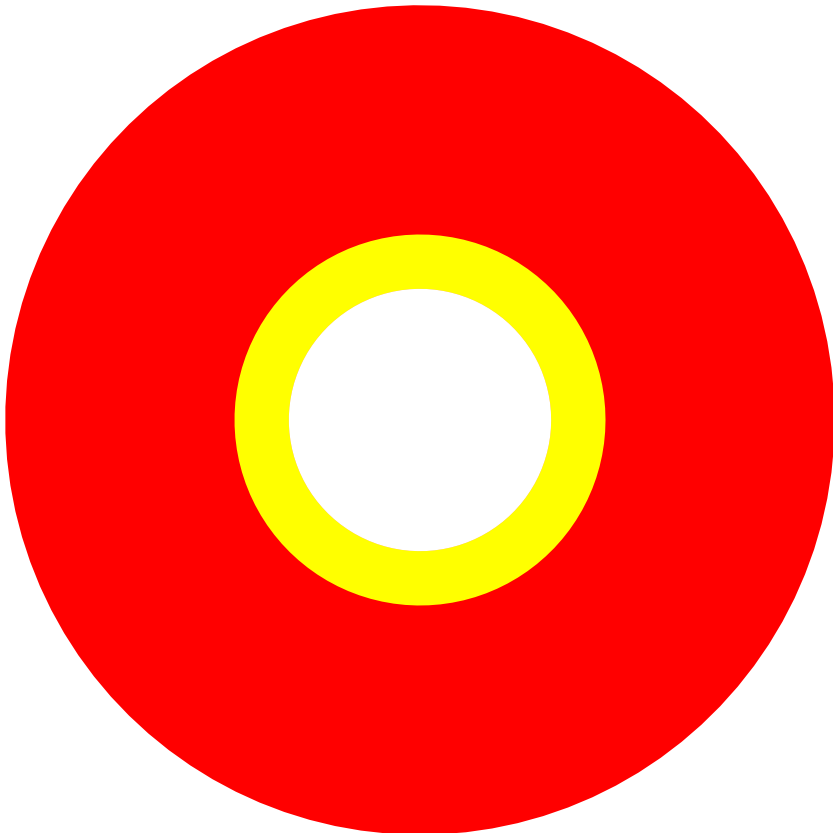


- cloud (10 , genomes<=2)
- shell (1)
- soft core (2 , genomes>=46)
- core (1 , genomes=49)



total gene clusters = 13   taxa = 49