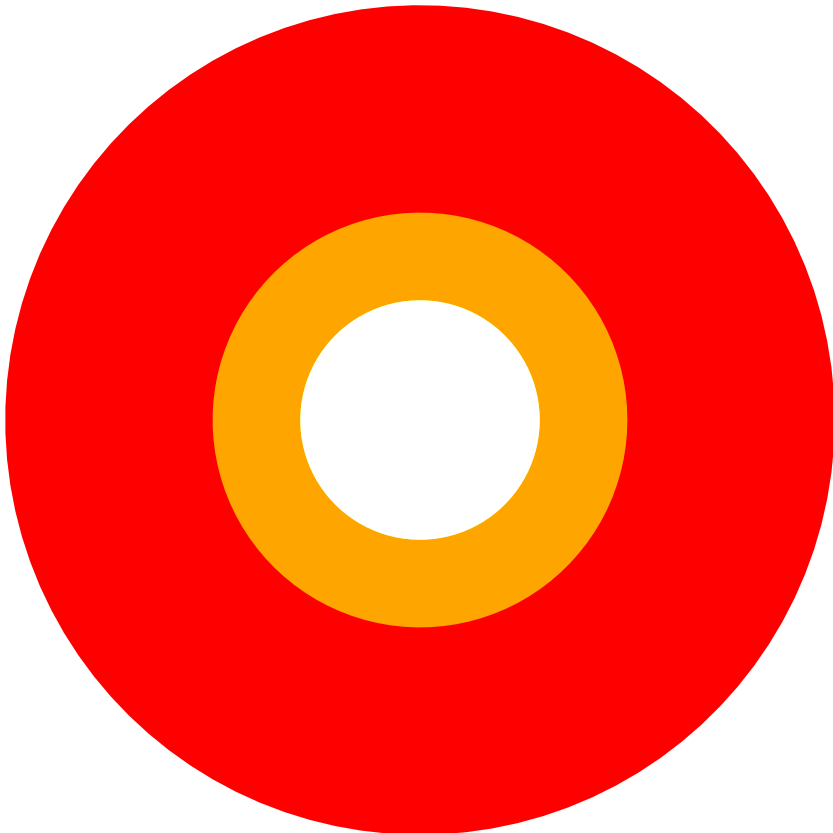


- cloud (12 , genomes<=2)
- shell (3)
- soft core (1 , genomes>=7)
- core (1 , genomes=8)



total gene clusters = 16 taxa = 8