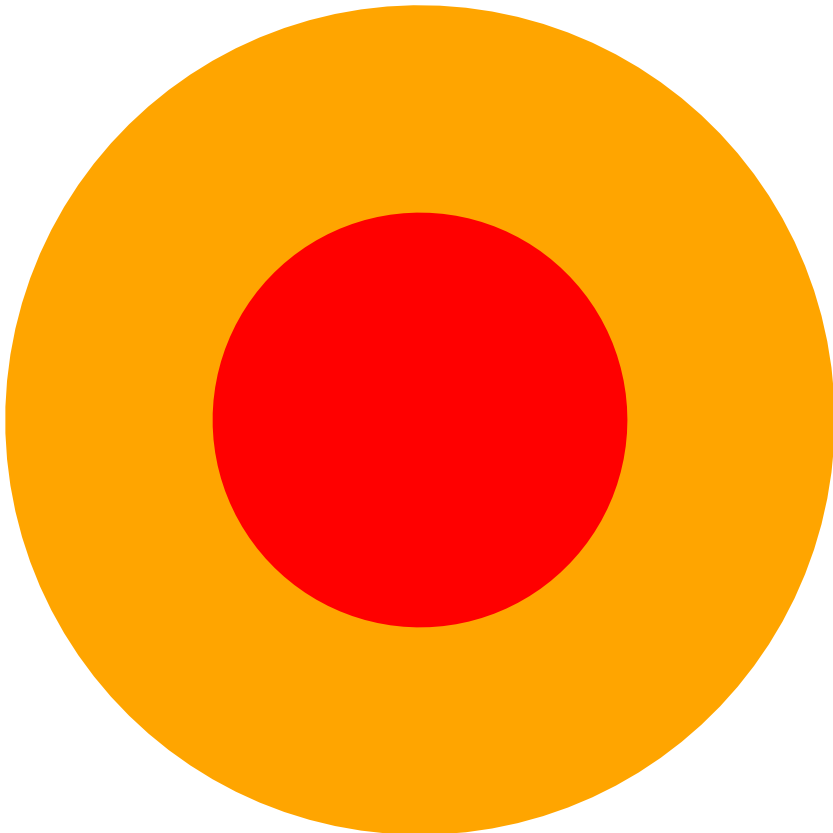


- cloud (1 , genomes<=2)
- shell (4)
- soft core (0 , genomes>=11)
- core (0 , genomes=12)



total gene clusters = 5   taxa = 12