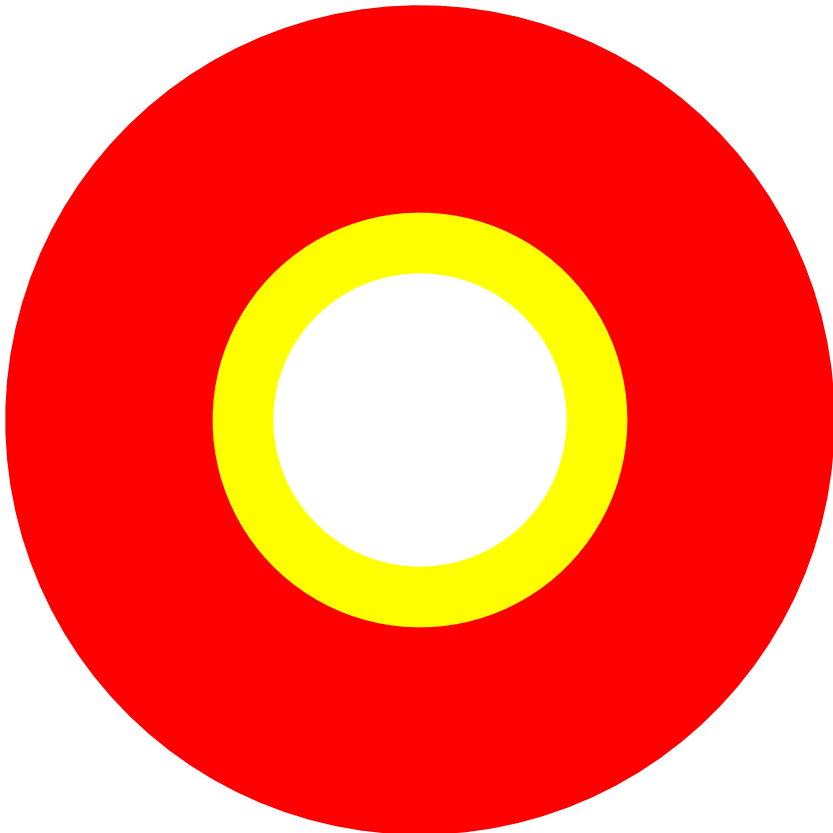


- cloud (8 , genomes<=2)
- shell (0)
- soft core (2 , genomes>=32)
- core (1 , genomes=34)



total gene clusters = 10   taxa = 34