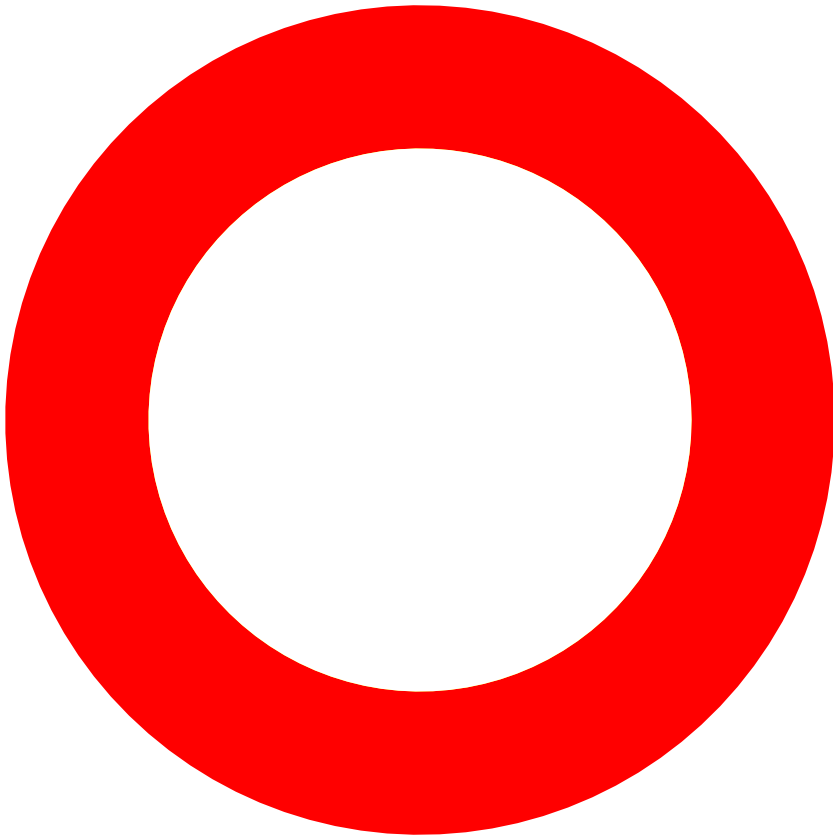


- cloud (7 , genomes<=10)
- shell (0)
- soft core (3 , genomes>=11)
- core (3 , genomes=12)



total gene clusters = 10   taxa = 12