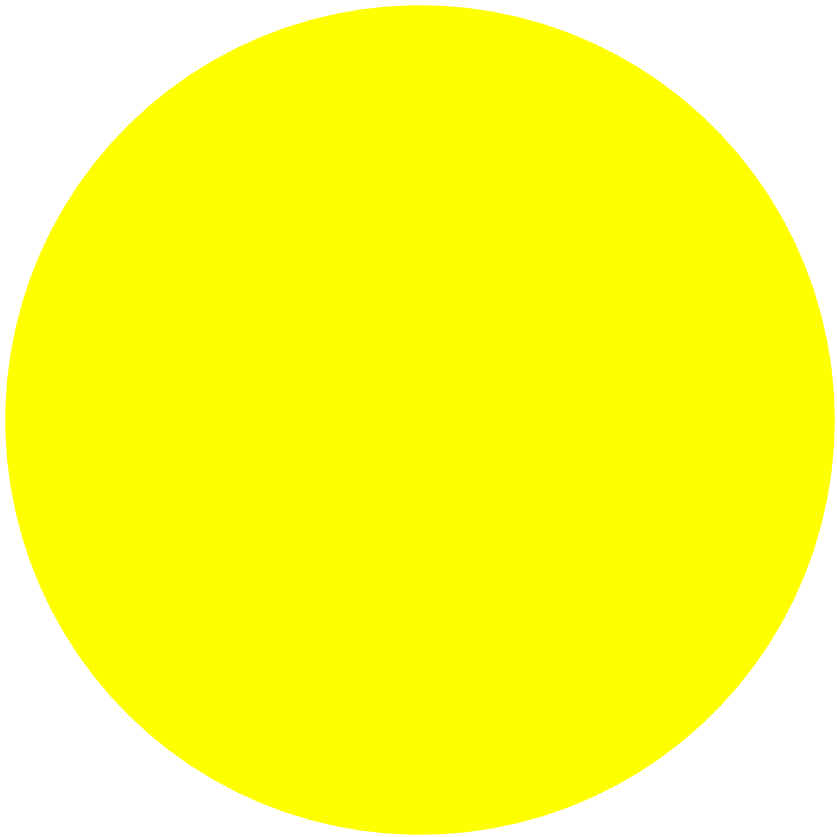


- cloud (0 , genomes<=2)
- shell (0)
- soft core (10 , genomes>=10)
- core (10 , genomes=11)



total gene clusters = 10   taxa = 11