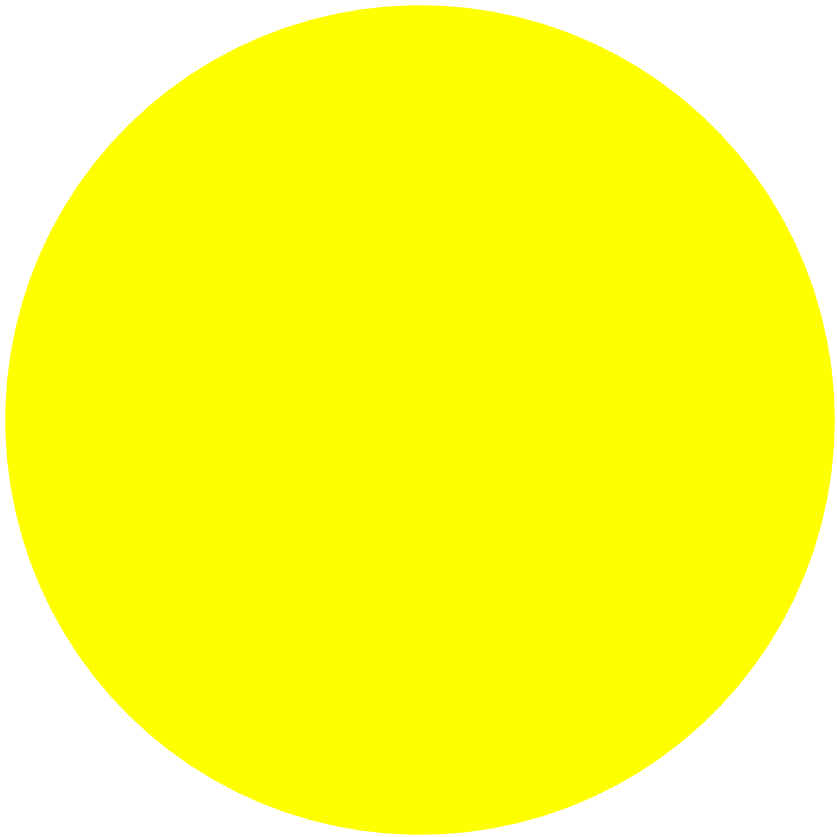


- cloud (0 , genomes<=2)
- shell (0)
- soft core (8 , genomes>=7)
- core (8 , genomes=8)



total gene clusters = 8   taxa = 8