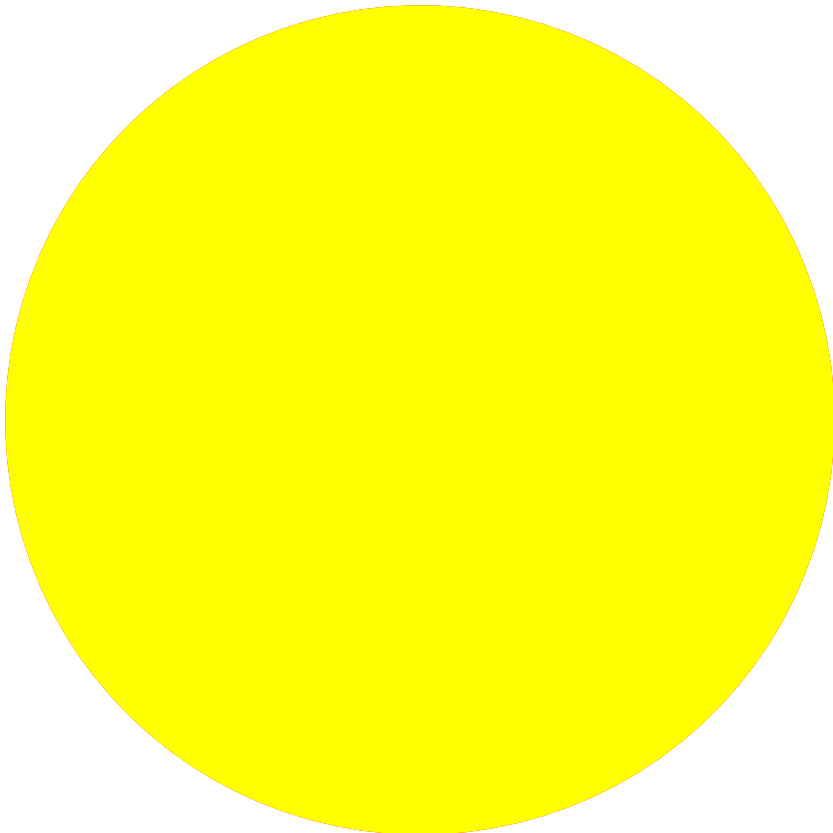


- cloud (1 , genomes<=2)
- shell (0)
- soft core (1 , genomes>=77)
- core (0 , genomes=82)



total gene clusters = 2   taxa = 82