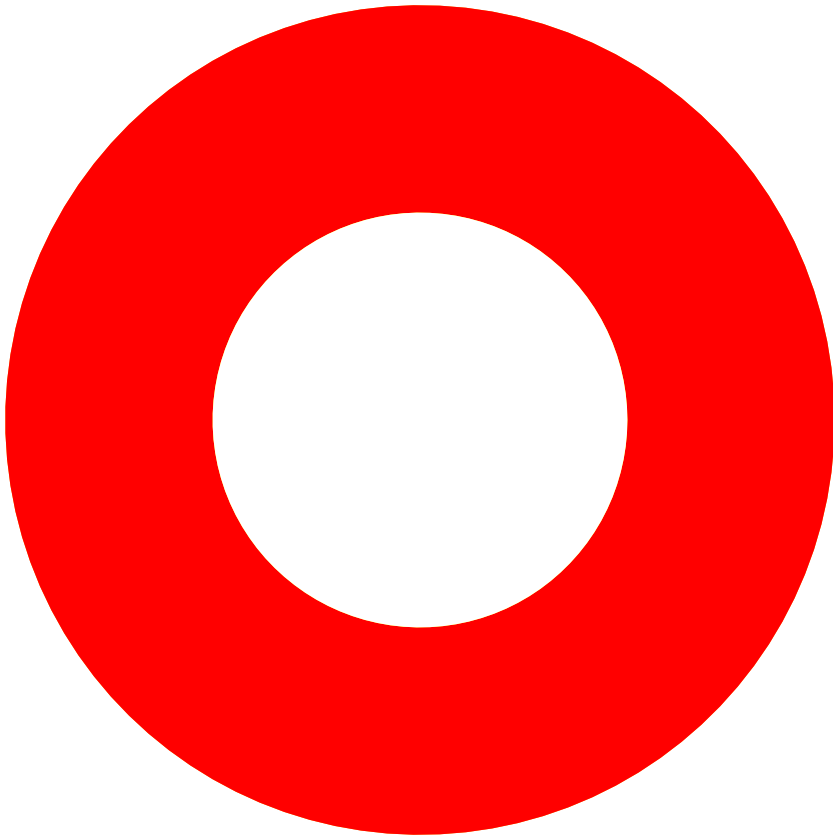


- cloud (4 , genomes<=2)
- shell (4)
- soft core (1 , genomes>=71)
- core (1 , genomes=75)



total gene clusters = 9   taxa = 75