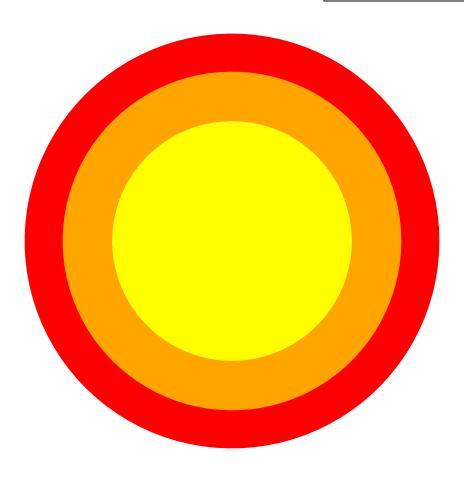
cloud (3, genomes<=2)</li>shell (2)soft core (1, genomes>=5)core (1, genomes=6)



total gene clusters = 6 taxa = 6