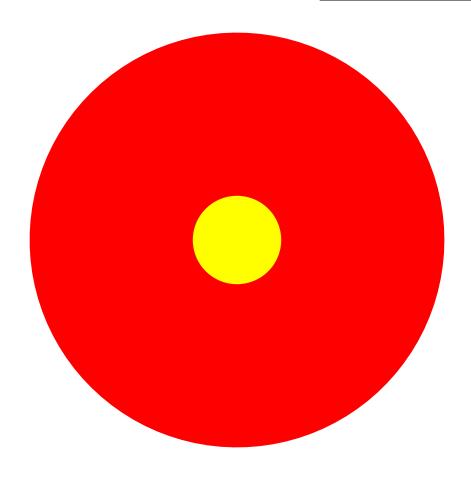
cloud (22, genomes<=2)</li>shell (0)soft core (1, genomes>=21)core (0, genomes=23)



total gene clusters = 23 taxa = 23