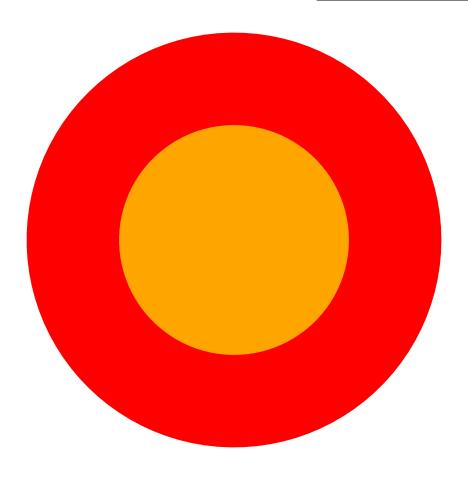
- cloud (140, genomes<=2)shell (43)soft core (0, genomes>=39)
- \Box core (0, genomes=42)



total gene clusters = 183 taxa = 42