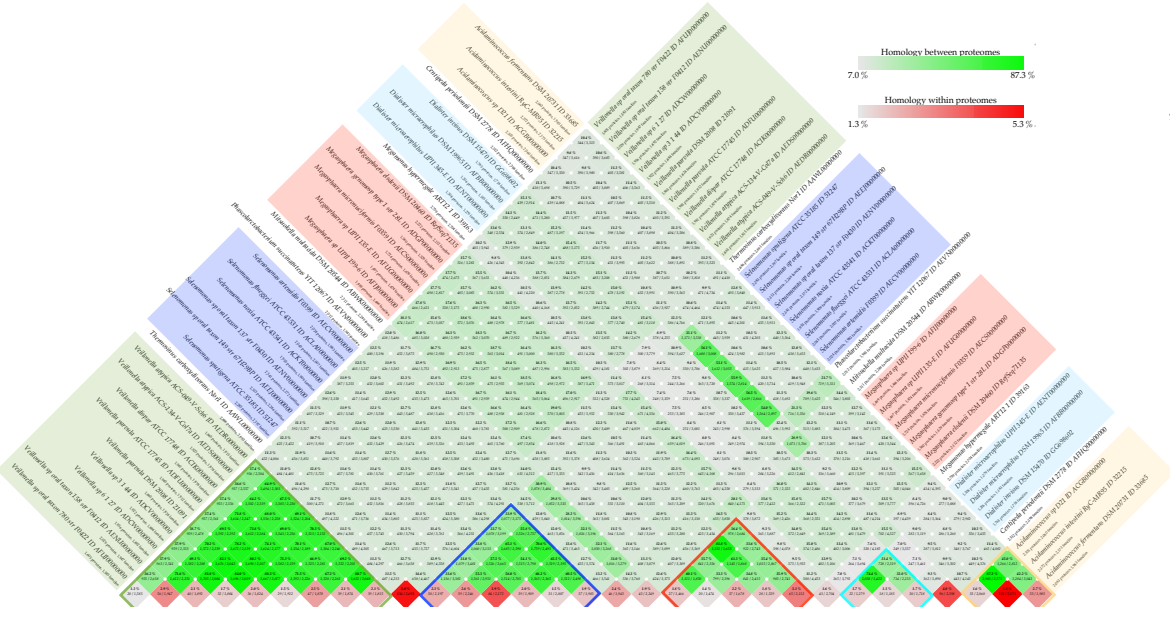
Homology between proteomes
7.0 %
87.3 %
Homology within proteomes
1.3 %
5.3 %