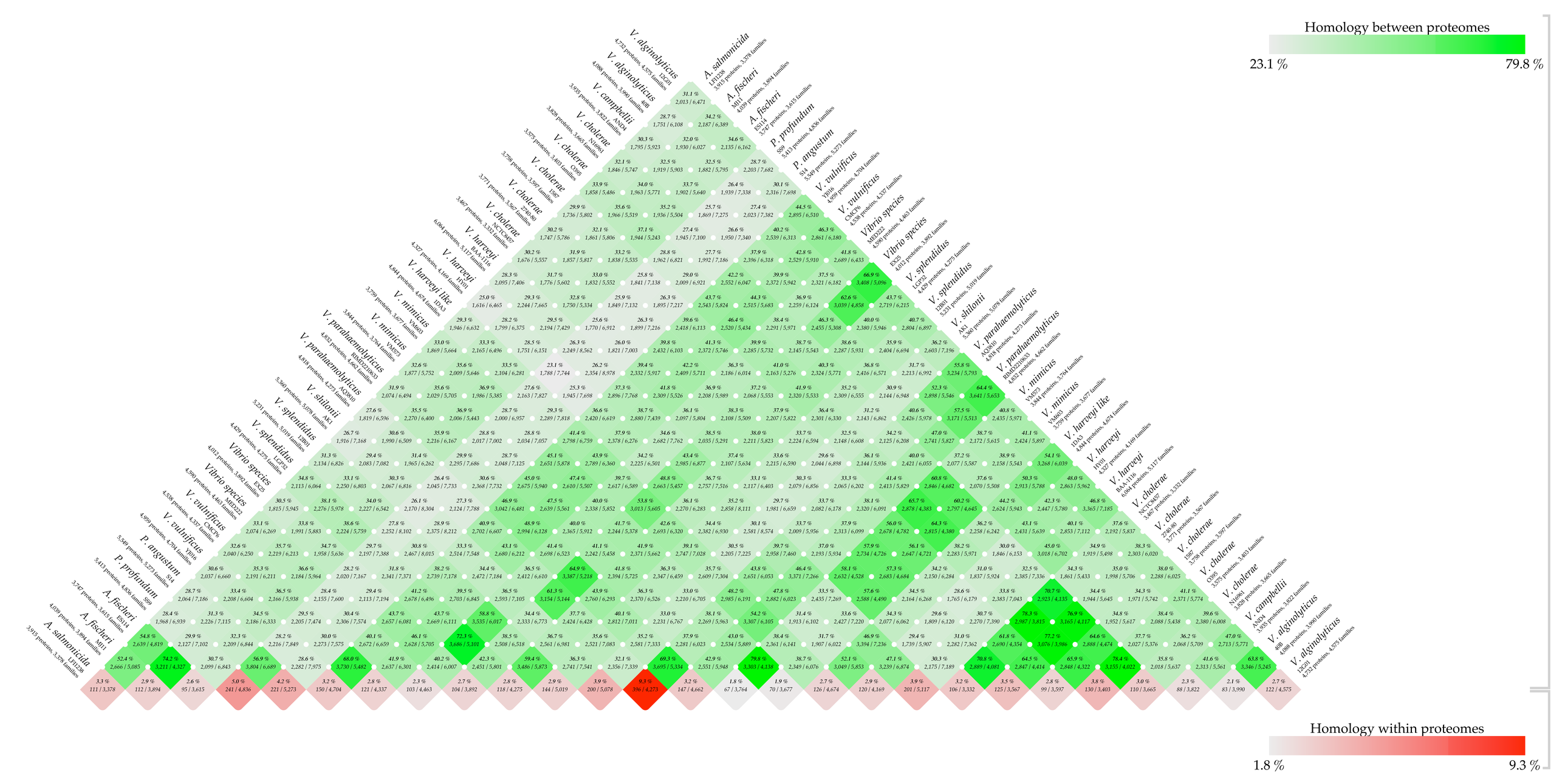
Homology between proteomes
23.1 %
79.8 %
Homology within proteomes
1.8 %
9.3 %