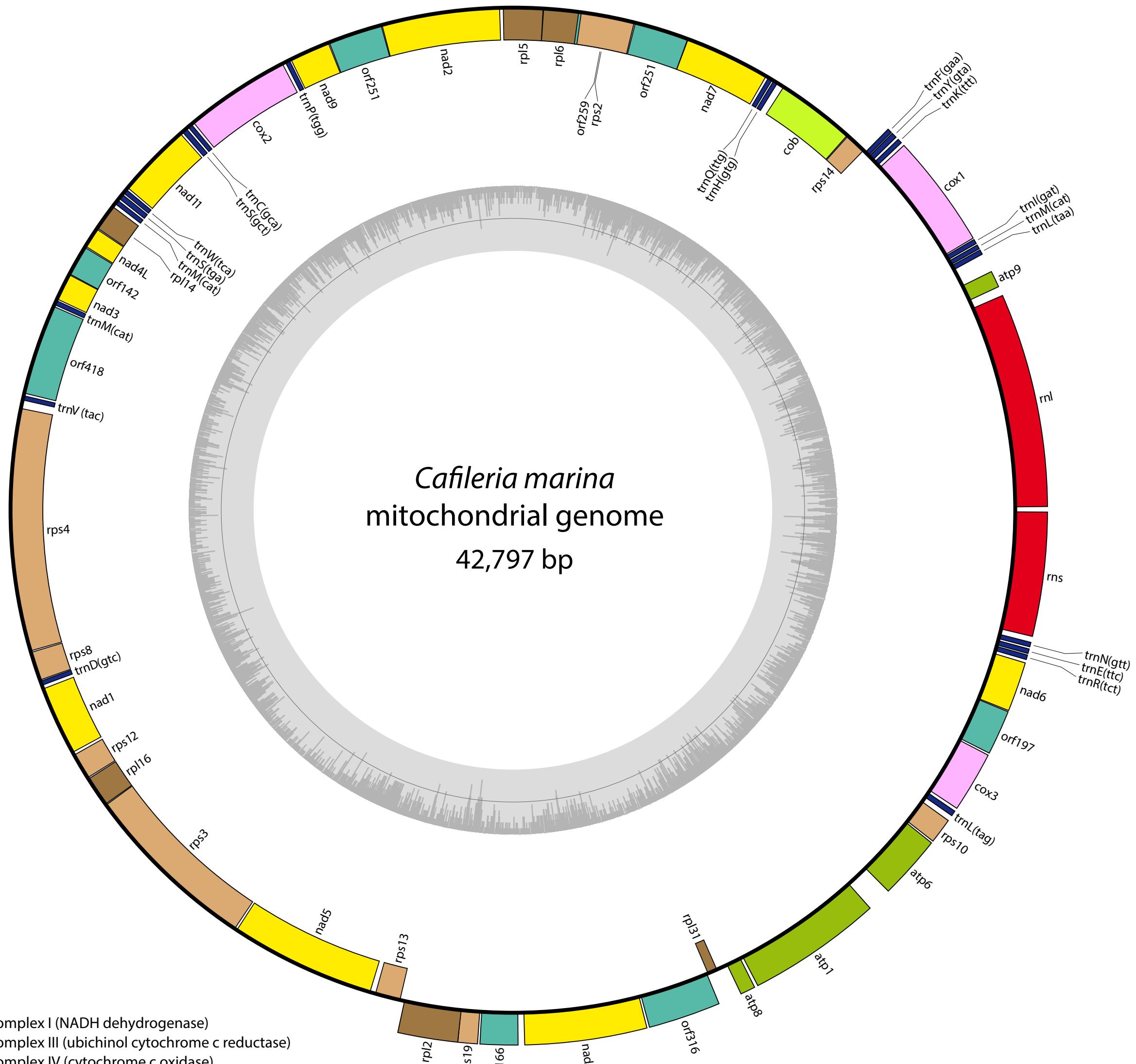


Cafileria marina
mitochondrial genome

42,797 bp



- [Yellow square] complex I (NADH dehydrogenase)
- [Light green square] complex III (ubiquinol cytochrome c reductase)
- [Pink square] complex IV (cytochrome c oxidase)
- [Dark green square] ATP synthase
- [Brown square] ribosomal proteins (SSU)
- [Tan square] ribosomal proteins (LSU)
- [Teal square] ORFs
- [Dark blue square] transfer RNAs: C(gca), D(gtc), E(ttc), F(gaa), H(gtg), I(gat), K(ttt), L(taa), L(tag), 3xM(cat), N(gtt), P(tgg), Q(ttg), R(tct), S(gct), V(tga), W(tca), Y(gta)
- [Red square] ribosomal RNAs