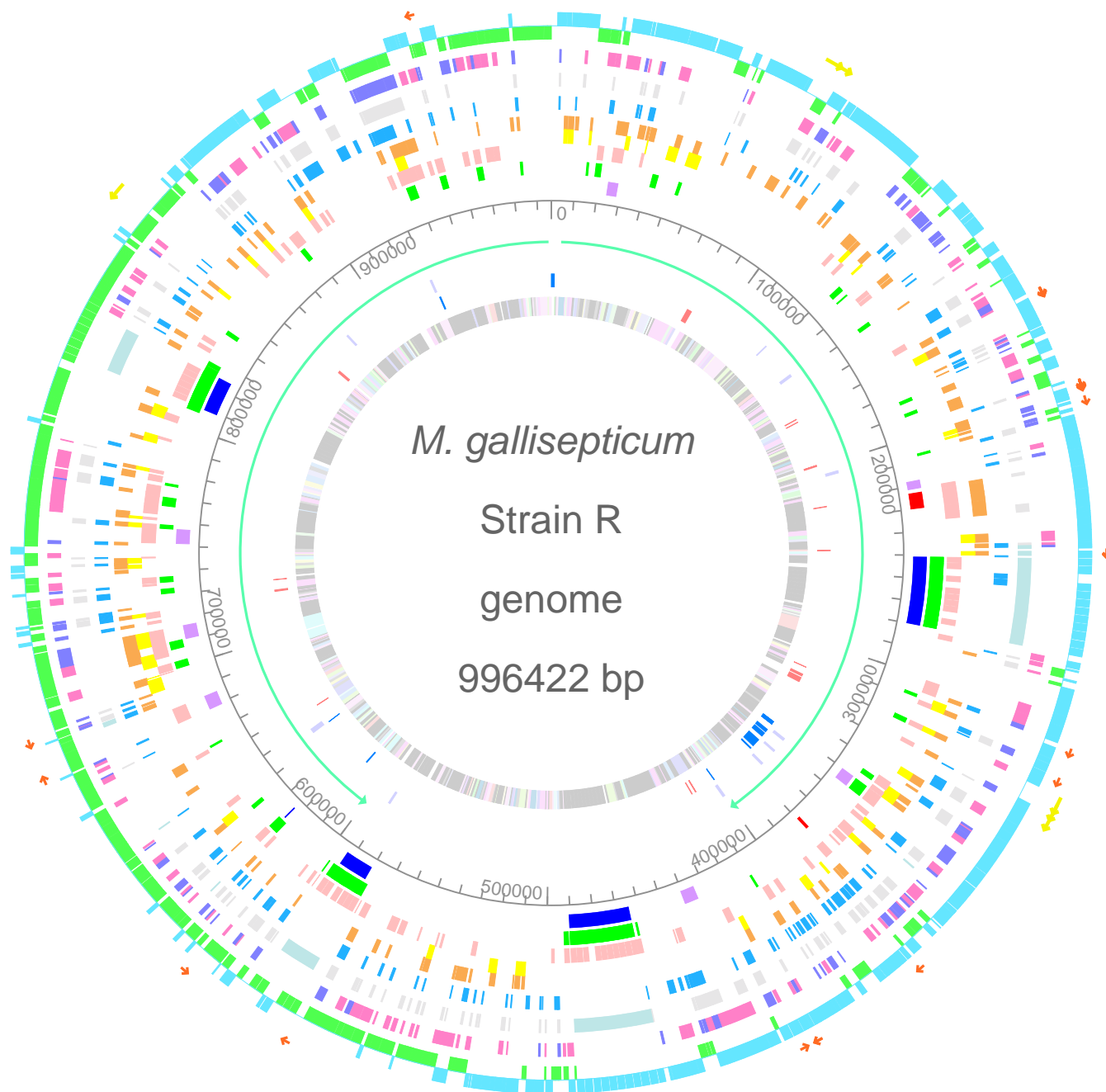


- Ribosomal RNAs
- tRNAs
- Genes in the direct strand
- Genes in the complementary strand
- Conserved hypotheticals
- Unique hypotheticals
- Genes with no homologues in other Mycoplasma spp.
- Genes which have homologues in only one or two Mycoplasma spp.
- Non-paralogous M. gallisepticum genes
- Proteins with >=2 transmembrane domains
- Proteins with ABC transporters and/or solute-binding motifs
- Proteins with lectin binding-like motifs
- Lipoproteins
- VlhA loci
- GapACrmA, CrmBC, CrmB-like
- high molecular weight proteins



- [J] Translation, ribosomal structure and biogenesis
- [K] Transcription
- [L] DNA replication, recombination and repair
- [D] Cell division and chromosome partitioning
- [O] Posttranslational modification, protein turnover, chaperones
- [M] Cell envelope biogenesis, outer membrane
- [N] Cell motility and secretion
- [P] Inorganic ion transport and metabolism
- [T] Signal transduction mechanisms

- [C] Energy production and conversion
- [G] Carbohydrate transport and metabolism
- [E] Amino acid transport and metabolism
- [F] Nucleotide transport and metabolism
- [H] Coenzyme metabolism
- [I] Lipid metabolism
- [Q] Secondary metabolites biosynthesis, transport and catabolism
- [R] General function prediction only
- [S] Function unknown

- Replication arrows
- Transposases
- GC rich >= 2,5 STD of the M. gall. genome
- AT rich >= 2,5 STD of the M. gall. genome