BCSL_114 (complete genomes)

- Genome annotation: Prokka
- Protein-coding gene annotations
- Pan-genome construction: Roary

59,989 orthologous sequence clusters

- Cluster filtering

8,954 orthologous sequence clusters

- Multiple sequence alignment: MAFFT

8,954 aligned sequence clusters

- Gene model construction: hmmbuild

HAMSTR_FULL_MGES_REMOVED