

The proposed plan for 2016

- Sequence
 - **January: Submit the REL-1509 chromosomes+MT to NCBI**
 - February: QC the assemblies at NCBI+get accessions, tell UCSC+Ensembl
 - February-September: Manual curation and improvement of protein coding regions
 - September: Submit updated chromosomes+lifted over annotation
- Annotation
 - January-February: QC+feedback of consensus gene sets
 - January-May: Manual annotation+checking of genes identified for the paper
 - **May: Load frozen paper version of annotations into public UCSC/Ensembl**
 - September: Liftover of updated annotation onto improved chromosomes
- Genomes paper
 - January-May: Work on analysis topics, figures, tables for the paper
 - May: Finalise analysis+section text+figures+tables for draft of paper
 - **June: Finalise paper for submission**
- July-September
 - Paper submitted and in review (reviewers can login to see the genomes in the public UCSC/Ensembl)