# 2019-02-27 Meeting notes

## Date

27 Feb 2019

## Attendees

- Clayton L Birkett

## Goals

## Discussion items

| Time | Item | Who | Notes |
|---|---|---|---|
| | GrainGenes | Taner | creating scripts (on github) to load JBrowse data<br>T3 can contribute scripts for our tracks |
| | data loading on T3 | | |
| | data loading on TriticumBase | Clay | T3/wheat new crop ontologies terms - accepted on github, will be published on crop ontology in a week or two<br>phenotype trials - working on bulk loading<br>locations - all loaded except sixteen trials that only have state and one with no location<br>genotype protocols |
| | WheatCAP | | user group conference call March 20th<br>summer workshop plans July 8-12 |
| | KnetMiner | | RefSeq v1.1<br>search by gene (without trait)<br>sorting genes by publication data<br>better ranking by score and p-value |
| | shared data files | | Eduard's 1000 wheat exomes |

## Action items

- ☑ Clayton L Birkett - can Eduard post shared data to SRA?
- ☐ Clayton L Birkett - ask Eduard if 1000 how to display in JBrowse (similar to Pangenome or other type of display?)