

2019-05-08 Meeting notes

Date

08 May 2019

Attendees

- [Clayton L Birkett](#)

Goals

Discussion items

Time	Item	Who	Notes
	GrainGenes	Taner	T3 logo, send icon to GrainGenes working with Docker
	Oat		<ul style="list-style-type: none">• genotype experiment - added CornellMetabolomics_2016 to replace CornellMetabolomics_2014• newly added oat lines to genotype• WI oat line synonyms
	Wheat	Clay	Select Lines by Genotype Experiment <ul style="list-style-type: none">• select more than one experiment• combine with the previous selection Download Portion of VCF file KASP Filter for Exome Capture
	New T3	David	download page (fixed) - output is coded as 0/1/2, marker names appended with UID
	New T3	Clay	genotype protocol - use the same as a project (T3 trial_code)

Action items

- ☐ [Clayton L Birkett](#) - look for a way to automate merging oat lines
- ☒ [Clayton L Birkett](#) - check combine lines code on each page.
- ☐ [Clayton L Birkett](#) - create line panels for 1000 exomes