2016-09-28 Meeting notes

Date

28 Sep 2016

Attendees

- Clayton L Birkett

Goals

Discussion items

Time	Item	Who	Notes
	PAG 2017	Clay	Managing Phenotypes (BMS, Fieldbook, GnpIS) or Application Interoperability (BRAPI, Flapjack, GOBII, BMS, KDDart, GnpIS SolR, WheatIS, QTLNetMiner)
	barley exome capture	Jean-Luc	Russell et al. 2016 Nature Genetics, could be used for imputation, loading in T3
	user group meeting	Clay	plan meeting for this Fall, coordinate with Kevin Smith
	breeders data farm merging	Clare	trait ontology (use general ontology or specific wheat ontology is available) breeding program vs data program
	Grain genes liason meeting		

Action items

- ☐ Clayton L Birkett change plot in training set optimization so that selected lines are not hidden
- ☑ Clayton L Birkett email Keven to set date for user group meeting
- ☑ Clayton L Birkett load barley hap map