2017-05-24 Meeting notes

Date

24 May 2017

Attendees

- Clayton L Birkett

Goals

Discussion items

Time	Item	Who	Notes
	JBrowse		- wheat pangenome - T3 markers track, gene presence track - limit of 2^29 or 536,000,000 bases for indexing vcf file using tabix - barley Morex assembly by Martin Mascher published, should be okay to publish on T3
	Knetminer		how should we link from T3 to Knetminer link from gene link to nearest gene all genes within specified distance of marker use markers sequence for BLAST so gene name not needed
	Plant Reactome		links from T3 - gene representation in pathway is easier to understand in new reactome release
	AgBioData paper		T3 installation documentation updated

Action items

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