

2017-06-28 Meeting notes

Date

28 Jun 2017

Attendees

- [Clayton L Birkett](#)

Goals

Discussion items

Time	Item	Who	Notes
	germplasm	Clay	• MCPD requires new fields ◦ optional or required? ◦ should we switch Breeding Programs to match FAO? • BrAPI calls • FAO plant genetic resources • Description of fields • Institute Code
	genome-specific primers in polyploids	Jean-Luc	Cristobal Uauy tool https://github.com/TGAC/bioruby-polyploid-tools
	wheat assembly	Clay	dividing chromosomes into _part1 and _part2
	SolR, WheatIS		create load file, no header entry_type - Experiment, Accession, Genetic Marker, Physical Marker, Phenotype database_name - Triticeae Toolbox db_id, db_version, description, url, speies, xref(not used), ... description of format - http://www.transplantdb.eu/node/4322

Action items

[Clayton L Birkett](#) - use MCPD codes for breeding codes so we don't need to add new fields to germplasm template, Clare will check if we can add entries not on list

[Clayton L Birkett](#) - polyploid tools not needed until the parents are loaded