

# QTLNetMiner

9/14/2016 Skype call with Keywan Hassani-Pak

He can add our data to their site.

We should use the IWGSC assembly. They are working on the NRGene assembly but don't plan to release it until January 2017.

T3 QTL Report

extend gene matches to within 1kb of gene

are introns included?

how to report multiple traits with same TO

include gene name in export, provide download link for complete data set

include information on VEP, variant effect predictor

need to use meta p-value and only send significant result

figure out how to link to QTLNetMiner

which trait ontology does QTLNetMiner use

Colaborate by email and have another skype call by next month.