BIG MISTAKE LOG

ABI Runs 17 and 18 used a sample input spreadsheet with the wrong plates!

It uses sample names from plates 1-4 instead of plates 17-20.

Runs 17 and 18 are plates 17-20 panels 2 and 3 respectively.

The plate names on the ABI and genemapper are:

Ots\_MCKR\_PT17-20\_2

Ots\_MCKR\_PT17-20\_3

The respective sheet in the excel notebook ABI\_input.xlsx is labeled with “error”. A corrected version is appended to the notebook labeled “corrected.” The correct input sheet is used on the other two 17-20 ABI runs, run 8 (panel 1) and run \_\_\_ (Oty3)

This cannot be fixed in the respective .fsa files because they are not human-readable. Instead the error must be fixed by renaming the samples after the finalized genotypes are output by genemapper.