1. For marker wells, the format is

<Sample type>\_Mk\_<labid>\_<markername>\_inputamount

1. For albumin wells with Dx sample, the format is

Dx\_ALB\_<labid>\_<inpputamount>

1. For albumin wells with follow up samples

FU\_ALB\_<labid>\_<dilution>

1. For albumin wells with MNC

MNC\_ALB \_<dilution>

1. For albumin wells with H2O

H2O\_ALB\_0