# METHODS AND SYSTEMS FOR EXPERIMENT DESIGN AND ANALYSIS

## BACKGROUND

**[0001]** Generally, users perform biological studies by gathering and comparing various sets of biological data. For example, a user may run various qPCR-based experiments to gather distinct types of biological data, such as genotyping data or gene expression data, about a gene of interest to the study. If the user wishes to compare various types of biological data, it is often done manually. The number of samples needed for one experiment may also be great and it is often difficult to isolate a particular sample for a user to examine and visualize so that a user may quickly and easily get valuable information from the data. Further, labs or facilities often operate several biological instruments. It may be difficult to track maintenance or calibration of the instruments

## SUMMARY

**[0002]** In one exemplary embodiment, a method for identifying a reaction site associated with an amplification curve from a plurality of amplification curves is provided. The method includes receiving amplification data from a plurality of reaction sites, wherein each reaction site contains a sample and generating a plurality of amplification curves from the amplification data. The method further includes displaying a first portion of the plurality of amplification curves on a display screen, and displaying a list of indications of reaction sites associated with the first portion of amplification curves alongside the first portion of amplification curves on the display screen. The method includes shifting from a view of the first portion of the plurality of amplification curves to display a view comprising a second portion of the plurality of amplification curves, wherein at least a portion of the second portion of the plurality of amplification curves are not included within the first portion of the plurality of amplification curves, and dynamically adjusting the list to display indications of reaction sites associated with viewable amplification curves included in the second portion of the plurality of amplification curves alongside the second portion of the plurality of amplification curves on the display screen. The list is configured to be scrollable.

**[0003]** In another exemplary embodiment, a non-transitory computer-readable storage medium encoded with processor-executable instructions, the instruction for identifying a reaction site associated with an amplification curve from a plurality of amplification curves, is provided. The instructions comprise instructions for receiving amplification data from a plurality of reaction sites, wherein each reaction site contains a sample, and generating a plurality of amplification

curves from the amplification data. The instructions further include instructions for displaying a first portion of the plurality of amplification curves on a display screen and displaying a list of indications of reaction sites associated with the first portion of amplification curves alongside the first portion of amplification curves on the display screen. The instructions further include instructions for shifting from a view of the first portion of the plurality of amplification curves to display a view comprising a second portion of the plurality of amplification curves, wherein at least a portion of the second portion of the plurality of amplification curves are not included within the first portion of the plurality of amplification curves, and dynamically adjusting the list to display indications of reaction sites associated with viewable amplification curves included in the second portion of the plurality of amplification curves alongside the second portion of the plurality of amplification curves on the display screen. The list is configured to be scrollable.

**[0004]** In yet another exemplary embodiment, a system for identifying a reaction site associated with an amplification curve from a plurality of amplification curves is provided. The system includes a processor; and a memory. The memory is encoded instructions, executable by the processor. The instructions include instructions for receiving amplification data from a plurality of reaction sites, wherein each reaction site contains a sample, and generating a plurality of amplification curves from the amplification data. The instructions further include instructions for displaying a first portion of the plurality of amplification curves on a display screen and displaying a list of indications of reaction sites associated with the first portion of amplification curves alongside the first portion of amplification curves on the display screen. The instructions further include instructions for shifting from a view of the first portion of the plurality of amplification curves to display a view comprising a second portion of the plurality of amplification curves, wherein at least a portion of the second portion of the plurality of amplification curves are not included within the first portion of the plurality of amplification curves, and dynamically adjusting the list to display indications of reaction sites associated with viewable amplification curves included in the second portion of the plurality of amplification curves alongside the second portion of the plurality of amplification curves on the display screen. The list is configured to be scrollable.

## DESCRIPTION OF THE FIGURES

**[0005]** FIG. 1 illustrates an example of a graphical user interface for displaying amplification curves according to various embodiments described herein;

**[0006]** FIG. 2 illustrates an example of a graphical user interface for viewing a selection of amplification curves according to various embodiments described herein;

**[0007]** FIG. 3 illustrates an example of a graphical user interface for identifying amplification curves according to various embodiments described herein;