

# CloneSelect Imager



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## Changes

## Build 1.3.73.1073

Changes and fixes made since version 1.3.56.1043 up to and including build 1.3.73.1073:

- 1. Added new licensable feature: Cell Migration to enable interpretation of cell migration.
- 2. A new pipeline named 'Very Low Contrast' has been added to improve confluence calculation when using very low contrast cell lines.
- **3.** Fixed bug. If the results folder for a plate is deleted or moved a meaningful message is now displayed instead of an exception alert.

### Build 1.3.56.1043

Changes and fixes made since version 1.3.51.990, up to and including build 1.3.56.1043:

- 1. Fixed a number of issues with the export of data.
- 2. Option to support multiple results locations has been added.
- 3. Two extra functions have been added to the automation interface.

#### Build 1.3.51.990

Changes and fixes made since version 1.3.45.961, up to and including build 1.3.51.990:

- 1. Fixed Batch Plate Imaging process so that user selected cell detection methods are used correctly.
- 2. Fixed data export of loci information to the Excel xml file format.
- **3.** Fixed system hang during imaging when enhance images is disabled and images are being stored in collections.
- 4. Fixed crash in whole well image display when contrast is altered.
- 5. Moving between time points on the loci tab now does not cause the zoom to reset.
- 6. The pick list on the loci tab is now not reset if the user moves to a different tab.
- 7. The pick list on the loci tab now tracks the user selected item.

### Build 1.3.45.961

Changes and fixes made since version 1.2.4.871, up to and including build 1.3.45.961:

- 1. A new cell detection method has been added to improve detection of low contrast cell lines.
- 2. A new cell detection method has been added to improve detection of colonies.
- 3. Barcode or plate name can be changed in results review.
- Time sequence images in bmp, png and jpg formats are exportable from the loci tab of image review.
- 5. Area histogram on the loci tab now uses  $\mu m^2$  instead of mm<sup>2</sup>.
- 6. Area and compactness histograms on the loci tab allow gating of large as well as small features.
- 7. Number of histogram bins can be set when exporting loci data.

- 8. Users can now give a descriptive name to a well on the confluence, cell number and loci tabs.
- 9. Wells can be excluded from loci processing on the thumbnail view tab.
- **10.** Images can be stored in a collection on disk instead of individual files. This decreases the time taken to move, backup and delete results.

#### Build 1.2.4.871

Changes and fixes made since version 1.2.2.848, up to and including build 1.2.4.871:

- 1. Fixed a problem when loci count analysis occurred outside of mask.
- 2. Added export of compactness and area data for every found feature to csv, xml and excel xml files.

#### Build 1.2.2.848

Changes and fixes made since version 1.1.0.818, up to and including build 1.2.2.848:

- 1. Changed automated imaging process to support image compression
- 2. Added 'Are you sure' prompt when pausing an ICCMS process
- 3. Enforced requirement for images to be saved to local disk
- 4. Added error report generation facility

#### Build 1.1.0.818

Changes and fixes made since version 1.0.1.807, up to and including build 1.1.0.818:

- 1. Resolved memory leak when loading image files
- 2. Added more decimal places for loci area
- 3. Minor performance improvements

#### Build 1.0.1.807

Changes and fixes made since version 1.0.1.795, up to and including build 1.0.1.807:

- 1. Is now compatible with the new OEM integration kit (WCF-based)
- 2. Fixes an issue where the plate row and column pitches were not used correctly
- 3. Ensures that the magnification caption is visible on the image display when plates are being imaged
- 4. Removes the black line drawn around the red mask area to make it less intrusive
- 5. Removes the Hardware model view from general display
- 6. Renames the ColdFire upload/download operations to make it more obvious what is happening
- 7. Fixes an issue where manual barcoding was not working correctly
- 8. Fixes a problem where dependant devices cannot connect if they are created in a specific order

### Build 1.0.1.795

Changes and fixes made since version 0.4.0.428, up to and including 1.0.1.795:

- 1. Full well view for all microplate formats
- 2. Improved auto focus with feedback
- 3. Improved growth curves display with selectable wells, well thumbnails and fitted curves
- 4. Summary page showing when the plate was imaged and growth trends
- 5. Faster loci counting with improved criteria display using histograms
- 6. Added 'back' button during plate focusing etc.
- 7. Store name of PC images are taken on to facilitate tracking data from multiple instruments
- 8. Selectable confluence generation algorithms
- 9. Added ability to remove cell estimation formulae
- 10. Ability to toggle confluence overlay display in focus screen
- **11.** Added ability to export graph images
- **12.** Added HTML plate report
- 13. Export clone size classes when loci counting
- 14. Added support for Genetix plate loader
- 15. Numerous usability and performance improvements

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