

MiniSeq Software Release Notes

MiniSeq System Suite Installer v1.0.1

MiniSeq Control Software v1.0.0

MiniSeq Service Software v1.0.1

MiniSeq RTA v2.8.6

BaseSpace Broker v2.8.1

MiniSeq Recipe Installer v1.0.0

Run Copy Service v1.0.20

Illumina DMA Driver v4.5.3

For MiniSeq Systems

Introduction

These Release Notes detail the key software components for the Illumina MiniSeq Software, highlighting known issues in the initial release.

Software Application	Prior Version	New Version
MiniSeq System Suite Installer	N/A	1.0.1
MiniSeq Control Software	N/A	1.0.0
MiniSeq Service Software	N/A	1.0.1
MiniSeq RTA	N/A	2.8.6
BaseSpace Broker	N/A	2.8.1
MiniSeq Recipe Installer	N/A	1.0.0
Run Copy Service	N/A	1.0.20
Illumina DMA Driver	N/A	4.5.3

This release supports the initial product launch of the MiniSeq System.

For more information about the MiniSeq System and how to use the software included with it, see the product and support pages at [illumina.com](http://www.illumina.com):

<http://www.illumina.com/systems/miniseq.html>

http://support.illumina.com/sequencing/sequencing_instruments/miniseq.html

I. MiniSeq System Suite Installer v.1.0

KNOWN ISSUES:

- None

II. MiniSeq Control Software v1.0

KNOWN ISSUES:

- When in BaseSpace Cloud or BaseSpace Onsite mode, indexes in the Run Setup screen are not editable. If editing is desired, the user can launch a browser, edit in BaseSpace, refresh the list in the Control Software, and reselect the run.

III. MiniSeq Service Software (MNSS) v1.0

KNOWN ISSUES:

- When selected, the exit button at the top of the screen leaves the instrument in an initialized state, potentially leading to instrument damage. To exit MiniSeq Service Software, use the Exit button in the bottom navigation bar. To close the Service Software and uninitialized the instrument, click the minimize button within the Service Software interface. Then right click the application icon in the Windows taskbar or press and hold with the touch screen to close the App.

IV. MiniSeq RTA v2.8.6

KNOWN ISSUES:

- Template generation may yield cluster locations that are within one tenth of a pixel of each other, leading to fastq read names that are identical. If downstream applications don't support this, only one read with that name should be retained in the fastq file.

V. BaseSpace Broker v2.8

KNOWN ISSUES:

- None

VI. MiniSeq Recipe Installer v1.0

KNOWN ISSUES:

- None

VII. Run Copy Service v1.0

KNOWN ISSUES:

- None

VIII. Illumina DMA Driver v4.5

KNOWN ISSUES:

- None

Revision History

Version	ER#	Originator	Description of Change
00	1006905	B. Chu	Initial version