

# BaseSpace Core Apps Software Release Notes

### MethylSeq v1.0

For BaseSpace

April 28, 2015

## illumına<sup>.</sup>

### Introduction

These Release Notes detail notable items for the MethylSeq App v1.0 release. This app is deployed on BaseSpace Cloud.

For more information about this app and how to use it, refer to the app User Guides, available from the details page of each app, and the BaseSpace Support Page, Documentation and Literature, on illumina.com.

http://support.illumina.com/sequencing/sequencing\_software/basespace/documentation.ilmn

The software package includes:

• MethylSeq v1.0

#### I. MethylSeq v1.0

**New Features:** 

• Initial Release In BaseSpace

**DEFECT REPAIRS** 

• None

KNOWN ISSUES:

- Bedgraph files output by the MethylSeq app are currently not viewable in the BaseSpace IGV.
  - As a workaround methylation levels can be viewed in a locally installed IGV by downloading the bedGraph file, and converting to a tdf or bigwig file format.