

Local Run Manager

DNA Enrichment Analysis Module

v2.0.0

Release Notes

09-FEB-2018

Introduction

These Release Notes detail notable items for the Local Run Manager DNA Enrichment Analysis Module v2.0.0 release.

Please note that the DNA Enrichment Analysis Module v2.0.0 requires Local Run Manager Framework v2.0.0 or higher and is not compatible with Local Run Manager Framework v1.3.1 or lower.

For more information about this analysis module and how to use it, refer to the Local Run Manager DNA Enrichment Analysis Module Workflow Guide, available from the Local Run Manager Support Page, Documentation & Literature, at illumina.com.

[HTTPS://SUPPORT.ILLUMINA.COM/SEQUENCING/SEQUENCING_SOFTWARE/LOCAL-RUN-MANAGER.HTML](https://support.illumina.com/sequencing/sequencing_software/local-run-manager.html)

NEW FEATURES:

- Added support for processing datasets generated from iSeq 100, MiniSeq, MiSeq, and NextSeq 550 sequencing systems.
- Added support for importing and exporting Sample Sheet files on the Run Setup page.
- Added support for the following Library Prep Kits:
 - IDT-ILMN TruSeq DNA-RNA UD Indexes 24 Indexes
 - IDT-ILMN TruSeq DNA-RNA UD Indexes 96 Indexes
 - Nextera DNA CD Indexes - 24
 - Nextera DNA CD Indexes - 96
- Added support for entering custom index sequences directly on the Run Setup page.
- Added support for the following special characters to be used as part of the "Run Name" and "Run Description" on the Run Setup page:
` ~ ! @ # \$ % - _ { }
- Increased the maximum number of samples on the Run Setup page from 384 to 1536.

DEFECT REPAIRS:

- Fixed an issue where the HTML report file incorrectly listed the software version numbers of Picard and GATK as "NA".

KNOWN ISSUES:

- None