



# **Sequencing Analysis Viewer**

## **Software Release Notes**

### **SAV 2.4.7**

**04-FEB-2019**



## Introduction

These Release Notes detail the key changes to the Sequencing Analysis Viewer (SAV) Software since the release of SAV version 2.4.5.

<b>Software Application</b>	<b>Prior Version</b>	<b>New Version</b>
<i>Sequencing Analysis Viewer</i>	2.4.5	2.4.7

Sequencing Analysis Viewer version 2.4.7 is used to visualize sequencing run data from MiniSeq, MiSeq, NextSeq, HiScanSQ, all HiSeq (2000, 2500, 3000, 4000, and HiSeq X), and NovaSeq. For data generated on Illumina GAIIx sequencing instruments, continue using SAV 1.8.37.

This SAV version requires Windows .NET Framework 4.5.1 and Visual C++ Redistributable for Visual Studio 2015. If your system is using Windows .NET Framework 4 or you are running a MiSeq with MiSeq Control Software v2.5 and lower, continue using SAV 1.8.37.

### NEW FEATURES:

- Added occupancy information to Analysis and Imaging tabs for iSeqs and MiniSeqs
- Added %PF as a metric on the Analysis tab
- Added automatic single-surface detection from the RunInfo tileset

### IMPROVEMENTS:

- Improved the consistency of cluster count and cluster density notations across SAV
- Made channel naming capitalization consistent with RunInfo.xml