

Raw data delivery note

SciLifeLab Stockholm

July 15, 2015

Project name

A.Blomberg_15_01

UPPNEX project id

b2011228

Flow cell id

AC6VGLANXX

Lane

7

Sequence data directory

/proj/b2011228/INBOX/A.Blomberg_15_01/P1887_101/150521_AC6VGLANXX

Sample

P1887_101 / Fves11. Ordered amount: 188 million read pairs.

Method

Clustered using cBot and sequenced on HiSeq2500 (HiSeq Control Software 2.2.58/RTA 1.18.64) with a 2x126 setup in RapidHighOutput mode. Bcl to Fastq conversion was performed using bcl2Fastq v1.8.3 from the CASAVA software suite. The quality scale is Sanger / phred33 / Illumina 1.8+.

Results

265.7 million read pairs. Average quality score 34.29 (89.90% bases \geq Q30).