

Institutionen för medicinska vetenskaper SNP&SEQ-teknologiplattformen



Report on sequencing results

The SNP&SEQ Technology Platform at Uppsala University REPORT issued by an Accredited Laboratory

Project name Allele-specific gene expression (ASE) in human white blood

cells with induced inflammation

Project code JA3

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Report ID JA3_130130_Report_on_sequencing_results

Report sent to Jacob Odeberg, <u>jacob@biotech.kth.se</u>

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Date of delivery January 30, 2013

Report issued by Olof Karlberg, Department of Medical Sciences, Molecular

Medicine, Uppsala University, 751 85 Uppsala, Sweden

seq@medsci.uu.se, www.sequencing.se

Sequencing performed at The SNP&SEQ Technology Platform in Uppsala, Uppsala

University, Dept. of Medical Sciences, Molecular Medicine, Uppsala University Hospital, entr. 70, 3rd floor, Res. Dept. 2,

751 85 Uppsala, Sweden

Sequencing performed using instrument type

HiSeq2000 Paired End 100bp

Result files Files delivered to UPPNEX account b2013015

/proj/b2013015/INBOX/130104_SN866_0197_AC1DLVACXX /proj/b2013015/INBOX/130104_SN866_0198_BC1DAYACXX

Additional files and folders The results are divided into one folder per sample.

For more information about the files and folders, see the file report.html, which also contains detailed information about

the run and all samples.

Comments Tile s 4 2113 had to be discarded from lane 4 on

130104_SN866_0197_AC1DLVACXX

Clustering and sequencing are done according to accredited methods. Library preparation for RNA sequencing is not an

accredited method

P:\Projekt\Pågående projekt\SEKVENSNING\JA3_Sarah-JayneReilly\Resultatrapport\JA3_130130_Report on sequencing results.docx
Utskrift: 2013-01-30 17:49 / karlol Sidan 1/1 BLA 0120-4