SciLifeLab

Project status note

SciLifeLab Stockholm

July 18, 2015

Project name

T.Backhaus_15_01 (Second re-run of T.Backhaus_14_02 - Pseudomonas putida silver)

UPPNEX project id

b2014338

Sequence data directories

/proj/b2014338/INBOX/T.Backhaus_15_01/

Flowcells Delivered

150702_AC6U4VANXX

Samples

ScilifeID	SubmittedID	BarcodeSeq	MSequenced	MOrdered
P2201_101	0,001A Ag+	GAATTCGT-GGCTCTGA	7.573077	2.925
P2201_102	0,01A Ag+	TCCGCGAA-ATAGAGGC	6.18103	2.925
P2201_103	0,04A Ag+	TCCGCGAA-CCTATCCT	6.345198	2.925
P2201_104	ctrlB Ag+	TCCGCGAA-GGCTCTGA	6.473671	2.925
P2201_105	0,001B Ag+	TCCGCGAA-AGGCGAAG	5.768181	2.925
P2201_106	0,01B Ag+	TCCGCGAA-TAATCTTA	6.600037	2.925
P2201_107	0,04B Ag+	GAATTCGT-AGGCGAAG	7.204352	2.925
P2201_108	ctrlD Ag+	TCCGCGAA-GTACTGAC	6.759836	2.925
P2201_109	0,001c Ag+	TCTCGCGC-TATAGCCT	5.879769	2.925
P2201_110	0,01c Ag+	TCTCGCGC-ATAGAGGC	6.746163	2.925
P2201_111	0,04c Ag+	TCTCGCGC-CCTATCCT	6.056807	2.925
P2201_112	ctrlC Ag+	TCTCGCGC-GGCTCTGA	6.690721	2.925
P2201_113	0,004A AgNP	TCTCGCGC-AGGCGAAG	5.803687	2.925
P2201_114	0,04A AgNP	TCTCGCGC-TAATCTTA	6.38459	2.925
P2201_115	0,12A AgNP	TCTCGCGC-CAGGACGT	6.731037	2.925
P2201_116	ctrlA AgNP	GAATTCGT-TAATCTTA	8.866899	2.925
P2201_117	0,004B AgNP	AGCGATAG-TATAGCCT	6.461851	2.925
P2201_118	0,04B AgNP	AGCGATAG-ATAGAGGC	5.732191	2.925
P2201_119	0,12B AgNP	AGCGATAG-CCTATCCT	7.191323	2.925
P2201_120	ctrlE AgNP	AGCGATAG-GGCTCTGA	7.126219	2.925

P2201_121	0,004C	AgNP	AGCGATAG-AGGCGAAG	6.732849	2.925
P2201_122	0,04C	AgNP	AGCGATAG-TAATCTTA	6.523513	2.925
P2201_123	0,12C	AgNP	AGCGATAG-CAGGACGT	7.876543	2.925
P2201_124	ctrlF	AgNP	AGCGATAG-GTACTGAC	7.375041	2.925

Information

Naming conventions

The data is delivered in fastq format using Illumina 1.8 quality scores. There will be one file for the forward reads and one file for the reverse reads (if the run was a paired-end run). The naming of the files follow the convention: [LANE]_[DATE]_[FLOWCELL]_[SCILIFE NAME]_[READ].fastq.gz

Data access at UPPMAX

Data from the sequencing will be uploaded to the UPPNEX (UPPMAX Next Generation sequence Cluster Storage, www.uppmax.uu.se), from which the user can access it. You can find the data in the INBOX folder of the UPPNEX project, which was created for you when your order was placed, e.g. /proj/b2013000/INBOX/J.Doe_13_01 If you have problems to access your data, please contact SciLifeLab genomics_support@scilifelab.se. If you have questions regarding UPPNEX, please contact support@uppmax.uu.se.

Acknowledgement

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