

## Project status note

### SciLifeLab Stockholm

**2014-08-11**

#### ***Project name***

A.Blomberg\_14\_03 (Debaryomyces/Skeletonema - Nextera XT)

#### ***UPPNEX project id***

b2011228

#### ***Sequence data directories***

/proj/b2011228/INBOX/A.Blomberg\_14\_03

#### ***Samples***

ScilifeID	Submitted D	BarcodeSeq	MSequenced	MOrdered	Status
P1260_1001	A1_Debaryo myces	TAAGGCGA- TAGATCGC	5.94	N/A	P
P1260_1002	B1_Debaryo myces	TAAGGCGA- CTCTCTAT	3.01	N/A	P
P1260_1003	C1_Debaryo myces	TAAGGCGA- TATCCTCT	5.54	N/A	P
P1260_1004	D1_Debaryo myces	TAAGGCGA- AGAGTAGA	4.01	N/A	P
P1260_1005	E1_Debaryo myces	TAAGGCGA- GTAAGGAG	5.01	N/A	P
P1260_1006	F1_Debaryo myces	TAAGGCGA- ACTGCATA	4.78	N/A	P
P1260_1007	G1_Debaryo myces	TAAGGCGA- AAGGAGTA	5.78	N/A	P
P1260_1008	H1_Debaryo myces	TAAGGCGA- CTAAGCCT	3.25	N/A	P
P1260_1009	A2_Debaryo myces	CGTACTAG- TAGATCGC	4.83	N/A	P
P1260_1010	B2_Debaryo myces	CGTACTAG- CTCTCTAT	3.23	N/A	P
P1260_1011	C2_Debaryo myces	CGTACTAG- TATCCTCT	1.95	N/A	P
P1260_1012	D2_Debaryo myces	CGTACTAG- AGAGTAGA	5.29	N/A	P

P1260_1013	E2_Debaryo myces	CGTACTAG- GTAAGGAG	4.57	N/A	P
P1260_1014	F2_Debaryo myces	CGTACTAG- ACTGCATA	5.74	N/A	P
P1260_1015	G2_Debaryo myces	CGTACTAG- AAGGAGTA	5.10	N/A	P
P1260_1016	H2_Debaryo myces	CGTACTAG- CTAAGCCT	6.57	N/A	P
P1260_1017	A3_Debaryo myces	AGGCAGAA- TAGATCGC	4.63	N/A	P
P1260_1018	B3_Debaryo myces	AGGCAGAA- CTCTCTAT	6.29	N/A	P
P1260_1019	C3_Debaryo myces	AGGCAGAA- TATCCTCT	4.66	N/A	P
P1260_1020	D3_skeleto nema	AGGCAGAA- AGAGTAGA	6.60	N/A	P
P1260_1021	E3_skeleto nema	AGGCAGAA- GTAAGGAG	17.78	N/A	P
P1260_1022	F3_skeleto nema	AGGCAGAA- ACTGCATA	11.03	N/A	P
P1260_1023	G3_skeleto nema	AGGCAGAA- AAGGAGTA	10.01	N/A	P
P1260_1024	H3_skeleto nema	AGGCAGAA- CTAAGCCT	5.20	N/A	P
P1260_1025	A4_skeleto nema	TCCTGAGC- TAGATCGC	25.60	N/A	P
P1260_1026	B4_skeleto nema	TCCTGAGC- CTCTCTAT	14.46	N/A	P
P1260_1027	C4_skeleto nema	TCCTGAGC- TATCCTCT	9.61	N/A	P
P1260_1028	D4_skeleto nema	TCCTGAGC- AGAGTAGA	15.14	N/A	P
P1260_1029	E4_skeleto nema	TCCTGAGC- GTAAGGAG	2.03	N/A	P
P1260_1030	F4_skeleto nema	TCCTGAGC- ACTGCATA	26.99	N/A	P
P1260_1031	G4_skeleto nema	TCCTGAGC- AAGGAGTA	5.79	N/A	P
P1260_1032	H4_skeleto nema	TCCTGAGC- CTAAGCCT	31.49	N/A	P
P1260_1033	A5_skeleto nema	CAGAGAGG- CTAAGCCT	28.04	N/A	P

## Comments

Project finished.

## ***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](http://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](mailto:genomics_support@scilifelab.se). If you have questions regarding UPPNEX, please contact [support@uppmax.uu.se](mailto:support@uppmax.uu.se).

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