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Sequence Data Delivery Note

Project VB2077 17.11.2020

The NGS libraries (30 samples) were single-read sequenced on an Illumina NextSeq 500 system using 75 bp read length.

Sample ID	Sample Name	Barcode i5	Barcode i7	Sequence file (R1/R2 in case of PE)	Number of Reads
R-2010185	1_1	AGGCTATA	ATTACTCG	ID-006413_S1_R1_001.fastq.gz	11.831.771
R-2010186	1_2	GCCTCTAT	ATTACTCG	ID-006414_S2_R1_001.fastq.gz	11.554.354
R-2010187	1_3	AGGATAGG	ATTACTCG	ID-006415_S3_R1_001.fastq.gz	11.566.711
R-2010188	1_4	TCAGAGCC	ATTACTCG	ID-006416_S4_R1_001.fastq.gz	11.394.009
R-2010189	1_5	CTTCGCCT	ATTACTCG	ID-006417_S5_R1_001.fastq.gz	11.575.416
R-2010190	1_6	TAAGATTA	TCCGGAGA	ID-006418_S6_R1_001.fastq.gz	12.231.108
R-2010191	1_7	ACGTCCTG	TCCGGAGA	ID-006419_S7_R1_001.fastq.gz	13.640.111
R-2010192	1_8	GTCAGTAC	TCCGGAGA	ID-006420_S8_R1_001.fastq.gz	12.515.989
R-2010193	1_9	ATAGAGAG	TCCGGAGA	ID-006421_S9_R1_001.fastq.gz	11.478.071
R-2010194	1_10	AGAGGATA	TCCGGAGA	ID-006422_S10_R1_001.fastq.gz	11.900.490
R-2010195	2_1	CTCCTTAC	CTGAAGCT	ID-006423_S11_R1_001.fastq.gz	12.361.921
R-2010196	2_2	TATGCAGT	CTGAAGCT	ID-006424_S12_R1_001.fastq.gz	13.003.672
R-2010197	2_3	TACTCCTT	CTGAAGCT	ID-006425_S13_R1_001.fastq.gz	11.417.069
R-2010198	2_4	AGGCTTAG	CTGAAGCT	ID-006426_S14_R1_001.fastq.gz	12.936.520
R-2010199	2_5	ATTAGACG	CTGAAGCT	ID-006427_S15_R1_001.fastq.gz	12.519.808
R-2010200	2_6	CGGAGAGA	CGGCTATG	ID-006428_S16_R1_001.fastq.gz	11.660.466
R-2010201	2_7	CTAGTCGA	CGGCTATG	ID-006429_S17_R1_001.fastq.gz	11.799.811
R-2010202	2_8	CTTAATAG	CGGCTATG	ID-006430_S18_R1_001.fastq.gz	13.672.938
R-2010203	2_9	ATAGCCTT	CGGCTATG	ID-006431_S19_R1_001.fastq.gz	11.205.089
R-2010204	2_10	TAAGGCTC	CGGCTATG	ID-006432_S20_R1_001.fastq.gz	11.350.925
R-2010205	3_1	TCGCATAA	TCCGCGAA	ID-006433_S21_R1_001.fastq.gz	11.544.070
R-2010206	3_2	AATTTGGT	TCCGCGAA	ID-006434_S22_R1_001.fastq.gz	13.260.062
R-2010207	3_3	GCTACGGT	TCCGCGAA	ID-006435_S23_R1_001.fastq.gz	13.638.425
R-2010208	3_4	CATCCCGT	TCCGCGAA	ID-006436_S24_R1_001.fastq.gz	12.967.228
R-2010209	3_5	CTTATAGT	TCCGCGAA	ID-006437_S25_R1_001.fastq.gz	14.460.722
R-2010210	3_6	AGGTCGTT	AGCGATAG	ID-006438_S26_R1_001.fastq.gz	11.555.018
R-2010211	3_7	ACCTTCTT	AGCGATAG	ID-006439_S27_R1_001.fastq.gz	11.414.454
R-2010212	3_8	TAGCTCTT	AGCGATAG	ID-006440_S28_R1_001.fastq.gz	12.128.143
R-2010213	3_9	CGAGACTT	AGCGATAG	ID-006441_S29_R1_001.fastq.gz	11.377.770
R-2010214	3_10	TTCGCTGT	AGCGATAG	ID-006442_S30_R1_001.fastq.gz	10.693.456

Summary of sequencing Run

Total Reads	475.005.304	
Cluster(PF)	422.925.394	
% Cluster	89,0	
Reads ident.	364.655.597	
% Index	86,2	

The NGS library pool was analysed on a Shimadzu MultiNA microchip electrophoresis system.

