

# Maryland Genomics Illumina Report



This report contains run information and quality metrics for an Illumina run generated by Maryland Genomics. If the project requires multiple flowcell runs, a separate report will be generated for each flowcell. For additional information or questions, please contact [IGS-Services@SOM.UMaryland.edu](mailto:IGS-Services@SOM.UMaryland.edu)

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303186	IL100305238	MCR.CVU.125	MCR.CVU.125	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-F04	ACCGTAAG-CATATCCA
3	XUMMR	MS100303363	IL100303407	MCR.CVU.295	MCR.CVU.295	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N5UD-F02	AAGGCCCT-TAACCAGT
3	XUMMR	MS100303325	IL100305473	MCR.CVU.261	MCR.CVU.261	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-A10	ACGTCGAG-TTCTCTA
3	XUMMR	MS100303165	IL100305217	MCR.CVU.104	MCR.CVU.104	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-A02	AAGCGACT-ACGAATCC
3	XUMMR	MS100303345	IL100305493	MCR.CVU.281	MCR.CVU.281	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-E12	CAGTGACG-TACCGCTC
3	XUMMR	MS100303088	IL100303620	MCR.CVU.29	MCR.CVU.29	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-D04	GTAACCGA-AAGACACC
3	XUMMR	MS100303210	IL100305262	MCR.CVU.149	MCR.CVU.149	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-F07	CTTTCCTT-GGCTTACT
3	XUMMR	MS100303198	IL100305250	MCR.CVU.137	MCR.CVU.137	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-B06	CTTGACGA-GCACACAA
3	XUMMR	MS100303269	IL100305417	MCR.CVU.205	MCR.CVU.205	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-A03	TTGTTGCA-ACGACGTC
3	XUMMR	MS100303122	IL100303654	MCR.CVU.63	MCR.CVU.63	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-F08	GAATCACC-GAACGAAG
3	XUMMR	MS100303258	IL100305406	MCR.CVU.194	MCR.CVU.194	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-F01	TAGTAATC-GCAGAGCC
3	XUMMR	MS100303237	IL100305289	MCR.CVU.176	MCR.CVU.176	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-A11	AAGGAAGG-ACCGGAGT
3	XUMMR	MS100303362	IL100303406	MCR.CVU.294	MCR.CVU.294	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N5UD-E02	AGCGCAAA-GTACTCGC
3	XUMMR	MS100303084	IL100303616	MCR.CVU.25	MCR.CVU.25	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-H03	GCGTTAGA-CAGCATAC
3	XUMMR	MS100303240	IL100305292	MCR.CVU.179	MCR.CVU.179	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-D11	ATTCGAGC-CTAGGTTG
3	XUMMR	MS100303331	IL100305479	MCR.CVU.267	MCR.CVU.267	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-G10	GTTCTGCG-TGTCTCT

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Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303288	IL100305436	MCR.CVU.224	MCR.CVU.224	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-D05	TATTTACC-CTCACGTC
3	XUMMR	MS100303130	IL100303662	MCR.CVU.71	MCR.CVU.71	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-F09	GGCAAGTT-CTCAGAAG
3	XUMMR	MS100303244	IL100305296	MCR.CVU.183	MCR.CVU.183	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-H11	CAGCTTCG-ACCCGTTG
3	XUMMR	MS100303330	IL100305478	MCR.CVU.266	MCR.CVU.266	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-F10	GATGGAAA-TTGTGCCC
3	XUMMR	MS100303222	IL100305274	MCR.CVU.161	MCR.CVU.161	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-B09	AAGGCGTA-CCATCCGC
3	XUMMR	MS100303270	IL100305418	MCR.CVU.206	MCR.CVU.206	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-B03	CCACACTT-GTATGACG
3	XUMMR	MS100303195	IL100305247	MCR.CVU.134	MCR.CVU.134	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-G05	ATTGGACA-TTGCTGGA
3	XUMMR	MS100303171	IL100305223	MCR.CVU.110	MCR.CVU.110	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-G02	CGCTCTTA-CATACCGT
3	XUMMR	MS100303113	IL100303645	MCR.CVU.54	MCR.CVU.54	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-E07	GTCGATTG-AAGACCGT
3	XUMMR	MS100303199	IL100305251	MCR.CVU.138	MCR.CVU.138	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-C06	GCCACGAC-CCTCGGGT
3	XUMMR	MS100303241	IL100305293	MCR.CVU.180	MCR.CVU.180	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-E11	CACCTGTA-TGTGTCAG
3	XUMMR	MS100303352	IL100303396	MCR.CVU.284	MCR.CVU.284	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N5UD-C01	GCCCAGTG-AATTGCGG
3	XUMMR	MS100303090	IL100303622	MCR.CVU.31	MCR.CVU.31	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-F04	CATGAGCA-GATCTTGC
3	XUMMR	MS100303227	IL100305279	MCR.CVU.166	MCR.CVU.166	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-G09	CAGCAGTC-GTTGGCGT
3	XUMMR	MS100303291	IL100305439	MCR.CVU.227	MCR.CVU.227	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-G05	CGTCTTGG-CGCCAGTC
3	XUMMR	MS100303125	IL100303657	MCR.CVU.66	MCR.CVU.66	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-A09	CGCAACTA-TCGGATTC
3	XUMMR	MS100303187	IL100305239	MCR.CVU.126	MCR.CVU.126	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-G04	TATGACGT-CGCTAATC
3	XUMMR	MS100303128	IL100303660	MCR.CVU.69	MCR.CVU.69	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-D09	CAATAGCC-TCTACGCA
3	XUMMR	MS100303095	IL100303627	MCR.CVU.36	MCR.CVU.36	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-C05	GGAGGAAT-GAACGGTT
3	XUMMR	MS100303153	IL100303685	MCR.CVU.95	MCR.CVU.95	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-E12	GGACATCA-TACCTGCA
3	XUMMR	MS100303140	IL100303672	MCR.CVU.81	MCR.CVU.81	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-H10	CTCGACTT-ATCTGACC
3	XUMMR	MS100303364	IL100303408	MCR.CVU.296	MCR.CVU.296	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N5UD-G02	GCGTACGG-CGAAACT
3	XUMMR	MS100303341	IL100305489	MCR.CVU.277	MCR.CVU.277	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-A12	TATACCTC-TAGTCACA

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303366	IL100303410	MCR.CVU.297	MCR.CVU.297	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated	N5UD-A03	ATAGCAGA-ATGTCTTA
3	XUMMR	MS100303257	IL100305405	MCR.CVU.193	MCR.CVU.193	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-E01	GAAGAGGG-CGGACGGA
3	XUMMR	MS100303268	IL100305416	MCR.CVU.204	MCR.CVU.204	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-H02	GAGCACTC-GCAACTTG
3	XUMMR	MS100303202	IL100305254	MCR.CVU.141	MCR.CVU.141	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-F06	TTGCGTTA-TTCCCGAA
3	XUMMR	MS100303292	IL100305440	MCR.CVU.228	MCR.CVU.228	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-H05	CGTAGCGA-CTAAACAA
3	XUMMR	MS100303118	IL100303650	MCR.CVU.59	MCR.CVU.59	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-B08	TGCGTAAC-AGCCTATC
3	XUMMR	MS100303206	IL100305258	MCR.CVU.145	MCR.CVU.145	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-B07	ACAAGGCA-TCGCGCAA
3	XUMMR	MS100303166	IL100305218	MCR.CVU.105	MCR.CVU.105	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-B02	TGATAGGC-GTCTGAGT
3	XUMMR	MS100303315	IL100305463	MCR.CVU.251	MCR.CVU.251	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-G08	TGAATCTT-CTGTTTAC
3	XUMMR	MS100303262	IL100305410	MCR.CVU.198	MCR.CVU.198	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-B02	AGAGCTGC-GTTGAAGG
3	XUMMR	MS100303314	IL100305462	MCR.CVU.250	MCR.CVU.250	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-F08	TACAAGTC-GAAGTGAA
3	XUMMR	MS100303164	IL100305216	MCR.CVU.103	MCR.CVU.103	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-H01	ATCCGCAG-ACACGACT
3	XUMMR	MS100303192	IL100305244	MCR.CVU.131	MCR.CVU.131	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-D05	CTAGGTTG-ATTCGAGC
3	XUMMR	MS100303334	IL100305482	MCR.CVU.270	MCR.CVU.270	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-B11	TTCCCTTT-CACCAAGC
3	XUMMR	MS100303303	IL100305451	MCR.CVU.239	MCR.CVU.239	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-C07	AATTAGAC-GTTCTTAT
3	XUMMR	MS100303194	IL100305246	MCR.CVU.133	MCR.CVU.133	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-F05	AGAACCAG-CCGACTCT
3	XUMMR	MS100303272	IL100305420	MCR.CVU.208	MCR.CVU.208	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-D03	ATGCTCCC-CAGCAGGG
3	XUMMR	MS100303368	IL100303412	401525_neg_ex01	neg_ex01	Genomic PE	neg_ex01	Kapa Evo Plus II	Dedicated	N5UD-C03	GCCCGTCT-GCAACAGC
3	XUMMR	MS100303070	IL100303602	MCR.CVU.10	MCR.CVU.10	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-B02	AGCCTATC-TGCGTAAC
3	XUMMR	MS100303249	IL100305301	MCR.CVU.188	MCR.CVU.188	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-E12	TGAGGACT-CACTAGAC
3	XUMMR	MS100303310	IL100305458	MCR.CVU.246	MCR.CVU.246	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-B08	GTTGAAGG-AGAGCTGC
3	XUMMR	MS100303231	IL100305283	MCR.CVU.170	MCR.CVU.170	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-C10	CTCGAAAT-CAGGTAAG
3	XUMMR	MS100303229	IL100305281	MCR.CVU.168	MCR.CVU.168	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N2UD-A10	AGGTAGGA-TGTTGCGC

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303146	IL100303678	MCR.CVU.88	MCR.CVU.88	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-F11	CAATGCGA-GACGAAC
3	XUMMR	MS100303316	IL100305464	MCR.CVU.252	MCR.CVU.252	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-H08	GCAACTTG-GAGCACTC
3	XUMMR	MS100303265	IL100305413	MCR.CVU.201	MCR.CVU.201	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-E02	GCGGGATA-AGAGACGG
3	XUMMR	MS100303207	IL100305259	MCR.CVU.146	MCR.CVU.146	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-C07	TCAGCGCC-ATGTTCT
3	XUMMR	MS100303126	IL100303658	MCR.CVU.67	MCR.CVU.67	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-B09	CACAGACT-CTGTACCA
3	XUMMR	MS100303077	IL100303609	MCR.CVU.17	MCR.CVU.17	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-A03	TCGGATTG-CGCAACTA
3	XUMMR	MS100303119	IL100303651	MCR.CVU.60	MCR.CVU.60	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-C08	CTTGCTAG-TCATCTCC
3	XUMMR	MS100303251	IL100305303	401523_zp_ex02	zp_ex02	Genomic PE	zp_ex02	Kapa Evo Plus II	Dedicated Dual	N2UD-G12	GAGTCGAT-CCTATGCA
3	XUMMR	MS100303282	IL100305430	MCR.CVU.218	MCR.CVU.218	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-F04	TTGTGCCC-GATGGAAA
3	XUMMR	MS100303104	IL100303636	MCR.CVU.45	MCR.CVU.45	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-D06	CTTACAGC-TGGTGAAG
3	XUMMR	MS100303246	IL100305298	MCR.CVU.185	MCR.CVU.185	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-B12	ACCAACAG-GTTTGCTC
3	XUMMR	MS100303204	IL100305256	MCR.CVU.143	MCR.CVU.143	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N2UD-H06	CAACCGAG-TACCTGTG
3	XUMMR	MS100303184	IL100305236	MCR.CVU.123	MCR.CVU.123	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-D04	AAGGAGAC-CTCACAA
3	XUMMR	MS100303190	IL100305242	MCR.CVU.129	MCR.CVU.129	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-B05	GTTTGCTC-ACCAACAG
3	XUMMR	MS100303290	IL100305438	MCR.CVU.226	MCR.CVU.226	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-F05	GTTAACAT-ACGCCGCA
3	XUMMR	MS100303254	IL100305402	MCR.CVU.190	MCR.CVU.190	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-B01	CAATCATA-TGGAGATT
3	XUMMR	MS100303252	IL100305304	401523_empty	empty	Genomic PE	empty	Kapa Evo Plus II	Dedicated Dual	N2UD-H12	TACCTGTG-CAACCGAG
3	XUMMR	MS100303343	IL100305491	MCR.CVU.279	MCR.CVU.279	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-C12	ACTCTTAG-CGTGGATT
3	XUMMR	MS100303302	IL100305450	MCR.CVU.238	MCR.CVU.238	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-B07	TGGAGATT-CAATCATA
3	XUMMR	MS100303266	IL100305414	MCR.CVU.202	MCR.CVU.202	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-F02	GAAGTGAA-TACAAGTC
3	XUMMR	MS100303110	IL100303642	MCR.CVU.51	MCR.CVU.51	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-B07	GTCTTGA-TCGTCTGA
3	XUMMR	MS100303361	IL100303405	MCR.CVU.293	MCR.CVU.293	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated	N5UD-D02	GTACCTGG-CACCTAAA
3	XUMMR	MS100303163	IL100305215	MCR.CVU.102	MCR.CVU.102	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N2UD-G01	ACGATATG-GACAATTC

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3	XUMMR	MS100303174	IL100305226	MCR.CVU.113	MCR.CVU.113	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-B03	GTCTGAGT-GACTTG TG
3	XUMMR	MS100303063	IL100303595	MCR.CVU.3	MCR.CVU.3	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-C01	TTCCAGGT-CAGTGCTT
3	XUMMR	MS100303301	IL100305449	MCR.CVU.237	MCR.CVU.237	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-A07	AAAGCTAA-TGTCGTAG
3	XUMMR	MS100303157	IL100305209	MCR.CVU.96	MCR.CVU.96	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-A01	CACTGTAG-AAGCGACT
3	XUMMR	MS100303155	IL100303687	401524_zp_ex02	zp_ex02	Genomic PE	zp_ex02	Kapa Evo Plus II	Dedicated Dual	N1UD-G12	GATAGCCA-CAACACAG
3	XUMMR	MS100303335	IL100305483	MCR.CVU.271	MCR.CVU.271	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-C11	GACGCTCG-CGGAGAGG
3	XUMMR	MS100303263	IL100305411	MCR.CVU.199	MCR.CVU.199	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-C02	CATACTTA-GCACGGGA
3	XUMMR	MS100303071	IL100303603	MCR.CVU.11	MCR.CVU.11	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-C02	TCATCTCC-CTTGCTAG
3	XUMMR	MS100303256	IL100305404	MCR.CVU.192	MCR.CVU.192	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-D01	GATGCGAC-ACTTTGGG
3	XUMMR	MS100303261	IL100305409	MCR.CVU.197	MCR.CVU.197	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-A02	GCGCTAAT-ACCCTGAC
3	XUMMR	MS100303209	IL100305261	MCR.CVU.148	MCR.CVU.148	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-E07	CCTTTTCA-TCTTGTTT
3	XUMMR	MS100303135	IL100303667	MCR.CVU.76	MCR.CVU.76	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-C10	AGCTAAGC-GTCGTTAC
3	XUMMR	MS100303307	IL100305455	MCR.CVU.243	MCR.CVU.243	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-G07	GCATGATC-GTGTGGAG
3	XUMMR	MS100303074	IL100303606	MCR.CVU.14	MCR.CVU.14	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-F02	GAACGAAG-GAATCACC
3	XUMMR	MS100303336	IL100305484	MCR.CVU.272	MCR.CVU.272	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-D11	CTCACGTC-TATTTACC
3	XUMMR	MS100303115	IL100303647	MCR.CVU.56	MCR.CVU.56	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N1UD-G07	GCCTTAAC-TAGGAGCT
3	XUMMR	MS100303277	IL100305425	MCR.CVU.213	MCR.CVU.213	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-A04	TTTCTCTA-ACGTCGAG
3	XUMMR	MS100303346	IL100305494	401522_neg_ex01	401522_neg_ex01	Genomic PE	neg_ex01	Kapa Evo Plus II	Dedicated Dual	N3UD-F12	AAGATTGA-CGAACCAC
3	XUMMR	MS100303094	IL100303626	MCR.CVU.35	MCR.CVU.35	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-B05	AGGATAGC-CCAAGGTT
3	XUMMR	MS100303083	IL100303615	MCR.CVU.24	MCR.CVU.24	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-G03	GTCCTAAG-AGCTACCA
3	XUMMR	MS100303081	IL100303613	MCR.CVU.21	MCR.CVU.21	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-E03	GCAATTCC-CTCGAACA
3	XUMMR	MS100303313	IL100305461	MCR.CVU.249	MCR.CVU.249	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-E08	AGAGACGG-GCGGGATA
3	XUMMR	MS100303309	IL100305457	MCR.CVU.245	MCR.CVU.245	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-A08	ACCCTGAC-GCGCTAAT

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303200	IL100305252	MCR.CVU.139	MCR.CVU.139	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-D06	TCTGGAAC-TAGCACCT
3	XUMMR	MS100303185	IL100305237	MCR.CVU.124	MCR.CVU.124	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-E04	AGTCAGGT-GTAACCAC
3	XUMMR	MS100303169	IL100305221	MCR.CVU.108	MCR.CVU.108	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N2UD-E02	GTCCGATC-GCCAATCC
3	XUMMR	MS100303099	IL100303631	MCR.CVU.40	MCR.CVU.40	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-G05	TCCACGTT-GGTTGAAC
3	XUMMR	MS100303359	IL100303403	MCR.CVU.291	MCR.CVU.291	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N5UD-B02	ACAGGAAA-TAGACAAT
3	XUMMR	MS100303212	IL100305264	MCR.CVU.151	MCR.CVU.151	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-H07	ACACGACT-ATCCGCAG
3	XUMMR	MS100303285	IL100305433	MCR.CVU.221	MCR.CVU.221	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-A05	TTTGGGAT-ACGTGAAC
3	XUMMR	MS100303284	IL100305432	MCR.CVU.220	MCR.CVU.220	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-H04	GTATAGTC-GAGACCAA
3	XUMMR	MS100303121	IL100303653	MCR.CVU.62	MCR.CVU.62	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-E08	TATGGCAC-TTGCGAGA
3	XUMMR	MS100303176	IL100305228	MCR.CVU.115	MCR.CVU.115	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-D03	TGTCACAC-TAATCTCG
3	XUMMR	MS100303150	IL100303682	MCR.CVU.92	MCR.CVU.92	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-B12	CACGCAAT-CCTTCCAT
3	XUMMR	MS100303136	IL100303668	MCR.CVU.77	MCR.CVU.77	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-D10	AAGACACC-GTAACCGA
3	XUMMR	MS100303139	IL100303671	MCR.CVU.80	MCR.CVU.80	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-G10	CTTCACTG-CTTAGGAC
3	XUMMR	MS100303358	IL100303402	MCR.CVU.290	MCR.CVU.290	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N5UD-A02	GTGACGTT-ATCTGGGA
3	XUMMR	MS100303253	IL100305401	MCR.CVU.189	MCR.CVU.189	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-A01	TGTCGTAG-AAAGCTAA
3	XUMMR	MS100303351	IL100303395	MCR.CVU.283	MCR.CVU.283	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N5UD-B01	GAACCTCT-GCGCAGAC
3	XUMMR	MS100303279	IL100305427	MCR.CVU.215	MCR.CVU.215	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-C04	TGGGAGTG-TAGACGGG
3	XUMMR	MS100303092	IL100303624	MCR.CVU.33	MCR.CVU.33	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N1UD-H04	ATCTGACC-CTCGACTT
3	XUMMR	MS100303075	IL100303607	MCR.CVU.15	MCR.CVU.15	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-G02	CGAATTGC-GTAAGGTG
3	XUMMR	MS100303215	IL100305267	MCR.CVU.154	MCR.CVU.154	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-C08	GGTGTGAG-AACACCAC
3	XUMMR	MS100303271	IL100305419	MCR.CVU.207	MCR.CVU.207	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-C03	CCCGTTTG-TACAGCAA
3	XUMMR	MS100303289	IL100305437	MCR.CVU.225	MCR.CVU.225	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-E05	TATATGGA-CTGCCAAG
3	XUMMR	MS100303156	IL100303688	401524_Empty	EMPTY	Genomic PE	Empty	Kapa Evo Plus II	Dedicated Dual	N1UD-H12	CCACAACA-GTACCACA

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303225	IL100305277	MCR.CVU.164	MCR.CVU.164	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-E09	TACGGCAG-GATATGAA
3	XUMMR	MS100303181	IL100305233	MCR.CVU.120	MCR.CVU.120	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-A04	ATTACCCA-CACTGTAG
3	XUMMR	MS100303148	IL100303680	MCR.CVU.90	MCR.CVU.90	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-H11	CTTCGGTT-AACCAGAG
3	XUMMR	MS100303281	IL100305429	MCR.CVU.217	MCR.CVU.217	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-E04	CGATATGG-TACGTCCC
3	XUMMR	MS100303347	IL100305495	401522_zp_ex02	401522_zp_ex02	Genomic PE	zp_ex02	Kapa Evo Plus II	Dedicated Dual	N3UD-G12	GTGTGTTT-TATTGTTC
3	XUMMR	MS100303196	IL100305248	MCR.CVU.135	MCR.CVU.135	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-H05	ACCCGTTG-CAGCTTCG
3	XUMMR	MS100303112	IL100303644	MCR.CVU.53	MCR.CVU.53	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-D07	TCCATTGC-TACGTCT
3	XUMMR	MS100303340	IL100305488	MCR.CVU.276	MCR.CVU.276	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-H11	CTAAACAA-CGTAGCGA
3	XUMMR	MS100303120	IL100303652	MCR.CVU.61	MCR.CVU.61	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-D08	AGCGAGAT-CCAGTATC
3	XUMMR	MS100303105	IL100303637	MCR.CVU.46	MCR.CVU.46	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-E06	TACCTGCA-GGACATCA
3	XUMMR	MS100303170	IL100305222	MCR.CVU.109	MCR.CVU.109	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-F02	GAGGACCA-ATGCCGGT
3	XUMMR	MS100303353	IL100303397	MCR.CVU.285	MCR.CVU.285	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N5UD-D01	TGACAGCT-CCTACGGG
3	XUMMR	MS100303355	IL100303399	MCR.CVU.287	MCR.CVU.287	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N5UD-F01	CTGGAGTA-AACCGAAC
3	XUMMR	MS100303211	IL100305263	MCR.CVU.150	MCR.CVU.150	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-G07	GACAATTG-ACGATATG
3	XUMMR	MS100303369	IL100303413	401525_zp_ex02	zp_ex02	Genomic PE	zp_ex02	Kapa Evo Plus II	Dedicated Dual	N5UD-D03	TTCGCCTG-ATCGTCTC
3	XUMMR	MS100303213	IL100305265	MCR.CVU.152	MCR.CVU.152	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-A08	CCTGTCAA-ATGGCTGT
3	XUMMR	MS100303142	IL100303674	MCR.CVU.84	MCR.CVU.84	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-B11	CCAAGGTT-AGGATAGC
3	XUMMR	MS100303219	IL100305271	MCR.CVU.158	MCR.CVU.158	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-G08	CATACCGT-CGCTCTTA
3	XUMMR	MS100303267	IL100305415	MCR.CVU.203	MCR.CVU.203	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-G02	CTGTTTAC-TGAATCTT
3	XUMMR	MS100303201	IL100305253	MCR.CVU.140	MCR.CVU.140	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-E06	CACTAGAC-TGAGGACT
3	XUMMR	MS100303068	IL100303600	MCR.CVU.8	MCR.CVU.8	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-H01	TACTCCAG-GGTATAGG
3	XUMMR	MS100303143	IL100303675	MCR.CVU.85	MCR.CVU.85	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-C11	GAACGGTT-GGAGGAAT
3	XUMMR	MS100303296	IL100305444	MCR.CVU.232	MCR.CVU.232	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-D06	GTAGATGC-GAGCAACA

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303221	IL100305273	MCR.CVU.160	MCR.CVU.160	Genomic PE	<i>Pseudomonas aeruginosa</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-A09	ATGGCTGT-CCTGTCAA
3	XUMMR	MS100303356	IL100303400	MCR.CVU.288	MCR.CVU.288	Genomic PE	<i>Acinetobacter baumannii</i>	Kapa Evo Plus II	Dedicated	N5UD-G01	GATCCGGG-ACCTGCTT
3	XUMMR	MS100303247	IL100305299	MCR.CVU.186	MCR.CVU.186	Genomic PE	<i>Pseudomonas aeruginosa</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-C12	CCTCGGGT-GCCACGAC
3	XUMMR	MS100303297	IL100305445	MCR.CVU.233	MCR.CVU.233	Genomic PE	<i>Staphylococcus aureus</i>	Kapa Evo Plus II	Dedicated Dual	N3UD-E06	TACCGCTC-CAGTGACG
3	XUMMR	MS100303109	IL100303641	MCR.CVU.50	MCR.CVU.50	Genomic PE	<i>Staphylococcus aureus</i>	Kapa Evo Plus II	Dedicated Dual	N1UD-A07	CGAATACG-TTACCGAC
3	XUMMR	MS100303097	IL100303629	MCR.CVU.38	MCR.CVU.38	Genomic PE	<i>Klebsiella pneumoniae</i>	Kapa Evo Plus II	Dedicated Dual	N1UD-E05	CCGCTTAA-GTCATCGT
3	XUMMR	MS100303123	IL100303655	MCR.CVU.64	MCR.CVU.64	Genomic PE	<i>Acinetobacter baumannii</i>	Kapa Evo Plus II	Dedicated Dual	N1UD-G08	GTAAGGTG-CGAATTGC
3	XUMMR	MS100303208	IL100305260	MCR.CVU.147	MCR.CVU.147	Genomic PE	<i>Pseudomonas aeruginosa</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-D07	AGTCACAT-CATTATGG
3	XUMMR	MS100303175	IL100305227	MCR.CVU.114	MCR.CVU.114	Genomic PE	<i>Staphylococcus aureus</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-C03	CCTAAACT-ACGAGGAG
3	XUMMR	MS100303217	IL100305269	MCR.CVU.156	MCR.CVU.156	Genomic PE	<i>Enterobacter cloacae</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-E08	GCCAATCC-GTCCGATC
3	XUMMR	MS100303137	IL100303669	MCR.CVU.78	MCR.CVU.78	Genomic PE	<i>Acinetobacter baumannii</i>	Kapa Evo Plus II	Dedicated Dual	N1UD-E10	CAACTCCA-GAATCCGT
3	XUMMR	MS100303149	IL100303681	MCR.CVU.91	MCR.CVU.91	Genomic PE	<i>Acinetobacter baumannii</i>	Kapa Evo Plus II	Dedicated Dual	N1UD-A12	CGGCATTA-GTCAGTCA
3	XUMMR	MS100303318	IL100305466	MCR.CVU.254	MCR.CVU.254	Genomic PE	<i>Pseudomonas aeruginosa</i>	Kapa Evo Plus II	Dedicated Dual	N3UD-B09	GTATGACG-CCACACTT
3	XUMMR	MS100303197	IL100305249	MCR.CVU.136	MCR.CVU.136	Genomic PE	<i>Pseudomonas aeruginosa</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-A06	ACCGGAGT-AAGGAAGG
3	XUMMR	MS100303214	IL100305266	MCR.CVU.153	MCR.CVU.153	Genomic PE	<i>Pseudomonas aeruginosa</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-B08	CCATCCGC-AAGGCGTA
3	XUMMR	MS100303138	IL100303670	MCR.CVU.79	MCR.CVU.79	Genomic PE	<i>Acinetobacter baumannii</i>	Kapa Evo Plus II	Dedicated Dual	N1UD-F10	GATCTTGC-CATGAGCA
3	XUMMR	MS100303179	IL100305231	MCR.CVU.118	MCR.CVU.118	Genomic PE	<i>Acinetobacter baumannii</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-G03	GTTGGCGT-CAGCAGTC
3	XUMMR	MS100303064	IL100303596	MCR.CVU.4	MCR.CVU.4	Genomic PE	<i>Staphylococcus aureus</i>	Kapa Evo Plus II	Dedicated Dual	N1UD-D01	TACGGTCT-TCCATTGC
3	XUMMR	MS100303278	IL100305426	MCR.CVU.214	MCR.CVU.214	Genomic PE	<i>Acinetobacter baumannii</i>	Kapa Evo Plus II	Dedicated Dual	N3UD-B04	CCAGCGAT-TGACTAGA
3	XUMMR	MS100303193	IL100305245	MCR.CVU.132	MCR.CVU.132	Genomic PE	<i>Staphylococcus aureus</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-E05	TGTGTCAG-CACCTGTA
3	XUMMR	MS100303337	IL100305485	MCR.CVU.273	MCR.CVU.273	Genomic PE	<i>Staphylococcus aureus</i>	Kapa Evo Plus II	Dedicated Dual	N3UD-E11	CTGCCAAG-TATATGGA
3	XUMMR	MS100303180	IL100305232	MCR.CVU.119	MCR.CVU.119	Genomic PE	<i>Acinetobacter baumannii</i>	Kapa Evo Plus II	Dedicated Dual	N2UD-H03	TAGCTGGC-TACCGGCT
3	XUMMR	MS100303114	IL100303646	MCR.CVU.55	MCR.CVU.55	Genomic PE	<i>Pseudomonas aeruginosa</i>	Kapa Evo Plus II	Dedicated Dual	N1UD-F07	ATAACGCC-CAGGTTCA



Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303141	IL100303673	MCR.CVU.82	MCR.CVU.82	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-A11	GTACACCT-TCCTCATG
3	XUMMR	MS100303312	IL100305460	MCR.CVU.248	MCR.CVU.248	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-D08	CGTATAAA-TTGCACCG
3	XUMMR	MS100303223	IL100305275	MCR.CVU.162	MCR.CVU.162	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-C09	ACGAGGAG-CCTAAACT
3	XUMMR	MS100303145	IL100303677	MCR.CVU.87	MCR.CVU.87	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-E11	GTCATCGT-CCGCTTAA
3	XUMMR	MS100303342	IL100305490	MCR.CVU.278	MCR.CVU.278	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-B12	CTCTTGAT-AGAAGTGG
3	XUMMR	MS100303274	IL100305422	MCR.CVU.210	MCR.CVU.210	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-F03	GTAGTTCG-GCATCAAG
3	XUMMR	MS100303365	IL100303409	MCR.CVU.298	MCR.CVU.298	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N5UD-H02	GCCAGATT-GCTGAGAA
3	XUMMR	MS100303073	IL100303605	MCR.CVU.13	MCR.CVU.13	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-E02	TTGCGAGA-TATGGCAC
3	XUMMR	MS100303093	IL100303625	MCR.CVU.34	MCR.CVU.34	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-A05	TCCTCATG-GTACACCT
3	XUMMR	MS100303129	IL100303661	MCR.CVU.70	MCR.CVU.70	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-E09	CTCGAACA-GCAATTCC
3	XUMMR	MS100303255	IL100305403	MCR.CVU.191	MCR.CVU.191	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-C01	GTTCTTAT-AATTAGAC
3	XUMMR	MS100303317	IL100305465	MCR.CVU.253	MCR.CVU.253	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-A09	ACGACGTC-TTGTTGCA
3	XUMMR	MS100303319	IL100305467	MCR.CVU.255	MCR.CVU.255	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-C09	TACAGCAA-CCCCTTTG
3	XUMMR	MS100303168	IL100305220	MCR.CVU.107	MCR.CVU.107	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-D02	ACCTCTTC-CTTGATA
3	XUMMR	MS100303080	IL100303612	MCR.CVU.20	MCR.CVU.20	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-D03	TCTACGCA-CAATAGCC
3	XUMMR	MS100303061	IL100303593	MCR.CVU.1	MCR.CVU.1	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-A01	TTACCGAC-CGAATACG
3	XUMMR	MS100303304	IL100305452	MCR.CVU.240	MCR.CVU.240	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-D07	ACTTTGGG-GATGCGAC
3	XUMMR	MS100303183	IL100305235	MCR.CVU.122	MCR.CVU.122	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-C04	CAGGTAAG-CTCGAAAT
3	XUMMR	MS100303106	IL100303638	MCR.CVU.47	MCR.CVU.47	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-F06	AGACGCTA-GGTGTACA
3	XUMMR	MS100303294	IL100305442	MCR.CVU.230	MCR.CVU.230	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-B06	AGAAGTGG-CTCTTGAT
3	XUMMR	MS100303350	IL100303394	MCR.CVU.282	MCR.CVU.282	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N5UD-A01	AGTTTCGA-ATCGTAGG
3	XUMMR	MS100303264	IL100305412	MCR.CVU.200	MCR.CVU.200	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-D02	TTGCACCG-CGTATAAA
3	XUMMR	MS100303205	IL100305257	MCR.CVU.144	MCR.CVU.144	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-A07	TGTTCCGC-AGGTAGGA

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Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method		Index Type	Index Number	Index Sequence
3	XUMMR	MS100303287	IL100305435	MCR.CVU.223	MCR.CVU.223	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N3UD-C05	CGGAGAGG-GACGCTCG
3	XUMMR	MS100303299	IL100305447	MCR.CVU.235	MCR.CVU.235	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N3UD-G06	TATTGTTC-GTGTGTTT
3	XUMMR	MS100303323	IL100305471	MCR.CVU.259	MCR.CVU.259	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N3UD-G09	CGATACAT-CGAGAACC
3	XUMMR	MS100303305	IL100305453	MCR.CVU.241	MCR.CVU.241	Genomic PE	Klebsiella pneumoniae	Kapa	Evo Plus II	Dedicated Dual	N3UD-E07	CGGACGGA-GAAGAGGG
3	XUMMR	MS100303230	IL100305282	MCR.CVU.169	MCR.CVU.169	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N2UD-B10	TCGCGCAA-ACAAGGCA
3	XUMMR	MS100303232	IL100305284	MCR.CVU.171	MCR.CVU.171	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N2UD-D10	CTCACAACTCAGAGAC
3	XUMMR	MS100303238	IL100305290	MCR.CVU.177	MCR.CVU.177	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N2UD-B11	GCACACAACTTGACGA
3	XUMMR	MS100303098	IL100303630	MCR.CVU.39	MCR.CVU.39	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N1UD-F05	GACGAACCTCAATGCGA
3	XUMMR	MS100303172	IL100305224	MCR.CVU.111	MCR.CVU.111	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N2UD-H02	CTGAGCTCATCAGAGC
3	XUMMR	MS100303320	IL100305468	MCR.CVU.256	MCR.CVU.256	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N3UD-D09	CAGCAGGGATGCTCCC
3	XUMMR	MS100303228	IL100305280	MCR.CVU.167	MCR.CVU.167	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N2UD-H09	TACCGGCTTAGCTGGC
3	XUMMR	MS100303124	IL100303656	MCR.CVU.65	MCR.CVU.65	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N1UD-H08	CGAGAGAA-GGAAGAGA
3	XUMMR	MS100303276	IL100305424	MCR.CVU.212	MCR.CVU.212	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N3UD-H03	GCCATGTAAACCTAT
3	XUMMR	MS100303067	IL100303599	MCR.CVU.7	MCR.CVU.7	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N1UD-G01	TAGGAGCTGCCTTAAC
3	XUMMR	MS100303329	IL100305477	MCR.CVU.265	MCR.CVU.265	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N3UD-E10	TACGTCCCGATATGG
3	XUMMR	MS100303328	IL100305476	MCR.CVU.264	MCR.CVU.264	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N3UD-D10	CTCTTCTACCTCGTA
3	XUMMR	MS100303167	IL100305219	MCR.CVU.106	MCR.CVU.106	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N2UD-C02	AACACCACGGTGTGAG
3	XUMMR	MS100303066	IL100303598	MCR.CVU.6	MCR.CVU.6	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N1UD-F01	CAGGTTCAATAACGCC
3	XUMMR	MS100303127	IL100303659	MCR.CVU.68	MCR.CVU.68	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N1UD-C09	TGGAAGCAGAGGTAC
3	XUMMR	MS100303102	IL100303634	MCR.CVU.43	MCR.CVU.43	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N1UD-B06	CCTTCCATCACGCAAT
3	XUMMR	MS100303182	IL100305234	MCR.CVU.121	MCR.CVU.121	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N2UD-B04	GACTTGTGTGCACGA
3	XUMMR	MS100303338	IL100305486	MCR.CVU.274	MCR.CVU.274	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N3UD-F11	ACGCCGCA-GTTAACAT
3	XUMMR	MS100303239	IL100305291	MCR.CVU.178	MCR.CVU.178	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N2UD-C11	GTCAGTATAGAAGCCT

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303220	IL100305272	MCR.CVU.159	MCR.CVU.159	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-H08	ATCAGAGC-CTGAGCTC
3	XUMMR	MS100303178	IL100305230	MCR.CVU.117	MCR.CVU.117	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-F03	AAGTGTGG-TGCCCATC
3	XUMMR	MS100303082	IL100303614	MCR.CVU.23	MCR.CVU.23	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-F03	CTCAGAAG-GGCAAGTT
3	XUMMR	MS100303242	IL100305294	MCR.CVU.181	MCR.CVU.181	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-F11	CCGACTCT-AGAACCAG
3	XUMMR	MS100303079	IL100303611	MCR.CVU.19	MCR.CVU.19	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-C03	GAGAGTAC-TGGAAGCA
3	XUMMR	MS100303173	IL100305225	MCR.CVU.112	MCR.CVU.112	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-A03	ACGAATCC-ATTACCCA
3	XUMMR	MS100303295	IL100305443	MCR.CVU.231	MCR.CVU.231	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-C06	CGTGGATT-ACTCTTAG
3	XUMMR	MS100303306	IL100305454	MCR.CVU.242	MCR.CVU.242	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-F07	GCAGAGCC-TAGTAATC
3	XUMMR	MS100303360	IL100303404	MCR.CVU.292	MCR.CVU.292	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N5UD-C02	GTGCTCTG-TTCTTCTT
3	XUMMR	MS100303116	IL100303648	MCR.CVU.57	MCR.CVU.57	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-H07	GGTATAGG-TACTCCAG
3	XUMMR	MS100303357	IL100303401	MCR.CVU.289	MCR.CVU.289	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N5UD-H01	AACACCTG-CCCATGCG
3	XUMMR	MS100303100	IL100303632	MCR.CVU.41	MCR.CVU.41	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-H05	AACCAGAG-CTTCGGTT
3	XUMMR	MS100303286	IL100305434	MCR.CVU.222	MCR.CVU.222	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-B05	CACCAAGC-TTCCCTTT
3	XUMMR	MS100303250	IL100305302	401523_neg_ex01	neg_ex01	Genomic PE	neg_ex01	Kapa Evo Plus II	Dedicated Dual	N2UD-F12	TTCCCGAA-TTGCGTTA
3	XUMMR	MS100303300	IL100305448	MCR.CVU.236	MCR.CVU.236	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-H06	GTTGTGTG-CGTCCGAC
3	XUMMR	MS100303344	IL100305492	MCR.CVU.280	MCR.CVU.280	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-D12	GAGCAACA-GTAGATGC
3	XUMMR	MS100303243	IL100305295	MCR.CVU.182	MCR.CVU.182	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-G11	TTGCTGGA-ATTGGACA
3	XUMMR	MS100303132	IL100303664	MCR.CVU.73	MCR.CVU.73	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-H09	CAGCATAC-GCGTTAGA
3	XUMMR	MS100303332	IL100305480	MCR.CVU.268	MCR.CVU.268	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-H10	GAGACCAA-GTATAGTC
3	XUMMR	MS100303370	IL100303414	401525_EMPTY	EMPTY	Genomic PE	EMPTY	Kapa Evo Plus II	Dedicated Dual	N5UD-E03	CTAGCTCC-CCTTGTGA
3	XUMMR	MS100303224	IL100305276	MCR.CVU.163	MCR.CVU.163	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-D09	TAATCTCG-TGTACAC
3	XUMMR	MS100303188	IL100305240	MCR.CVU.127	MCR.CVU.127	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-H04	TTGGGTAC-CTTCCAAC
3	XUMMR	MS100303245	IL100305297	MCR.CVU.184	MCR.CVU.184	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-A12	TCCCACGA-TTCAATAG

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303324	IL100305472	MCR.CVU.260	MCR.CVU.260	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-H09	AACCCCTAT-GCCATGTA
3	XUMMR	MS100303152	IL100303684	MCR.CVU.94	MCR.CVU.94	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-D12	TGGTGAAG-CTTACAGC
3	XUMMR	MS100303260	IL100305408	MCR.CVU.196	MCR.CVU.196	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-H01	ACGTTGTA-TCGACCTA
3	XUMMR	MS100303235	IL100305287	MCR.CVU.174	MCR.CVU.174	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-G10	CGCTAATC-TATGACGT
3	XUMMR	MS100303085	IL100303617	MCR.CVU.26	MCR.CVU.26	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-A04	CAAGGTAC-CGTATCTC
3	XUMMR	MS100303087	IL100303619	MCR.CVU.28	MCR.CVU.28	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-C04	GTCGTTAC-AGCTAAGC
3	XUMMR	MS100303147	IL100303679	MCR.CVU.89	MCR.CVU.89	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-G11	GGTTGAAC-TCCACGTT
3	XUMMR	MS100303076	IL100303608	MCR.CVU.16	MCR.CVU.16	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-H02	GGAAGAGA-CGAGAGAA
3	XUMMR	MS100303339	IL100305487	MCR.CVU.275	MCR.CVU.275	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-G11	CGCCAGTC-CGTCTTGG
3	XUMMR	MS100303158	IL100305210	MCR.CVU.97	MCR.CVU.97	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-B01	GTGCACGA-TGATAGGC
3	XUMMR	MS100303218	IL100305270	MCR.CVU.157	MCR.CVU.157	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-F08	ATGCCGGT-GAGGACCA
3	XUMMR	MS100303159	IL100305211	MCR.CVU.98	MCR.CVU.98	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-C01	ATGTTCTC-TCAGCGCC
3	XUMMR	MS100303191	IL100305243	MCR.CVU.130	MCR.CVU.130	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-C05	AGAAGCCT-GTCAGTAT
3	XUMMR	MS100303233	IL100305285	MCR.CVU.172	MCR.CVU.172	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-E10	GTAACCAC-AGTCAGGT
3	XUMMR	MS100303103	IL100303635	MCR.CVU.44	MCR.CVU.44	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-C06	AGGAACAC-GGAATGTC
3	XUMMR	MS100303354	IL100303398	MCR.CVU.286	MCR.CVU.286	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated	N5UD-E01	CATCACCC-TGCTATAT
3	XUMMR	MS100303321	IL100305469	MCR.CVU.257	MCR.CVU.257	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-E09	GATAAATG-GCTCAATA
3	XUMMR	MS100303117	IL100303649	MCR.CVU.58	MCR.CVU.58	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-A08	TCTAGGAG-AGTGACCT
3	XUMMR	MS100303203	IL100305255	MCR.CVU.142	MCR.CVU.142	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N2UD-G06	CCTATGCA-GAGTCGAT
3	XUMMR	MS100303322	IL100305470	MCR.CVU.258	MCR.CVU.258	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-F09	GCATCAAG-GTAGTTCG
3	XUMMR	MS100303065	IL100303597	MCR.CVU.5	MCR.CVU.5	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-E01	AAGACCGT-GTCGATTG
3	XUMMR	MS100303108	IL100303640	MCR.CVU.49	MCR.CVU.49	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-H06	GTACCACA-CCACAACA
3	XUMMR	MS100303131	IL100303663	MCR.CVU.72	MCR.CVU.72	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-G09	AGCTACCA-GTCCTAAG

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method	Index Type	Index Number	Index Sequence
3	XUMMR	MS100303236	IL100305288	MCR.CVU.175	MCR.CVU.175	Genomic PE	Klebsiella pneumoniae	Kapa Evo Plus II	Dedicated Dual	N2UD-H10	CTTCCAAC-TTGGGTAC
3	XUMMR	MS100303111	IL100303643	MCR.CVU.52	MCR.CVU.52	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-C07	CAGTGTCTT-TTCCAGGT
3	XUMMR	MS100303160	IL100305212	MCR.CVU.99	MCR.CVU.99	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-D01	CATTATGG-AGTCACAT
3	XUMMR	MS100303101	IL100303633	MCR.CVU.42	MCR.CVU.42	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-A06	GTCAGTCA-CGGCATT
3	XUMMR	MS100303072	IL100303604	MCR.CVU.12	MCR.CVU.12	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-D02	CCAGTATC-AGCGAGAT
3	XUMMR	MS100303293	IL100305441	MCR.CVU.229	MCR.CVU.229	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-A06	TAGTCACA-TATACCTC
3	XUMMR	MS100303348	IL100305496	401522_empty	401522_empty	Genomic PE	empty	Kapa Evo Plus II	Dedicated Dual	N3UD-H12	CGTCCGAC-GTTGTGTG
3	XUMMR	MS100303226	IL100305278	MCR.CVU.165	MCR.CVU.165	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N2UD-F09	TGCCCCATC-AAGTGTGG
3	XUMMR	MS100303189	IL100305241	MCR.CVU.128	MCR.CVU.128	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N2UD-A05	TTCAATAG-TCCCCAGA
3	XUMMR	MS100303133	IL100303665	MCR.CVU.74	MCR.CVU.74	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-A10	CGTATCTC-CAAGGTAC
3	XUMMR	MS100303333	IL100305481	MCR.CVU.269	MCR.CVU.269	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N3UD-A11	ACGTGAAC-TTTGGGAT
3	XUMMR	MS100303308	IL100305456	MCR.CVU.244	MCR.CVU.244	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-H07	TCGACCTA-ACGTTGTA
3	XUMMR	MS100303089	IL100303621	MCR.CVU.30	MCR.CVU.30	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N1UD-E04	GAATCCGT-CAACTCCA
3	XUMMR	MS100303216	IL100305268	MCR.CVU.155	MCR.CVU.155	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N2UD-D08	CTTGACATA-ACCTCTTC
3	XUMMR	MS100303273	IL100305421	MCR.CVU.209	MCR.CVU.209	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N3UD-E03	GCTCAATA-GATAAATG
3	XUMMR	MS100303280	IL100305428	MCR.CVU.216	MCR.CVU.216	Genomic PE	Acinetobacter baumannii	Kapa Evo Plus II	Dedicated Dual	N3UD-D04	CCCTCGTA-CTCTTCTA
3	XUMMR	MS100303134	IL100303666	MCR.CVU.75	MCR.CVU.75	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-B10	TTACGTGC-AGACCTTG
3	XUMMR	MS100303096	IL100303628	MCR.CVU.37	MCR.CVU.37	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-D05	GACGTCAT-CCAGTTGA
3	XUMMR	MS100303107	IL100303639	MCR.CVU.48	MCR.CVU.48	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-G06	CAACACAG-GATAGCCA
3	XUMMR	MS100303069	IL100303601	MCR.CVU.9	MCR.CVU.9	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-A02	AGTGACCT-TCTAGGAG
3	XUMMR	MS100303086	IL100303618	MCR.CVU.27	MCR.CVU.27	Genomic PE	Staphylococcus aureus	Kapa Evo Plus II	Dedicated Dual	N1UD-B04	AGACCTTG-TTACGTGC
3	XUMMR	MS100303154	IL100303686	401524_NEG-ex01	neg_ex01	Genomic PE	Neg	Kapa Evo Plus II	Dedicated Dual	N1UD-F12	GGTGTAAC-AGACGCTA
3	XUMMR	MS100303078	IL100303610	MCR.CVU.18	MCR.CVU.18	Genomic PE	Pseudomonas aeruginosa	Kapa Evo Plus II	Dedicated Dual	N1UD-B03	CTGTACCA-CACAGACT

Table 1: Sample summary table for samples from project XUMMR

Lane	Project	Master Sample ID	Library ID	Sample Name	External ID	Type	Species	Method		Index Type	Index Number	Index Sequence
3	XUMMR	MS100303234	IL100305286	MCR.CVU.173	MCR.CVU.173	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N2UD-F10	CATATCCA-ACCGTAAG
3	XUMMR	MS100303062	IL100303594	MCR.CVU.2	MCR.CVU.2	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N1UD-B01	TCGTCTGA-GTCCTTGA
3	XUMMR	MS100303177	IL100305229	MCR.CVU.116	MCR.CVU.116	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N2UD-E03	GATATGAA-TACGGCAG
3	XUMMR	MS100303248	IL100305300	MCR.CVU.187	MCR.CVU.187	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N2UD-D12	TAGCACCT-TCTGGAAC
3	XUMMR	MS100303144	IL100303676	MCR.CVU.86	MCR.CVU.86	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N1UD-D11	CCAGTTGA-GACGTCAT
3	XUMMR	MS100303283	IL100305431	MCR.CVU.219	MCR.CVU.219	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N3UD-G04	TGTCCTCT-GTTCTCG
3	XUMMR	MS100303162	IL100305214	MCR.CVU.101	MCR.CVU.101	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N2UD-F01	GGCTTACT-CTTTCCT
3	XUMMR	MS100303311	IL100305459	MCR.CVU.247	MCR.CVU.247	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N3UD-C08	GCACGGGA-CATACTTA
3	XUMMR	MS100303275	IL100305423	MCR.CVU.211	MCR.CVU.211	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N3UD-G03	CGAGAACC-CGATACAT
3	XUMMR	MS100303161	IL100305213	MCR.CVU.100	MCR.CVU.100	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N2UD-E01	TCTTGTTT-CCTTTCAC
3	XUMMR	MS100303326	IL100305474	MCR.CVU.262	MCR.CVU.262	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N3UD-B10	TGACTAGA-CCAGCGAT
3	XUMMR	MS100303367	IL100303411	MCR.CVU.299	MCR.CVU.299	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated	N5UD-B03	GAGATGAT-TCACGCCT
3	XUMMR	MS100303091	IL100303623	MCR.CVU.32	MCR.CVU.32	Genomic PE	Acinetobacter baumannii	Kapa	Evo Plus II	Dedicated Dual	N1UD-G04	CTTAGGAC-CTTCACTG
3	XUMMR	MS100303259	IL100305407	MCR.CVU.195	MCR.CVU.195	Genomic PE	Pseudomonas aeruginosa	Kapa	Evo Plus II	Dedicated Dual	N3UD-G01	GTGTGGAG-GCATGATC
3	XUMMR	MS100303298	IL100305446	MCR.CVU.234	MCR.CVU.234	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N3UD-F06	CGAACCAC-AAGATTGA
3	XUMMR	MS100303327	IL100305475	MCR.CVU.263	MCR.CVU.263	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N3UD-C10	TAGACGGG-TGGGAGTG
3	XUMMR	MS100303151	IL100303683	MCR.CVU.93	MCR.CVU.93	Genomic PE	Staphylococcus aureus	Kapa	Evo Plus II	Dedicated Dual	N1UD-C12	GGAATGTC-AGGAACAC