Supplementary Table 1: Characteristics for Blastn analysis and identity of isolates sequenced

S/N	Sample No /Type	Query Length	Sequence Nucleotide obtained (FASTA format)	Similarity/ E-score	Gene Bank Accession Number/ Identical with	Identity Of Isolate Obtained
1	7	716	>7_907R_G03_21 GGGGGGGGCGGACTTAACGCGTTAGCTCCGGAAGCCA CGCTCAAGGGGCACAACCTCAAGTCGACATCGTTTACG GCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCC ACGCTTTCGCACTGAGCGTCAGTCTTCGTCCAGGGGG CCGCCTTCGCCACCGGTATTCCTCCAACTCTACGCATT TCACCGCTACACCTGGAATTCTACCCCCCTCTACGAGA CTCAGCTTGCCAGTATCAGATGCAGTTCCCAGGTTGAG CCCGGGGATTTCACATCTGACTTACAAACCGCCTGCGT GCGCTTTACGCCCAGTAATTCCGATTAACGCTTGCACC CTCCGTATTACCGCGGCTGCTGGCACGGAGTTAGCCG GTGCTTCTTCTGCGGGTAACGTCAATGACAAAGGTATT AACTTTACTCCCTTCCTCCCCGTGAAAGTACTTTACAAC CCAAGGCCTTCTTCATACCGCGGATGGTGCATCAGCTT GCGCCCATTGTGCAATATTCCCCACTGCTGCCTCCCGA GGAGTCTGGACCGGTCTCAGTCCGTGTGGTGGTCATC CTCTCAGACCAGTAGGGATCGCGCCTAGGTGAGGCTTA CCCCACCTACTAGCTAATCCCATCTGGGACATCCGATG GCAAAGGCCGAAGACCCCTCTTTGGTCTTGCGACGTTA TGCG	95.73/0	MH266079.1 .	***Escherichia coli***
2		826	>10_907R_A04_01 TGGGGGGGTGGATTTAACGCGTTAGCTCCGGAAGCCA CGCCTCAAGGGCACAACCTCCAAATCGACATCGTTTAC GCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCC ACGCTTTCGCACCTGAGCGTCAGTCTTTGTCCAGGGGG CCGCCTTCGCCACCGGTATTCCTCCAATCTCTACGCAT TTCACCGCTACACCTGGAATTCTACCCCCCTCTACAAG ACTCTAGCGCCAGTTTCAAATGCATTCCCAGGTTAGCC CGGGGATTTCACATCGACTTAAACCGCCTGCGTGCGCT TTACGCCCAGTAATTCCGATTAACGCTTGCACCCTCCG TATTACCGCGGCTGCTGGCACGGAGTTAGCCGGTGCTT CTTCTGCGGGTAACGTCAATGAAGGGTATTAACCTACG CCTTCCTCCCCCTGAAAGTCTTTACAACCCAAGGCCTT CTTCAACACGCGGCATGGCTGCATCAGGCTTGCGCCC ATTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAGT CTGGCCGTGTCTCAGTTCCAGTGTGGTGTCATCCTCTC AGACCAGCTAGAGATCGTCGCCTAGGTGAGCCTTACCC ACCTACTAGCTAATCCCATTGGGCATCTGATGGCTGAG GCCCGAAGGTCCCCCTTTGGTTGGACTTATGCGGTATT AGCTACCGTTTCCAGTAGTTATCCCCCTCATCGGCGTT CCCAGACATTACTCACCCGTCCGCCGCTCGTCACCAGA AGCAAGCTCTCTCCGCCCGAA	94.51	MT912763.1 OR MT990398.1	***Klebsiella quasivariicola*** Klebsiella pneumoniae
3		719	>11_907R_B04_04 TTCCCAGGGCGGGTCGATTAACGCGTTAGCTCCGGAA GCCACCCTCAGGGCAACAACCTCCAATCGACATCGTTT ACAGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTC CCCACGCTTTCGCACCTGAGCGTCAGTCTTTGTCCAGG GGGCCGCCTTCGCCACCGGTATTCCTCCACATCTCTAC GCATTTCACCGCTACACATGGAATTCTACCCCCCTCTAC AAGACTCTAGCTACCAGTMTAGATGCATTCCCGGTTAA GCCCGGGGATTTCACATCTACTAATAACCGCCTGCGTG CGCTTTACGCCCAGTAATTCCGATTAACGCTTGCACCC TCCGTATTACCGCGGCTGCTGGCACGGAGTTAGCCGG TGCTTCTTCTGTCGGTAACGTCAATGAGTATTAACTAAC CCTTCCTCCCGACTGAAAGTACTTTACAACCCAGGCCT TCTTCATACACGCGGCATGGGCTGCTCAGGCTTGCGCC CATTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGAG TCTGGGGCCGTGTCTCAGTCCCAGTGTGGCTGATCATC CTCTCAGACCAGCTAGGGGATCGTCGCCTAGGTAGCCT TACCTTACCTACTAGCTAATCCCATATGGGTCATCTGAG GCGCAAGGCCCGAAGGTCCCCCTTTGGTCCAAGACAT ATGG	94.41/0	CP064826.1	***Morganella morganii subsp. morganii***
4		867	>12_907R_C04_07 TCCAGGGGCGGGGTCGGATTTACGCGTTAGCTCCGAA GCCACCTCAGCAACCTCAAATCGACATCGTTTACAGCG TGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACG CTTTCGCACCTGAGCGTCAGTCTTTGTCCAGGGGGCCG CCTTCGCCACCGGTATTCCTCCACATCTCTACGCATTTC ACCGCTACACATGGAATTCTACCCCCCTCTACAGACTC TAGCACCAGTTAGATGCCATTCGCTAGTTAAGCCGGGG ATTTCACATCTAACTTTTACCGCCTGCGTGCGCTTTACG CCCAGTAATTCCGATTAACGCTTGCACCCTCCGTATTAC CGCGGCTGCTGGCACGGAGTTAGCCGGTGCTTCTTCT GGGTAACGTCAATGTGATGTATTAACTACGCCTTCCTCC CCTGAAAGTACTTTACAACCCTAGGCCTTCTTCATACAG CGGCATGGCTGCATCAGGCTTGCGCCCATTGTGCAATA TTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGT CTCAGTCCCGTGTGGCTGATCATCCTCTCAGACCAGTA GGATCGCGCCTAGGTGAGCCATTACCTCACCTATCTAA TCCCATATGGGTGTCCGAAGCGCAGGCCCGAAGACCC	90.92/0	MH532483.1	***Providencia stuartii***

			TGCTTTGTCCTAAGAATATGCGGATTAGCCACCGTTTCC AGTATATCCCCTTATTCGGGCAGATCCCATACATACTCA CGTCGCGGTCGTCAGGAGAGCAGCTTCCTGTTACCCTC ATAGTAAGTCGCCCCGTCATGTAAAACTCTAGG			
5		883	>13_907R_E03_15 AGGGGGGGTGATTTACGCGTTAGCTCCGGAAGCCACG CCTCAAGGGCACAACCTCCAAATCGACATCGTTTACGG CGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCA CGCTTTCGCACCTGAGCGTCAGTCTTTGTCCAGGGGGC CGCCTTCGCCACCGGTATTCCTCCAGATCTCTACGCAT TTCACCGCTACACCTGGAATTCTACCCCCCCTCTACAAG ACTCTAGCCTGCCAGTTTCGAATGCAGTTCCCAGGTTG AGCCCGGGGATTTCACATCCGACTTGACAGACCGCCTG CGTGCGCTTTACGCCCAGTAATTCCGATTAACGCTTGC ACCCTCCGTATTACCGCGGCTGCTGGCACGGAGTTAG CCGGTGCTTCTTCTGCGGGTAACGTCAATCGMAGGTTA TTAACCTACGCCTTCCTCCCCGCTGAAAGTGCTTTACAA CCCGAAGGCCTTCTTCACACACGCGGCATGGCTGCATC AGGCTTGCGCCCATTGTGCAATATTCCCCACTGCTGCC TCCCGTAGGAGTCTGGACCGTGTCTCAGTTCCAGTGTG GCTGGTCATCCTCTCAGACCAGCTAGGGATCGTCGCCT AGGTGAGCCGTTACCCCACCTACTAGCTAATCCCATCT GGGCACATCTGATGGCATGAGGCCCGAAGGTCCCCCA CTTTGGTCTTGCGACTTATGCGGTATTAGCTACCGTTTC CAGTAGTTATCCCCCTCCATCAGGCAGTTTCCCAGACA TTACTCACCCGTCCGCCGCTCGTCACCCGAGAGCAAG CTCTCTGTGCTACCGCTCGACTTTGCATGTGTTAGGCC TGCCGCCAGCGTTCAATCTGAGCCATGATCAAACTCT	98.85	MT990399.1	***Klebsiella pneumoniae***
6	15	479	>15_907R_D04_10 ATCAGGGCGGGGTCGATTTACGCGTTAGCTCCAGAAG CCACGGTTCAAGACCACAACCTCTAAATCGACATCGTTT ACAGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTC CCCACGCTTTCGCACCTGAGCGTCAGTCTTTGTCCAGG GGGCCGCCTTCGCCACCGGTATTCCTCCACATCTCTAC GCATTTCACCGCTACACATGGAATTCTACCCCCCTCTCA AGACTCTAGCCAACCAGTTTCAGATGCAATTCCCAAGTT AAGCTCGGGGCTTTCACATCTGACTTAATTGACCGCCT GCGTGCGCTTTACGCCCAGTAATTCCGATTAACGCTTG CACCCTCCGTATTACCGCGGCTGCTGGACGGAGTTAG CCGGTGCTTCTTCTGGGGGGAACGTCAATTGGTAAGG GATGGGCTTATCACCTTCCTCCCGGTAAAAGACTTTGG AGCCTAAGGG	96.10	MG663472.1	***Salmonella enterica subsp. enterica***