

Supplementary Table 2. Core genome single nucleotide polymorphisms (SNPs) variation between *K. pneumoniae* isolates from the same patients (T0 versus latest follow up times) and between each other

ST101

	RM2.10. T0	RM2.12. T0	RM2.12 .T4	RM2.31 .T0	RM2.31 .T4	RM2.55 .T0	RM2.55 .T8	RM2.56 .T0	RM2.56 .T4	RM2.69 .T0	RM2.69 .T4	RM2.71 .T0	RM2.72 .T0	RM2.72.T8
RM2.10.T0	0	10	10	7	6	7	9	14	18	11	10	11	12	16
RM2.12.T0	10	0	2	11	10	11	13	18	22	15	14	19	4	8
RM2.12.T4	10	2	0	11	10	11	13	18	22	13	14	19	4	8
RM2.31.T0	7	11	11	0	5	6	8	15	19	12	11	16	13	17
RM2.31.T4	6	10	10	5	0	5	7	14	18	11	10	13	12	12
RM2.55.T0	7	11	11	6	5	0	8	15	19	12	11	16	13	17
RM2.55.T8	9	13	13	8	7	8	0	17	21	14	13	18	15	19
RM2.56.T0	14	18	18	15	14	15	17	0	4	17	18	23	18	24
RM2.56.T4	18	22	22	19	18	19	21	4	0	21	22	27	22	28
RM2.69.T0	11	15	13	12	11	12	14	17	21	0	5	20	15	21
RM2.69.T4	10	14	14	11	10	11	13	18	22	5	0	19	16	20
RM2.71.T0	11	19	19	16	13	16	18	23	27	20	19	0	21	23
RM2.72.T0	12	4	4	13	12	13	15	18	22	15	16	21	0	8
RM2.72.T8	16	8	8	17	12	17	19	24	28	21	20	23	8	0

ST512

	RM2.05. T0	RM2.15. T0	RM2.15 .T4	RM2.19 .T0	RM2.19 .T4	RM2.20 .T0	RM2.20 .T4	RM2.21 .T0	RM2.21 .T4	RM2.21 .T8	RM2.21 .T0	RM2.67 .T4
RM2.05.T0	0	24	79	73	76	84	83	81	80	80	88	79
RM2.15.T0	24	0	75	69	72	80	79	77	76	76	84	75
RM2.15.T4	79	75	0	48	43	23	6	18	19	19	73	58
RM2.19.T0	73	69	48	0	9	53	52	50	49	49	67	58
RM2.19.T4	76	72	43	9	0	48	47	45	44	44	70	55
RM2.20.T0	84	80	23	53	48	0	27	25	26	26	78	63
RM2.20.T4	83	79	6	52	47	27	0	22	23	23	77	62
RM2.21.T0	81	77	18	50	45	25	22	0	4	7	75	60
RM2.21.T4	80	76	19	49	44	26	23	4	0	3	72	59
RM2.21.T8	80	76	19	49	44	26	23	7	3	0	72	59
RM2.67.T0	88	84	73	67	70	78	77	75	72	72	0	59
RM2.67.T4	79	75	58	58	55	63	62	60	59	59	59	0