

Table S3
Major tandem repeat tracts

replicon	location	unit repeat length (bp)	B31	N40	JD1	297
chrM	<i>bb0210 (Imp1)</i>	162	7 ^a	2	6	nd ^c
chrM	<i>bb0546</i>	60	5	3	5	nd
chrM	<i>bb0801</i>	33	12	10	12	nd
lp17	13154 in B31	21	8	20	9	15
lp21	3618 in B31	63	176	– ^b	126 ^d	–
lp28-4	<i>B31_bbi16/vraA</i>	27	22	31	24	35
lp28-5	12056 in JD1	133	–	16	26	6
lp38	5938 in B31	17	7	6	–	–
lp38	10287 in B31	7	12	119	–	–
lp54	11362 in B31	8	2	7	3	10
lp54	20128 in B31	11	7	5	3	3
lp54	27258 in B31	10	4	3	3	4

Table S3 footnotes:

- a. values rounded to nearest whole integer
- b. "–", orthologous sequence not present in this strain
- c. nd, not sequenced
- d. in variable part of chromosome