

Table S1 Genome related information of strains RD2P54^T, M1R5S18^T, and M1R5S59^T

No.	Strains	Total reads after filtering (Mb)	Average length of reads (bp)	Total base after filtering (Gb)	Completeness (%)	Contamination (%)	Assembly size (bp)	Contigs	G+C (%)	CDS
1	RD2P54 ^T	9.64	149	1.45	99.59	1.38	4,193,932	46	71.3	3,513
2	M1R5S18 ^T	14.94	149	2.24	99.95	1.16	3,898,385	29	71.3	3,315
3	M1R5S59 ^T	12.30	149	1.84	99.95	1.03	3,959,197	20	70.6	3,356