Table S3. Assembly statistics for the 20 IATS serotype isolates.

The assemblies were performed using the A5 assembly pipeline.

	Platform	Scaffolds	Genome Size	Longest Scaffold	N50	% reads passing EC	Median cov
O1	HiSeq	145	6703016	422485	111875	98.37	99
O2	HiSeq	133	6750246	356749	96874	98.51	128
O3	HiSeq	140	6357593	420867	91877	98.51	139
O4	HiSeq	236	6568643	455691	96234	98.40	134
O5	HiSeq	194	6214018	264927	76249	98.41	144
O6	MiSeq	78	6659911	626676	209526	98.00	127
O7	HiSeq	224	6765780	312849	172697	98.44	124
O8	HiSeq	112	6647277	338467	148018	98.41	131
O9	HiSeq	111	6378192	436537	153192	98.43	134
O10	HiSeq	180	6426301	320194	84223	98.49	137
O11	HiSeq	207	6658142	334986	86674	98.47	132
O12	HiSeq	217	6448608	296739	75294	98.46	136
O13	HiSeq	258	6524980	239302	63465	98.42	136
O14	HiSeq	261	6527572	239303	66103	98.42	136
O15	HiSeq	130	6482228	459823	143401	98.43	132
O16	HiSeq	129	6813102	423952	178313	98.48	128
O17	MiSeq	99	6993770	403929	158754	98.02	148
O18	HiSeq	123	6603159	350348	145684	98.55	133
O19	HiSeq	197	6700599	270540	78080	98.51	132
O20	HiSeq	167	6534632	269982	96164	98.51	133