

Reference sequence (1): AP006557.1_SARS_coronavirus_TWH_genomic_RNA_complete_genome
Identities normalised by aligned length.
Colored by: identity

Table with columns: ID, Name, cov, pid, and sequence alignment. The table lists 114 SARS coronavirus genomic RNA complete genomes with their coverage and identity percentages, followed by a detailed sequence alignment with a reference sequence.

Table with columns: ID, Name, cov, pid, and sequence alignment. This table continues the list of SARS coronavirus genomic RNA complete genomes, showing IDs 1 through 114 and their corresponding coverage and identity data.

Accession	Sequence ID	From	Patent	Genome Type	Consensus	Cov	Pid	321	4	400
111	CQ918584.1	Sequence_1	From Patent	NC_0200409684.2	100.0%	95.0%	95.0%	GTGTTCAGTGCATCATCAGCATACTAGGTTTCTGTC	CGGGTGTGACCGAAGGTAAGTGGAGAGCC	TGTTCTTGTGTT
112	CQ918585.1	Sequence_2	From Patent	NC_0200409684.2	100.0%	95.0%	95.0%	GTGTTCAGTGCATCATCAGCATACTAGGTTTCTGTC	CGGGTGTGACCGAAGGTAAGTGGAGAGCC	TGTTCTTGTGTT
113	CQ918598.1	Sequence_15	From Patent	NC_0200409684.2	100.0%	100.0%	100.0%	GTGTTCAGTGCATCATCAGCATACTAGGTTTCTGTC	CGGGTGTGACCGAAGGTAAGTGGAGAGCC	TGTTCTTGTGTT
114	NC_0204718.3	SARS_coronavirus_complete_genome			consensus/100%	100.0%	100.0%	GTGTTCAGTGCATCATCAGCATACTAGGTTTCTGTC	CGGGTGTGACCGAAGGTAAGTGGAGAGCC	TGTTCTTGTGTT
					consensus/90%		GACCGAAGGTAAGT.....		
					consensus/80%			GTGTTCAGTGCATCATCAGCATACTAGGTTTCTGTC	CGGGTGTGACCGAAGGTAAGTGGAGAGCC	TGTTCTTGTGTT
					consensus/70%			GTGTTCAGTGCATCATCAGCATACTAGGTTTCTGTC	CGGGTGTGACCGAAGGTAAGTGGAGAGCC	TGTTCTTGTGTT
1	AP006557.1	SARS_coronavirus_TWH_genomic_RNA_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
2	AP006558.1	SARS_coronavirus_TWJ_genomic_RNA_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
3	AP006559.1	SARS_coronavirus_TWK_genomic_RNA_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
4	AP006560.1	SARS_coronavirus_TWS_genomic_RNA_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
5	AP006561.1	SARS_coronavirus_TWY_genomic_RNA_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
6	AY274119.3	Severe_acute_respiratory_syndrome-related_coronavirus_isolate_Tor2_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
7	AY278487.3	SARS_coronavirus_BJ02_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
8	AY278488.2	SARS_coronavirus_BJ01_complete_genome			93.7%	93.7%		TCAACGAGAAAAACA		
9	AY278489.2	SARS_coronavirus_BJ03_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
10	AY278490.3	SARS_coronavirus_BJ03_complete_genome			98.7%	98.7%		TCAACGAGAAAAACA		
11	AY278491.2	SARS_coronavirus_HKU-39849_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
12	AY278554.2	SARS_coronavirus_CUHK-WI_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
13	AY278741.1	SARS_coronavirus_Urbani_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
14	AY279354.2	SARS_coronavirus_BJ04_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
15	AY282752.2	SARS_coronavirus_CUHK-Su10_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
16	AY283794.1	SARS_coronavirus_Sin2500_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
17	AY283795.1	SARS_coronavirus_Sin2677_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
18	AY283796.1	SARS_coronavirus_Sin2679_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
19	AY283797.1	SARS_coronavirus_Sin2748_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
20	AY283798.2	SARS_coronavirus_Sin2774_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
21	AY291315.1	SARS_coronavirus_Frankfurt_1_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
22	AY291451.1	SARS_coronavirus_TW1_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
23	AY297028.1	SARS_coronavirus_ZJ01_complete_genome			95.3%	95.3%		TCAACGAGAAAAACA		
24	AY304486.1	SARS_coronavirus_S23_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
25	AY304488.1	SARS_coronavirus_S216_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
26	AY304495.1	SARS_coronavirus_G250_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
27	AY310120.1	SARS_coronavirus_FRA_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
28	AY313906.1	SARS_coronavirus_GD69_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
29	AY321118.1	SARS_coronavirus_TW_C_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
30	AY323977.2	SARS_coronavirus_HSR_1_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
31	AY338174.1	SARS_coronavirus_Taiwan_TC1_complete_genome			77.0%	77.0%		TCAACGAGAAAAACA		
32	AY338175.1	SARS_coronavirus_Taiwan_TC2_complete_genome			77.0%	77.0%		TCAACGAGAAAAACA		
33	AY345986.1	SARS_coronavirus_CUHK-AG01_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
34	AY345987.1	SARS_coronavirus_CUHK-AG02_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
35	AY345988.1	SARS_coronavirus_CUHK-AG03_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
36	AY348314.1	SARS_coronavirus_Taiwan_TC3_complete_genome			77.0%	77.0%		TCAACGAGAAAAACA		
37	AY350750.1	SARS_coronavirus_PUM01_complete_genome			95.7%	95.7%		TCAACGAGAAAAACA		
38	AY351680.1	SARS_coronavirus_ZW1_complete_genome			100.0%	99.7%		TCAACGAGAAAAACA		
39	AY357075.1	SARS_coronavirus_PUM02_complete_genome			95.3%	95.3%		TCAACGAGAAAAACA		
40	AY357076.1	SARS_coronavirus_PUM03_complete_genome			95.3%	95.3%		TCAACGAGAAAAACA		
41	AY362698.1	SARS_coronavirus_TW_C2_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
42	AY362699.1	SARS_coronavirus_TW_C3_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
43	AY390556.1	SARS_coronavirus_G202_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
44	AY394850.1	SARS_coronavirus_WH1_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
45	AY394977.1	SARS_coronavirus_G2-A_partial_genome			83.0%	83.0%		TCAACGAGAAAAACA		
46	AY394978.1	SARS_coronavirus_G2-B_partial_genome			76.0%	76.0%		TCAACGAGAAAAACA		
47	AY394979.1	SARS_coronavirus_G2-C_partial_genome			82.7%	82.7%		TCAACGAGAAAAACA		
48	AY394980.1	SARS_coronavirus_G2-D_partial_genome			84.7%	84.3%		TCAACGAGAAAAACA		
49	AY394981.1	SARS_coronavirus_H2-B1A_partial_genome			84.7%	84.3%		TCAACGAGAAAAACA		
50	AY394982.1	SARS_coronavirus_H2-B1B_partial_genome			82.7%	82.7%		TCAACGAGAAAAACA		
51	AY394983.1	SARS_coronavirus_H2-B2_A_complete_genome			82.7%	82.7%		TCAACGAGAAAAACA		
52	AY394984.1	SARS_coronavirus_H2-B2_B_partial_genome			84.3%	84.3%		TCAACGAGAAAAACA		
53	AY394985.1	SARS_coronavirus_H2-B2_FB_complete_genome			16.7%	16.7%		TCAACGAGAAAAACA		
54	AY394986.1	SARS_coronavirus_H2-B2_Cb_complete_genome			83.0%	83.0%		TCAACGAGAAAAACA		
55	AY394987.1	SARS_coronavirus_H2-B2_Fc_complete_genome			86.0%	86.0%		TCAACGAGAAAAACA		
56	AY394988.1	SARS_coronavirus_JMD_partial_genome			71.3%	71.0%		TCAACGAGAAAAACA		
57	AY394989.1	SARS_coronavirus_H2S2-B_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
58	AY394990.1	SARS_coronavirus_H2S2-E_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
59	AY394991.1	SARS_coronavirus_H2S2-Fc_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
60	AY394992.1	SARS_coronavirus_H2S2-C_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
61	AY394993.1	SARS_coronavirus_H2S2-L2_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
62	AY394994.1	SARS_coronavirus_H2S2-Bc_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
63	AY394995.1	SARS_coronavirus_H2S2-Cc_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
64	AY394996.1	SARS_coronavirus_ZS-B_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
65	AY394997.1	SARS_coronavirus_ZS-A_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
66	AY394998.1	SARS_coronavirus_LC1_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
67	AY394999.1	SARS_coronavirus_LC2_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
68	AY395000.1	SARS_coronavirus_LC3_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
69	AY395001.1	SARS_coronavirus_LC4_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
70	AY395002.1	SARS_coronavirus_LC5_complete_genome			95.0%	95.0%		TCAACGAGAAAAACA		
71	AY395003.1	SARS_coronavirus_ZS-C_complete_genome			83.0%	83.0%		TCAACGAGAAAAACA		
72	AY395004.1	SARS_coronavirus_H2S2-Bb_partial_genome			84.3%	84.3%		TCAACGAGAAAAACA		
73	AY427439.1	SARS_coronavirus_AS_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
74	AY461660.1	SARS_coronavirus_SoJ_complete_genome			95.0%	94.7%		TCAACGAGAAAAACA		
75	AY463059.1	SARS_coronavirus_ShanghaiQXC1_complete_genome			73.7%	73.7%		TCAACGAGAAAAACA		
76	AY463060.1	SARS_coronavirus_ShanghaiQXC2_complete_genome			73.7%	73.7%		TCAACGAGAAAAACA		
77	AY485277.1	SARS_coronavirus_Sin11-01_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
78	AY485278.1	SARS_coronavirus_Sin03-11_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
79	AY502923.1	SARS_coronavirus_TW10_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
80	AY502924.1	SARS_coronavirus_TW11_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
81	AY502925.1	SARS_coronavirus_TW2_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
82	AY502926.1	SARS_coronavirus_TW3_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
83	AY502927.1	SARS_coronavirus_TW4_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
84	AY502928.1	SARS_coronavirus_TW5_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
85	AY502929.1	SARS_coronavirus_TW6_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
86	AY502930.1	SARS_coronavirus_TW7_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
87	AY502931.1	SARS_coronavirus_TW8_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
88	AY502932.1	SARS_coronavirus_TW9_complete_genome			100.0%	100.0%		TCAACGAGAAAAACA		
89	AY508704.1	SARS_coronavirus_SNS1_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
90	AY559081.1	SARS_coronavirus_Sin842_complete_genome			95.7%	95.3%		TCAACGAGAAAAACA		
91	AY559082.1	SARS_coronavirus_Sin852_complete_genome			99.7%	96.3%		TCAACGAGAAAAACA		
92	AY559083.1	SARS_coronavirus_Sin3408_complete_genome			88.7%	85.3%		TCAACGAGAAAAACA		
93	AY559084.1	SARS_coronavirus_Sin765V_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
94	AY559085.1	SARS_coronavirus_Sin848_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
95	AY559086.1	SARS_coronavirus_Sin849_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
96	AY559087.1	SARS_coronavirus_Sin3725V_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
97	AY559088.1	SARS_coronavirus_SinP1_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
98	AY559089.1	SARS_coronavirus_SinP2_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
99	AY559090.1	SARS_coronavirus_SinP3_complete_genome			95.3%	92.7%		TCAACGAGAAAAACA		
100	AY559091.1	SARS_coronavirus_SinP4_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
101	AY559092.1	SARS_coronavirus_SinP5_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
102	AY559093.1	SARS_coronavirus_Sin845_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
103	AY559094.1	SARS_coronavirus_Sin846_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
104	AY559095.1	SARS_coronavirus_Sin847_complete_genome			96.7%	96.0%		TCAACGAGAAAAACA		
105	AY559096.1	SARS_coronavirus_Sin850_complete_genome			97.3%	95.7%		TCAACGAGAAAAACA		
106	AY559097.1	SARS_coronavirus_Sin3408L_complete_genome			94.7%	94.7%		TCAACGAGAAAAACA		
107	AY595412.1	SARS_coronavirus_LLJ-2004_complete_genome			93.0%	93.0%		TCAACGAGAAAAACA		
108	AY654624.1	SARS_coronavirus_TJF_complete_genome			99.7%	99.0%				

69	AY395001.1_SARS_coronavirus_LC4_complete_genome	95.0%	95.0%	-----
70	AY395002.1_SARS_coronavirus_LC5_complete_genome	95.0%	95.0%	-----
71	AY395003.1_SARS_coronavirus_ZS-C_complete_genome	83.0%	83.0%	-----
72	AY395004.1_SARS_coronavirus_HSZ2-Bb_partial_genome	84.3%	84.3%	-----
73	AY427439.1_SARS_coronavirus_AS_complete_genome	94.7%	94.7%	-----
74	AY461660.1_SARS_coronavirus_SoD_complete_genome	95.0%	94.7%	-----
75	AY463059.1_SARS_coronavirus_ShanghaiQXC1_complete_genome	73.7%	73.7%	-----
76	AY463060.1_SARS_coronavirus_ShanghaiQXC2_complete_genome	73.7%	73.7%	-----
77	AY485277.1_SARS_coronavirus_Sin01-11_complete_genome	100.0%	100.0%	-----
78	AY485278.1_SARS_coronavirus_Sin03-11_complete_genome	100.0%	100.0%	-----
79	AY502923.1_SARS_coronavirus_TW10_complete_genome	100.0%	100.0%	-----
80	AY502924.1_SARS_coronavirus_TW11_complete_genome	100.0%	100.0%	-----
81	AY502925.1_SARS_coronavirus_TW2_complete_genome	100.0%	100.0%	-----
82	AY502926.1_SARS_coronavirus_TW3_complete_genome	100.0%	100.0%	-----
83	AY502927.1_SARS_coronavirus_TW4_complete_genome	100.0%	100.0%	-----
84	AY502928.1_SARS_coronavirus_TW5_complete_genome	100.0%	100.0%	-----
85	AY502929.1_SARS_coronavirus_TW6_complete_genome	100.0%	100.0%	-----
86	AY502930.1_SARS_coronavirus_TW7_complete_genome	100.0%	100.0%	-----
87	AY502931.1_SARS_coronavirus_TW8_complete_genome	100.0%	100.0%	-----
88	AY502932.1_SARS_coronavirus_TW9_complete_genome	100.0%	100.0%	-----
89	AY508724.1_SARS_coronavirus_NS-1_complete_genome	94.7%	94.7%	-----
90	AY559081.1_SARS_coronavirus_Sin842_complete_genome	95.7%	95.3%	-----
91	AY559082.1_SARS_coronavirus_Sin852_complete_genome	99.7%	96.3%	-----
92	AY559083.1_SARS_coronavirus_Sin408_complete_genome	88.7%	85.3%	-----
93	AY559084.1_SARS_coronavirus_Sin3765V_complete_genome	94.7%	94.7%	-----
94	AY559085.1_SARS_coronavirus_Sin848_complete_genome	94.7%	94.7%	-----
95	AY559086.1_SARS_coronavirus_Sin849_complete_genome	94.7%	94.7%	-----
96	AY559087.1_SARS_coronavirus_Sin3725V_complete_genome	94.7%	94.7%	-----
97	AY559088.1_SARS_coronavirus_SinP1_complete_genome	94.7%	94.7%	-----
98	AY559089.1_SARS_coronavirus_SinP2_complete_genome	94.7%	94.7%	-----
99	AY559090.1_SARS_coronavirus_SinP3_complete_genome	95.3%	92.7%	-----
100	AY559091.1_SARS_coronavirus_SinP4_complete_genome	94.7%	94.7%	-----
101	AY559092.1_SARS_coronavirus_SinP5_complete_genome	94.7%	94.7%	-----
102	AY559093.1_SARS_coronavirus_Sin845_complete_genome	94.7%	94.7%	-----
103	AY559094.1_SARS_coronavirus_Sin846_complete_genome	94.7%	94.7%	-----
104	AY559095.1_SARS_coronavirus_Sin847_complete_genome	96.7%	96.0%	-----
105	AY559096.1_SARS_coronavirus_Sin850_complete_genome	97.3%	95.7%	-----
106	AY559097.1_SARS_coronavirus_Sin3408L_complete_genome	94.7%	94.7%	-----
107	AY594412.1_SARS_coronavirus_LiJ-2004_complete_genome	93.0%	93.0%	-----
108	AY654624.1_SARS_coronavirus_TJF_complete_genome	99.7%	99.0%	-----
109	AY714217.1_SARS_Coronavirus_CDC_200301157_complete_genome	100.0%	100.0%	-----
110	CQ903009.1_Sequence_13_from_Patent_WO2004094667	100.0%	99.7%	-----
111	CQ918584.1_Sequence_1_from_Patent_WO2004096842	95.0%	95.0%	-----
112	CQ918595.1_Sequence_2_from_Patent_WO2004096842	95.0%	95.0%	-----
113	CQ918598.1_Sequence_15_from_Patent_WO2004096842	100.0%	100.0%	-----
114	NC_004718.3_SARS_coronavirus_complete_genome	100.0%	100.0%	-----
	consensus/100%			-----
	consensus/90%			-----
	consensus/80%			-----
	consensus/70%			-----

		cov	pid	561				584
1	AP006557.1_SARS_coronavirus_TWH_genomic_RNA_complete_genome	100.0%	100.0%	-----				-----
2	AP006558.1_SARS_coronavirus_TWJ_genomic_RNA_complete_genome	100.0%	100.0%	-----				-----
3	AP006559.1_SARS_coronavirus_TWK_genomic_RNA_complete_genome	100.0%	100.0%	-----				-----
4	AP006560.1_SARS_coronavirus_TWS_genomic_RNA_complete_genome	100.0%	100.0%	-----				-----
5	AP006561.1_SARS_coronavirus_TWY_genomic_RNA_complete_genome	100.0%	100.0%	-----				-----
6	AY274119.3_Severe_acute_respiratory_syndrome-related_coronavirus_isolate_Tor2_complete_genome	100.0%	100.0%	-----				-----
7	AY278487.1_SARS_coronavirus_B01_complete_genome	100.0%	100.0%	-----				-----
8	AY278488.2_SARS_coronavirus_B01_complete_genome	93.7%	93.7%	-----				-----
9	AY278489.2_SARS_coronavirus_GD01_complete_genome	94.7%	94.7%	-----				-----
10	AY278490.3_SARS_coronavirus_B03_complete_genome	98.7%	98.7%	-----				-----
11	AY278491.2_SARS_coronavirus_HKU-39849_complete_genome	100.0%	100.0%	-----				-----
12	AY27854.1_SARS_coronavirus_CUH-K1_complete_genome	95.0%	95.0%	-----				-----
13	AY278741.1_SARS_coronavirus_Urbani_complete_genome	100.0%	100.0%	-----				-----
14	AY279354.2_SARS_coronavirus_B04_complete_genome	94.7%	94.7%	-----				-----
15	AY282752.2_SARS_coronavirus_CUH-K-Su10_complete_genome	95.0%	95.0%	-----				-----
16	AY283794.1_SARS_coronavirus_Sin2500_complete_genome	94.7%	94.7%	-----				-----
17	AY283795.1_SARS_coronavirus_Sin2677_complete_genome	94.7%	94.7%	-----				-----
18	AY283796.1_SARS_coronavirus_Sin2679_complete_genome	94.7%	94.7%	-----				-----
19	AY283797.1_SARS_coronavirus_Sin2748_complete_genome	94.7%	94.7%	-----				-----
20	AY283798.2_SARS_coronavirus_Sin2774_complete_genome	94.7%	94.7%	-----				-----
21	AY291315.1_SARS_coronavirus_Frankfurt_1_complete_genome	100.0%	100.0%	-----				-----
22	AY291451.1_SARS_coronavirus_TW1_complete_genome	100.0%	100.0%	-----				-----
23	AY297028.1_SARS_coronavirus_Z01_complete_genome	95.3%	95.3%	-----				-----
24	AY304486.1_SARS_coronavirus_S23_complete_genome	95.0%	95.0%	-----				-----
25	AY304488.1_SARS_coronavirus_S216_complete_genome	95.0%	95.0%	-----				-----
26	AY304495.1_SARS_coronavirus_G250_complete_genome	95.0%	95.0%	-----				-----
27	AY310120.1_SARS_coronavirus_FNA_complete_genome	100.0%	100.0%	-----				-----
28	AY313906.1_SARS_coronavirus_GD69_complete_genome	100.0%	100.0%	-----				-----
29	AY321118.1_SARS_coronavirus_TWC_complete_genome	100.0%	100.0%	-----				-----
30	AY323977.2_SARS_coronavirus_HSR_1_complete_genome	100.0%	100.0%	-----				-----
31	AY338174.1_SARS_coronavirus_Taiwan_TC1_complete_genome	77.0%	77.0%	-----				-----
32	AY338175.1_SARS_coronavirus_Taiwan_TC2_complete_genome	77.0%	77.0%	-----				-----
33	AY345986.1_SARS_coronavirus_CUH-K-AG01_complete_genome	95.0%	95.0%	-----				-----
34	AY345987.1_SARS_coronavirus_CUH-K-AG02_complete_genome	95.0%	95.0%	-----				-----
35	AY345988.1_SARS_coronavirus_CUH-K-AG03_complete_genome	95.0%	95.0%	-----				-----
36	AY348314.1_SARS_coronavirus_Taiwan_TC3_complete_genome	77.0%	77.0%	-----				-----
37	AY350750.1_SARS_coronavirus_PUMC01_complete_genome	95.7%	95.7%	-----				-----
38	AY351680.1_SARS_coronavirus_ZMY_1_complete_genome	100.0%	99.7%	-----				-----
39	AY357075.1_SARS_coronavirus_PUMC02_complete_genome	95.3%	95.3%	-----				-----
40	AY357076.1_SARS_coronavirus_PUMC03_complete_genome	95.3%	95.3%	-----				-----
41	AY362698.1_SARS_coronavirus_TW2_complete_genome	100.0%	100.0%	-----				-----
42	AY362699.1_SARS_coronavirus_TW3_complete_genome	100.0%	100.0%	-----				-----
43	AY390556.1_SARS_coronavirus_G202_complete_genome	100.0%	100.0%	-----				-----
44	AY394850.2_SARS_coronavirus_WUO_complete_genome	100.0%	100.0%	-----				-----
45	AY394977.1_SARS_coronavirus_G2-A_partial_genome	83.0%	83.0%	-----				-----
46	AY394978.1_SARS_coronavirus_G2-B_partial_genome	76.0%	76.0%	-----				-----
47	AY394979.1_SARS_coronavirus_G2-C_partial_genome	82.7%	82.7%	-----				-----
48	AY394980.1_SARS_coronavirus_G2-D_partial_genome	84.7%	84.3%	-----				-----
49	AY394981.1_SARS_coronavirus_HGZBL1-A_partial_genome	83.0%	83.0%	-----				-----
50	AY394982.1_SARS_coronavirus_HGZBL1-B_partial_genome	82.7%	82.7%	-----				-----
51	AY394983.1_SARS_coronavirus_HGZ2-A_complete_genome	82.7%	82.7%	-----				-----
52	AY394984.1_SARS_coronavirus_HSZ-A_partial_genome	84.3%	84.3%	-----				-----
53	AY394985.1_SARS_coronavirus_HSZ-Bb_partial_genome	16.7%	16.7%	-----	TGGTTCGAGAAATGGACGCGATTC			-----
54	AY394986.1_SARS_coronavirus_HSZ-Cb_partial_genome	83.0%	83.0%	-----				-----
55	AY394987.1_SARS_coronavirus_HSZ2-Fb_complete_genome	86.0%	86.0%	-----				-----
56	AY394988.1_SARS_coronavirus_JMD_partial_genome	71.3%	71.0%	-----				-----
57	AY394989.1_SARS_coronavirus_HZS-D_complete_genome	95.0%	95.0%	-----				-----
58	AY394990.1_SARS_coronavirus_HZS2-E_complete_genome	95.0%	95.0%	-----				-----
59	AY394991.1_SARS_coronavirus_HZS2-Fc_complete_genome	95.0%	95.0%	-----				-----
60	AY394992.1_SARS_coronavirus_HZS2-C_complete_genome	95.0%	95.0%	-----				-----
61	AY394993.1_SARS_coronavirus_HGZBL2_complete_genome	95.0%	95.0%	-----				-----
62	AY394994.1_SARS_coronavirus_HSZ-Bc_complete_genome	95.0%	95.0%	-----				-----
63	AY394995.1_SARS_coronavirus_HSZ-Cc_complete_genome	95.0%	95.0%	-----				-----
64	AY394996.1_SARS_coronavirus_ZS-B_complete_genome	95.0%	95.0%	-----				-----
65	AY394997.1_SARS_coronavirus_ZS-A_complete_genome	95.0%	95.0%	-----				-----
66	AY394998.1_SARS_coronavirus_LC1_complete_genome	95.0%	95.0%	-----				-----
67	AY394999.1_SARS_coronavirus_LC2_complete_genome	95.0%	95.0%	-----				-----
68	AY395000.1_SARS_coronavirus_LC3_complete_genome	95.0%	95.0%	-----				-----
69	AY395001.1_SARS_coronavirus_LC4_complete_genome	95.0%	95.0%	-----				-----
70	AY395002.1_SARS_coronavirus_LC5_complete_genome	95.0%	95.0%	-----				-----
71	AY395003.1_SARS_coronavirus_ZS-C_complete_genome	83.0%	83.0%	-----				-----
72	AY395004.1_SARS_coronavirus_HSZ2-Bb_partial_genome	84.3%	84.3%	-----				-----
73	AY427439.1_SARS_coronavirus_AS_complete_genome	94.7%	94.7%	-----				-----
74	AY461660.1_SARS_coronavirus_SoD_complete_genome	95.0%	94.7%	-----				-----
75	AY463059.1_SARS_coronavirus_ShanghaiQXC1_complete_genome	73.7%	73.7%	-----				-----
76	AY463060.1_SARS_coronavirus_ShanghaiQXC2_complete_genome	73.7%	73.7%	-----				-----
77	AY485277.1_SARS_coronavirus_Sin01-11_complete_genome	100.0%	100.0%	-----				-----
78	AY485278.1_SARS_coronavirus_Sin03-11_complete_genome	100.0%	100.0%	-----				-----
79	AY502923.1_SARS_coronavirus_TW10_complete_genome	100.0%	100.0%	-----				-----
80	AY502924.1_SARS_coronavirus_TW11_complete_genome	100.0%	100.0%	-----				-----
81	AY502925.1_SARS_coronavirus_TW2_complete_genome	100.0%	100.0%	-----				-----
82	AY502926.1_SARS_coronavirus_TW3_complete_genome	100.0%	100.0%	-----				-----
83	AY502927.1_SARS_coronavirus_TW4_complete_genome	100.0%	100.0%	-----				-----
84	AY502928.1_SARS_coronavirus_TW5_complete_genome	100.0%	100.0%	-----				-----
85	AY502929.1_SARS_coronavirus_TW6_complete_genome	100.0%	100.0%	-----				-----
86	AY502930.1_SARS_coronavirus_TW7_complete_genome	100.0%	100.0%	-----				-----
87	AY502931.1_SARS_coronavirus_TW8_complete_genome	100.0%	100.0%	-----				-----
88	AY502932.1_SARS_coronavirus_TW9_complete_genome	100.0%	100.0%	-----				-----
89	AY508724.1_SARS_coronavirus_NS-1_complete_genome	94.7%	94.7%	-----				-----
90	AY559081.1_SARS_coronavirus_Sin842_complete_genome	95.7%	95.3%	-----				-----
91	AY559082.1_SARS_coronavirus_Sin852_complete_genome	99.7%	96.3%	-----				-----
92	AY559083.1_SARS_coronavirus_Sin408_complete_genome	88.7%	85.3%	-----				-----
93	AY559084.1_SARS_coronavirus_Sin3765V_complete_genome	94.7%	94.7%	-----				-----
94	AY559085.1_SARS_coronavirus_Sin848_complete_genome	94.7%	94.7%	-----				-----
95	AY559086.1_SARS_coronavirus_Sin849_complete_genome	94.7%	94.7%	-----				-----
96	AY559087.1_SARS_coronavirus_Sin3725V_complete_genome	94.7%	94.7%	-----				-----
97	AY559088.1_SARS_coronavirus_SinP1_complete_genome	94.7%	94.7%	-----				-----
98	AY559089.1_SARS_coronavirus_SinP2_complete_genome	94.7%	94.7%	-----				-----
99	AY559090.1_SARS_coronavirus_SinP3_complete_genome	95.3%	92.7%	-----				-----
100	AY559091.1_SARS_coronavirus_SinP4_complete_genome	94.7%	94.7%	-----				-----
101	AY559092.1_SARS_coronavirus_SinP5_complete_genome	94.7%	94.7%	-----				-----
102	AY559093.1_SARS_coronavirus_Sin845_complete_genome	94.7%	94.7%	-----				-----
103	AY559094.1_SARS_coronavirus_Sin846_complete_genome	94.7%	94.7%	-----				-----
104	AY559095.1_SARS_coronavirus_Sin847_complete_genome	96.7%	96.0%	-----				-----
105	AY559096.1_SARS_coronavirus_Sin850_complete_genome	97.3%	95.7%	-----				-----
106	AY559097.1_SARS_coronavirus_Sin3408L_complete_genome	94.7%	94.7%	-----				-----
107	AY594412.1_SARS_coronavirus_LiJ-2004_complete_genome	93.0%	93.0%	-----				-----

108	AY654624.1_SARS_coronavirus_TJF_complete_genome	99.7%	99.0%	-----
109	AY714217.1_SARS_Coronavirus_CDC*200301157_complete_genome	100.0%	100.0%	-----
110	CQ903009.1_Sequence_13_from_Patent_WO2004094667	100.0%	99.7%	-----
111	CQ918584.1_Sequence_1_from_Patent_WO2004096842	95.0%	95.0%	-----
112	CQ918585.1_Sequence_2_from_Patent_WO2004096842	95.0%	95.0%	-----
113	CQ918598.1_Sequence_15_from_Patent_WO2004096842	100.0%	100.0%	-----
114	NC_004718.3_SARS_coronavirus_complete_genome	100.0%	100.0%	-----
	consensus/100%		
	consensus/90%		
	consensus/80%		
	consensus/70%		

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