

Visus Test Table

NCBI Id	Genome_lenth	Number of aligned reads	Coverage (bases)
NC_001422.1	5386	4834017	5386
NC_001802.1	9181	4412255	9047
NC_022518.1	9472	20130	6670
NC_001362.1	8323	10327	4465
NC_001357.1	7857	7430	5222
NC_001560.1	11161	6527	1656
NC_006273.2	235646	2103	742
NC_001500.1	6296	1952	1975
NC_001501.1	8332	1937	7684
NC_009823.1	9711	763	110
NC_001416.1	48502	711	4486
NC_019723.1	47090	651	3095
NC_029853.1	8183	640	3908
NC_019711.1	47288	614	2707
NC_001479.1	7835	570	693
NC_001493.2	134226	435	275
NC_023437.1	7934	353	104
NC_001499.1	5894	343	1423
NC_001782.1	1801	239	149
NC_031032.1	24320	227	183
NC_001502.1	5833	208	1565
NC_019491.1	291144	180	1075
NC_002641.1	159160	164	304
NC_028696.2	41338	142	2121
NC_016447.1	219474	137	102
NC_019445.1	44784	134	3330
NC_032111.1	163005	133	263
NC_009127.1	295146	128	441
NC_004813.1	57930	103	1194
NC_009758.1	8926	86	97
NC_011357.1	62147	84	2323

NCBI Id	Genome_lenth	Number of aligned reads	Coverage (bases)
NC_028095.1	1705	83	107
NC_011356.1	54896	79	2136
NC_008310.2	6485	76	97
NC_005856.1	94800	72	1234
NC_008580.1	8380	70	89
NC_013221.1	2896	68	126
NC_004107.1	3323	65	588
NC_001716.2	153080	60	712
NC_001664.3	159321	56	485
NC_022749.1	39758	54	797
NC_001722.1	10359	53	163
NC_029302.1	6038	47	96
NC_002229.3	177874	47	334
NC_034266.1	204237	45	302
NC_024450.1	204054	45	308
NC_028656.1	65955	43	1181
NC_015253.1	127065	41	110
NC_027339.1	38389	40	1240
NC_022098.1	2473870	36	382
NC_004102.1	9646	34	101
NC_007821.1	6068	33	125
NC_001954.1	8454	31	530
NC_009514.1	47021	29	350
NC_008912.1	3141	22	152
NC_028449.1	63569	22	185
NC_004914.3	62706	21	292
NC_019768.1	41468	20	262
NC_002167.1	39732	19	325
NC_031129.1	103445	19	570
NC_000898.1	162114	16	267
NC_024771.1	151601	14	98
NC_021857.1	41475	14	245

NCBI Id	Genome_lenth	Number of aligned reads	Coverage (bases)
NC_005344.1	39043	14	323
NC_003287.2	6407	14	387
NC_001875.2	131995	12	185
NC_006641.1	15959	10	105
NC_007815.2	8197	8	218
NC_002577.1	164270	7	111
NC_001669.1	5243	7	239
NC_021858.1	1908524	6	83
AC_000007.1	35937	6	95
NC_002166.1	40751	6	116
NC_001405.1	35937	6	123
AC_000017.1	36001	6	148
NC_027984.1	57677	6	394
NC_019767.1	40155	5	90
NC_008925.1	3821	5	93
NC_008291.1	198050	5	94
NC_001330.1	6087	5	98
NC_009240.1	96944	5	99
NC_021190.1	46150	5	284
NC_018874.1	211518	4	80
NC_019769.1	38964	4	190
NC_020231.1	233501	3	84
NC_024364.1	131952	3	84
NC_019704.1	44375	3	90
NC_007646.1	135135	3	100
NC_001806.2	152222	3	185
NC_019716.1	44510	3	277
NC_004560.1	5784	2	80
NC_019442.1	60523	2	90
NC_011345.1	155122	2	92
NC_022337.1	29467	2	95
NC_001702.1	8135	2	122

NCBI Id	Genome_lenth	Number of aligned reads	Coverage (bases)
NC_011048.1	19282	2	190
NC_012743.2	62715	2	190
NC_018839.1	29729	2	190
NC_001266.1	159857	1	77
NC_022334.1	29266	1	82
NC_018847.1	29516	1	84
NC_027396.1	33756	1	89
NC_001420.2	5577	1	90
NC_007856.1	5486	1	90
NC_027983.1	167435	1	90
NC_027362.1	29394	1	93
NC_031119.1	29347	1	93
NC_000924.1	61670	1	95
NC_018841.1	29574	1	95
NC_018842.1	29348	1	95
NC_018845.1	29371	1	95
NC_018852.1	29506	1	95
NC_022338.1	29514	1	95
NC_027359.1	29491	1	95
NC_027371.1	29741	1	95
NC_027401.1	29751	1	95
NC_027626.1	29347	1	95
NC_027628.1	29517	1	95
NC_028943.1	29237	1	95
NC_007817.1	5486	1	96