

Organism	No. of genes (N_{genes})	Average fold occurrence (C)	No. of folds (N_{folds})	No. of generations (t)	Initial no. of folds (N_0)	Rate of fold transfer (r)	Generations per transfer ($1/r$)
<i>M. genitalium</i>	481	2.4	200	280	20	0.64	1.6
<i>M. pneumoniae</i>	688	2.5	275	413	20	0.62	1.6
<i>R. prowazekii</i>	834	2.6	321	514	40	0.55	1.8
<i>C. trachomatis</i>	894	2.6	344	550	60	0.52	1.9
<i>T. pallidum</i>	1031	2.8	368	662	30	0.51	2.0
<i>C. pneumoniae</i>	1052	2.7	390	662	60	0.5	2.0
<i>A. aeolicus</i>	1522	4.3	354	1180	100	0.21	4.8
<i>H. pylori</i>	1553	3.3	470	1081	40	0.4	2.5
<i>B. burgdorferi</i>	1638	2.9	565	1074	120	0.41	2.4
<i>H. influenzae</i>	1709	3.7	462	1249	90	0.3	3.3
<i>M. jannaschii</i>	1715	4.8	357	1357	70	0.21	4.8
<i>M. thermoautotrophicum</i>	1869	5	374	1495	80	0.2	5.0
<i>P. horikoshii</i>	2064	4.6	449	1616	70	0.24	4.2
<i>A. fulgidus</i>	2420	5.8	417	2001	90	0.16	5.3
<i>Synechocystis sp.</i>	3169	5.7	556	2613	110	0.17	5.9
<i>M. tuberculosis</i>	3918	8	490	3426	160	0.096	10.4
<i>B. subtilis</i>	4100	7	586	3447	190	0.12	8.3
<i>E. coli</i>	4289	7	613	3605	170	0.12	8.3
<i>S. cerevisiae</i>	6269	10.9	575	5693	200	0.066	15.1
<i>C. elegans</i>	19099	31.6	604	18482	280	0.018	55.6