

Amino acid	% abundance	ln (% abunda	Cost (ATP)	Cost (ATP/tin	ln (% abunda	Organism	Genomic GC	R % abundan	R ln(% abund
A	5.51	1.70656462	11.7	11.7	1.292	S. cerevisiae	0.38	-0.52	-0.6
C	1.28	0.24686008	24.7	741	1.082				
D	5.84	1.7647308	12.7	114.3	1.082				
E	6.57	1.88251383	15.3	76.5	1.082				
F	4.48	1.49962305	52	208	1.565				
G	5.01	1.61143592	11.7	11.7	1.292				
H	2.13	0.75612198	38.3	536.2	1.082				
I	6.55	1.87946505	32.3	64.6	2.046				
K	7.35	1.99470031	30.3	242.4	1.565				
L	9.56	2.25758773	27.3	54.6	2.368				
M	2.07	0.72754861	34.3	445.9	0.601				
N	6.1	1.80828877	14.7	147	1.565				
P	4.31	1.4609379	20.3	60.9	1.292				
Q	3.92	1.36609165	16.3	130.4	1.082				
R	4.44	1.49065438	27.3	109.2	1.885				
S	8.96	2.19277023	11.7	70.2	2.18				
T	5.82	1.76130026	18.7	112.2	1.775				
V	5.6	1.7227666	23.3	46.6	1.775				
W	1.04	3.92E-02	74.3	891.6	0.118				
Y	3.34	1.20597081	50	350	1.565				
A	6.33	1.8453	11.7	11.7	1.178	S. pombe	0.36	-0.52	-0.61
C	1.53	0.42527	24.7	741	1.058				
D	5.32	1.67147	12.7	114.3	1.058				
E	6.49	1.87026	15.3	76.5	1.058				
F	4.59	1.52388	52	208	1.632				
G	4.99	1.60744	11.7	11.7	1.178				
H	2.25	0.81093	38.3	536.2	1.058				
I	6.13	1.81319	32.3	64.6	2.126				
K	6.39	1.85473	30.3	242.4	1.632				
L	9.83	2.28544	27.3	54.6	2.387				
M	2.05	0.71784	34.3	445.9	0.611				
N	5.18	1.64481	14.7	147	1.632				

R % abundan R ln(% abund R ln(% abundance), observed vs predicted

-0.76 -0.91 0.77

-0.77 -0.92 0.78

P	4.69	1.54543	20.3	60.9	1.178			
Q	3.79	1.33237	16.3	130.4	1.058			
R	4.84	1.57691	27.3	109.2	1.813			
S	9.39	2.23965	11.7	70.2	2.157			
T	5.6	1.72277	18.7	112.2	1.752			
V	6.01	1.79342	23.3	46.6	1.752			
W	1.11	0.10436	74.3	891.6	3.80E-02			
Y	3.4	1.22378	50	350	1.632			
A	8.7	2.16332303	11.7	11.7	1.827	N. crassa	0.5	-0.66
C	1.13	0.12221763	24.7	741	1.139			-0.67
D	5.63	1.72810944	12.7	114.3	1.139			
E	6.46	1.86562932	15.3	76.5	1.139			
F	3.37	1.21491274	52	208	1.145			
G	7.2	1.97408103	11.7	11.7	1.827			
H	2.45	0.89608803	38.3	536.2	1.139			
I	4.42	1.4861397	32.3	64.6	1.551			
K	5.08	1.62531126	30.3	242.4	1.145			
L	8.35	2.12226154	27.3	54.6	2.24			
M	2.15	0.76546784	34.3	445.9	0.449			
N	3.7	1.30833282	14.7	147	1.145			
P	6.52	1.87487438	20.3	60.9	1.827			
Q	4.26	1.44926916	16.3	130.4	1.139			
R	6.17	1.81969884	27.3	109.2	2.235			
S	8.29	2.11504997	11.7	70.2	2.238			
T	6.11	1.80992677	18.7	112.2	1.833			
V	5.98	1.78842057	23.3	46.6	1.833			
W	1.35	0.30010459	74.3	891.6	0.444			
Y	2.56	0.94000726	50	350	1.145			
A	4.93	1.59533899	11.7	11.7	1.032	C. albicans	0.34	-0.51
C	1.17	0.15700375	24.7	741	1.024			-0.59
D	5.71	1.74221902	12.7	114.3	1.024			
E	6.23	1.82937633	15.3	76.5	1.024			

-0.83

-0.93

0.77

-0.75

-0.9

0.75

F	4.52	1.50851199	52	208	1.709			
G	4.97	1.60341984	11.7	11.7	1.032			
H	2.14	0.76080583	38.3	536.2	1.024			
I	7.13	1.96431123	32.3	64.6	2.219			
K	7.2	1.97408103	30.3	242.4	1.709			
L	9.3	2.2300144	27.3	54.6	2.407			
M	1.87	0.62593843	34.3	445.9	0.616			
N	6.69	1.90061387	14.7	147	1.709			
P	4.49	1.5018527	20.3	60.9	1.032			
Q	4.42	1.4861397	16.3	130.4	1.024			
R	3.72	1.31372367	27.3	109.2	1.721			
S	9.29	2.22893855	11.7	70.2	2.123			
T	6.16	1.81807678	18.7	112.2	1.717			
V	5.51	1.70656462	23.3	46.6	1.717			
W	0.98	-2.02E-02	74.3	891.6	-6.90E-02			
Y	3.47	1.24415459	50	350	1.709			
A	9.38	2.23857976	11.7	11.7	1.88	M. grisea		
C	1.28	0.24686008	24.7	741	1.139		0.51	
D	5.7	1.74046618	12.7	114.3	1.139			-0.64
E	5.9	1.77495235	15.3	76.5	1.139			-0.66
F	3.47	1.24415459	52	208	1.09			
G	7.38	1.99877364	11.7	11.7	1.88			
H	2.33	0.84586827	38.3	536.2	1.139			
I	4.37	1.47476301	32.3	64.6	1.488			
K	4.78	1.56444055	30.3	242.4	1.09			
L	8.52	2.14241634	27.3	54.6	2.222			
M	2.2	0.78845736	34.3	445.9	0.421			
N	3.54	1.26412673	14.7	147	1.09			
P	6.34	1.84687877	20.3	60.9	1.88			
Q	4.1	1.41098697	16.3	130.4	1.139			
R	6.55	1.87946505	27.3	109.2	2.27			
S	8	2.07944154	11.7	70.2	2.237			
T	5.86	1.7681496	18.7	112.2	1.832			

-0.81

-0.92

0.8

V	6.2	1.82454929	23.3	46.6	1.832			
W	1.47	0.3852624	74.3	891.6	0.47			
Y	2.52	0.9242589	50	350	1.09			
A	8.23	2.10778601	11.7	11.7	1.903	F. graminearu	0.52	-0.63
C	1.28	0.24686008	24.7	741	1.138			-0.64
D	5.94	1.78170913	12.7	114.3	1.138			
E	6.21	1.8261609	15.3	76.5	1.138			
F	3.76	1.32441896	52	208	1.066			
G	6.7	1.90210753	11.7	11.7	1.903			
H	2.38	0.86710049	38.3	536.2	1.138			
I	5.11	1.6311994	32.3	64.6	1.459			
K	5.13	1.63510566	30.3	242.4	1.066			
L	8.68	2.16102153	27.3	54.6	2.213			
M	2.27	0.81977983	34.3	445.9	0.408			
N	3.87	1.35325451	14.7	147	1.066			
P	5.87	1.76985463	20.3	60.9	1.903			
Q	4.03	1.39376638	16.3	130.4	1.138			
R	5.75	1.74919986	27.3	109.2	2.285			
S	8.14	2.09679018	11.7	70.2	2.237			
T	6.1	1.80828877	18.7	112.2	1.831			
V	6.12	1.8115621	23.3	46.6	1.831			
W	1.51	0.41210965	74.3	891.6	0.48			
Y	2.8	1.02961942	50	350	1.066			
A	8.56	2.14710019	11.7	11.7	1.848	A. nidulans	0.5	-0.61
C	1.24	0.21511138	24.7	741	1.139			-0.63
D	5.57	1.71739505	12.7	114.3	1.139			
E	6.19	1.82293509	15.3	76.5	1.139			
F	3.7	1.30833282	52	208	1.124			
G	6.77	1.91250109	11.7	11.7	1.848			
H	2.37	0.86288996	38.3	536.2	1.139			
I	5	1.60943791	32.3	64.6	1.527			
K	4.55	1.51512723	30.3	242.4	1.124			

-0.84

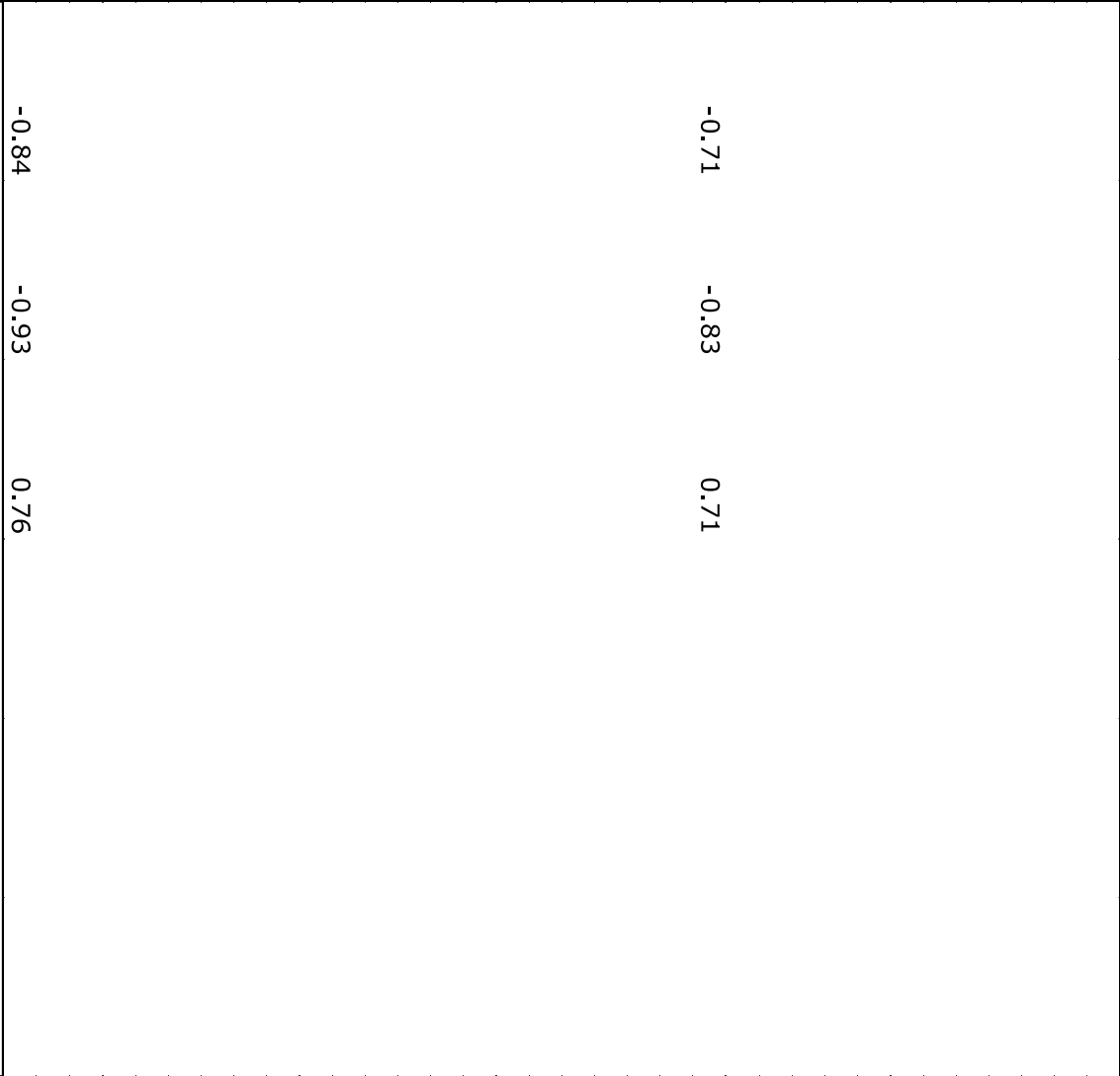
-0.94

0.76

-0.82

-0.93

0.8



C	2.03	0.70803579	24.7	741	1.051		
D	5.3	1.66770682	12.7	114.3	1.051		
E	6.5	1.87180218	15.3	76.5	1.051		
F	4.83	1.57484647	52	208	1.651		
G	5.34	1.67522565	11.7	11.7	1.144		
H	2.32	0.84156719	38.3	536.2	1.051		
I	6.14	1.81482474	32.3	64.6	2.149		
K	6.42	1.85941812	30.3	242.4	1.651		
L	8.65	2.15755932	27.3	54.6	2.392		
M	2.61	0.95935022	34.3	445.9	0.613		
N	4.91	1.59127394	14.7	147	1.651		
P	4.91	1.59127394	20.3	60.9	1.144		
Q	4.11	1.41342303	16.3	130.4	1.051		
R	5.21	1.65057986	27.3	109.2	1.792		
S	8.04	2.08442908	11.7	70.2	2.149		
T	5.84	1.7647308	18.7	112.2	1.744		
V	6.19	1.82293509	23.3	46.6	1.744		
W	1.1	9.53E-02	74.3	891.6	1.30E-02		
Y	3.16	1.15057203	50	350	1.651		
A	6.18	1.82131827	11.7	11.7	1.249	C. briggsae	
C	1.95	0.66782937	24.7	741	1.073		0.37
D	5.29	1.66581825	12.7	114.3	1.073		-0.59
E	6.81	1.91839212	15.3	76.5	1.073		-0.67
F	4.66	1.53901545	52	208	1.591		
G	5.37	1.68082791	11.7	11.7	1.249		
H	2.32	0.84156719	38.3	536.2	1.073		
I	5.89	1.773256	32.3	64.6	2.077		
K	6.46	1.86562932	30.3	242.4	1.591		
L	8.46	2.13534917	27.3	54.6	2.375		
M	2.62	0.96317432	34.3	445.9	0.606		
N	4.78	1.56444055	14.7	147	1.591		
P	5.05	1.61938824	20.3	60.9	1.249		
Q	4.13	1.41827741	16.3	130.4	1.073		

-0.84

-0.94

0.77

R	5.49	1.70292826	27.3	109.2	1.858			
S	8.01	2.08069076	11.7	70.2	2.172			
T	5.73	1.74571553	18.7	112.2	1.766			
V	6.09	1.80664808	23.3	46.6	1.766			
W	1.15	0.13976194	74.3	891.6	8.80E-02			
Y	3.09	1.12817109	50	350	1.591			
A	7.46	2.00955541	11.7	11.7	1.495	D. melanogas	0.42	-0.68
C	1.87	0.62593843	24.7	741	1.115			-0.76
D	5.19	1.6467337	12.7	114.3	1.115			
E	6.41	1.85785927	15.3	76.5	1.115			
F	3.55	1.2669476	52	208	1.428			
G	6.18	1.82131827	11.7	11.7	1.495			
H	2.68	0.9858168	38.3	536.2	1.115			
I	4.87	1.58309394	32.3	64.6	1.884			
K	5.55	1.71379793	30.3	242.4	1.428			
L	9.15	2.21375388	27.3	54.6	2.329			
M	2.36	0.85866162	34.3	445.9	0.566			
N	4.65	1.53686722	14.7	147	1.428			
P	5.46	1.69744879	20.3	60.9	1.495			
Q	5.19	1.6467337	16.3	130.4	1.115			
R	5.59	1.72097929	27.3	109.2	2.016			
S	8.26	2.11142459	11.7	70.2	2.214			
T	5.59	1.72097929	18.7	112.2	1.808			
V	5.93	1.78002421	23.3	46.6	1.808			
W	1	0	74.3	891.6	0.253			
Y	2.97	1.08856195	50	350	1.428			
A	7.43	2.00552586	11.7	11.7	1.602	A. gambiae	0.45	-0.63
C	2.01	0.69813472	24.7	741	1.127			-0.72
D	5.25	1.65822808	12.7	114.3	1.127			
E	6.24	1.83098018	15.3	76.5	1.127			
F	3.76	1.32441896	52	208	1.346			
G	6.36	1.85002838	11.7	11.7	1.602			

-0.84

-0.94

0.78

-0.84

-0.94

0.83

H	2.65	0.97455964	38.3	536.2	1.127			
I	5.23	1.65441128	32.3	64.6	1.788			
K	5.53	1.71018782	30.3	242.4	1.346			
L	9.32	2.23216263	27.3	54.6	2.304			
M	2.24	0.80647587	34.3	445.9	0.538			
N	4.38	1.47704872	14.7	147	1.346			
P	5.13	1.63510566	20.3	60.9	1.602			
Q	4.32	1.4632554	16.3	130.4	1.127			
R	6.02	1.79508726	27.3	109.2	2.086			
S	7.46	2.00955541	11.7	70.2	2.226			
T	5.83	1.763017	18.7	112.2	1.821			
V	6.41	1.85785927	23.3	46.6	1.821			
W	1.12	0.11332869	74.3	891.6	0.319			
Y	3.14	1.14422228	50	350	1.346			
A	6.16	1.81807678	11.7	11.7	1.177	A. thaliana	0.36	-0.56
C	1.92	0.65232519	24.7	741	1.058			-0.65
D	5.37	1.68082791	12.7	114.3	1.058			
E	6.68	1.89911799	15.3	76.5	1.058			
F	4.32	1.4632554	52	208	1.632			
G	6.31	1.84213568	11.7	11.7	1.177			
H	2.32	0.84156719	38.3	536.2	1.058			
I	5.35	1.67709656	32.3	64.6	2.127			
K	6.39	1.85473427	30.3	242.4	1.632			
L	9.54	2.25549349	27.3	54.6	2.387			
M	2.43	0.88789126	34.3	445.9	0.611			
N	4.39	1.47932923	14.7	147	1.632			
P	4.73	1.5539252	20.3	60.9	1.177			
Q	3.5	1.25276297	16.3	130.4	1.058			
R	5.5	1.70474809	27.3	109.2	1.813			
S	9.02	2.19944433	11.7	70.2	2.157			
T	5.13	1.63510566	18.7	112.2	1.751			
V	6.66	1.89611949	23.3	46.6	1.751			
W	1.29	0.25464222	74.3	891.6	3.70E-02			

-0.78

-0.91

0.73

Y	2.89	1.0612565	50	350	1.632			
A	6.97	1.94161523	11.7	11.7	1.402	H. sapiens	0.4	-0.62
C	2.31	0.83724753	24.7	741	1.101			-0.73
D	4.68	1.54329811	12.7	114.3	1.101			
E	6.95	1.93874166	15.3	76.5	1.101			
F	3.64	1.29198368	52	208	1.494			
G	6.7	1.90210753	11.7	11.7	1.402			
H	2.64	0.97077892	38.3	536.2	1.101			
I	4.35	1.47017585	32.3	64.6	1.962			
K	5.72	1.74396881	30.3	242.4	1.494			
L	9.87	2.28949985	27.3	54.6	2.348			
M	2.2	0.78845736	34.3	445.9	0.585			
N	3.58	1.2753628	14.7	147	1.494			
P	6.37	1.85159947	20.3	60.9	1.402			
Q	4.71	1.54968791	16.3	130.4	1.101			
R	5.68	1.73695123	27.3	109.2	1.956			
S	8.25	2.1102132	11.7	70.2	2.2			
T	5.34	1.67522565	18.7	112.2	1.794			
V	6	1.79175947	23.3	46.6	1.794			
W	1.27	0.2390169	74.3	891.6	0.193			
Y	2.63	0.96698385	50	350	1.494			
A	6.7	1.90210753	11.7	11.7	1.448	M. musculus	0.41	-0.6
C	2.47	0.90421815	24.7	741	1.108			-0.72
D	4.75	1.55814462	12.7	114.3	1.108			
E	6.79	1.91545094	15.3	76.5	1.108			
F	3.91	1.36353737	52	208	1.462			
G	6.4	1.85629799	11.7	11.7	1.448			
H	2.76	1.01523068	38.3	536.2	1.108			
I	4.57	1.51951321	32.3	64.6	1.924			
K	5.8	1.75785792	30.3	242.4	1.462			
L	9.99	2.30158459	27.3	54.6	2.339			
M	2.18	0.77932488	34.3	445.9	0.576			

-0.78

-0.89

0.74

-0.77

-0.89

0.79

N	3.64	1.29198368	14.7	147	1.462			
P	5.86	1.7681496	20.3	60.9	1.448			
Q	4.67	1.54115907	16.3	130.4	1.108			
R	5.53	1.71018782	27.3	109.2	1.986			
S	8.18	2.10169215	11.7	70.2	2.207			
T	5.43	1.69193913	18.7	112.2	1.801			
V	6.18	1.82131827	23.3	46.6	1.801			
W	1.22	0.19885086	74.3	891.6	0.223			
Y	2.85	1.04731899	50	350	1.462			
A	6.89	1.93007109	11.7	11.7	1.481	R. norvegicus	0.42	-0.61
C	2.28	0.82417544	24.7	741	1.113			-0.72
D	4.76	1.56024767	12.7	114.3	1.113			
E	6.83	1.92132467	15.3	76.5	1.113			
F	3.7	1.30833282	52	208	1.439			
G	6.49	1.87026253	11.7	11.7	1.481			
H	2.62	0.96317432	38.3	536.2	1.113			
I	4.36	1.47247206	32.3	64.6	1.897			
K	5.65	1.73165555	30.3	242.4	1.439			
L	10	2.30258509	27.3	54.6	2.332			
M	2.23	0.80200159	34.3	445.9	0.57			
N	3.55	1.2669476	14.7	147	1.439			
P	6.04	1.79840401	20.3	60.9	1.481			
Q	4.7	1.54756251	16.3	130.4	1.113			
R	5.64	1.72988407	27.3	109.2	2.007			
S	8.37	2.12465389	11.7	70.2	2.212			
T	5.46	1.69744879	18.7	112.2	1.806			
V	6.27	1.83577636	23.3	46.6	1.806			
W	1.24	0.21511138	74.3	891.6	0.244			
Y	2.66	0.97832612	50	350	1.439			
A	6.56	1.8809906	11.7	11.7	1.642	F. rubripes	0.45	-0.62
C	2.38	0.86710049	24.7	741	1.131			-0.73
D	5.08	1.62531126	12.7	114.3	1.131			

-0.77

-0.89

0.78

-0.79

-0.9

0.84

E	6.49	1.87026253	15.3	76.5	1.131		
F	3.8	1.33500107	52	208	1.313		
G	6.44	1.86252854	11.7	11.7	1.642		
H	2.73	1.00430161	38.3	536.2	1.131		
I	4.52	1.50851199	32.3	64.6	1.749		
K	5.42	1.69009582	30.3	242.4	1.313		
L	9.8	2.28238239	27.3	54.6	2.294		
M	2.28	0.82417544	34.3	445.9	0.525		
N	3.81	1.33762919	14.7	147	1.313		
P	5.65	1.73165555	20.3	60.9	1.642		
Q	4.64	1.53471437	16.3	130.4	1.131		
R	5.78	1.75440368	27.3	109.2	2.112		
S	8.31	2.11745961	11.7	70.2	2.23		
T	5.54	1.7119945	18.7	112.2	1.824		
V	6.55	1.87946505	23.3	46.6	1.824		
W	1.25	0.22314355	74.3	891.6	0.343		
Y	2.85	1.04731899	50	350	1.313		
A	5.81	1.75958057	11.7	11.7	1.157	P. falciparum	
C	2.39	0.87129337	24.7	741	1.054	0.19	
D	5.41	1.68824909	12.7	114.3	1.054		-0.24
E	6.24	1.83098018	15.3	76.5	1.054		-0.43
F	4.11	1.41342303	52	208	1.643		
G	5.72	1.74396881	11.7	11.7	1.157		
H	2.53	0.9282193	38.3	536.2	1.054		
I	5.58	1.71918878	32.3	64.6	2.14		
K	6.44	1.86252854	30.3	242.4	1.643		
L	8.82	2.17702187	27.3	54.6	2.39		
M	2.48	0.90825856	34.3	445.9	0.613		
N	5.16	1.64093658	14.7	147	1.643		
P	4.68	1.54329811	20.3	60.9	1.157		
Q	4.1	1.41098697	16.3	130.4	1.054		
R	5.06	1.62136648	27.3	109.2	1.8		
S	8.04	2.08442908	11.7	70.2	2.152		

-0.36

-0.59

0.82

T	6.11	1.80992677	18.7	112.2	1.747			
V	6.7	1.90210753	23.3	46.6	1.747			
W	1.2	0.18232156	74.3	891.6	2.30E-02			
Y	3.25	1.178655	50	350	1.643			
A	8.61	2.15292432	11.7	11.7	1.825	C. intestinalis	0.36	-0.71
C	1.27	0.2390169	24.7	741	1.139			
D	5.55	1.71379793	12.7	114.3	1.139			
E	6.14	1.81482474	15.3	76.5	1.139			
F	3.7	1.30833282	52	208	1.147			
G	6.7	1.90210753	11.7	11.7	1.825			
H	2.42	0.88376754	38.3	536.2	1.139			
I	4.91	1.59127394	32.3	64.6	1.554			
K	4.62	1.53039471	30.3	242.4	1.147			
L	9.12	2.2104698	27.3	54.6	2.241			
M	2.11	0.74668795	34.3	445.9	0.45			
N	3.58	1.2753628	14.7	147	1.147			
P	6.05	1.80005827	20.3	60.9	1.825			
Q	4.06	1.40118297	16.3	130.4	1.139			
R	6.34	1.84687877	27.3	109.2	2.233			
S	8.4	2.12823171	11.7	70.2	2.238			
T	5.88	1.77155676	18.7	112.2	1.833			
V	6.19	1.82293509	23.3	46.6	1.833			
W	1.45	0.37156356	74.3	891.6	0.442			
Y	2.78	1.02245093	50	350	1.147			
A	8.22	2.10657021	11.7	11.7	1.773	A. fumigatus	0.5	-0.61
C	1.32	0.27763174	24.7	741	1.139			-0.63
D	5.55	1.71379793	12.7	114.3	1.139			
E	6.05	1.80005827	15.3	76.5	1.139			
F	3.86	1.35066718	52	208	1.197			
G	6.88	1.92861865	11.7	11.7	1.773			
H	2.42	0.88376754	38.3	536.2	1.139			
I	5.16	1.64093658	32.3	64.6	1.612			

-0.81

-0.91

0.77

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-0.93

0.81

K	4.64	1.53471437	30.3	242.4	1.197			
L	9.22	2.22137504	27.3	54.6	2.257			
M	2.21	0.79299252	34.3	445.9	0.474			
N	3.73	1.31640823	14.7	147	1.197			
P	5.81	1.75958057	20.3	60.9	1.773			
Q	3.97	1.3787661	16.3	130.4	1.139			
R	5.94	1.78170913	27.3	109.2	2.199			
S	8.2	2.10413415	11.7	70.2	2.237			
T	5.95	1.78339122	18.7	112.2	1.832			
V	6.31	1.84213568	23.3	46.6	1.832			
W	1.53	0.42526774	74.3	891.6	0.416			
Y	2.95	1.08180517	50	350	1.197			
A	11	2.39789527	11.7	11.7	2.023	C. neoforman	0.49	-0.62
C	1.78	0.57661336	24.7	741	1.129			
D	4.91	1.59127394	12.7	114.3	1.129			
E	6.29	1.83896107	15.3	76.5	1.129			
F	3.29	1.19088757	52	208	0.929			
G	6.39	1.85473427	11.7	11.7	2.023			
H	2.48	0.90825856	38.3	536.2	1.129			
I	3.63	1.28923265	32.3	64.6	1.3			
K	2.76	1.01523068	30.3	242.4	0.929			
L	10.3	2.3321439	27.3	54.6	2.165			
M	1.83	0.60431597	34.3	445.9	0.331			
N	2.56	0.94000726	14.7	147	0.929			
P	5.5	1.70474809	20.3	60.9	2.023			
Q	4.3	1.45861502	16.3	130.4	1.129			
R	8.74	2.16791019	27.3	109.2	2.366			
S	7.96	2.074429	11.7	70.2	2.228			
T	5.57	1.71739505	18.7	112.2	1.823			
V	6.74	1.90805993	23.3	46.6	1.823			
W	1.52	0.41871034	74.3	891.6	0.532			
Y	2.3	0.83290912	50	350	0.929			

-0.83

-0.93

0.8

A	6.7	1.90210753	11.7	11.7	1.577	C. merolae	0.55	-0.53	-0.63
C	1.34	0.29266961	24.7	741	1.125				
D	5.67	1.73518912	12.7	114.3	1.125				
E	6.67	1.89761986	15.3	76.5	1.125				
F	4.32	1.4632554	52	208	1.366				
G	5.55	1.71379793	11.7	11.7	1.577				
H	2.21	0.79299252	38.3	536.2	1.125				
I	5.48	1.7011051	32.3	64.6	1.811				
K	6.45	1.86408013	30.3	242.4	1.366				
L	9.77	2.27931647	27.3	54.6	2.311				
M	2.05	0.71783979	34.3	445.9	0.545				
N	4.69	1.54543258	14.7	147	1.366				
P	4.56	1.51732262	20.3	60.9	1.577				
Q	4.22	1.43983513	16.3	130.4	1.125				
R	5	1.60943791	27.3	109.2	2.069				
S	9.1	2.20827441	11.7	70.2	2.224				
T	5.71	1.74221902	18.7	112.2	1.818				
V	6.18	1.82131827	23.3	46.6	1.818				
W	1.09	8.62E-02	74.3	891.6	0.304				
Y	3.15	1.14740245	50	350	1.366				
A	7.85	2.06051353	11.7	11.7	1.911	K. waltii	0.44	-0.56	-0.63
C	1.37	0.31481074	24.7	741	1.138				
D	5.63	1.72810944	12.7	114.3	1.138				
E	6.55	1.87946505	15.3	76.5	1.138				
F	3.93	1.36863943	52	208	1.058				
G	5.95	1.78339122	11.7	11.7	1.911				
H	2.39	0.87129337	38.3	536.2	1.138				
I	5.04	1.61740608	32.3	64.6	1.45				
K	5.65	1.73165555	30.3	242.4	1.058				
L	10	2.30258509	27.3	54.6	2.21				
M	2.18	0.77932488	34.3	445.9	0.404				
N	4.09	1.40854497	14.7	147	1.058				
P	4.86	1.58103844	20.3	60.9	1.911				

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-0.93

0.78

Q	4.26	1.44926916	16.3	130.4	1.138			
R	5.85	1.76644166	27.3	109.2	2.291			
S	7.96	2.074429	11.7	70.2	2.236			
T	5.44	1.69377906	18.7	112.2	1.831			
V	6.49	1.87026253	23.3	46.6	1.831			
W	1.12	0.11332869	74.3	891.6	0.484			
Y	3.26	1.18172719	50	350	1.058			
A	5.52	1.70837786	11.7	11.7	1.316	A. gossypii	0.52	-0.58
C	1.17	0.15700375	24.7	741	1.086			-0.65
D	6.15	1.81645208	12.7	114.3	1.086			
E	6.72	1.90508816	15.3	76.5	1.086			
F	4.18	1.43031125	52	208	1.55			
G	5.23	1.65441128	11.7	11.7	1.316			
H	2.12	0.75141609	38.3	536.2	1.086			
I	6.51	1.87333946	32.3	64.6	2.029			
K	7.28	1.98513086	30.3	242.4	1.55			
L	9.3	2.2300144	27.3	54.6	2.364			
M	2.2	0.78845736	34.3	445.9	0.598			
N	6.02	1.79508726	14.7	147	1.55			
P	4.25	1.44691898	20.3	60.9	1.316			
Q	3.93	1.36863943	16.3	130.4	1.086			
R	4.43	1.48839958	27.3	109.2	1.901			
S	8.69	2.16217294	11.7	70.2	2.185			
T	5.87	1.76985463	18.7	112.2	1.779			
V	5.85	1.76644166	23.3	46.6	1.779			
W	0.98	-2.02E-02	74.3	891.6	0.134			
Y	3.48	1.24703229	50	350	1.55			
A	5.58	1.71918878	11.7	11.7	1.323	C. glabrata	0.39	-0.54
C	1.25	0.22314355	24.7	741	1.088			-0.61
D	6.02	1.79508726	12.7	114.3	1.088			
E	6.65	1.89461686	15.3	76.5	1.088			
F	4.4	1.48160454	52	208	1.545			

-0.82

-0.94

0.74

-0.78

-0.92

0.76

G	5.09	1.62727783	11.7	11.7	1.323		
H	2.19	0.78390154	38.3	536.2	1.088		
I	6.33	1.84530024	32.3	64.6	2.023		
K	7.04	1.95160817	30.3	242.4	1.545		
L	9.72	2.27418562	27.3	54.6	2.363		
M	2.08	0.73236789	34.3	445.9	0.597		
N	5.54	1.7119945	14.7	147	1.545		
P	4.31	1.4609379	20.3	60.9	1.323		
Q	4.19	1.43270073	16.3	130.4	1.088		
R	4.36	1.47247206	27.3	109.2	1.906		
S	8.86	2.18154677	11.7	70.2	2.186		
T	5.84	1.7647308	18.7	112.2	1.781		
V	6	1.79175947	23.3	46.6	1.781		
W	1.09	8.62E-02	74.3	891.6	0.14		
Y	3.34	1.20597081	50	350	1.545		
A	7.81	2.05540496	11.7	11.7	1.794	K. lactis	
C	1.22	0.19885086	24.7	741	1.139		0.39
D	5.97	1.78674693	12.7	114.3	1.139		-0.53
E	6.47	1.86717611	15.3	76.5	1.139		-0.6
F	3.88	1.35583515	52	208	1.177		
G	6.37	1.85159947	11.7	11.7	1.794		
H	2.42	0.88376754	38.3	536.2	1.139		
I	4.9	1.58923521	32.3	64.6	1.589		
K	5.85	1.76644166	30.3	242.4	1.177		
L	8.71	2.16447179	27.3	54.6	2.251		
M	2.24	0.80647587	34.3	445.9	0.465		
N	4.02	1.3912819	14.7	147	1.177		
P	5.46	1.69744879	20.3	60.9	1.794		
Q	4.19	1.43270073	16.3	130.4	1.139		
R	5.09	1.62727783	27.3	109.2	2.213		
S	8.25	2.1102132	11.7	70.2	2.238		
T	6.16	1.81807678	18.7	112.2	1.832		
V	6.71	1.90359895	23.3	46.6	1.832		

-0.78

-0.91

0.77

W	1.2	0.18232156	74.3	891.6	0.427			
Y	2.98	1.0919233	50	350	1.177			
A	5.14	1.63705308	11.7	11.7	1.191	Y. lipolytica	0.49	-0.64
C	1.17	0.15700375	24.7	741	1.061			-0.66
D	6.14	1.81482474	12.7	114.3	1.061			
E	6.56	1.8809906	15.3	76.5	1.061			
F	4.47	1.49738841	52	208	1.624			
G	5.21	1.65057986	11.7	11.7	1.191			
H	2.08	0.73236789	38.3	536.2	1.061			
I	7.05	1.95302762	32.3	64.6	2.117			
K	7.35	1.99470031	30.3	242.4	1.624			
L	9.61	2.26280422	27.3	54.6	2.385			
M	1.99	0.68813464	34.3	445.9	0.61			
N	6.63	1.8916048	14.7	147	1.624			
P	4.23	1.44220199	20.3	60.9	1.191			
Q	3.8	1.33500107	16.3	130.4	1.061			
R	4.02	1.3912819	27.3	109.2	1.821			
S	8.88	2.18380156	11.7	70.2	2.16			
T	5.46	1.69744879	18.7	112.2	1.754			
V	5.42	1.69009582	23.3	46.6	1.754			
W	0.99	-1.01E-02	74.3	891.6	4.70E-02			
Y	3.6	1.28093385	50	350	1.624			
A	2.03	0.71	11.7	11.7	-6.20E-02	D. hansenii	0.36	-0.5
C	1.78	0.58	24.7	741	0.67			-0.58
D	6.46	1.87	12.7	114.3	0.67			
E	7.13	1.96	15.3	76.5	0.67			
F	4.25	1.45	52	208	2.095			
G	2.94	1.08	11.7	11.7	-6.20E-02			
H	2.43	0.89	38.3	536.2	0.67			
I	9.12	2.21	32.3	64.6	2.686			
K	11.7	2.46	30.3	242.4	2.095			
L	7.51	2.02	27.3	54.6	2.487			

-0.85

-0.95

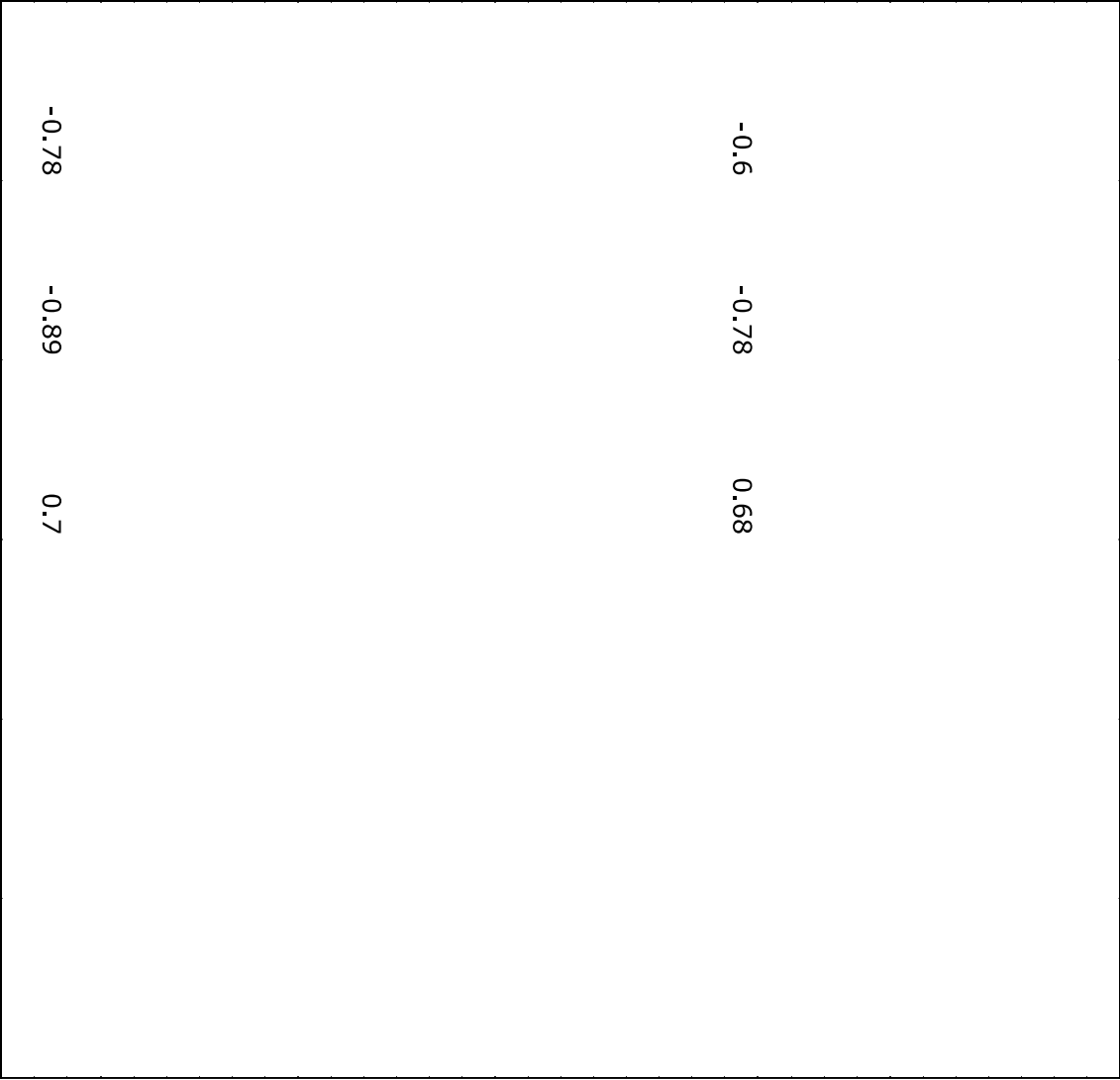
0.75

-0.75

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0.76

M	2.21	0.79	34.3	445.9	0.454			
N	14.1	2.65	14.7	147	2.095			
P	2.09	0.74	20.3	60.9	-6.20E-02			
Q	2.78	1.02	16.3	130.4	0.67			
R	2.67	0.98	27.3	109.2	1.063			
S	6.42	1.86	11.7	70.2	1.768			
T	4.16	1.43	18.7	112.2	1.363			
V	3.94	1.37	23.3	46.6	1.363			
W	0.49	-0.71	74.3	891.6	-0.971			
Y	5.64	1.73	50	350	2.095			
A	5.47	1.69927862	11.7	11.7	0.895	M. jannaschii	0.31	-0.29
C	1.27	0.2390169	24.7	741	0.989			-0.44
D	5.54	1.7119945	12.7	114.3	0.989			
E	8.68	2.16102153	15.3	76.5	0.989			
F	4.22	1.43983513	52	208	1.775			
G	6.34	1.84687877	11.7	11.7	0.895			
H	1.43	0.35767444	38.3	536.2	0.989			
I	10.4	2.34180581	32.3	64.6	2.298			
K	10.4	2.34180581	30.3	242.4	1.775			
L	9.48	2.24918432	27.3	54.6	2.423			
M	2.2	0.78845736	34.3	445.9	0.613			
N	5.29	1.66581825	14.7	147	1.775			
P	3.36	1.21194097	20.3	60.9	0.895			
Q	1.45	0.37156356	16.3	130.4	0.989			
R	3.85	1.34807315	27.3	109.2	1.636			
S	4.48	1.49962305	11.7	70.2	2.087			
T	4.04	1.39624469	18.7	112.2	1.682			
V	6.86	1.92570744	23.3	46.6	1.682			
W	0.72	-0.3285041	74.3	891.6	-0.173			
Y	4.36	1.47247206	50	350	1.775			
A	7.33	1.99197552	11.7	11.7	1.814	M. thermoaut	0.5	-0.49
C	1.2	0.18232156	24.7	741	1.139			-0.57



D	5.91	1.77664583	12.7	114.3	1.139		
E	8.14	2.09679018	15.3	76.5	1.139		
F	3.64	1.29198368	52	208	1.158		
G	7.97	2.07568449	11.7	11.7	1.814		
H	1.87	0.62593843	38.3	536.2	1.139		
I	7.7	2.04122033	32.3	64.6	1.566		
K	4.56	1.51732262	30.3	242.4	1.158		
L	9.47	2.24812891	27.3	54.6	2.244		
M	2.93	1.07500242	34.3	445.9	0.455		
N	3.31	1.19694819	14.7	147	1.158		
P	4.3	1.45861502	20.3	60.9	1.814		
Q	1.9	0.64185389	16.3	130.4	1.139		
R	6.79	1.91545094	27.3	109.2	2.226		
S	6.14	1.81482474	11.7	70.2	2.238		
T	4.96	1.60140574	18.7	112.2	1.832		
V	7.74	2.04640169	23.3	46.6	1.832		
W	0.84	-0.1743534	74.3	891.6	0.437		
Y	3.22	1.16938136	50	350	1.158		
A	7.84	2.05923883	11.7	11.7	1.775	A. fulgidus	
C	1.17	0.15700375	24.7	741	1.139		0.49
D	4.89	1.5871923	12.7	114.3	1.139		-0.4
E	8.9	2.18605128	15.3	76.5	1.139		-0.46
F	4.59	1.52388002	52	208	1.195		
G	7.24	1.97962121	11.7	11.7	1.775		
H	1.51	0.41210965	38.3	536.2	1.139		
I	7.22	1.97685495	32.3	64.6	1.61		
K	6.86	1.92570744	30.3	242.4	1.195		
L	9.5	2.2512918	27.3	54.6	2.257		
M	2.62	0.96317432	34.3	445.9	0.473		
N	3.21	1.16627094	14.7	147	1.195		
P	3.86	1.35066718	20.3	60.9	1.775		
Q	1.78	0.57661336	16.3	130.4	1.139		
R	5.77	1.75267208	27.3	109.2	2.2		

-0.73

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S	5.51	1.70656462	11.7	70.2	2.237			
T	4.16	1.42551507	18.7	112.2	1.832			
V	8.61	2.15292432	23.3	46.6	1.832			
W	1.03	2.96E-02	74.3	891.6	0.417			
Y	3.64	1.29198368	50	350	1.195			
A	6.37	1.85159947	11.7	11.7	1.478	P. horikoshii	0.42	-0.32
C	0.63	-0.4620355	24.7	741	1.113			-0.36
D	4.26	1.44926916	12.7	114.3	1.113			
E	8.29	2.11504997	15.3	76.5	1.113			
F	4.6	1.5260563	52	208	1.44			
G	6.97	1.94161523	11.7	11.7	1.478			
H	1.49	0.39877612	38.3	536.2	1.113			
I	8.78	2.17247641	32.3	64.6	1.899			
K	7.74	2.04640169	30.3	242.4	1.44			
L	10.3	2.3321439	27.3	54.6	2.333			
M	2.4	0.87546874	34.3	445.9	0.57			
N	3.53	1.26129787	14.7	147	1.44			
P	4.5	1.5040774	20.3	60.9	1.478			
Q	1.63	0.48858002	16.3	130.4	1.113			
R	5.45	1.69561561	27.3	109.2	2.005			
S	5.85	1.76644166	11.7	70.2	2.211			
T	4.51	1.50629715	18.7	112.2	1.806			
V	7.55	2.02154756	23.3	46.6	1.806			
W	1.17	0.15700375	74.3	891.6	0.242			
Y	3.84	1.34547237	50	350	1.44			
A	6.67	1.89761986	11.7	11.7	1.609	P. abysssi	0.45	-0.33
C	0.55	-0.597837	24.7	741	1.128			-0.35
D	4.6	1.5260563	12.7	114.3	1.128			
E	8.84	2.17928688	15.3	76.5	1.128			
F	4.35	1.47017585	52	208	1.341			
G	7.26	1.98237983	11.7	11.7	1.609			
H	1.5	0.40546511	38.3	536.2	1.128			

-0.71

-0.83

0.67

-0.71

-0.83

0.63

I	8.49	2.138889	32.3	64.6	1.781			
K	7.8	2.05412373	30.3	242.4	1.341			
L	10.2	2.32238772	27.3	54.6	2.303			
M	2.39	0.87129337	34.3	445.9	0.536			
N	3.33	1.2029723	14.7	147	1.341			
P	4.25	1.44691898	20.3	60.9	1.609			
Q	1.66	0.5068176	16.3	130.4	1.128			
R	5.72	1.74396881	27.3	109.2	2.09			
S	4.97	1.60341984	11.7	70.2	2.227			
T	4.2	1.43508453	18.7	112.2	1.821			
V	8.07	2.08815348	23.3	46.6	1.821			
W	1.17	0.15700375	74.3	891.6	0.323			
Y	3.82	1.34025042	50	350	1.341			
A	9.69	2.27109443	11.7	11.7	2.07	A. pernix		
C	0.78	-0.2484614	24.7	741	1.123		0.56	
D	4.22	1.43983513	12.7	114.3	1.123			-0.44
E	7.27	1.98375629	15.3	76.5	1.123			
F	2.87	1.05431203	52	208	0.87			
G	8.84	2.17928688	11.7	11.7	2.07			
H	1.61	0.47623418	38.3	536.2	1.123			
I	5.53	1.71018782	32.3	64.6	1.232			
K	3.94	1.37118072	30.3	242.4	0.87			
L	11	2.39789527	27.3	54.6	2.144			
M	2.21	0.79299252	34.3	445.9	0.295			
N	1.97	0.67803354	14.7	147	0.87			
P	5.52	1.70837786	20.3	60.9	2.07			
Q	1.76	0.56531381	16.3	130.4	1.123			
R	7.8	2.05412373	27.3	109.2	2.398			
S	6.66	1.89611949	11.7	70.2	2.222			
T	4.34	1.46787435	18.7	112.2	1.817			
V	9.03	2.20055237	23.3	46.6	1.817			
W	1.34	0.29266961	74.3	891.6	0.549			
Y	3.46	1.24126859	50	350	0.87			

-0.71

-0.82

0.73

A	6.95	1.93874166	11.7	11.7	1.665	T. acidophilur	0.46	-0.42	-0.45
C	0.6	-0.5108256	24.7	741	1.133				
D	5.74	1.74745921	12.7	114.3	1.133				
E	5.97	1.78674693	15.3	76.5	1.133				
F	4.7	1.54756251	52	208	1.294				
G	7.25	1.98100147	11.7	11.7	1.665				
H	1.64	0.49469624	38.3	536.2	1.133				
I	9.01	2.19833507	32.3	64.6	1.726				
K	5.63	1.72810944	30.3	242.4	1.294				
L	8.38	2.12584791	27.3	54.6	2.288				
M	3.2	1.16315081	34.3	445.9	0.517				
N	4.25	1.44691898	14.7	147	1.294				
P	3.96	1.37624403	20.3	60.9	1.665				
Q	2.15	0.76546784	16.3	130.4	1.133				
R	5.49	1.70292826	27.3	109.2	2.127				
S	7.57	2.02419307	11.7	70.2	2.232				
T	4.77	1.56234631	18.7	112.2	1.826				
V	7.15	1.96711236	23.3	46.6	1.826				
W	0.85	-0.1625189	74.3	891.6	0.356				
Y	4.63	1.53255687	50	350	1.294				
A	6.35	1.84845481	11.7	11.7	1.382	T. volcanium	0.4	-0.39	-0.44
C	0.62	-0.4780358	24.7	741	1.098				
D	5.48	1.7011051	12.7	114.3	1.098				
E	6.35	1.84845481	15.3	76.5	1.098				
F	4.73	1.5539252	52	208	1.507				
G	6.95	1.93874166	11.7	11.7	1.382				
H	1.52	0.41871034	38.3	536.2	1.098				
I	9.22	2.22137504	32.3	64.6	1.977				
K	6.89	1.93007109	30.3	242.4	1.507				
L	8.76	2.17019591	27.3	54.6	2.352				
M	2.71	0.99694864	34.3	445.9	0.588				
N	4.75	1.55814462	14.7	147	1.507				

-0.8

-0.89

0.66

-0.77

-0.88

0.71

P	3.76	1.32441896	20.3	60.9	1.382			
Q	2.09	0.73716407	16.3	130.4	1.098			
R	4.67	1.54115907	27.3	109.2	1.943			
S	7.52	2.01756614	11.7	70.2	2.197			
T	4.8	1.56861592	18.7	112.2	1.791			
V	7.15	1.96711236	23.3	46.6	1.791			
W	0.83	-0.1863296	74.3	891.6	0.18			
Y	4.75	1.55814462	50	350	1.507			
A	13.1	2.57261223	11.7	11.7	2.386	Halobacterium	0.66	-0.55
C	0.73	-0.3147107	24.7	741	1.032			-0.56
D	9.02	2.19944433	12.7	114.3	1.032			
E	6.69	1.90061387	15.3	76.5	1.032			
F	3.11	1.13462273	52	208	0.372			
G	8.52	2.14241634	11.7	11.7	2.386			
H	2.24	0.80647587	38.3	536.2	1.032			
I	3.58	1.2753628	32.3	64.6	0.665			
K	1.64	0.49469624	30.3	242.4	0.372			
L	8.34	2.12106322	27.3	54.6	1.955			
M	1.74	0.55388511	34.3	445.9	-4.40E-02			
N	2.12	0.75141609	14.7	147	0.372			
P	4.69	1.54543258	20.3	60.9	2.386			
Q	2.6	0.95551145	16.3	130.4	1.032			
R	6.49	1.87026253	27.3	109.2	2.616			
S	5.24	1.6563215	11.7	70.2	2.131			
T	6.83	1.92132467	18.7	112.2	1.726			
V	9.61	2.26280422	23.3	46.6	1.726			
W	1.06	5.83E-02	74.3	891.6	0.616			
Y	2.52	0.9242589	50	350	0.372			
A	5.6	1.7227666	11.7	11.7	1.164	S. solfataricu	0.36	-0.32
C	0.61	-0.4942963	24.7	741	1.055			-0.38
D	4.68	1.54329811	12.7	114.3	1.055			
E	6.81	1.91839212	15.3	76.5	1.055			

-0.68

-0.83

0.69

-0.72

-0.85

0.71

F	4.44	1.49065438	52	208	1.64			
G	6.43	1.86097454	11.7	11.7	1.164			
H	1.29	0.25464222	38.3	536.2	1.055			
I	9.43	2.2438961	32.3	64.6	2.136			
K	7.74	2.04640169	30.3	242.4	1.64			
L	10.3	2.3321439	27.3	54.6	2.389			
M	2.2	0.78845736	34.3	445.9	0.612			
N	4.95	1.59938758	14.7	147	1.64			
P	3.8	1.33500107	20.3	60.9	1.164			
Q	2.09	0.73716407	16.3	130.4	1.055			
R	4.71	1.54968791	27.3	109.2	1.804			
S	6.7	1.90210753	11.7	70.2	2.154			
T	4.72	1.5518088	18.7	112.2	1.748			
V	7.47	2.010895	23.3	46.6	1.748			
W	1.05	4.88E-02	74.3	891.6	2.80E-02			
Y	4.82	1.57277393	50	350	1.64			
A	6.55	1.87946505	11.7	11.7	1.424	P. furiosus		
C	0.59	-0.5276327	24.7	741	1.105		0.41	
D	4.36	1.47247206	12.7	114.3	1.105			-0.31
E	8.9	2.18605128	15.3	76.5	1.105			
F	4.43	1.48839958	52	208	1.478			
G	7.12	1.96290773	11.7	11.7	1.424			
H	1.51	0.41210965	38.3	536.2	1.105			
I	8.7	2.16332303	32.3	64.6	1.943			
K	8.1	2.09186406	30.3	242.4	1.478			
L	10.1	2.31253542	27.3	54.6	2.344			
M	2.2	0.78845736	34.3	445.9	0.581			
N	3.48	1.24703229	14.7	147	1.478			
P	4.28	1.45395301	20.3	60.9	1.424			
Q	1.75	0.55961579	16.3	130.4	1.105			
R	5.33	1.67335124	27.3	109.2	1.97			
S	4.94	1.59736533	11.7	70.2	2.203			
T	4.41	1.48387469	18.7	112.2	1.798			

-0.7

-0.83

0.64

V	7.9	2.06686276	23.3	46.6	1.798			
W	1.23	0.20701417	74.3	891.6	0.208			
Y	3.99	1.38379123	50	350	1.478			
A	5.56	1.71559811	11.7	11.7	0.989	S. tokodaii	0.33	-0.3
C	0.65	-0.4307829	24.7	741	1.013			-0.39
D	4.63	1.53255687	12.7	114.3	1.013			
E	7.02	1.94876322	15.3	76.5	1.013			
F	4.54	1.51292701	52	208	1.731			
G	6.28	1.83736998	11.7	11.7	0.989			
H	1.3	0.26236426	38.3	536.2	1.013			
I	9.92	2.29455292	32.3	64.6	2.245			
K	8	2.07944154	30.3	242.4	1.731			
L	10.3	2.3321439	27.3	54.6	2.412			
M	2.14	0.76080583	34.3	445.9	0.616			
N	4.87	1.58309394	14.7	147	1.731			
P	3.92	1.36609165	20.3	60.9	0.989			
Q	2.08	0.73236789	16.3	130.4	1.013			
R	4.15	1.42310833	27.3	109.2	1.694			
S	6.68	1.89911799	11.7	70.2	2.112			
T	4.76	1.56024767	18.7	112.2	1.706			
V	7.21	1.97546895	23.3	46.6	1.706			
W	1.01	9.95E-03	74.3	891.6	-0.102			
Y	4.89	1.5871923	50	350	1.731			
A	9.89	2.29152415	11.7	11.7	1.886	P. aerophilum	0.51	-0.38
C	0.87	-0.1392621	24.7	741	1.139			-0.39
D	4.29	1.45628673	12.7	114.3	1.139			
E	6.99	1.94448056	15.3	76.5	1.139			
F	3.63	1.28923265	52	208	1.084			
G	7.67	2.03731662	11.7	11.7	1.886			
H	1.51	0.41210965	38.3	536.2	1.139			
I	6.29	1.83896107	32.3	64.6	1.481			
K	5.68	1.73695123	30.3	242.4	1.084			

-0.7

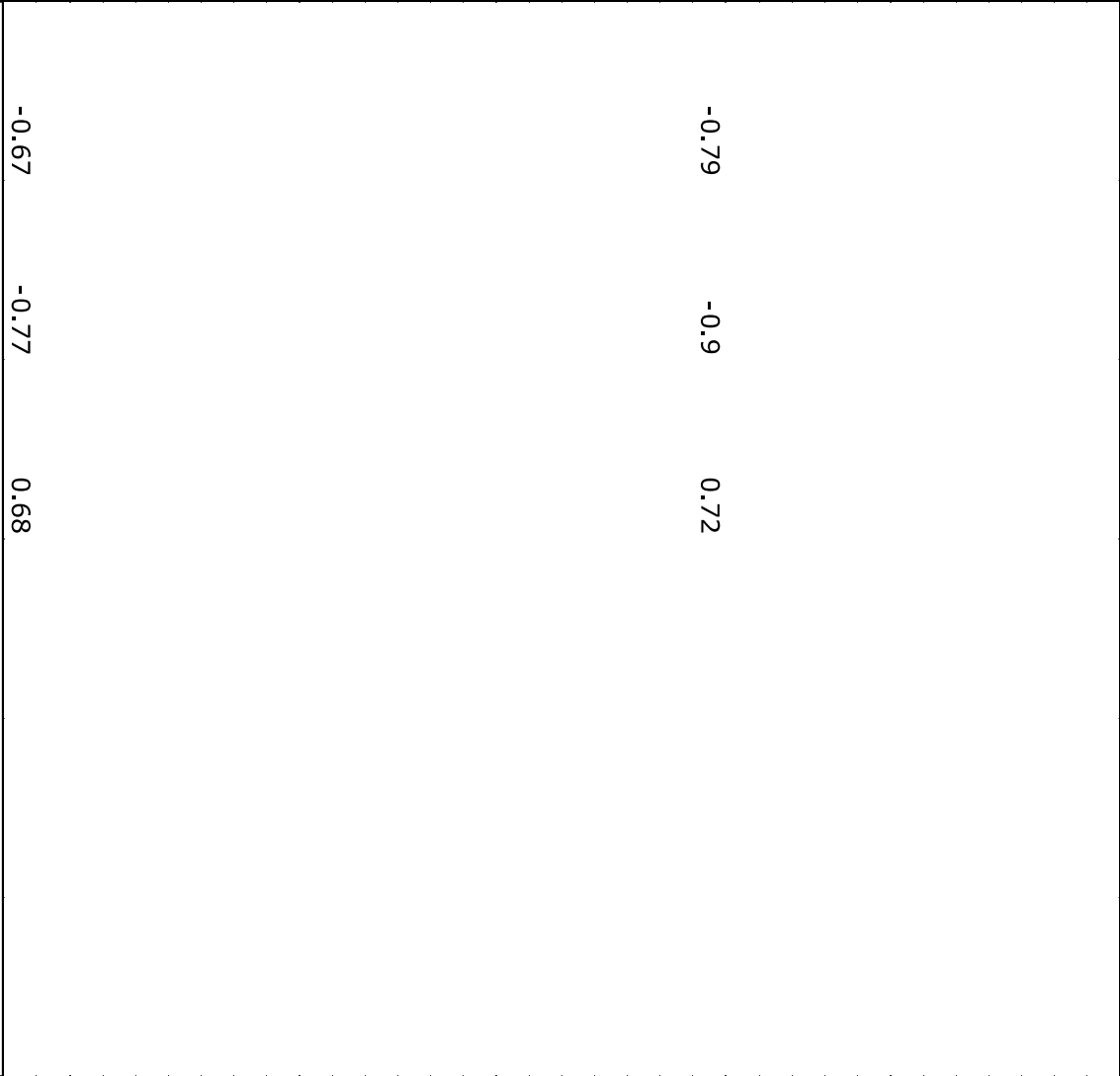
-0.85

0.7

-0.71

-0.83

0.69



C	1.31	0.27002714	24.7	741	1.088			
D	5.79	1.75613229	12.7	114.3	1.088			
E	9.99	2.30158459	15.3	76.5	1.088			
F	2.87	1.05431203	52	208	0.634			
G	8.05	2.08567209	11.7	11.7	2.236			
H	1.93	0.65752	38.3	536.2	1.088			
I	4.82	1.57277393	32.3	64.6	0.962			
K	4.02	1.3912819	30.3	242.4	0.634			
L	10	2.30258509	27.3	54.6	2.057			
M	1.9	0.64185389	34.3	445.9	0.143			
N	1.91	0.64710324	14.7	147	0.634			
P	5.46	1.69744879	20.3	60.9	2.236			
Q	1.4	0.33647224	16.3	130.4	1.088			
R	8.34	2.12106322	27.3	109.2	2.511			
S	4.62	1.53039471	11.7	70.2	2.187			
T	4.59	1.52388002	18.7	112.2	1.781			
V	10.4	2.34180581	23.3	46.6	1.781			
W	1.23	0.20701417	74.3	891.6	0.597			
Y	2.82	1.03673689	50	350	0.634			
A	7.07	1.95586048	11.7	11.7	1.459	M. mazel		
C	1.24	0.21511138	24.7	741	1.11		0.41	
D	5.25	1.65822808	12.7	114.3	1.11		-0.49	-0.57
E	8.16	2.09924417	15.3	76.5	1.11			
F	4.29	1.45628673	52	208	1.454			
G	7.22	1.97685495	11.7	11.7	1.459			
H	1.67	0.51282363	38.3	536.2	1.11			
I	7.65	2.03470565	32.3	64.6	1.915			
K	6.81	1.91839212	30.3	242.4	1.454			
L	9.37	2.2375131	27.3	54.6	2.337			
M	2.49	0.91228271	34.3	445.9	0.574			
N	4.35	1.47017585	14.7	147	1.454			
P	4	1.38629436	20.3	60.9	1.459			
Q	2.48	0.90825856	16.3	130.4	1.11			

-0.79

-0.9

0.73

R	4.67	1.54115907	27.3	109.2	1.993			
S	6.75	1.90954251	11.7	70.2	2.209			
T	5.11	1.6311994	18.7	112.2	1.803			
V	6.86	1.92570744	23.3	46.6	1.803			
W	0.93	-7.26E-02	74.3	891.6	0.23			
Y	3.53	1.26129787	50	350	1.454			
A	6.69	1.90061387	11.7	11.7	1.424	M. burtonii		
C	1.22	0.19885086	24.7	741	1.105		0.41	-0.5
D	6.29	1.83896107	12.7	114.3	1.105			-0.58
E	7.3	1.98787435	15.3	76.5	1.105			
F	4.06	1.40118297	52	208	1.479			
G	6.96	1.94017947	11.7	11.7	1.424			
H	1.95	0.66782937	38.3	536.2	1.105			
I	8.46	2.13534917	32.3	64.6	1.944			
K	6.45	1.86408013	30.3	242.4	1.479			
L	8.98	2.19499988	27.3	54.6	2.344			
M	2.96	1.08518927	34.3	445.9	0.581			
N	4.6	1.5260563	14.7	147	1.479			
P	3.61	1.28370777	20.3	60.9	1.424			
Q	2.45	0.89608803	16.3	130.4	1.105			
R	4.19	1.43270073	27.3	109.2	1.97			
S	6.8	1.91692261	11.7	70.2	2.203			
T	5.41	1.68824909	18.7	112.2	1.798			
V	7.26	1.98237983	23.3	46.6	1.798			
W	0.86	-0.1508229	74.3	891.6	0.207			
Y	3.39	1.22082992	50	350	1.479			
A	5.2	1.64865863	11.7	11.7	0.912	N. equitans	0.32	-0.21
C	0.82	-0.1984509	24.7	741	0.993			-0.36
D	4.95	1.59938758	12.7	114.3	0.993			
E	7.85	2.06051353	15.3	76.5	0.993			
F	4.42	1.4861397	52	208	1.767			
G	5.29	1.66581825	11.7	11.7	0.912			

-0.78

-0.9

0.71

-0.59

-0.81

0.7

H	1.34	0.29266961	38.3	536.2	0.993		
I	10.4	2.34180581	32.3	64.6	2.289		
K	10.8	2.37954613	30.3	242.4	1.767		
L	10.4	2.34180581	27.3	54.6	2.421		
M	1.69	0.52472853	34.3	445.9	0.614		
N	5.24	1.6563215	14.7	147	1.767		
P	4.02	1.3912819	20.3	60.9	0.912		
Q	2.21	0.79299252	16.3	130.4	0.993		
R	3.91	1.36353737	27.3	109.2	1.647		
S	4.62	1.53039471	11.7	70.2	2.092		
T	4.1	1.41098697	18.7	112.2	1.686		
V	5.92	1.77833645	23.3	46.6	1.686		
W	0.96	-4.08E-02	74.3	891.6	-0.16		
Y	5.64	1.72988407	50	350	1.767		
A	5.61	1.72455072	11.7	11.7	1.174	P. torridus	-0.4
C	0.61	-0.4942963	24.7	741	1.057	0.36	
D	5.86	1.7681496	12.7	114.3	1.057		
E	5.66	1.73342389	15.3	76.5	1.057		
F	4.85	1.57897871	52	208	1.634		
G	6.51	1.87333946	11.7	11.7	1.174		
H	1.48	0.39204209	38.3	536.2	1.057		
I	11	2.39789527	32.3	64.6	2.129		
K	6.9	1.93152141	30.3	242.4	1.634		
L	8.62	2.15408509	27.3	54.6	2.387		
M	3.06	1.11841492	34.3	445.9	0.612		
N	6.61	1.88858365	14.7	147	1.634		
P	3.48	1.24703229	20.3	60.9	1.174		
Q	1.46	0.37843644	16.3	130.4	1.057		
R	4.42	1.4861397	27.3	109.2	1.811		
S	7.43	2.00552586	11.7	70.2	2.156		
T	4.37	1.47476301	18.7	112.2	1.751		
V	5.73	1.74571553	23.3	46.6	1.751		
W	0.73	-0.3147107	74.3	891.6	3.50E-02		

-0.69

-0.82

0.71

Y	5.44	1.69377906	50	350	1.634			
A	10.6	2.360854	11.7	11.7	2.234	H. marismortu	0.61	-0.58
C	0.73	-0.3147107	24.7	741	1.089			
D	8.46	2.13534917	12.7	114.3	1.089			
E	8.15	2.09801793	15.3	76.5	1.089			
F	3.22	1.16938136	52	208	0.636			
G	8.39	2.12704052	11.7	11.7	2.234			
H	1.97	0.67803354	38.3	536.2	1.089			
I	4.27	1.45161383	32.3	64.6	0.965			
K	1.92	0.65232519	30.3	242.4	0.636			
L	8.79	2.17361471	27.3	54.6	2.058			
M	1.89	0.63657683	34.3	445.9	0.144			
N	2.51	0.92028275	14.7	147	0.636			
P	4.57	1.51951321	20.3	60.9	2.234			
Q	3.03	1.10856262	16.3	130.4	1.089			
R	6.09	1.80664808	27.3	109.2	2.51			
S	5.8	1.75785792	11.7	70.2	2.187			
T	6.82	1.91985947	18.7	112.2	1.782			
V	8.83	2.17815502	23.3	46.6	1.782			
W	1.12	0.11332869	74.3	891.6	0.596			
Y	2.66	0.97832612	50	350	0.636			
A	7.42	2.00417906	11.7	11.7	1.911	T. kodakaraei	0.52	-0.37
C	0.53	-0.6348783	24.7	741	1.138			-0.36
D	4.73	1.5539252	12.7	114.3	1.138			
E	8.82	2.17702187	15.3	76.5	1.138			
F	4.32	1.4632554	52	208	1.058			
G	7.58	2.0255132	11.7	11.7	1.911			
H	1.59	0.46373402	38.3	536.2	1.138			
I	6.96	1.94017947	32.3	64.6	1.45			
K	6.86	1.92570744	30.3	242.4	1.058			
L	10.4	2.34180581	27.3	54.6	2.21			
M	2.33	0.84586827	34.3	445.9	0.404			

-0.74

-0.87

0.67

-0.73

-0.84

0.57

N	3.21	1.16627094	14.7	147	1.058			
P	4.38	1.47704872	20.3	60.9	1.911			
Q	1.82	0.5988365	16.3	130.4	1.138			
R	5.83	1.763017	27.3	109.2	2.291			
S	5.04	1.61740608	11.7	70.2	2.236			
T	4.61	1.52822786	18.7	112.2	1.831			
V	8.28	2.11384297	23.3	46.6	1.831			
W	1.29	0.25464222	74.3	891.6	0.484			
Y	3.83	1.3428648	50	350	1.058			
A	8.2	2.10413415	11.7	11.7	1.292	H. influenzae	0.38	-0.49
C	1.03	2.96E-02	24.7	741	1.082			-0.57
D	4.98	1.60542989	12.7	114.3	1.082			
E	6.49	1.87026253	15.3	76.5	1.082			
F	4.46	1.49514877	52	208	1.565			
G	6.64	1.89311196	11.7	11.7	1.292			
H	2.05	0.71783979	38.3	536.2	1.082			
I	7.08	1.95727391	32.3	64.6	2.046			
K	6.33	1.84530024	30.3	242.4	1.565			
L	10.5	2.35137526	27.3	54.6	2.368			
M	2.4	0.87546874	34.3	445.9	0.601			
N	4.89	1.5871923	14.7	147	1.565			
P	3.72	1.31372367	20.3	60.9	1.292			
Q	4.63	1.53255687	16.3	130.4	1.082			
R	4.5	1.5040774	27.3	109.2	1.885			
S	5.84	1.7647308	11.7	70.2	2.18			
T	5.2	1.64865863	18.7	112.2	1.775			
V	6.7	1.90210753	23.3	46.6	1.775			
W	1.12	0.11332869	74.3	891.6	0.118			
Y	3.14	1.1442228	50	350	1.565			
A	5.57	1.71739505	11.7	11.7	0.921	M. genitalium	0.32	-0.33
C	0.82	-0.1984509	24.7	741	0.995			-0.45
D	4.92	1.59330853	12.7	114.3	0.995			

-0.8

-0.93

0.69

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-0.84

0.73

E	5.68	1.73695123	15.3	76.5	0.995		
F	6.09	1.80664808	52	208	1.764		
G	4.62	1.53039471	11.7	11.7	0.921		
H	1.58	0.45742485	38.3	536.2	0.995		
I	8.24	2.10900034	32.3	64.6	2.284		
K	9.5	2.2512918	30.3	242.4	1.764		
L	10.6	2.360854	27.3	54.6	2.42		
M	1.52	0.41871034	34.3	445.9	0.614		
N	7.5	2.01490302	14.7	147	1.764		
P	2.99	1.09527339	20.3	60.9	0.921		
Q	4.73	1.5539252	16.3	130.4	0.995		
R	3.1	1.13140211	27.3	109.2	1.652		
S	6.64	1.89311196	11.7	70.2	2.094		
T	5.4	1.68639895	18.7	112.2	1.689		
V	6.14	1.81482474	23.3	46.6	1.689		
W	0.96	-4.08E-02	74.3	891.6	-0.154		
Y	3.23	1.17248214	50	350	1.764		
A	6.67	1.89761986	11.7	11.7	1.387	M. pneumoniae	
C	0.75	-0.2876821	24.7	741	1.099		0.4
D	4.96	1.60140574	12.7	114.3	1.099		-0.42
E	5.67	1.73518912	15.3	76.5	1.099		-0.47
F	5.59	1.72097929	52	208	1.504		
G	5.52	1.70837786	11.7	11.7	1.387		
H	1.8	0.58778667	38.3	536.2	1.099		
I	6.6	1.88706965	32.3	64.6	1.974		
K	8.56	2.14710019	30.3	242.4	1.504		
L	10.3	2.3321439	27.3	54.6	2.351		
M	1.57	0.45107562	34.3	445.9	0.588		
N	6.22	1.82776991	14.7	147	1.504		
P	3.5	1.25276297	20.3	60.9	1.387		
Q	5.36	1.67896398	16.3	130.4	1.099		
R	3.48	1.24703229	27.3	109.2	1.946		
S	6.48	1.86872051	11.7	70.2	2.197		

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-0.88

0.69

T	5.96	1.78507048	18.7	112.2	1.792			
V	6.47	1.86717611	23.3	46.6	1.792			
W	1.19	0.17395331	74.3	891.6	0.183			
Y	3.22	1.16938136	50	350	1.504			
A	8.48	2.13771045	11.7	11.7	1.725	Synechocysti	0.47	-0.51
C	1	0	24.7	741	1.137			-0.55
D	5.02	1.61342993	12.7	114.3	1.137			
E	6.04	1.79840401	15.3	76.5	1.137			
F	4.01	1.38879124	52	208	1.242			
G	7.37	1.99741771	11.7	11.7	1.725			
H	1.86	0.62057649	38.3	536.2	1.137			
I	6.28	1.83736998	32.3	64.6	1.665			
K	4.18	1.43031125	30.3	242.4	1.242			
L	11.4	2.43361336	27.3	54.6	2.272			
M	2.01	0.69813472	34.3	445.9	0.495			
N	4.04	1.39624469	14.7	147	1.242			
P	5.14	1.63705308	20.3	60.9	1.725			
Q	5.55	1.71379793	16.3	130.4	1.137			
R	5.05	1.61938824	27.3	109.2	2.166			
S	5.81	1.75958057	11.7	70.2	2.235			
T	5.5	1.70474809	18.7	112.2	1.83			
V	6.69	1.90061387	23.3	46.6	1.83			
W	1.55	0.43825493	74.3	891.6	0.389			
Y	2.91	1.06815308	50	350	1.242			
A	9.48	2.24918432	11.7	11.7	1.853	E. coli	0.51	-0.52
C	1.17	0.15700375	24.7	741	1.139			-0.58
D	5.13	1.63510566	12.7	114.3	1.139			
E	5.74	1.74745921	15.3	76.5	1.139			
F	3.89	1.35840916	52	208	1.119			
G	7.36	1.99605993	11.7	11.7	1.853			
H	2.27	0.81977983	38.3	536.2	1.139			
I	6	1.79175947	32.3	64.6	1.521			

-0.8

-0.93

0.72

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-0.93

0.71

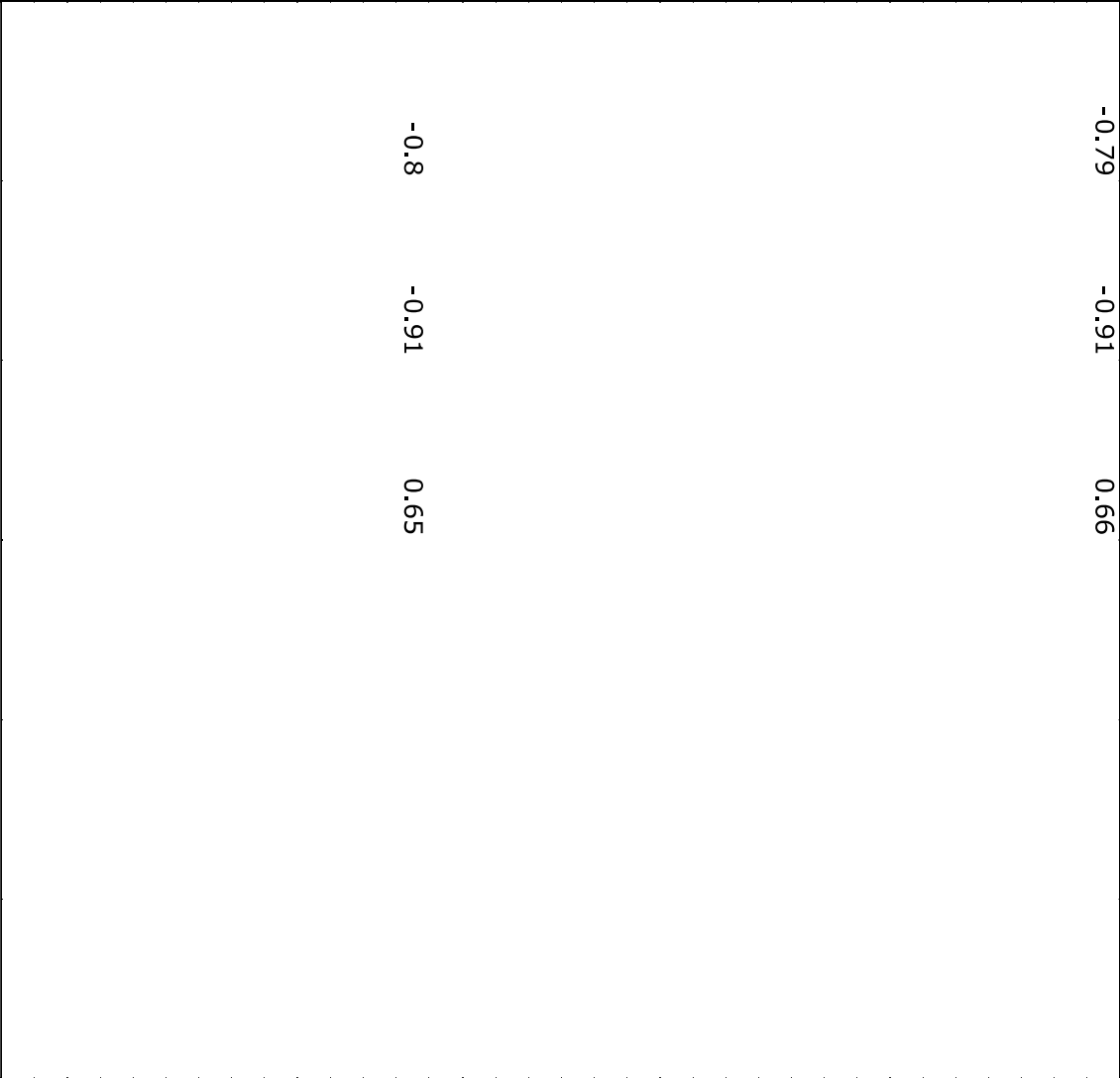
K	4.4	1.48160454	30.3	242.4	1.119			
L	10.6	2.360854	27.3	54.6	2.231			
M	2.85	1.04731899	34.3	445.9	0.436			
N	3.95	1.37371558	14.7	147	1.119			
P	4.42	1.4861397	20.3	60.9	1.853			
Q	4.43	1.48839958	16.3	130.4	1.139			
R	5.53	1.71018782	27.3	109.2	2.252			
S	5.82	1.76130026	11.7	70.2	2.238			
T	5.4	1.68639895	18.7	112.2	1.832			
V	7.05	1.95302762	23.3	46.6	1.832			
W	1.52	0.41871034	74.3	891.6	0.457			
Y	2.84	1.04380405	50	350	1.119			
A	6.83	1.92132467	11.7	11.7	1.329	H. pylori		
C	1.09	8.62E-02	24.7	741	1.089		0.39	
D	4.77	1.56234631	12.7	114.3	1.089			-0.54
E	6.86	1.92570744	15.3	76.5	1.089			
F	5.41	1.68824909	52	208	1.541			
G	5.76	1.75093748	11.7	11.7	1.329			
H	2.12	0.75141609	38.3	536.2	1.089			
I	7.19	1.97269117	32.3	64.6	2.018			
K	8.94	2.19053559	30.3	242.4	1.541			
L	11.2	2.41591378	27.3	54.6	2.362			
M	2.22	0.7975072	34.3	445.9	0.596			
N	5.85	1.76644166	14.7	147	1.541			
P	3.28	1.18784342	20.3	60.9	1.329			
Q	3.72	1.31372367	16.3	130.4	1.089			
R	3.45	1.23837423	27.3	109.2	1.909			
S	6.82	1.91985947	11.7	70.2	2.187			
T	4.39	1.47932923	18.7	112.2	1.782			
V	5.61	1.72455072	23.3	46.6	1.782			
W	0.7	-0.3566749	74.3	891.6	0.144			
Y	3.69	1.30562646	50	350	1.541			

-0.69

-0.88

0.74

A	7.67	2.03731662	11.7	11.7	1.555	B. subtilis	0.44	-0.47	-0.52
C	0.8	-0.2231436	24.7	741	1.122				
D	5.17	1.64287269	12.7	114.3	1.122				
E	7.23	1.97823904	15.3	76.5	1.122				
F	4.49	1.5018527	52	208	1.384				
G	6.9	1.93152141	11.7	11.7	1.555				
H	2.27	0.81977983	38.3	536.2	1.122				
I	7.36	1.99605993	32.3	64.6	1.831				
K	7.05	1.95302762	30.3	242.4	1.384				
L	9.64	2.26592111	27.3	54.6	2.316				
M	2.78	1.02245093	34.3	445.9	0.551				
N	3.94	1.37118072	14.7	147	1.384				
P	3.68	1.30291275	20.3	60.9	1.555				
Q	3.83	1.3428648	16.3	130.4	1.122				
R	4.12	1.41585316	27.3	109.2	2.055				
S	6.29	1.83896107	11.7	70.2	2.221				
T	5.42	1.69009582	18.7	112.2	1.816				
V	6.74	1.90805993	23.3	46.6	1.816				
W	1.03	2.96E-02	74.3	891.6	0.29				
Y	3.48	1.24703229	50	350	1.384				
A	7.38	1.99877364	11.7	11.7	1.563	B. halodurans	0.44	-0.47	-0.5
C	0.73	-0.3147107	24.7	741	1.123				
D	5.07	1.62334082	12.7	114.3	1.123				
E	7.84	2.05923883	15.3	76.5	1.123				
F	4.45	1.4929041	52	208	1.377				
G	7.01	1.9473377	11.7	11.7	1.563				
H	2.41	0.87962675	38.3	536.2	1.123				
I	6.93	1.93585981	32.3	64.6	1.824				
K	5.86	1.7681496	30.3	242.4	1.377				
L	9.98	2.30058309	27.3	54.6	2.314				
M	2.76	1.01523068	34.3	445.9	0.549				
N	3.62	1.28647403	14.7	147	1.377				
P	3.81	1.33762919	20.3	60.9	1.563				



Q	4.11	1.41342303	16.3	130.4	1.123			
R	4.84	1.57691472	27.3	109.2	2.06			
S	5.61	1.72455072	11.7	70.2	2.222			
T	5.55	1.71379793	18.7	112.2	1.817			
V	7.44	2.00687085	23.3	46.6	1.817			
W	1.13	0.12221763	74.3	891.6	0.295			
Y	3.38	1.21787571	50	350	1.377			
A	4.49	1.5018527	11.7	11.7	0.685	B. burgdoferi	0.28	-0.24
C	0.66	-0.4155154	24.7	741	0.928			-0.43
D	5.18	1.64480506	12.7	114.3	0.928			
E	6.77	1.91250109	15.3	76.5	0.928			
F	6.31	1.84213568	52	208	1.864			
G	5.2	1.64865863	11.7	11.7	0.685			
H	1.22	0.19885086	38.3	536.2	0.928			
I	10.7	2.37024374	32.3	64.6	2.405			
K	10.2	2.32238772	30.3	242.4	1.864			
L	10.3	2.3321439	27.3	54.6	2.443			
M	1.9	0.64185389	34.3	445.9	0.597			
N	7.27	1.98375629	14.7	147	1.864			
P	2.52	0.9242589	20.3	60.9	0.685			
Q	2.26	0.81536481	16.3	130.4	0.928			
R	3.22	1.16938136	27.3	109.2	1.507			
S	7.47	2.010895	11.7	70.2	2.027			
T	3.94	1.37118072	18.7	112.2	1.621			
V	5.35	1.67709656	23.3	46.6	1.621			
W	0.5	-0.6931472	74.3	891.6	-0.339			
Y	4.23	1.44220199	50	350	1.864			
A	5.89	1.773256	11.7	11.7	1.545	A. aeolicus	0.43	-0.29
C	0.79	-0.2357223	24.7	741	1.121			-0.39
D	4.31	1.4609379	12.7	114.3	1.121			
E	9.63	2.26488323	15.3	76.5	1.121			
F	5.14	1.63705308	52	208	1.391			

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G	6.75	1.90954251	11.7	11.7	1.545		
H	1.54	0.43178242	38.3	536.2	1.121		
I	7.33	1.99197552	32.3	64.6	1.84		
K	9.4	2.24070969	30.3	242.4	1.391		
L	10.5	2.35137526	27.3	54.6	2.318		
M	1.93	0.65752	34.3	445.9	0.554		
N	3.59	1.2781522	14.7	147	1.391		
P	4.07	1.403643	20.3	60.9	1.545		
Q	2.04	0.71294981	16.3	130.4	1.121		
R	4.92	1.59330853	27.3	109.2	2.048		
S	4.79	1.56653041	11.7	70.2	2.22		
T	4.21	1.43746265	18.7	112.2	1.814		
V	7.93	2.07065304	23.3	46.6	1.814		
W	0.93	-7.26E-02	74.3	891.6	0.284		
Y	4.13	1.41827741	50	350	1.391		
A	13.2	2.58021683	11.7	11.7	2.376	M. tuberculosis	-0.55
C	0.88	-0.1278334	24.7	741	1.037		
D	5.8	1.75785792	12.7	114.3	1.037		
E	4.68	1.54329811	15.3	76.5	1.037		
F	2.95	1.08180517	52	208	0.391		
G	9.98	2.30058309	11.7	11.7	2.376		
H	2.23	0.80200159	38.3	536.2	1.037		
I	4.26	1.44926916	32.3	64.6	0.686		
K	2.03	0.70803579	30.3	242.4	0.391		
L	9.75	2.27726729	27.3	54.6	1.963		
M	1.84	0.60976557	34.3	445.9	-3.10E-02		
N	2.53	0.9282193	14.7	147	0.391		
P	5.8	1.75785792	20.3	60.9	2.376		
Q	3.09	1.12817109	16.3	130.4	1.037		
R	7.32	1.99061033	27.3	109.2	2.609		
S	5.48	1.7011051	11.7	70.2	2.135		
T	5.92	1.77833645	18.7	112.2	1.73		
V	8.56	2.14710019	23.3	46.6	1.73		

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-0.83

0.78

W	1.46	0.37843644	74.3	891.6	0.615			
Y	2.08	0.73236789	50	350	0.391			
A	13.2	2.58021683	11.7	11.7	2.376	M. tuberculosis	0.66	-0.51
C	0.93	-7.26E-02	24.7	741	1.037			-0.56
D	5.79	1.75613229	12.7	114.3	1.037			
E	4.66	1.53901545	15.3	76.5	1.037			
F	2.91	1.06815308	52	208	0.391			
G	9.77	2.27931647	11.7	11.7	2.376			
H	2.26	0.81536481	38.3	536.2	1.037			
I	4.21	1.43746265	32.3	64.6	0.686			
K	2.05	0.71783979	30.3	242.4	0.391			
L	9.7	2.27212589	27.3	54.6	1.963			
M	1.99	0.68813464	34.3	445.9	-3.10E-02			
N	2.47	0.90421815	14.7	147	0.391			
P	5.88	1.77155676	20.3	60.9	2.376			
Q	3.12	1.137833	16.3	130.4	1.037			
R	7.51	2.01623547	27.3	109.2	2.609			
S	5.56	1.71559811	11.7	70.2	2.135			
T	5.91	1.77664583	18.7	112.2	1.73			
V	8.46	2.13534917	23.3	46.6	1.73			
W	1.48	0.39204209	74.3	891.6	0.615			
Y	2.06	0.72270598	50	350	0.391			
A	11.7	2.45958884	11.7	11.7	2.123	M. leprae	0.58	-0.53
C	0.94	-6.19E-02	24.7	741	1.115			-0.58
D	5.89	1.773256	12.7	114.3	1.115			
E	5.08	1.62531126	15.3	76.5	1.115			
F	2.97	1.08856195	52	208	0.8			
G	8.42	2.13060983	11.7	11.7	2.123			
H	2.22	0.7975072	38.3	536.2	1.115			
I	4.83	1.57484647	32.3	64.6	1.152			
K	2.67	0.98207847	30.3	242.4	0.8			
L	10	2.30258509	27.3	54.6	2.119			

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M	1.97	0.67803354	34.3	445.9	0.252			
N	2.65	0.97455964	14.7	147	0.8			
P	5.36	1.67896398	20.3	60.9	2.123			
Q	3.28	1.18784342	16.3	130.4	1.115			
R	6.94	1.93730178	27.3	109.2	2.434			
S	5.96	1.78507048	11.7	70.2	2.213			
T	6.09	1.80664808	18.7	112.2	1.808			
V	9.2	2.21920348	23.3	46.6	1.808			
W	1.38	0.3220835	74.3	891.6	0.567			
Y	2.24	0.80647587	50	350	0.8			
A	10.1	2.31253542	11.7	11.7	1.94	T. pallidum	0.53	-0.49
C	1.91	0.64710324	24.7	741	1.136			-0.61
D	4.52	1.50851199	12.7	114.3	1.136			
E	5.97	1.78674693	15.3	76.5	1.136			
F	4.45	1.4929041	52	208	1.025			
G	6.96	1.94017947	11.7	11.7	1.94			
H	2.75	1.01160091	38.3	536.2	1.136			
I	4.9	1.58923521	32.3	64.6	1.412			
K	3.97	1.3787661	30.3	242.4	1.025			
L	10.1	2.31253542	27.3	54.6	2.199			
M	2.09	0.73716407	34.3	445.9	0.386			
N	2.48	0.90825856	14.7	147	1.025			
P	4.2	1.43508453	20.3	60.9	1.94			
Q	3.84	1.34547237	16.3	130.4	1.136			
R	7.43	2.00552586	27.3	109.2	2.31			
S	6.62	1.89009537	11.7	70.2	2.235			
T	5.3	1.66770682	18.7	112.2	1.83			
V	8.25	2.1102132	23.3	46.6	1.83			
W	0.97	-3.05E-02	74.3	891.6	0.497			
Y	3.03	1.10856262	50	350	1.025			
A	7.52	2.01756614	11.7	11.7	1.452	C. trachomati	0.41	-0.48
C	1.62	0.48242615	24.7	741	1.109			-0.61

-0.73

-0.87

0.82

-0.77

-0.92

0.81

D	4.52	1.50851199	12.7	114.3	1.109		
E	6.59	1.88555335	15.3	76.5	1.109		
F	4.83	1.57484647	52	208	1.459		
G	6.34	1.84687877	11.7	11.7	1.452		
H	2.3	0.83290912	38.3	536.2	1.109		
I	6.61	1.88858365	32.3	64.6	1.921		
K	5.76	1.75093748	30.3	242.4	1.459		
L	11.2	2.41591378	27.3	54.6	2.338		
M	2.05	0.71783979	34.3	445.9	0.576		
N	3.5	1.25276297	14.7	147	1.459		
P	4.37	1.47476301	20.3	60.9	1.452		
Q	4.18	1.43031125	16.3	130.4	1.109		
R	4.84	1.57691472	27.3	109.2	1.988		
S	8.11	2.09309787	11.7	70.2	2.208		
T	5.11	1.6311994	18.7	112.2	1.802		
V	6.42	1.85941812	23.3	46.6	1.802		
W	0.95	-5.13E-02	74.3	891.6	0.225		
Y	3.07	1.12167756	50	350	1.459		
A	6.04	1.79840401	11.7	11.7	0.743	R. prowadz	kii
C	1.09	8.62E-02	24.7	741	0.945		
D	4.83	1.57484647	12.7	114.3	0.945		
E	5.77	1.75267208	15.3	76.5	0.945		
F	4.88	1.58514522	52	208	1.841		
G	5.41	1.68824909	11.7	11.7	0.743		
H	1.91	0.64710324	38.3	536.2	0.945		
I	10.8	2.37954613	32.3	64.6	2.377		
K	8.38	2.12584791	30.3	242.4	1.841		
L	10.1	2.31253542	27.3	54.6	2.438		
M	2.14	0.76080583	34.3	445.9	0.603		
N	6.64	1.89311196	14.7	147	1.841		
P	3.14	1.1442228	20.3	60.9	0.743		
Q	3.14	1.1442228	16.3	130.4	0.945		
R	3.39	1.22082992	27.3	109.2	1.542		

-0.67

-0.86

0.76

S	6.75	1.90954251	11.7	70.2	2.044			
T	5.21	1.65057986	18.7	112.2	1.639			
V	5.6	1.7227666	23.3	46.6	1.639			
W	0.71	-0.3424903	74.3	891.6	-0.293			
Y	3.89	1.35840916	50	350	1.841			
A	6.77	1.91250109	11.7	11.7	0.847	C. jejuni	0.31	-0.33
C	1.22	0.19885086	24.7	741	0.975			-0.51
D	5.24	1.6563215	12.7	114.3	0.975			
E	6.98	1.94304892	15.3	76.5	0.975			
F	6.03	1.79674701	52	208	1.797			
G	5.6	1.7227666	11.7	11.7	0.847			
H	1.64	0.49469624	38.3	536.2	0.975			
I	8.68	2.16102153	32.3	64.6	2.324			
K	9.49	2.25023861	30.3	242.4	1.797			
L	10.8	2.37954613	27.3	54.6	2.428			
M	2.2	0.78845736	34.3	445.9	0.611			
N	6.3	1.84054963	14.7	147	1.797			
P	2.67	0.98207847	20.3	60.9	0.847			
Q	3.12	1.137833	16.3	130.4	0.975			
R	2.98	1.0919233	27.3	109.2	1.607			
S	6.45	1.86408013	11.7	70.2	2.074			
T	4.06	1.40118297	18.7	112.2	1.669			
V	5.28	1.6639261	23.3	46.6	1.669			
W	0.65	-0.4307829	74.3	891.6	-0.21			
Y	3.69	1.30562646	50	350	1.797			
A	6.98	1.94304892	11.7	11.7	1.415	C. pneumonia	0.41	-0.46
C	1.59	0.46373402	24.7	741	1.103			-0.59
D	4.51	1.50629715	12.7	114.3	1.103			
E	6.61	1.88858365	15.3	76.5	1.103			
F	4.74	1.55603714	52	208	1.485			
G	6.23	1.82937633	11.7	11.7	1.415			
H	2.38	0.86710049	38.3	536.2	1.103			

-0.64

-0.84

0.73

-0.76

-0.91

0.81

I	6.92	1.93441577	32.3	64.6	1.951		
K	6.14	1.81482474	30.3	242.4	1.485		
L	11.3	2.42480273	27.3	54.6	2.346		
M	1.93	0.65752	34.3	445.9	0.583		
N	3.81	1.33762919	14.7	147	1.485		
P	4.48	1.49962305	20.3	60.9	1.415		
Q	4.03	1.39376638	16.3	130.4	1.103		
R	4.56	1.51732262	27.3	109.2	1.964		
S	8.02	2.08193842	11.7	70.2	2.202		
T	5.27	1.66203036	18.7	112.2	1.796		
V	6.12	1.8115621	23.3	46.6	1.796		
W	1.01	9.95E-03	74.3	891.6	0.201		
Y	3.26	1.18172719	50	350	1.485		
A	5.85	1.76644166	11.7	11.7	1.677	T. maritima	
C	0.7	-0.3566749	24.7	741	1.134		0.46
D	4.96	1.60140574	12.7	114.3	1.134		-0.35
E	8.92	2.18829595	15.3	76.5	1.134		-0.4
F	5.19	1.6467337	52	208	1.284		
G	6.9	1.93152141	11.7	11.7	1.677		
H	1.58	0.45742485	38.3	536.2	1.134		
I	7.18	1.97129938	32.3	64.6	1.714		
K	7.61	2.02946317	30.3	242.4	1.284		
L	10	2.30258509	27.3	54.6	2.285		
M	2.4	0.87546874	34.3	445.9	0.513		
N	3.6	1.28093385	14.7	147	1.284		
P	3.99	1.38379123	20.3	60.9	1.677		
Q	2.01	0.69813472	16.3	130.4	1.134		
R	5.53	1.71018782	27.3	109.2	2.135		
S	5.65	1.73165555	11.7	70.2	2.232		
T	4.52	1.50851199	18.7	112.2	1.827		
V	8.6	2.1517622	23.3	46.6	1.827		
W	1.1	9.53E-02	74.3	891.6	0.363		
Y	3.57	1.2725656	50	350	1.284		

-0.72

-0.86

0.64

A	12.1	2.49320545	11.7	11.7	2.406	D. radioduran	0.67	-0.5	-0.53
C	0.67	-0.4004776	24.7	741	1.022				
D	5.05	1.61938824	12.7	114.3	1.022				
E	5.72	1.74396881	15.3	76.5	1.022				
F	3.15	1.14740245	52	208	0.332				
G	9.19	2.21811594	11.7	11.7	2.406				
H	2.09	0.73716407	38.3	536.2	1.022				
I	3.27	1.18478999	32.3	64.6	0.62				
K	2.73	1.00430161	30.3	242.4	0.332				
L	11.6	2.4510051	27.3	54.6	1.939				
M	1.89	0.63657683	34.3	445.9	-7.40E-02				
N	2.41	0.87962675	14.7	147	0.332				
P	6.06	1.8017098	20.3	60.9	2.406				
Q	4.12	1.41585316	16.3	130.4	1.022				
R	7.38	1.99877364	27.3	109.2	2.63				
S	5.2	1.64865863	11.7	70.2	2.121				
T	5.81	1.75958057	18.7	112.2	1.716				
V	7.68	2.03861955	23.3	46.6	1.716				
W	1.38	0.3220835	74.3	891.6	0.616				
Y	2.3	0.83290912	50	350	0.332				
A	10.1	2.31253542	11.7	11.7	1.899	N. meningiditi	0.52	-0.54	-0.59
C	1.05	4.88E-02	24.7	741	1.138				
D	5.27	1.66203036	12.7	114.3	1.138				
E	6.15	1.81645208	15.3	76.5	1.138				
F	4.1	1.41098697	52	208	1.07				
G	7.76	2.04898233	11.7	11.7	1.899				
H	2.21	0.79299252	38.3	536.2	1.138				
I	5.86	1.7681496	32.3	64.6	1.464				
K	5.65	1.73165555	30.3	242.4	1.07				
L	9.83	2.28543893	27.3	54.6	2.215				
M	2.44	0.89199804	34.3	445.9	0.411				
N	4.11	1.41342303	14.7	147	1.07				

-0.69

-0.86

0.73

-0.8

-0.94

0.68

P	4.17	1.42791604	20.3	60.9	1.899			
Q	4.01	1.38879124	16.3	130.4	1.138			
R	5.49	1.70292826	27.3	109.2	2.283			
S	5.55	1.71379793	11.7	70.2	2.237			
T	5.24	1.6563215	18.7	112.2	1.831			
V	6.76	1.91102289	23.3	46.6	1.831			
W	1.17	0.15700375	74.3	891.6	0.479			
Y	2.97	1.08856195	50	350	1.07			
A	10.2	2.32238772	11.7	11.7	1.935	X. fastidiosa	0.53	-0.52
C	1.19	0.17395331	24.7	741	1.137			-0.59
D	5.3	1.66770682	12.7	114.3	1.137			
E	5.06	1.62136648	15.3	76.5	1.137			
F	3.46	1.24126859	52	208	1.032			
G	7.58	2.0255132	11.7	11.7	1.935			
H	2.74	1.00795792	38.3	536.2	1.137			
I	5.33	1.67335124	32.3	64.6	1.42			
K	3.55	1.2669476	30.3	242.4	1.032			
L	10.7	2.37024374	27.3	54.6	2.202			
M	2.41	0.87962675	34.3	445.9	0.39			
N	3.34	1.20597081	14.7	147	1.032			
P	4.93	1.59533899	20.3	60.9	1.935			
Q	4.22	1.43983513	16.3	130.4	1.137			
R	6.7	1.90210753	27.3	109.2	2.306			
S	5.97	1.78674693	11.7	70.2	2.235			
T	5.8	1.75785792	18.7	112.2	1.83			
V	7.33	1.99197552	23.3	46.6	1.83			
W	1.46	0.37843644	74.3	891.6	0.495			
Y	2.61	0.95935022	50	350	1.032			
A	9.14	2.21266039	11.7	11.7	1.73	V. cholerae	0.47	-0.52
C	1.05	4.88E-02	24.7	741	1.137			-0.58
D	5.01	1.61143592	12.7	114.3	1.137			
E	6.19	1.82293509	15.3	76.5	1.137			

-0.75

-0.9

0.79

-0.79

-0.93

0.69

F	4.07	1.403643	52	208	1.237		
G	6.68	1.89911799	11.7	11.7	1.73		
H	2.4	0.87546874	38.3	536.2	1.137		
I	6.04	1.79840401	32.3	64.6	1.659		
K	4.93	1.59533899	30.3	242.4	1.237		
L	10.8	2.37954613	27.3	54.6	2.27		
M	2.73	1.00430161	34.3	445.9	0.493		
N	3.9	1.36097655	14.7	147	1.237		
P	4.01	1.38879124	20.3	60.9	1.73		
Q	5.17	1.64287269	16.3	130.4	1.137		
R	4.94	1.59736533	27.3	109.2	2.17		
S	6.33	1.84530024	11.7	70.2	2.236		
T	5.19	1.6467337	18.7	112.2	1.83		
V	7.01	1.9473377	23.3	46.6	1.83		
W	1.32	0.27763174	74.3	891.6	0.392		
Y	2.96	1.08518927	50	350	1.237		
A	11.6	2.4510051	11.7	11.7	2.405	P. aeruginosa	
C	1	0	24.7	741	1.023		
D	5.32	1.6714733	12.7	114.3	1.023		
E	6.08	1.8050047	15.3	76.5	1.023		
F	3.55	1.2669476	52	208	0.335		
G	8.45	2.13416644	11.7	11.7	2.405		
H	2.17	0.77472717	38.3	536.2	1.023		
I	4.17	1.42791604	32.3	64.6	0.623		
K	2.86	1.05082163	30.3	242.4	0.335		
L	12.4	2.51769647	27.3	54.6	1.94		
M	2.06	0.72270598	34.3	445.9	-7.20E-02		
N	2.64	0.97077892	14.7	147	0.335		
P	5.07	1.62334082	20.3	60.9	2.405		
Q	4.25	1.44691898	16.3	130.4	1.023		
R	7.62	2.03077637	27.3	109.2	2.629		
S	5.51	1.70656462	11.7	70.2	2.122		
T	4.16	1.42551507	18.7	112.2	1.716		

-0.68

-0.86

0.71

V	6.88	1.92861865	23.3	46.6	1.716			
W	1.48	0.39204209	74.3	891.6	0.616			
Y	2.53	0.9282193	50	350	0.335			
A	4.47	1.49738841	11.7	11.7	0.552	Buchnera sp.	0.26	-0.26
C	1.21	0.19062036	24.7	741	0.886			-0.45
D	4.32	1.4632554	12.7	114.3	0.886			
E	5.51	1.70656462	15.3	76.5	0.886			
F	5.04	1.61740608	52	208	1.914			
G	5.43	1.69193913	11.7	11.7	0.552			
H	2.11	0.74668795	38.3	536.2	0.886			
I	11.5	2.44234704	32.3	64.6	2.466			
K	9.88	2.29051251	30.3	242.4	1.914			
L	9.9	2.29253476	27.3	54.6	2.454			
M	2.17	0.77472717	34.3	445.9	0.58			
N	7.16	1.96850998	14.7	147	1.914			
P	3	1.09861229	20.3	60.9	0.552			
Q	3.2	1.16315081	16.3	130.4	0.886			
R	3.79	1.33236602	27.3	109.2	1.426			
S	7.27	1.98375629	11.7	70.2	1.985			
T	4.56	1.51732262	18.7	112.2	1.58			
V	4.85	1.57897871	23.3	46.6	1.58			
W	0.9	-0.1053605	74.3	891.6	-0.447			
Y	3.61	1.28370777	50	350	1.914			
A	7.73	2.04510886	11.7	11.7	1.283	L. monocytog	0.38	-0.47
C	0.6	-0.5108256	24.7	741	1.08			-0.5
D	5.44	1.69377906	12.7	114.3	1.08			
E	7.43	2.00552586	15.3	76.5	1.08			
F	4.53	1.51072194	52	208	1.57			
G	6.66	1.89611949	11.7	11.7	1.283			
H	1.77	0.57097955	38.3	536.2	1.08			
I	7.83	2.05796251	32.3	64.6	2.053			
K	7.16	1.96850998	30.3	242.4	1.57			

-0.58

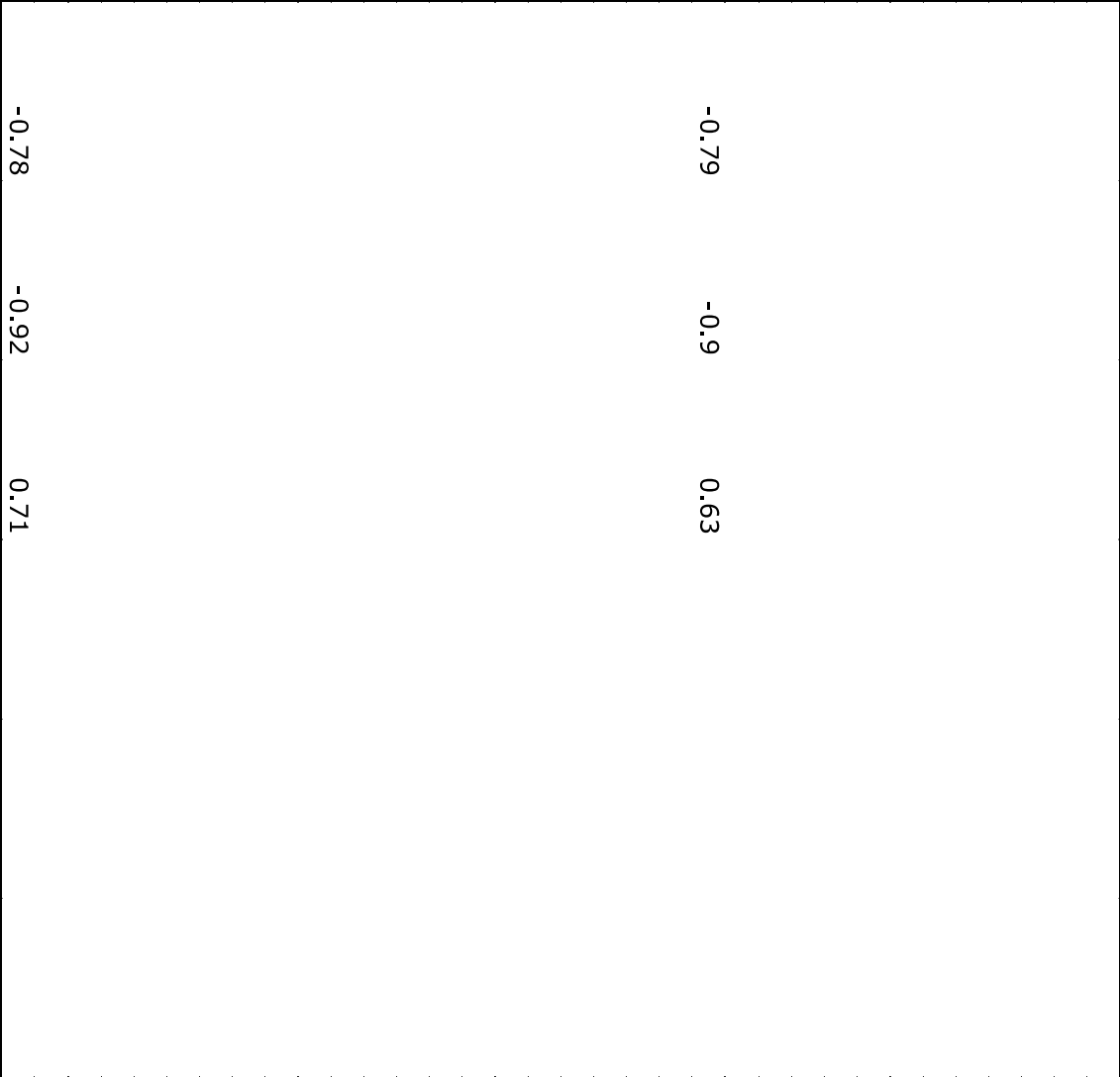
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-0.9

0.64



C	1.15	0.13976194	24.7	741	1.138		
D	5.21	1.65057986	12.7	114.3	1.138		
E	5.6	1.7227666	15.3	76.5	1.138		
F	3.85	1.34807315	52	208	1.063		
G	7.36	1.99605993	11.7	11.7	1.906		
H	2.29	0.82855182	38.3	536.2	1.138		
I	5.9	1.77495235	32.3	64.6	1.456		
K	4.32	1.4632554	30.3	242.4	1.063		
L	10.7	2.37024374	27.3	54.6	2.212		
M	2.75	1.01160091	34.3	445.9	0.407		
N	3.78	1.32972401	14.7	147	1.063		
P	4.47	1.49738841	20.3	60.9	1.906		
Q	4.37	1.47476301	16.3	130.4	1.138		
R	5.7	1.74046618	27.3	109.2	2.287		
S	5.79	1.75613229	11.7	70.2	2.237		
T	5.47	1.69927862	18.7	112.2	1.831		
V	7.06	1.95444505	23.3	46.6	1.831		
W	1.53	0.42526774	74.3	891.6	0.482		
Y	2.87	1.05431203	50	350	1.063		
A	9.19	2.21811594	11.7	11.7	1.735	Y. pestis	
C	1.04	3.92E-02	24.7	741	1.137		0.48
D	5.11	1.6311994	12.7	114.3	1.137		-0.53
E	5.48	1.7011051	15.3	76.5	1.137		-0.58
F	3.82	1.34025042	52	208	1.233		
G	7.22	1.97685495	11.7	11.7	1.735		
H	2.28	0.82417544	38.3	536.2	1.137		
I	6.21	1.8261609	32.3	64.6	1.654		
K	4.38	1.47704872	30.3	242.4	1.233		
L	10.9	2.38876279	27.3	54.6	2.269		
M	2.63	0.96698385	34.3	445.9	0.491		
N	4.05	1.39871688	14.7	147	1.233		
P	4.38	1.47704872	20.3	60.9	1.735		
Q	4.87	1.58309394	16.3	130.4	1.137		

-0.79

-0.94

0.73

R	5.41	1.68824909	27.3	109.2	2.173			
S	6.24	1.83098018	11.7	70.2	2.236			
T	5.4	1.68639895	18.7	112.2	1.83			
V	6.89	1.93007109	23.3	46.6	1.83			
W	1.36	0.3074847	74.3	891.6	0.395			
Y	3	1.09861229	50	350	1.233			
A	6.39	1.85473427	11.7	11.7	0.992	S. aureus Mu	0.33	-0.46
C	0.63	-0.4620355	24.7	741	1.014			-0.52
D	5.83	1.763017	12.7	114.3	1.014			
E	6.52	1.87487438	15.3	76.5	1.014			
F	4.48	1.49962305	52	208	1.73			
G	6.01	1.79342475	11.7	11.7	0.992			
H	2.31	0.83724753	38.3	536.2	1.014			
I	8.58	2.14943391	32.3	64.6	2.243			
K	7.56	2.02287119	30.3	242.4	1.73			
L	9.11	2.20937271	27.3	54.6	2.412			
M	2.64	0.97077892	34.3	445.9	0.616			
N	5.68	1.73695123	14.7	147	1.73			
P	3.2	1.16315081	20.3	60.9	0.992			
Q	4.16	1.42551507	16.3	130.4	1.014			
R	3.52	1.25846099	27.3	109.2	1.696			
S	6.14	1.81482474	11.7	70.2	2.113			
T	5.8	1.75785792	18.7	112.2	1.707			
V	6.68	1.89911799	23.3	46.6	1.707			
W	0.74	-0.3011051	74.3	891.6	-9.90E-02			
Y	3.92	1.36609165	50	350	1.73			
A	6.43	1.86097454	11.7	11.7	0.99	S. aureus N3	0.33	-0.46
C	0.63	-0.4620355	24.7	741	1.014			-0.52
D	5.82	1.76130026	12.7	114.3	1.014			
E	6.49	1.87026253	15.3	76.5	1.014			
F	4.48	1.49962305	52	208	1.73			
G	6.04	1.79840401	11.7	11.7	0.99			

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H	2.32	0.84156719	38.3	536.2	1.014		
I	8.6	2.1517622	32.3	64.6	2.244		
K	7.53	2.01889504	30.3	242.4	1.73		
L	9.13	2.2115657	27.3	54.6	2.412		
M	2.64	0.97077892	34.3	445.9	0.616		
N	5.65	1.73165555	14.7	147	1.73		
P	3.22	1.16938136	20.3	60.9	0.99		
Q	4.16	1.42551507	16.3	130.4	1.014		
R	3.5	1.25276297	27.3	109.2	1.695		
S	6.13	1.81319475	11.7	70.2	2.112		
T	5.79	1.75613229	18.7	112.2	1.707		
V	6.71	1.90359895	23.3	46.6	1.707		
W	0.74	-0.3011051	74.3	891.6	-0.101		
Y	3.89	1.35840916	50	350	1.73		
A	7.81	2.05540496	11.7	11.7	1.31	S. pyogenes	
C	0.63	-0.4620355	24.7	741	1.085		0.39
D	5.78	1.75440368	12.7	114.3	1.085		-0.48
E	6.38	1.8531681	15.3	76.5	1.085		-0.52
F	4.4	1.48160454	52	208	1.553		
G	6.45	1.86408013	11.7	11.7	1.31		
H	2.05	0.71783979	38.3	536.2	1.085		
I	7.4	2.00148	32.3	64.6	2.032		
K	7.04	1.95160817	30.3	242.4	1.553		
L	10.1	2.31253542	27.3	54.6	2.365		
M	2.56	0.94000726	34.3	445.9	0.599		
N	4.36	1.47247206	14.7	147	1.553		
P	3.35	1.20896035	20.3	60.9	1.31		
Q	4.3	1.45861502	16.3	130.4	1.085		
R	3.96	1.37624403	27.3	109.2	1.897		
S	6.18	1.82131827	11.7	70.2	2.184		
T	5.87	1.76985463	18.7	112.2	1.778		
V	6.7	1.90210753	23.3	46.6	1.778		
W	0.84	-0.1743534	74.3	891.6	0.131		

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Y	3.67	1.30019166	50	350	1.553			
A	12.3	2.50959926	11.7	11.7	2.279	M. loti	-0.51	-0.54
C	0.86	-0.1508229	24.7	741	1.075		0.63	
D	5.67	1.73518912	12.7	114.3	1.075			
E	5.37	1.68082791	15.3	76.5	1.075			
F	3.84	1.34547237	52	208	0.564			
G	8.61	2.15292432	11.7	11.7	2.279			
H	2.06	0.72270598	38.3	536.2	1.075			
I	5.38	1.68268837	32.3	64.6	0.882			
K	3.69	1.30562646	30.3	242.4	0.564			
L	9.84	2.28645571	27.3	54.6	2.03			
M	2.49	0.91228271	34.3	445.9	9.40E-02			
N	2.71	0.99694864	14.7	147	0.564			
P	5.06	1.62136648	20.3	60.9	2.279			
Q	3.11	1.13462273	16.3	130.4	1.075			
R	6.98	1.94304892	27.3	109.2	2.541			
S	5.69	1.73871025	11.7	70.2	2.173			
T	5.3	1.66770682	18.7	112.2	1.768			
V	7.34	1.99333884	23.3	46.6	1.768			
W	1.36	0.3074847	74.3	891.6	0.605			
Y	2.21	0.79299252	50	350	0.564			
A	12.1	2.49320545	11.7	11.7	2.268	S. meliloti	0.62	-0.54
C	0.86	-0.1508229	24.7	741	1.078			
D	5.48	1.7011051	12.7	114.3	1.078			
E	6.03	1.79674701	15.3	76.5	1.078			
F	3.91	1.36353737	52	208	0.582			
G	8.52	2.14241634	11.7	11.7	2.268			
H	2.05	0.71783979	38.3	536.2	1.078			
I	5.47	1.69927862	32.3	64.6	0.902			
K	3.43	1.23256026	30.3	242.4	0.582			
L	10	2.30258509	27.3	54.6	2.037			
M	2.51	0.92028275	34.3	445.9	0.106			

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N	2.67	0.98207847	14.7	147	0.582			
P	4.95	1.59938758	20.3	60.9	2.268			
Q	2.9	1.06471074	16.3	130.4	1.078			
R	7.19	1.97269117	27.3	109.2	2.534			
S	5.66	1.73342389	11.7	70.2	2.177			
T	5.15	1.63899672	18.7	112.2	1.772			
V	7.44	2.00687085	23.3	46.6	1.772			
W	1.3	0.26236426	74.3	891.6	0.603			
Y	2.27	0.81977983	50	350	0.582			
A	11.5	2.44234704	11.7	11.7	2.165	A. tumefaciar	0.59	-0.52
C	0.83	-0.1863296	24.7	741	1.106			-0.55
D	5.54	1.7119945	12.7	114.3	1.106			
E	5.79	1.75613229	15.3	76.5	1.106			
F	4.05	1.39871688	52	208	0.741			
G	8.25	2.1102132	11.7	11.7	2.165			
H	2.02	0.70309751	38.3	536.2	1.106			
I	5.75	1.74919986	32.3	64.6	1.084			
K	3.9	1.36097655	30.3	242.4	0.741			
L	9.86	2.28848617	27.3	54.6	2.097			
M	2.68	0.9858168	34.3	445.9	0.214			
N	2.98	1.0919233	14.7	147	0.741			
P	4.82	1.57277393	20.3	60.9	2.165			
Q	3.12	1.137833	16.3	130.4	1.106			
R	6.64	1.89311196	27.3	109.2	2.463			
S	5.98	1.78842057	11.7	70.2	2.205			
T	5.37	1.68082791	18.7	112.2	1.799			
V	7.28	1.98513086	23.3	46.6	1.799			
W	1.25	0.22314355	74.3	891.6	0.579			
Y	2.3	0.83290912	50	350	0.741			
A	13.2	2.58021683	11.7	11.7	2.377	M. bovis	0.66	-0.51
C	0.88	-0.1278334	24.7	741	1.037			-0.55
D	5.8	1.75785792	12.7	114.3	1.037			

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E	4.66	1.53901545	15.3	76.5	1.037		
F	2.95	1.08180517	52	208	0.39		
G	10	2.30258509	11.7	11.7	2.377		
H	2.23	0.80200159	38.3	536.2	1.037		
I	4.25	1.44691898	32.3	64.6	0.685		
K	2.03	0.70803579	30.3	242.4	0.39		
L	9.75	2.27726729	27.3	54.6	1.962		
M	1.84	0.60976557	34.3	445.9	-3.10E-02		
N	2.52	0.9242589	14.7	147	0.39		
P	5.81	1.75958057	20.3	60.9	2.377		
Q	3.08	1.1249296	16.3	130.4	1.037		
R	7.32	1.99061033	27.3	109.2	2.609		
S	5.48	1.7011051	11.7	70.2	2.135		
T	5.92	1.77833645	18.7	112.2	1.73		
V	8.57	2.14826773	23.3	46.6	1.73		
W	1.46	0.37843644	74.3	891.6	0.615		
Y	2.06	0.72270598	50	350	0.39		
A	13.7	2.61739583	11.7	11.7	2.562	S. coelicolor	
C	0.77	-0.2613648	24.7	741	0.924		0.72
D	6.12	1.8115621	12.7	114.3	0.924		-0.49
E	5.69	1.73871025	15.3	76.5	0.924		-0.51
F	2.65	0.97455964	52	208	-2.00E-02		
G	9.66	2.26799365	11.7	11.7	2.562		
H	2.34	0.85015093	38.3	536.2	0.924		
I	2.86	1.05082163	32.3	64.6	0.227		
K	2.05	0.71783979	30.3	242.4	-2.00E-02		
L	10.2	2.32238772	27.3	54.6	1.795		
M	1.57	0.45107562	34.3	445.9	-0.349		
N	1.69	0.52472853	14.7	147	-2.00E-02		
P	6.2	1.82454929	20.3	60.9	2.562		
Q	2.63	0.96698385	16.3	130.4	0.924		
R	8.36	2.12345843	27.3	109.2	2.74		
S	4.97	1.60341984	11.7	70.2	2.023		

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T	6.15	1.81645208	18.7	112.2	1.617			
V	8.67	2.15986879	23.3	46.6	1.617			
W	1.51	0.41210965	74.3	891.6	0.596			
Y	2.05	0.71783979	50	350	-2.00E-02			
A	4.89	1.5871923	11.7	11.7	0.486	U. urealyticus	0.26	-0.27
C	0.66	-0.4155154	24.7	741	0.865			-0.4
D	5.68	1.73695123	12.7	114.3	0.865			
E	5.75	1.74919986	15.3	76.5	0.865			
F	5.15	1.63899672	52	208	1.937			
G	4.13	1.41827741	11.7	11.7	0.486			
H	1.73	0.54812141	38.3	536.2	0.865			
I	10.3	2.3321439	32.3	64.6	2.494			
K	9.94	2.29656702	30.3	242.4	1.937			
L	10	2.30258509	27.3	54.6	2.458			
M	1.68	0.51879379	34.3	445.9	0.57			
N	9.06	2.20386912	14.7	147	1.937			
P	2.65	0.97455964	20.3	60.9	0.486			
Q	3.83	1.3428648	16.3	130.4	0.865			
R	2.77	1.01884732	27.3	109.2	1.386			
S	5.98	1.78842057	11.7	70.2	1.963			
T	5.01	1.61143592	18.7	112.2	1.558			
V	5.36	1.67896398	23.3	46.6	1.558			
W	0.87	-0.1392621	74.3	891.6	-0.502			
Y	4.44	1.49065438	50	350	1.937			
A	9.44	2.24495598	11.7	11.7	1.858	S. flexneri	0.51	-0.52
C	1.2	0.18232156	24.7	741	1.139			-0.58
D	5.03	1.61541998	12.7	114.3	1.139			
E	5.85	1.76644166	15.3	76.5	1.139			
F	3.78	1.32972401	52	208	1.113			
G	7.31	1.98924327	11.7	11.7	1.858			
H	2.33	0.84586827	38.3	536.2	1.139			
I	5.83	1.763017	32.3	64.6	1.514			

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K	4.49	1.5018527	30.3	242.4	1.113			
L	10.6	2.360854	27.3	54.6	2.229			
M	2.86	1.05082163	34.3	445.9	0.433			
N	3.81	1.33762919	14.7	147	1.113			
P	4.37	1.47476301	20.3	60.9	1.858			
Q	4.45	1.4929041	16.3	130.4	1.139			
R	5.88	1.77155676	27.3	109.2	2.255			
S	5.82	1.76130026	11.7	70.2	2.238			
T	5.34	1.67522565	18.7	112.2	1.832			
V	7.06	1.95444505	23.3	46.6	1.832			
W	1.54	0.43178242	74.3	891.6	0.459			
Y	2.85	1.04731899	50	350	1.113			
A	7.36	1.99605993	11.7	11.7	1.138	L. lactis		
C	0.45	-0.7985077	24.7	741	1.049		0.35	
D	5.28	1.6639261	12.7	114.3	1.049			-0.46
E	6.98	1.94304892	15.3	76.5	1.049			-0.45
F	4.76	1.56024767	52	208	1.654			
G	6.58	1.88403475	11.7	11.7	1.138			
H	1.79	0.58221562	38.3	536.2	1.049			
I	7.7	2.04122033	32.3	64.6	2.153			
K	7.41	2.00283044	30.3	242.4	1.654			
L	9.88	2.29051251	27.3	54.6	2.392			
M	2.52	0.9242589	34.3	445.9	0.614			
N	5.2	1.64865863	14.7	147	1.654			
P	3.22	1.16938136	20.3	60.9	1.138			
Q	3.69	1.30562646	16.3	130.4	1.049			
R	3.62	1.28647403	27.3	109.2	1.788			
S	6.62	1.89009537	11.7	70.2	2.148			
T	5.72	1.74396881	18.7	112.2	1.743			
V	6.58	1.88403475	23.3	46.6	1.743			
W	1	0	74.3	891.6	9.00E-03			
Y	3.54	1.26412673	50	350	1.654			

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A	6.45	1.86408013	11.7	11.7	0.967	R. conirlii	0.32	-0.37	-0.52
C	1.04	3.92E-02	24.7	741	1.008				
D	4.95	1.59938758	12.7	114.3	1.008				
E	6.15	1.81645208	15.3	76.5	1.008				
F	4.73	1.5539252	52	208	1.741				
G	5.46	1.69744879	11.7	11.7	0.967				
H	1.83	0.60431597	38.3	536.2	1.008				
I	9.97	2.29958058	32.3	64.6	2.258				
K	8.37	2.12465389	30.3	242.4	1.741				
L	10	2.30258509	27.3	54.6	2.415				
M	2.27	0.81977983	34.3	445.9	0.616				
N	6.43	1.86097454	14.7	147	1.741				
P	3.3	1.19392247	20.3	60.9	0.967				
Q	3.2	1.16315081	16.3	130.4	1.008				
R	3.41	1.22671229	27.3	109.2	1.681				
S	6.74	1.90805993	11.7	70.2	2.106				
T	5.28	1.6639261	18.7	112.2	1.701				
V	5.66	1.73342389	23.3	46.6	1.701				
W	0.76	-0.2744368	74.3	891.6	-0.118				
Y	3.81	1.33762919	50	350	1.741				
A	13.7	2.61739583	11.7	11.7	2.424	C. crescentus	0.67	-0.5	-0.53
C	0.74	-0.3011051	24.7	741	1.013				
D	5.79	1.75613229	12.7	114.3	1.013				
E	5.4	1.68639895	15.3	76.5	1.013				
F	3.54	1.26412673	52	208	0.296				
G	8.96	2.19277023	11.7	11.7	2.424				
H	1.8	0.58778667	38.3	536.2	1.013				
I	4.37	1.47476301	32.3	64.6	0.579				
K	3.52	1.25846099	30.3	242.4	0.296				
L	10	2.30258509	27.3	54.6	1.925				
M	2.26	0.81536481	34.3	445.9	-0.102				
N	2.36	0.85866162	14.7	147	0.296				
P	5.48	1.7011051	20.3	60.9	2.424				

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Q	3.21	1.16627094	16.3	130.4	1.013			
R	7.34	1.99333884	27.3	109.2	2.642			
S	5.1	1.62924054	11.7	70.2	2.112			
T	5.22	1.6524974	18.7	112.2	1.706			
V	7.54	2.02022218	23.3	46.6	1.706			
W	1.41	0.3435897	74.3	891.6	0.616			
Y	2.1	0.74193735	50	350	0.296			
A	8.05	2.08567209	11.7	11.7	1.449	Nostoc sp.	0.41	-0.52
C	0.93	-7.26E-02	24.7	741	1.108			-0.55
D	4.79	1.56653041	12.7	114.3	1.108			
E	6.08	1.8050047	15.3	76.5	1.108			
F	3.95	1.37371558	52	208	1.461			
G	6.69	1.90061387	11.7	11.7	1.449			
H	1.83	0.60431597	38.3	536.2	1.108			
I	6.93	1.93585981	32.3	64.6	1.923			
K	4.67	1.54115907	30.3	242.4	1.461			
L	11	2.39789527	27.3	54.6	2.339			
M	1.79	0.58221562	34.3	445.9	0.576			
N	4.56	1.51732262	14.7	147	1.461			
P	4.69	1.54543258	20.3	60.9	1.449			
Q	5.55	1.71379793	16.3	130.4	1.108			
R	5.02	1.61342993	27.3	109.2	1.986			
S	6.35	1.84845481	11.7	70.2	2.207			
T	5.83	1.763017	18.7	112.2	1.802			
V	6.67	1.89761986	23.3	46.6	1.802			
W	1.42	0.35065687	74.3	891.6	0.223			
Y	3.07	1.12167756	50	350	1.461			
A	9.75	2.27726729	11.7	11.7	1.984	T. elongatus	0.54	-0.47
C	1.13	0.12221763	24.7	741	1.133			-0.53
D	4.45	1.4929041	12.7	114.3	1.133			
E	5.85	1.76644166	15.3	76.5	1.133			
F	3.52	1.25846099	52	208	0.976			

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G	7.14	1.96571278	11.7	11.7	1.984		
H	2.2	0.78845736	38.3	536.2	1.133		
I	5.54	1.7119945	32.3	64.6	1.355		
K	3.11	1.13462273	30.3	242.4	0.976		
L	12	2.48490665	27.3	54.6	2.182		
M	1.92	0.65232519	34.3	445.9	0.358		
N	2.91	1.06815308	14.7	147	0.976		
P	5.77	1.75267208	20.3	60.9	1.984		
Q	5.78	1.75440368	16.3	130.4	1.133		
R	6.62	1.89009537	27.3	109.2	2.339		
S	5.17	1.64287269	11.7	70.2	2.232		
T	5.49	1.70292826	18.7	112.2	1.826		
V	7	1.94591015	23.3	46.6	1.826		
W	1.7	0.53062825	74.3	891.6	0.516		
Y	2.85	1.04731899	50	350	0.976		
A	6.5	1.87180218	11.7	11.7	1.261	T. tengconge	
C	0.76	-0.2744368	24.7	741	1.076		
D	4.94	1.59736533	12.7	114.3	1.076		
E	8.4	2.12823171	15.3	76.5	1.076		
F	4.37	1.47476301	52	208	1.583		
G	6.84	1.92278773	11.7	11.7	1.261		
H	1.44	0.36464311	38.3	536.2	1.076		
I	8.95	2.19165353	32.3	64.6	2.069		
K	8.86	2.18154677	30.3	242.4	1.583		
L	9.53	2.25444472	27.3	54.6	2.374		
M	2.52	0.9242589	34.3	445.9	0.605		
N	4.35	1.47017585	14.7	147	1.583		
P	3.56	1.26976055	20.3	60.9	1.261		
Q	2.33	0.84586827	16.3	130.4	1.076		
R	4.35	1.47017585	27.3	109.2	1.866		
S	5.25	1.65822808	11.7	70.2	2.174		
T	4.59	1.52388002	18.7	112.2	1.769		
V	7.54	2.02022218	23.3	46.6	1.769		

-0.71

-0.87

0.68

W	0.8	-0.2231436	74.3	891.6	9.70E-02			
Y	4.02	1.3912819	50	350	1.583			
A	4.56	1.51732262	11.7	11.7	0.628	B. floridanus	0.27	-0.3
C	1.57	0.45107562	24.7	741	0.91			-0.48
D	4.58	1.521699	12.7	114.3	0.91			
E	4.43	1.48839958	15.3	76.5	0.91			
F	4.45	1.4929041	52	208	1.886			
G	5.7	1.74046618	11.7	11.7	0.628			
H	2.58	0.94777894	38.3	536.2	0.91			
I	11.4	2.43361336	32.3	64.6	2.432			
K	7.16	1.96850998	30.3	242.4	1.886			
L	10	2.30258509	27.3	54.6	2.448			
M	2.5	0.91629073	34.3	445.9	0.591			
N	6.88	1.92861865	14.7	147	1.886			
P	3.18	1.1568812	20.3	60.9	0.628			
Q	3.89	1.35840916	16.3	130.4	0.91			
R	4.08	1.40609699	27.3	109.2	1.472			
S	7.02	1.94876322	11.7	70.2	2.009			
T	4.86	1.58103844	18.7	112.2	1.604			
V	5.75	1.74919986	23.3	46.6	1.604			
W	1.03	2.96E-02	74.3	891.6	-0.385			
Y	4.23	1.44220199	50	350	1.886			
A	6.9	1.93152141	11.7	11.7	1.2	P. marinus CC	0.36	-0.47
C	1.22	0.19885086	24.7	741	1.063			-0.56
D	4.93	1.59533899	12.7	114.3	1.063			
E	6.43	1.86097454	15.3	76.5	1.063			
F	4.06	1.40118297	52	208	1.619			
G	6.87	1.92716411	11.7	11.7	1.2			
H	1.81	0.59332685	38.3	536.2	1.063			
I	7.81	2.05540496	32.3	64.6	2.111			
K	6.6	1.88706965	30.3	242.4	1.619			
L	11.5	2.44234704	27.3	54.6	2.383			

-0.63

-0.82

0.81

-0.75

-0.89

0.7

M	1.94	0.66268797	34.3	445.9	0.61			
N	4.97	1.60341984	14.7	147	1.619			
P	4.25	1.44691898	20.3	60.9	1.2			
Q	3.66	1.29746315	16.3	130.4	1.063			
R	4.92	1.59330853	27.3	109.2	1.827			
S	7.71	2.04251819	11.7	70.2	2.162			
T	4.69	1.54543258	18.7	112.2	1.756			
V	5.7	1.74046618	23.3	46.6	1.756			
W	1.46	0.37843644	74.3	891.6	5.40E-02			
Y	2.44	0.89199804	50	350	1.619			
A	9.63	2.26488323	11.7	11.7	1.862	P. marinus MI	0.51	-0.51
C	1.32	0.27763174	24.7	741	1.139			-0.61
D	5.07	1.62334082	12.7	114.3	1.139			
E	5.83	1.763017	15.3	76.5	1.139			
F	3.25	1.178655	52	208	1.11			
G	7.84	2.05923883	11.7	11.7	1.862			
H	2.17	0.77472717	38.3	536.2	1.139			
I	4.9	1.58923521	32.3	64.6	1.51			
K	3.51	1.25561604	30.3	242.4	1.11			
L	12.5	2.52572864	27.3	54.6	2.228			
M	2.13	0.75612198	34.3	445.9	0.431			
N	3.21	1.16627094	14.7	147	1.11			
P	5.17	1.64287269	20.3	60.9	1.862			
Q	4.88	1.58514522	16.3	130.4	1.139			
R	6.5	1.87180218	27.3	109.2	2.258			
S	6.78	1.9139771	11.7	70.2	2.238			
T	4.71	1.54968791	18.7	112.2	1.832			
V	6.79	1.91545094	23.3	46.6	1.832			
W	1.72	0.54232429	74.3	891.6	0.461			
Y	1.98	0.68309685	50	350	1.11			
A	5.35	1.67709656	11.7	11.7	0.864	P. marinus CC	0.31	-0.38
C	1.16	0.14842001	24.7	741	0.98			-0.49

-0.71

-0.87

0.79

-0.68

-0.85

0.7

D	5.06	1.62136648	12.7	114.3	0.98		
E	6.62	1.89009537	15.3	76.5	0.98		
F	4.89	1.5871923	52	208	1.789		
G	6.28	1.83736998	11.7	11.7	0.864		
H	1.5	0.40546511	38.3	536.2	0.98		
I	9.22	2.22137504	32.3	64.6	2.315		
K	8.56	2.14710019	30.3	242.4	1.789		
L	10.7	2.37024374	27.3	54.6	2.426		
M	1.83	0.60431597	34.3	445.9	0.612		
N	6.56	1.8809906	14.7	147	1.789		
P	3.62	1.28647403	20.3	60.9	0.864		
Q	2.99	1.09527339	16.3	130.4	0.98		
R	3.92	1.36609165	27.3	109.2	1.617		
S	7.72	2.04381436	11.7	70.2	2.079		
T	4.46	1.49514877	18.7	112.2	1.673		
V	5.25	1.65822808	23.3	46.6	1.673		
W	1.25	0.22314355	74.3	891.6	-0.198		
Y	2.87	1.05431203	50	350	1.789		
A	7.87	2.06305806	11.7	11.7	1.77	W. succinoge	
C	0.94	-6.19E-02	24.7	741	1.138		0.48
D	4.33	1.46556754	12.7	114.3	1.138		-0.41
E	8.25	2.1102132	15.3	76.5	1.138		-0.5
F	4.8	1.56861592	52	208	1.2		
G	7.09	1.95868534	11.7	11.7	1.77		
H	2.07	0.72754861	38.3	536.2	1.138		
I	7.12	1.96290773	32.3	64.6	1.616		
K	6.76	1.91102289	30.3	242.4	1.2		
L	11.5	2.44234704	27.3	54.6	2.258		
M	2.47	0.90421815	34.3	445.9	0.476		
N	3.43	1.23256026	14.7	147	1.2		
P	3.72	1.31372367	20.3	60.9	1.77		
Q	3.18	1.1568812	16.3	130.4	1.138		
R	5	1.60943791	27.3	109.2	2.196		

-0.73

-0.89

0.67

S	6.95	1.93874166	11.7	70.2	2.237			
T	4.24	1.44456327	18.7	112.2	1.832			
V	5.86	1.7681496	23.3	46.6	1.832			
W	0.98	-2.02E-02	74.3	891.6	0.414			
Y	3.31	1.19694819	50	350	1.2			
A	7.96	2.074429	11.7	11.7	1.523	P. luminescen	0.43	-0.53
C	1.1	9.53E-02	24.7	741	1.119			-0.58
D	5.23	1.65441128	12.7	114.3	1.119			
E	5.91	1.77664583	15.3	76.5	1.119			
F	3.92	1.36609165	52	208	1.407			
G	6.8	1.91692261	11.7	11.7	1.523			
H	2.38	0.86710049	38.3	536.2	1.119			
I	6.9	1.93152141	32.3	64.6	1.86			
K	5.19	1.6467337	30.3	242.4	1.407			
L	10.5	2.35137526	27.3	54.6	2.323			
M	2.42	0.88376754	34.3	445.9	0.56			
N	4.66	1.53901545	14.7	147	1.407			
P	4.2	1.43508453	20.3	60.9	1.523			
Q	4.72	1.5518088	16.3	130.4	1.119			
R	5.12	1.63315444	27.3	109.2	2.034			
S	6.49	1.87026253	11.7	70.2	2.217			
T	5.41	1.68824909	18.7	112.2	1.812			
V	6.36	1.85002838	23.3	46.6	1.812			
W	1.35	0.30010459	74.3	891.6	0.271			
Y	3.26	1.18172719	50	350	1.407			
A	8.56	2.14710019	11.7	11.7	1.638	V. parahaemc	0.45	-0.55
C	1.04	3.92E-02	24.7	741	1.131			-0.59
D	5.5	1.70474809	12.7	114.3	1.131			
E	6.54	1.87793717	15.3	76.5	1.131			
F	4.19	1.43270073	52	208	1.317			
G	6.77	1.91250109	11.7	11.7	1.638			
H	2.24	0.80647587	38.3	536.2	1.131			

-0.82

-0.94

0.74

-0.81

-0.94

0.69

I	6.17	1.81969884	32.3	64.6	1.752		
K	5.38	1.68268837	30.3	242.4	1.317		
L	10.1	2.31253542	27.3	54.6	2.295		
M	2.74	1.00795792	34.3	445.9	0.526		
N	4.26	1.44926916	14.7	147	1.317		
P	3.89	1.35840916	20.3	60.9	1.638		
Q	4.53	1.51072194	16.3	130.4	1.131		
R	4.47	1.49738841	27.3	109.2	2.11		
S	6.58	1.88403475	11.7	70.2	2.229		
T	5.4	1.68639895	18.7	112.2	1.824		
V	7.19	1.97269117	23.3	46.6	1.824		
W	1.23	0.20701417	74.3	891.6	0.341		
Y	3.04	1.11185752	50	350	1.317		
A	8.8	2.17475172	11.7	11.7	1.696	V. vulnificus	-0.58
C	1.07	6.77E-02	24.7	741	1.135	0.47	
D	5.26	1.66013103	12.7	114.3	1.135		
E	6.36	1.85002838	15.3	76.5	1.135		
F	4.18	1.43031125	52	208	1.268		
G	6.68	1.89911799	11.7	11.7	1.696		
H	2.36	0.85866162	38.3	536.2	1.135		
I	6.06	1.8017098	32.3	64.6	1.695		
K	5.18	1.64480506	30.3	242.4	1.268		
L	10.4	2.34180581	27.3	54.6	2.28		
M	2.75	1.01160091	34.3	445.9	0.506		
N	4.11	1.41342303	14.7	147	1.268		
P	3.84	1.34547237	20.3	60.9	1.696		
Q	4.89	1.5871923	16.3	130.4	1.135		
R	4.69	1.54543258	27.3	109.2	2.147		
S	6.55	1.87946505	11.7	70.2	2.234		
T	5.26	1.66013103	18.7	112.2	1.828		
V	7.03	1.95018671	23.3	46.6	1.828		
W	1.29	0.25464222	74.3	891.6	0.373		
Y	3.07	1.12167756	50	350	1.268		

-0.8

-0.93

0.69

A	11.5	2.44234704	11.7	11.7	2.489	Thermo-thern	0.69	-0.35	-0.35
C	0.39	-0.9416085	24.7	741	0.976				
D	3.58	1.2753628	12.7	114.3	0.976				
E	8.61	2.15292432	15.3	76.5	0.976				
F	3.78	1.32972401	52	208	0.157				
G	9.29	2.22893855	11.7	11.7	2.489				
H	1.87	0.62593843	38.3	536.2	0.976				
I	2.68	0.9858168	32.3	64.6	0.424				
K	3.62	1.28647403	30.3	242.4	0.157				
L	14.5	2.67414865	27.3	54.6	1.868				
M	1.56	0.44468582	34.3	445.9	-0.208				
N	1.55	0.43825493	14.7	147	0.157				
P	6.46	1.86562932	20.3	60.9	2.489				
Q	2.46	0.90016135	16.3	130.4	0.976				
R	8.46	2.13534917	27.3	109.2	2.688				
S	3.39	1.22082992	11.7	70.2	2.075				
T	3.76	1.32441896	18.7	112.2	1.669				
V	8.14	2.09679018	23.3	46.6	1.669				
W	1.31	0.27002714	74.3	891.6	0.611				
Y	2.87	1.05431203	50	350	0.157				
A	11.5	2.44234704	11.7	11.7	2.491	Thermus-ther	0.7	-0.35	-0.35
C	0.39	-0.9416085	24.7	741	0.974				
D	3.59	1.2781522	12.7	114.3	0.974				
E	8.65	2.15755932	15.3	76.5	0.974				
F	3.76	1.32441896	52	208	0.151				
G	9.29	2.22893855	11.7	11.7	2.491				
H	1.86	0.62057649	38.3	536.2	0.974				
I	2.67	0.98207847	32.3	64.6	0.417				
K	3.61	1.28370777	30.3	242.4	0.151				
L	14.5	2.67414865	27.3	54.6	1.866				
M	1.55	0.43825493	34.3	445.9	-0.213				
N	1.55	0.43825493	14.7	147	0.151				

-0.61

-0.78

0.62

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-0.78

0.62

P	6.52	1.87487438	20.3	60.9	2.491			
Q	2.44	0.89199804	16.3	130.4	0.974			
R	8.5	2.14006616	27.3	109.2	2.69			
S	3.41	1.22671229	11.7	70.2	2.073			
T	3.75	1.32175584	18.7	112.2	1.668			
V	8.14	2.09679018	23.3	46.6	1.668			
W	1.32	0.27763174	74.3	891.6	0.611			
Y	2.87	1.05431203	50	350	0.151			
A	7.36	1.99605993	11.7	11.7	1.34	S. thermophil	0.39	-0.47
C	0.58	-0.5447272	24.7	741	1.091			-0.5
D	5.79	1.75613229	12.7	114.3	1.091			
E	6.79	1.91545094	15.3	76.5	1.091			
F	4.63	1.53255687	52	208	1.535			
G	6.55	1.87946505	11.7	11.7	1.34			
H	1.91	0.64710324	38.3	536.2	1.091			
I	7.41	2.00283044	32.3	64.6	2.01			
K	6.95	1.93874166	30.3	242.4	1.535			
L	9.94	2.29656702	27.3	54.6	2.36			
M	2.56	0.94000726	34.3	445.9	0.595			
N	4.62	1.53039471	14.7	147	1.535			
P	3.27	1.18478999	20.3	60.9	1.34			
Q	3.76	1.32441896	16.3	130.4	1.091			
R	4.14	1.42069579	27.3	109.2	1.916			
S	6.27	1.83577636	11.7	70.2	2.189			
T	5.69	1.73871025	18.7	112.2	1.784			
V	7.05	1.95302762	23.3	46.6	1.784			
W	0.84	-0.1743534	74.3	891.6	0.151			
Y	3.78	1.32972401	50	350	1.535			
A	7.37	1.99741771	11.7	11.7	1.34	S. thermophil	0.39	-0.47
C	0.58	-0.5447272	24.7	741	1.091			-0.5
D	5.82	1.76130026	12.7	114.3	1.091			
E	6.8	1.91692261	15.3	76.5	1.091			

-0.8

-0.91

0.67

-0.8

-0.91

0.67

F	4.6	1.5260563	52	208	1.534		
G	6.53	1.87640694	11.7	11.7	1.34		
H	1.91	0.64710324	38.3	536.2	1.091		
I	7.41	2.00283044	32.3	64.6	2.01		
K	6.96	1.94017947	30.3	242.4	1.534		
L	9.93	2.29556048	27.3	54.6	2.36		
M	2.57	0.9439059	34.3	445.9	0.595		
N	4.62	1.53039471	14.7	147	1.534		
P	3.25	1.178655	20.3	60.9	1.34		
Q	3.76	1.32441896	16.3	130.4	1.091		
R	4.16	1.42551507	27.3	109.2	1.916		
S	6.27	1.83577636	11.7	70.2	2.189		
T	5.68	1.73695123	18.7	112.2	1.784		
V	7.07	1.95586048	23.3	46.6	1.784		
W	0.84	-0.1743534	74.3	891.6	0.151		
Y	3.77	1.327075	50	350	1.534		
A	12	2.48490665	11.7	11.7	2.467	S. thermophil	
C	0.87	-0.1392621	24.7	741	0.989		
D	4.86	1.58103844	12.7	114.3	0.989		
E	6.6	1.88706965	15.3	76.5	0.989		
F	3.1	1.13140211	52	208	0.205		
G	8.98	2.19499988	11.7	11.7	2.467		
H	1.97	0.67803354	38.3	536.2	0.989		
I	4.38	1.47704872	32.3	64.6	0.477		
K	2.5	0.91629073	30.3	242.4	0.205		
L	10.9	2.38876279	27.3	54.6	1.888		
M	2.25	0.81093022	34.3	445.9	-0.171		
N	2.07	0.72754861	14.7	147	0.205		
P	5.79	1.75613229	20.3	60.9	2.467		
Q	3.42	1.22964055	16.3	130.4	0.989		
R	8.36	2.12345843	27.3	109.2	2.673		
S	4.3	1.45861502	11.7	70.2	2.088		
T	4.96	1.60140574	18.7	112.2	1.682		

-0.68

-0.84

0.71

V	8.49	2.138889	23.3	46.6	1.682			
W	1.45	0.37156356	74.3	891.6	0.613			
Y	2.51	0.92028275	50	350	0.205			
A	9.23	2.22245905	11.7	11.7	1.91	G. kaustophil	0.52	-0.49
C	0.86	-0.1508229	24.7	741	1.138			
D	4.85	1.57897871	12.7	114.3	1.138			
E	7.41	2.00283044	15.3	76.5	1.138			
F	4.16	1.42551507	52	208	1.059			
G	7.23	1.97823904	11.7	11.7	1.91			
H	2.32	0.84156719	38.3	536.2	1.138			
I	6.36	1.85002838	32.3	64.6	1.451			
K	5.62	1.72633166	30.3	242.4	1.059			
L	9.94	2.29656702	27.3	54.6	2.211			
M	2.66	0.97832612	34.3	445.9	0.404			
N	3.08	1.1249296	14.7	147	1.059			
P	4.27	1.45161383	20.3	60.9	1.91			
Q	3.66	1.29746315	16.3	130.4	1.138			
R	6.09	1.80664808	27.3	109.2	2.29			
S	4.91	1.59127394	11.7	70.2	2.236			
T	4.97	1.60341984	18.7	112.2	1.831			
V	7.62	2.03077637	23.3	46.6	1.831			
W	1.29	0.25464222	74.3	891.6	0.484			
Y	3.36	1.21194097	50	350	1.059			

-0.78

-0.9

0.64