

Amino acid	% abundance	In (% abundance)	Cost (ATP)	Cost (ATP/tin)	In (% abundance)	Organism	Genomic GC c	R % abundance	R ln(% abundance)
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	1.632					

R % 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
U	6.01	1.79342	23.3	46.6	1.752
V	1.11	0.10436	74.3	891.6	3.80E-02
W	3.4	1.22378	50	350	1.632
X	8.7	2.16332303	11.7	11.7	1.827
Y	1.13	0.12221763	24.7	74.1	N. crassa
Z	5.63	1.72810944	12.7	114.3	1.139
A	6.46	1.86562932	15.3	76.5	1.139
C	3.37	1.21491274	52	208	1.145
D	7.2	1.97408103	11.7	11.7	1.827
E	2.45	0.89608803	38.3	536.2	1.139
F	4.42	1.4861397	32.3	64.6	1.551
G	5.08	1.62531126	30.3	242.4	1.145
H	8.35	2.12226154	27.3	54.6	2.24
I	2.15	0.76546784	34.3	445.9	0.449
K	3.7	1.30833282	14.7	147	1.145
L	6.52	1.87487438	20.3	60.9	1.827
M	4.26	1.44926916	16.3	130.4	1.139
N	6.17	1.81969884	27.3	109.2	2.235
P	8.29	2.11504997	11.7	70.2	2.238
Q	6.11	1.80992677	18.7	112.2	1.833
R	5.98	1.78842057	23.3	46.6	1.833
S	1.35	0.30010459	74.3	891.6	0.444
T	2.56	0.94000726	50	350	1.145
U	4.93	1.59533899	11.7	11.7	C. albicans
V	1.17	0.15700375	24.7	74.1	0.34
W	5.71	1.74221902	12.7	114.3	-0.51
X	6.23	1.82937633	15.3	76.5	-0.59

-0.83

-0.93

0.77

-0.75

-0.9

0.75

-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			
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			
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			

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-0.93

0.8

L	9.16	2.21484618	27.3	54.6	2.233	
M	2.04	0.71294981	34.3	445.9	0.439	
N	3.67	1.30019166	14.7	147	1.124	
P	6	1.79175947	20.3	60.9	1.848	
Q	4.02	1.3912819	16.3	130.4	1.139	
R	6.24	1.83098018	27.3	109.2	2.248	
S	8.38	2.12584791	11.7	70.2	2.238	
T	5.97	1.78674693	18.7	112.2	1.833	
V	6.11	1.80992677	23.3	46.6	1.833	
W	1.46	0.37843644	74.3	891.6	0.454	
Y	2.89	1.0612565	50	350	1.124	
A	5.04	1.61740608	11.7	11.7	1.722	E. cuniculi
C	2.04	0.71294981	24.7	741	1.137	
D	5.46	1.69744879	12.7	114.3	1.137	
E	8.1	2.09186406	15.3	76.5	1.137	
F	4.78	1.56444055	52	208	1.244	
G	6.52	1.87487438	11.7	11.7	1.722	
H	1.92	0.65232519	38.3	536.2	1.137	
I	6.67	1.89761986	32.3	64.6	1.667	
K	7.08	1.95727391	30.3	242.4	1.244	
L	9.52	2.25339485	27.3	54.6	2.272	
M	3	1.09861229	34.3	445.9	0.496	
N	3.89	1.35840916	14.7	147	1.244	
P	3.38	1.21787571	20.3	60.9	1.722	
Q	2.27	0.81977983	16.3	130.4	1.137	
R	6.71	1.90359895	27.3	109.2	2.165	
S	7.99	2.07819076	11.7	70.2	2.235	
T	4.09	1.40854497	18.7	112.2	1.83	
V	7.04	1.95160817	23.3	46.6	1.83	
W	0.78	-0.2484614	74.3	891.6	0.388	
Y	3.62	1.28647403	50	350	1.244	
A	6.29	1.83896107	11.7	1.144	C. elegans	
				0.35	-0.58	
				-0.67		

-0.84 -0.93 0.76

-0.71 -0.83 0.71

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
U	6.19	1.82293509	23.3	46.6	1.744
V	1.1	9.53E-02	74.3	891.6	1.30E-02
W	3.16	1.15057203	50	350	1.651
X					
Y					
Z					
A	6.18	1.82131827	11.7	11.7	1.249 C. briggsae
B	1.95	0.66782937	24.7	741	1.073
C	5.29	1.66581825	12.7	114.3	1.073
D	6.81	1.91839212	15.3	76.5	1.073
E	4.66	1.53901545	52	208	1.591
F	5.37	1.68082791	11.7	11.7	1.249
G	2.32	0.84156719	38.3	536.2	1.073
H	5.89	1.773256	32.3	64.6	2.077
I	6.46	1.86562932	30.3	242.4	1.591
K	8.46	2.13534917	27.3	54.6	2.375
L	2.62	0.96317432	34.3	445.9	0.606
M	4.78	1.56444055	14.7	147	1.591
N	5.05	1.61938824	20.3	60.9	1.249
P	4.13	1.41827741	16.3	130.4	1.073

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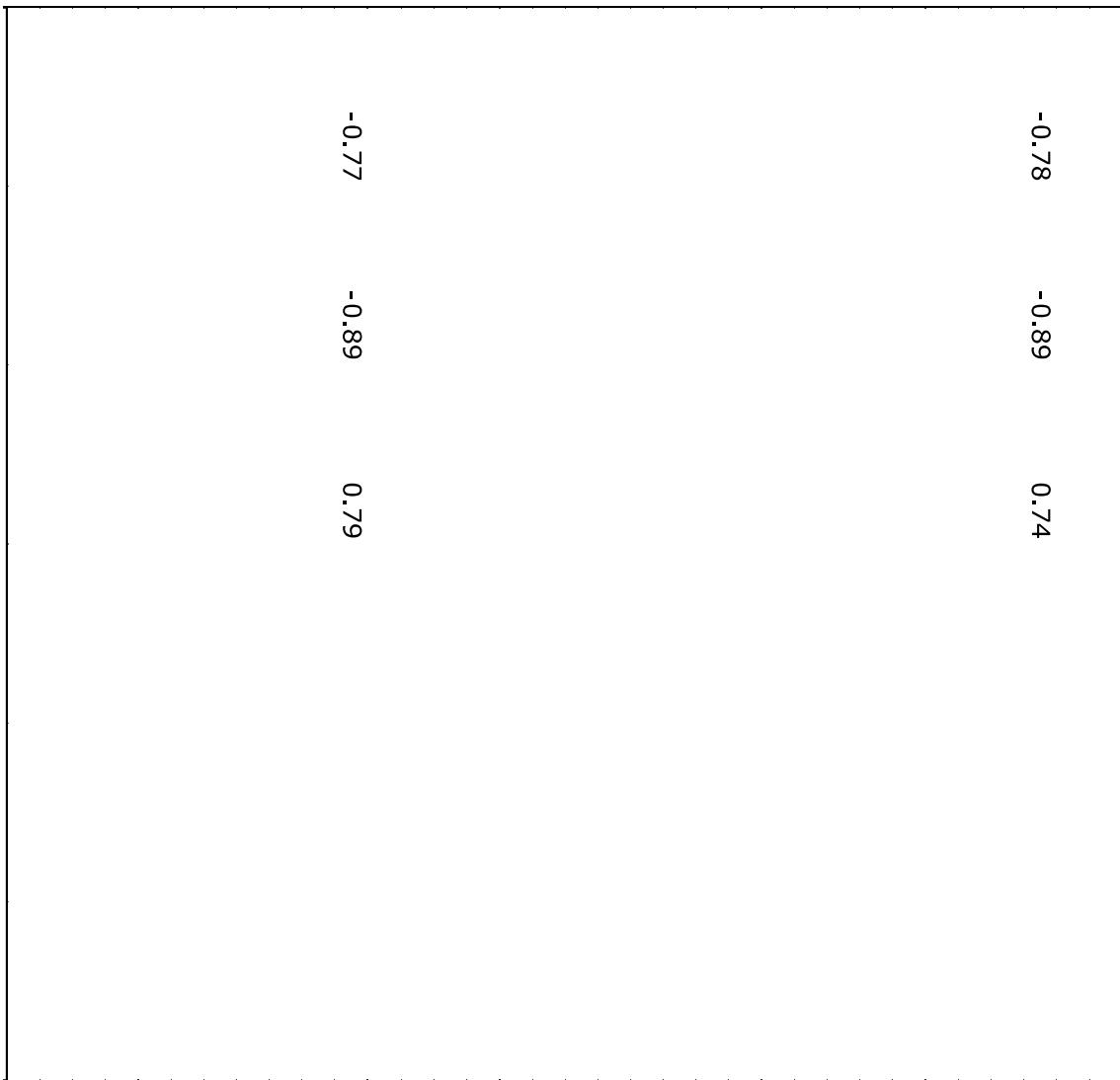
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		2.89	1.0612565	50	350	1.632
Y	A	6.97	1.94161523	11.7	11.7	1.402 H. sapiens
	C	2.31	0.83724753	24.7	741	0.4
	D	4.68	1.54329811	12.7	114.3	-0.62
	E	6.95	1.93874166	15.3	76.5	-0.73
	F	3.64	1.29198368	52	208	
	G	6.7	1.90210753	11.7	11.7	
	H	2.64	0.97077892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



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	1.691193913	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	<i>R. norvegicus</i>
C	2.28	0.82417544	24.7	741	1.113	
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	1.26669476	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
C	2.38	0.86710049	24.7	741	1.131	
D	5.08	1.62531126	12.7	114.3	1.131	

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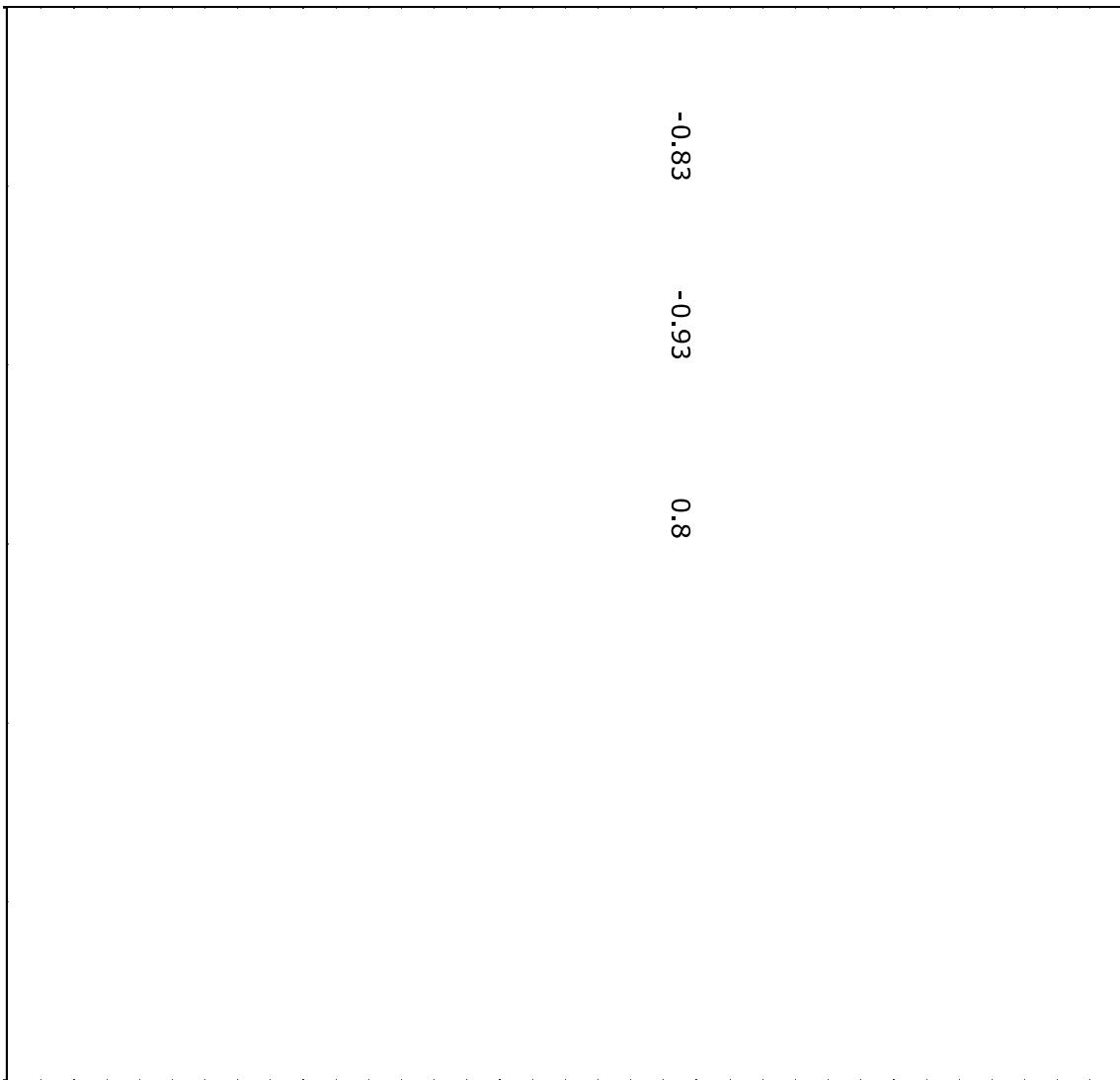
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-0.8 -0.93 0.78

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

-0.82

-0.94

0.74

-0.78

-0.92

0.76

-0.78

-0.91

0.77

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

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-0.9

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
C	1.27	0.2390169	24.7	741	0.989
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
C	1.2	0.18232156	24.7	741	1.139

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

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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				
J	5.63	1.72810944	30.3	242.4	1.294				
K	8.38	2.12584791	27.3	54.6	2.288				
L	3.2	1.16315081	34.3	445.9	0.517				
M	4.25	1.44691898	14.7	147	1.294				
N	3.96	1.37624403	20.3	60.9	1.665				
P	2.15	0.76546784	16.3	130.4	1.133				
Q	5.49	1.70292826	27.3	109.2	2.127				
R	7.57	2.02419307	11.7	70.2	2.232				
S	4.77	1.56234631	18.7	112.2	1.826				
T	7.15	1.96711236	23.3	46.6	1.826				
U	0.85	-0.1625189	74.3	891.6	0.356				
V	4.63	1.53255687	50	350	1.294				
W									
Y									
Z									
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				
J	6.89	1.93007109	30.3	242.4	1.507				
K	8.76	2.17019591	27.3	54.6	2.352				
L	2.71	0.99694864	34.3	445.9	0.588				
M	4.75	1.55814462	14.7	147	1.507				

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

S. solfataricu

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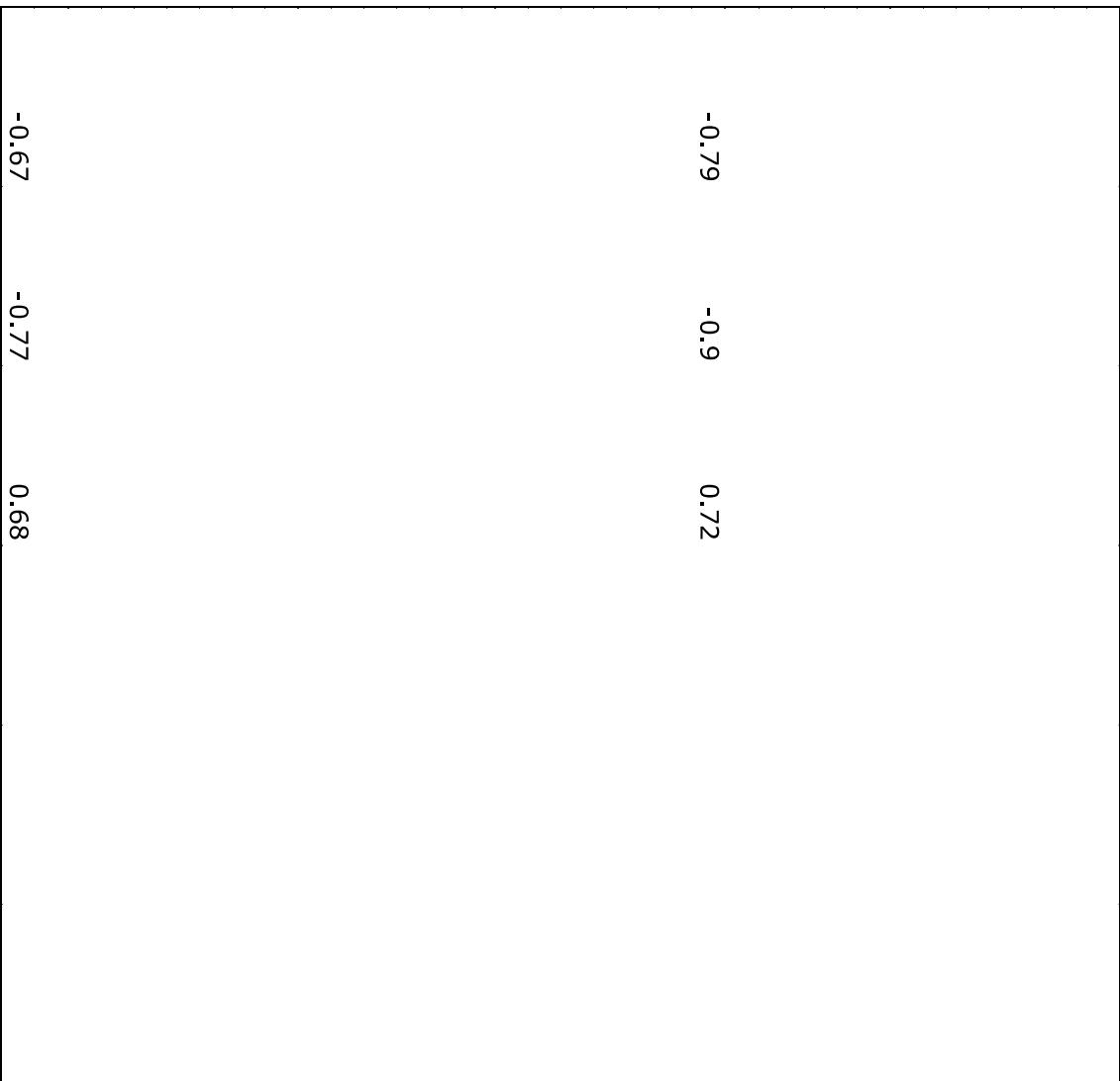
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L	10.5	2.35137526	27.3	54.6	2.219	
M	1.92	0.65232519	34.3	445.9	0.418	
N	2.6	0.95551145	14.7	147	1.084	
P	4.98	1.60542989	20.3	60.9	1.886	
Q	2.08	0.73236789	16.3	130.4	1.139	
R	6.54	1.87793717	27.3	109.2	2.274	
S	4.93	1.59533899	11.7	70.2	2.237	
T	4.4	1.48160454	18.7	112.2	1.832	
V	9.31	2.23108909	23.3	46.6	1.832	
W	1.47	0.3852624	74.3	891.6	0.472	
Y	4.3	1.45861502	50	350	1.084	
A	6.88	1.92861865	11.7	11.7	1.516	M. acetivoran
C	1.25	0.22314355	24.7	741	1.118	
D	5.34	1.67522565	12.7	114.3	1.118	
E	7.96	2.074429	15.3	76.5	1.118	
F	4.44	1.49065438	52	208	1.413	
G	7.23	1.97823904	11.7	11.7	1.516	
H	1.66	0.5068176	38.3	536.2	1.118	
I	7.35	1.99470031	32.3	64.6	1.866	
K	6.54	1.87793717	30.3	242.4	1.413	
L	9.39	2.23964529	27.3	54.6	2.325	
M	2.45	0.89608803	34.3	445.9	0.561	
N	4.48	1.49962305	14.7	147	1.413	
P	3.99	1.38379123	20.3	60.9	1.516	
Q	2.54	0.93216408	16.3	130.4	1.118	
R	4.48	1.49962305	27.3	109.2	2.03	
S	6.9	1.93152141	11.7	70.2	2.216	
T	5.43	1.69193913	18.7	112.2	1.811	
V	6.82	1.91985947	23.3	46.6	1.811	
W	1.05	4.88E-02	74.3	891.6	0.266	
Y	3.72	1.31372367	50	350	1.413	
A	8.34	2.12106322	11.7	11.7	0.61	M. kandleri
					-0.44	
					-0.48	



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
U	10.4	2.34180581	23.3	46.6	1.781
V	1.23	0.20701417	74.3	891.6	0.597
W	2.82	1.03673689	50	350	0.634
M. mazei					
A	7.07	1.95586048	11.7	11.7	1.459
C	1.24	0.21511138	24.7	741	1.11
D	5.25	1.65822808	12.7	114.3	1.11
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

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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		
D	6.29	1.83896107	12.7	114.3	1.105		
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	
C	0.82	-0.1984509	24.7	741	0.993		
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		

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

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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
C	1.03	2.96E-02	24.7	741	1.082	
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
C	0.82	-0.1984509	24.7	741	0.995	
D	4.92	1.59330853	12.7	114.3	0.995	

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E	5.68	1.733695123	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
J	9.5	2.2512918	30.3	242.4	1.764
K	10.6	2.360854	27.3	54.6	2.42
L	1.52	0.41871034	34.3	445.9	0.614
M	7.5	2.01490302	14.7	147	1.764
N	2.99	1.09527339	20.3	60.9	0.921
P	4.73	1.5539252	16.3	130.4	0.995
Q	3.1	1.13140211	27.3	109.2	1.652
R	6.64	1.89311196	11.7	70.2	2.094
S	5.4	1.68639895	18.7	112.2	1.689
T	6.14	1.81482474	23.3	46.6	1.689
V	0.96	-4.08E-02	74.3	891.6	-0.154
W	3.23	1.17248214	50	350	1.764
Y					
A	6.67	1.89761986	11.7	11.7	1.387
C	0.75	-0.2876821	24.7	741	1.099
D	4.96	1.60140574	12.7	114.3	1.099
E	5.67	1.73518912	15.3	76.5	1.099
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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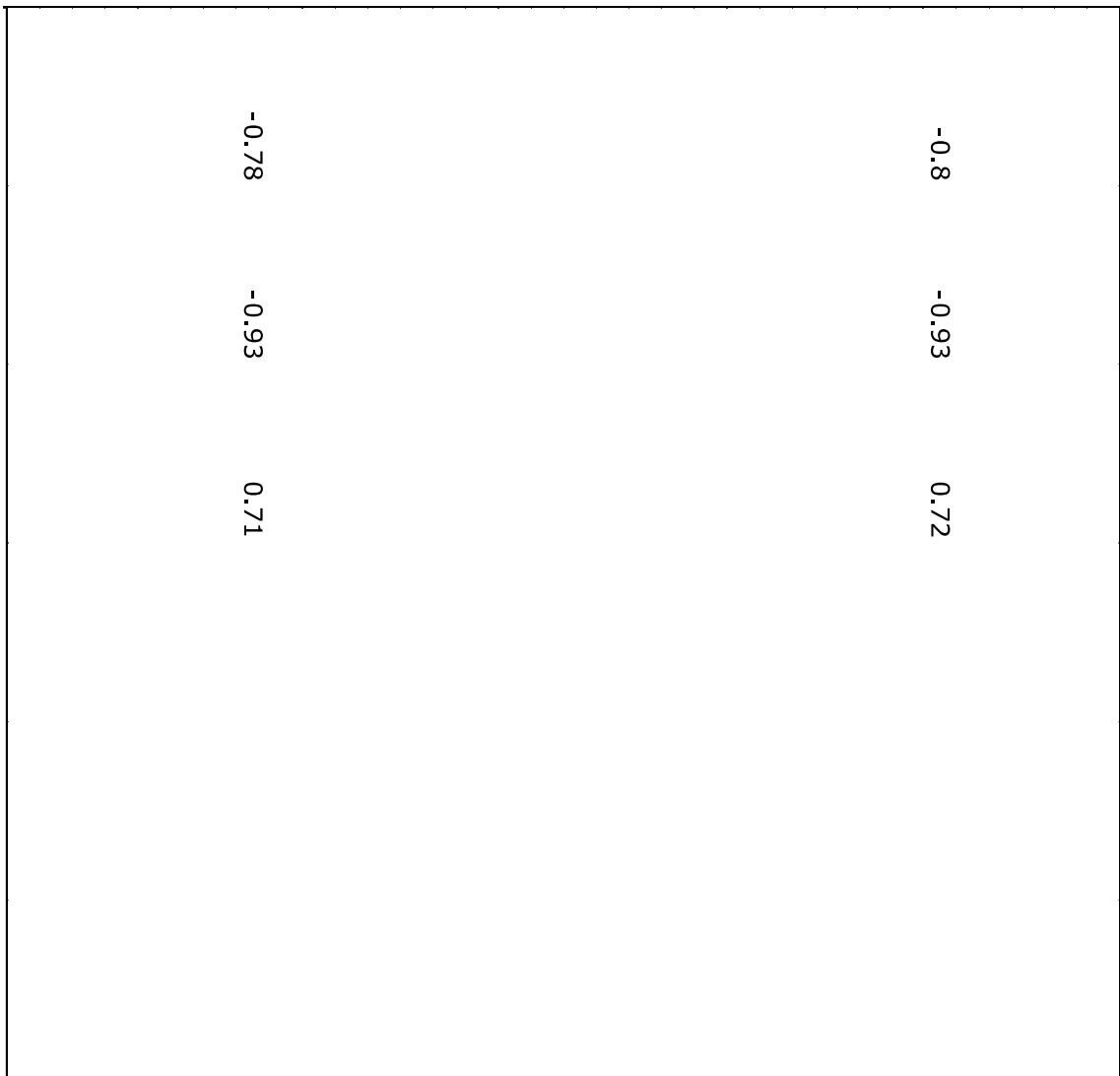
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A	7.67	2.03731662	11.7	11.7	1.555	B. subtilis	0.44	-0.47	-0.52
C	0.8	-0.2231436	24.7	741	1.122				
D	5.17	1.64287269	12.7	114.3	1.122				
E	7.23	1.97823904	15.3	76.5	1.122				
F	4.49	1.15018527	52	208	1.384				
G	6.9	1.93152141	11.7	11.7	1.555				
H	2.27	0.81977983	38.3	536.2	1.122				
I	7.36	1.99605993	32.3	64.6	1.831				
J	7.05	1.95302762	30.3	242.4	1.384				
K	9.64	2.26592111	27.3	54.6	2.316				
L	2.78	1.02245093	34.3	445.9	0.551				
M	3.94	1.37118072	14.7	147	1.384				
N	3.68	1.30291275	20.3	60.9	1.555				
P	3.83	1.34286448	16.3	130.4	1.122				
Q	4.12	1.41585316	27.3	109.2	2.055				
R	6.29	1.83896107	11.7	70.2	2.221				
S	5.42	1.69009582	18.7	112.2	1.816				
T	6.74	1.90805993	23.3	46.6	1.816				
V	1.03	2.96E-02	74.3	891.6	0.29				
W	3.48	1.24703229	50	350	1.384				
Y									
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				

-0.79 -0.91 0.66

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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. burgdorferi	0.28	-0.24
C	0.66	-0.4155154	24.7	741	0.928			
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			
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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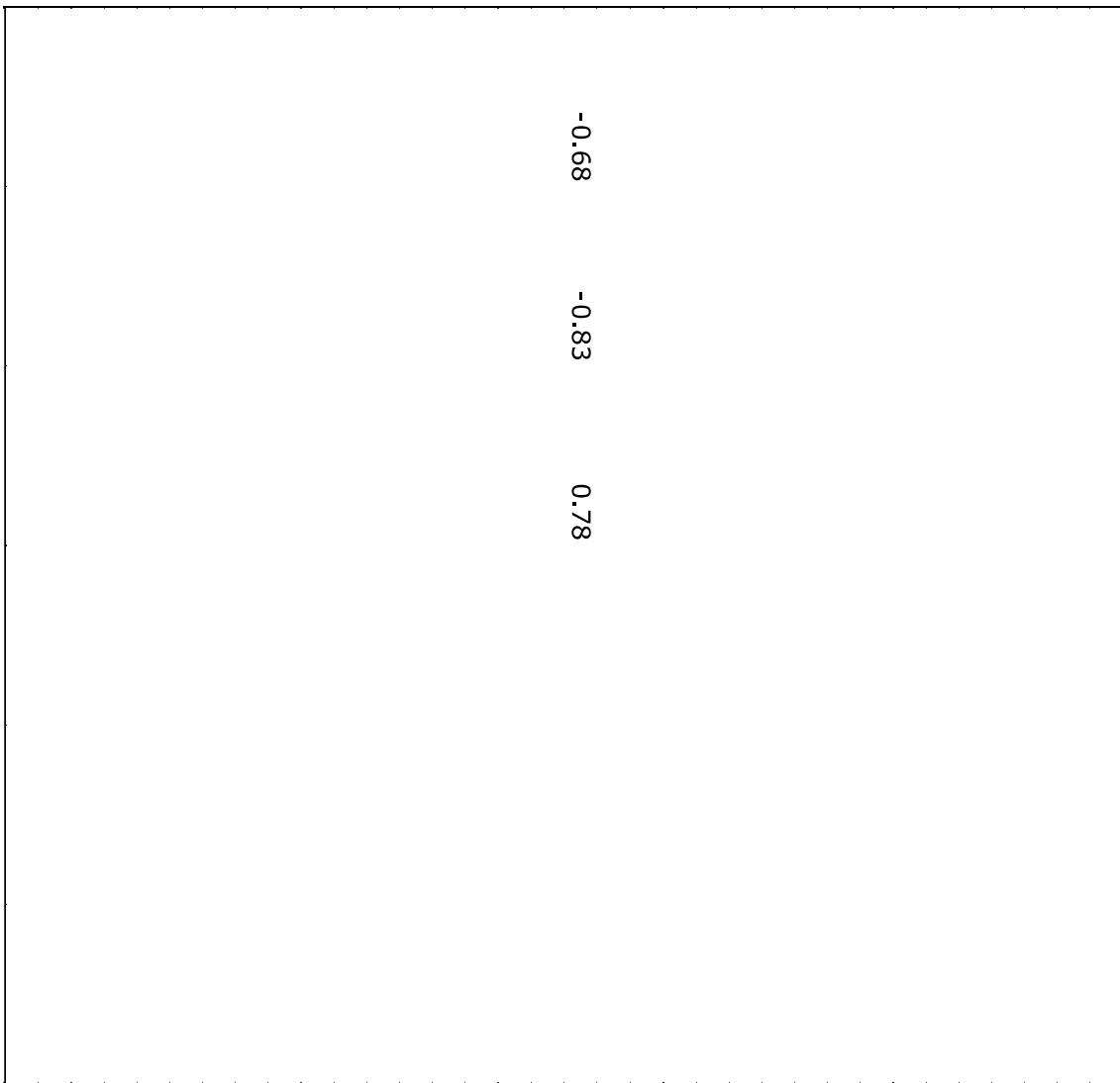
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G	6.75	1.90954251	11.7	11.7	1.545
H	1.54	0.43178242	38.3	536.2	1.121
I	7.33	1.99197552	32.3	64.6	1.84
K	9.4	2.24070969	30.3	242.4	1.391
L	10.5	2.35137526	27.3	54.6	2.318
M	1.93	0.65752	34.3	445.9	0.554
N	3.59	1.2781522	14.7	147	1.391
P	4.07	1.403643	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
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



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
C	1.91	0.64710324	24.7	741	1.136	
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
C	1.62	0.48242615	24.7	741	1.109	

-0.77 -0.92 0.81

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

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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. meningigit	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				

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-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			
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.. monocy tog	0.38	-0.47
C	0.6	-0.5108256	24.7	741	1.08			
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			

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L	9.5	2.2512918	27.3	54.6	2.37
M	2.74	1.00795792	34.3	445.9	0.602
N	4.62	1.53039471	14.7	147	1.57
P	3.47	1.24415459	20.3	60.9	1.283
Q	3.45	1.23837423	16.3	130.4	1.08
R	3.65	1.29472717	27.3	109.2	1.88
S	5.79	1.75613229	11.7	70.2	2.179
T	6.11	1.80992677	18.7	112.2	1.773
V	7.03	1.95018671	23.3	46.6	1.773
W	0.93	-7.26E-02	74.3	891.6	0.112
Y	3.44	1.23547147	50	350	1.57
A	7.67	2.03731662	11.7	11.7	1.251
C	0.59	-0.5276327	24.7	741	1.074
D	5.47	1.69927862	12.7	114.3	1.074
E	7.51	2.01623547	15.3	76.5	1.074
F	4.52	1.50851199	52	208	1.589
G	6.58	1.88403475	11.7	11.7	1.251
H	1.76	0.56531381	38.3	536.2	1.074
I	7.81	2.05540496	32.3	64.6	2.075
K	7.35	1.99470031	30.3	242.4	1.589
L	9.42	2.24283509	27.3	54.6	2.375
M	2.71	0.99694864	34.3	445.9	0.605
N	4.72	1.5518088	14.7	147	1.589
P	3.41	1.22671229	20.3	60.9	1.251
Q	3.43	1.23256026	16.3	130.4	1.074
R	3.7	1.30833282	27.3	109.2	1.86
S	5.8	1.75785792	11.7	70.2	2.172
T	6.08	1.8050047	18.7	112.2	1.767
V	6.96	1.94017947	23.3	46.6	1.767
W	0.94	-6.19E-02	74.3	891.6	9.00E-02
Y	3.46	1.24126859	50	350	1.589
A	9.71	2.27315628	11.7	1.906	S. typhi
				0.52	-0.51
				-0.57	

-0.78 -0.92 0.71

-0.79 -0.9 0.63

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	1.474746301	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
U	7.06	1.95444505	23.3	46.6	1.831
V	1.53	0.42526774	74.3	891.6	0.482
W	2.87	1.05431203	50	350	1.063
Y					
pestis					
	9.19	2.21811594	11.7	11.7	1.735
A	1.04	3.92E-02	24.7	741	1.137
C	5.11	1.6311994	12.7	114.3	1.137
D	5.48	1.7011051	15.3	76.5	1.137
E	3.82	1.34025042	52	208	1.233
F	7.22	1.97685495	11.7	11.7	1.735
G	2.28	0.82417544	38.3	536.2	1.137
H	6.21	1.8261609	32.3	64.6	1.654
I	4.38	1.47704872	30.3	242.4	1.233
K	10.9	2.38876279	27.3	54.6	2.269
L	2.63	0.96698385	34.3	445.9	0.491
M	4.05	1.39871688	14.7	147	1.233
N	4.38	1.47704872	20.3	60.9	1.735
P	4.87	1.58309394	16.3	130.4	1.137

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		3.67	1.30019166	50	350	1.553
A	C	12.3	2.50959926	11.7	11.7	2.279
C	D	0.86	-0.1508229	24.7	741	M. loti
D	E	5.67	1.73518912	12.7	114.3	1.075
E	F	5.37	1.68082791	15.3	76.5	1.075
F	G	3.84	1.34547237	52	208	0.564
G	H	8.61	2.15292432	11.7	11.7	2.279
H	I	2.06	0.72270598	38.3	536.2	1.075
I	J	5.38	1.68268837	32.3	64.6	0.882
J	K	3.69	1.30562646	30.3	242.4	0.564
K	L	9.84	2.28645571	27.3	54.6	2.03
L	M	2.49	0.91228271	34.3	445.9	9.40E-02
M	N	2.71	0.99694864	14.7	147	0.564
N	P	5.06	1.62136648	20.3	60.9	2.279
P	Q	3.11	1.13462273	16.3	130.4	1.075
Q	R	6.98	1.94304892	27.3	109.2	2.541
R	S	5.69	1.73871025	11.7	70.2	2.173
S	T	5.3	1.66770682	18.7	112.2	1.768
T	U	7.34	1.99333884	23.3	46.6	1.768
U	V	1.36	0.3074847	74.3	891.6	0.605
V	W	2.21	0.79299252	50	350	0.564
Y	Z					
A	C	12.1	2.49320545	11.7	11.7	2.268
C	D	0.86	-0.1508229	24.7	741	S. meliloti
D	E	5.48	1.7011051	12.7	114.3	1.078
E	F	6.03	1.79674701	15.3	76.5	1.078
F	G	3.91	1.36353737	52	208	0.582
G	H	8.52	2.14241634	11.7	11.7	2.268
H	I	2.05	0.71783979	38.3	536.2	1.078
I	J	5.47	1.69927862	32.3	64.6	0.902
J	K	3.43	1.23256026	30.3	242.4	0.582
K	L	10	2.30258509	27.3	54.6	2.037
L	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. tumefaciens
C	0.83	-0.1863296	24.7	741	1.106
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
C	0.88	-0.1278334	24.7	741	1.037
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
J	2.03	0.70803579	30.3	242.4	0.39
K	9.75	2.27726729	27.3	54.6	1.962
L	1.84	0.60976557	34.3	445.9	-3.10E-02
M	2.52	0.9242589	14.7	147	0.39
N	5.81	1.75958057	20.3	60.9	2.377
P	3.08	1.1249296	16.3	130.4	1.037
Q	7.32	1.99061033	27.3	109.2	2.609
R	5.48	1.7011051	11.7	70.2	2.135
S	5.92	1.77833645	18.7	112.2	1.73
T	8.57	2.14826773	23.3	46.6	1.73
V	1.46	0.37843644	74.3	891.6	0.615
W	2.06	0.722270598	50	350	0.39
Y					
A	13.7	2.61739583	11.7	11.7	2.562
C	0.77	-0.2613648	24.7	741	0.924
D	6.12	1.8115621	12.7	114.3	0.924
E	5.69	1.73871025	15.3	76.5	0.924
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

S. coelicolor

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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. urealyticum	0.26
C	0.66	-0.4155154	24.7	741	0.865		
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		
U	5.36	1.67896398	23.3	46.6	1.558		
V	0.87	-0.1392621	74.3	891.6	-0.502		
W	4.44	1.49065438	50	350	1.937		
Y	9.44	2.24495598	11.7	11.7	1.858	S. flexneri	0.51
A	1.2	0.18232156	24.7	741	1.139		-0.52
C	5.03	1.61541998	12.7	114.3	1.139		
D	5.85	1.76644166	15.3	76.5	1.139		
E	3.78	1.32972401	52	208	1.113		
F	7.31	1.98924327	11.7	11.7	1.858		
G	2.33	0.84586827	38.3	536.2	1.139		
H	5.83	1.763017	32.3	64.6	1.514		

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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
J	3.11	1.13462273	30.3	242.4	0.976
K	12	2.48490665	27.3	54.6	2.182
L	1.92	0.65232519	34.3	445.9	0.358
M	2.91	1.06815308	14.7	147	0.976
N	5.77	1.75267208	20.3	60.9	1.984
P	5.78	1.75440368	16.3	130.4	1.133
Q	6.62	1.89009537	27.3	109.2	2.339
R	5.17	1.64287269	11.7	70.2	2.232
S	5.49	1.70292826	18.7	112.2	1.826
T	7	1.94591015	23.3	46.6	1.826
V	1.7	0.53062825	74.3	891.6	0.516
W	2.85	1.04731899	50	350	0.976
A	6.5	1.87180218	11.7	11.7	1.261
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
U	7.54	2.02022218	23.3	46.6	1.769

T. tengconge

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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			
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	0.9477894	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			
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			

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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
C	1.32	0.27763174	24.7	741	1.139
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
C	1.16	0.14842001	24.7	741	0.98

-0.68 -0.85 0.7

-0.71 -0.87 0.79

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

-0.73

-0.89

0.67

-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
C	1.07	6.77E-02	24.7	741	V. vulnificus
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-therm	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

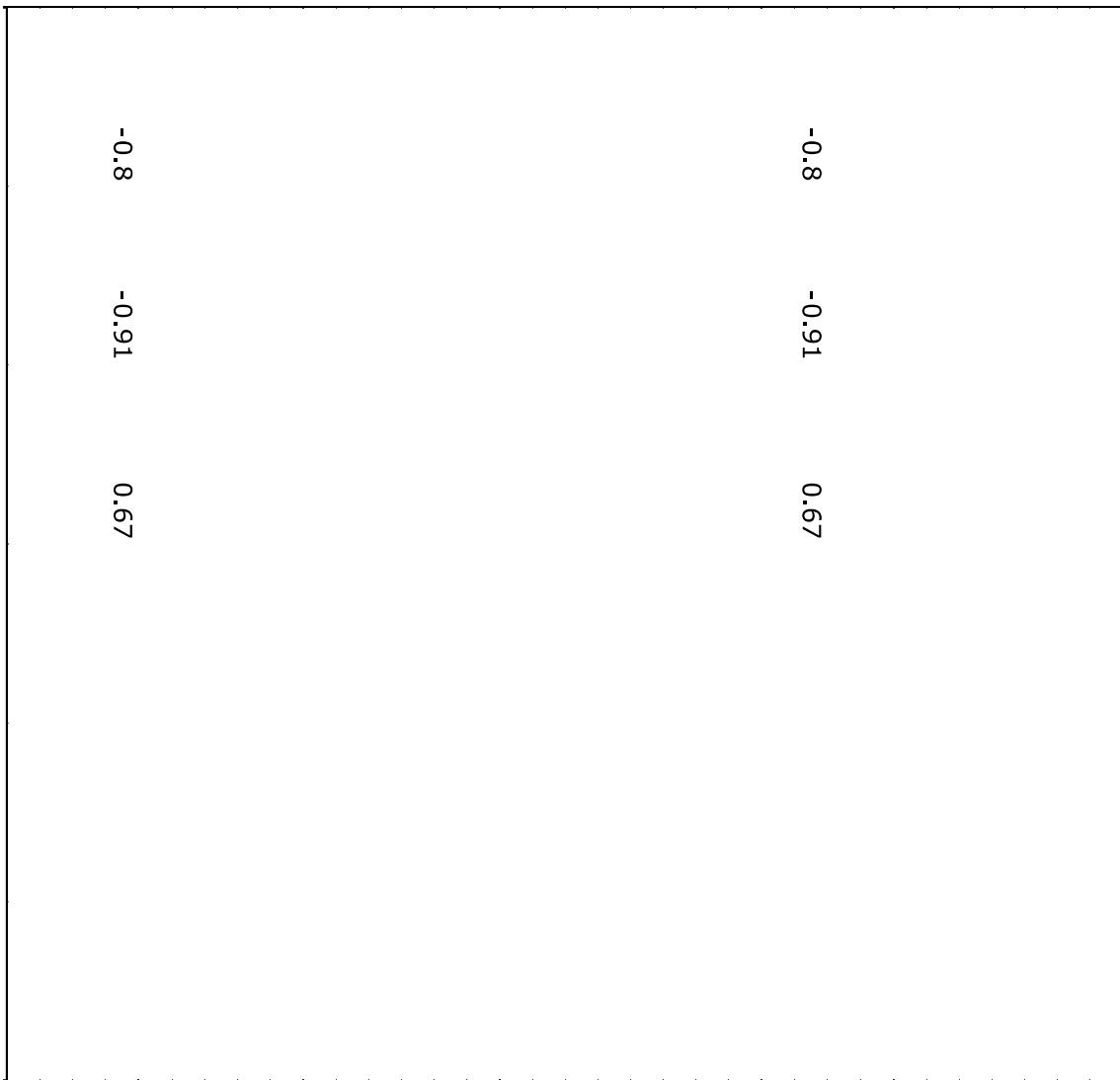
0.62

-0.61

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



-0.68

-0.84

0.71

-0.78

-0.9

0.64