Bacteria - Correlation of amino acid frequency and proteome factors for genetic code: Thraustochytrium mitochondrial

	Pearson correlation coefficient					Spearman correlation coefficient					00
∑ -	-0.28	-0.1	0.17	0.15	Σ-	-0.22	-0.16	0.18	0.17	1.0	10
≥ -	0.18	0.27	0.25	0.24	≥ -	0.22	0.37	0.35	0.33		
U -	-0.11	-0.077	0.12	0.13	U -	-0.062	-0.15	0.2	0.18	- 0.7	' 5
Δ -	0.16	0.31	-0.12	-0.23	Δ -	0.26	0.35	-0.21	-0.28		
ш -	-0.061	-0.21	0.14	0.21	ш -	-0.081	-0.26	0.11	0.23	0.5	- 0
L -	0.094	-0.46	0.45	0.45	L -	0.035	-0.61	0.61	0.61	- 0.5	0.50
ェ -	0.022	0.26	-0.032	-0.14	ェ -	0.088	0.34	0.0091	-0.11		
⊻ -	-0.1	-0.82	0.83	0.83	⊻ -	-0.22	-0.83	0.83	0.83	- 0.2	25
z -	0.11	-0.76	0.77	0.77	z -	-0.018	-0.83	0.83	0.83		
acid Q	-0.094	-0.044	0.34	0.3	ο -	-0.097	-0.098	0.4	0.35	0.0	
Amino acid Y Q	0.24	-0.54	0.54	0.54	> -	0.1	-0.62	0.62	0.62	- 0.0	10
	-0.082	-0.82	0.82	0.82		-0.16	-0.86	0.86	0.86		
∢ -	0.0085	0.83	0.82	0.82	∢ -	0.098	0.86	0.86	0.86	0).25
ს -	0.1	0.65	0.64	0.64	ს -	0.23	0.73	0.73	0.73		
۵ -	0.023	0.64	0.64	0.64	<u>a</u> -	0.19	0.77	0.77	0.77		. = 0
⊢ -	0.19	0.13	0.067	0.0046	- -	0.19	0.15	0.053	0.014	0).50
> -	-0.11	0.46	-0.035	-0.22	> -	-0.013	0.53	-0.088	-0.25		
	-0.057	0.15	-0.085	-0.11		-0.053	0.21	-0.094	-0.16	0).75
∝ -	-0.073	0.69	0.7	0.7	∝ -	-0.017	0.76	0.76	0.76		
ω -	0.094	-0.39	0.051	0.2	ω -	0.017	-0.49	0.13	0.27		00
	Length -	- OS	- Codon+GC -	Codon+GC+Cost -		Length -	- OS	Codon+GC -	Codon+GC+Cost -		00

Factor

Factor