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

				11461101111111		Horiariai				
Pearson correlation coefficient						Spearman correlation coefficient				1.00
Σ-	-0.28	-0.1	0.17	0.17	Σ-	-0.22	-0.16	0.18	0.18	1.00
≥ -	0.18	0.27	-0.042	-0.054	≥ -	0.22	0.37	-0.0047	-0.02	
O -	-0.11	-0.077	0.12	0.12	υ -	-0.062	-0.15	0.2	0.2	- 0.75
Ω-	0.16	0.31	-0.12	-0.14	Δ-	0.26	0.35	-0.21	-0.22	
ш -	-0.061	-0.21	0.14	0.15	ш -	-0.081	-0.26	0.11	0.13	0.50
ш-	0.094	-0.46	0.45	0.45	ш-	0.035	-0.61	0.61	0.61	- 0.50
ェ -	0.022	0.26	-0.032	-0.044	ェ -	0.088	0.34	0.0091	-0.0037	
⊻ -	-0.1	-0.82	0.75	0.75	⊻ -	-0.22	-0.83	0.81	0.81	- 0.25
z -	0.11	-0.76	0.77	0.77	z -	-0.018	-0.83	0.83	0.83	
o acid Q	-0.094	-0.044	0.34	0.34	ο -	-0.097	-0.098	0.4	0.4	0.00
Amino acid Y Q	0.24	-0.54	0.54	0.54	> -	0.1	-0.62	0.62	0.62	- 0.00
	-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	
d -	0.023	0.64	0.64	0.64		0.19	0.77	0.77	0.77	0.50
⊢ -	0.19	0.13	0.067	0.061	⊢ -	0.19	0.15	0.053	0.05	0.30
> -	-0.11	0.46	-0.035	-0.056	> -	-0.013	0.53	-0.088	-0.11	
	-0.057	0.15	-0.14	-0.14	-	-0.053	0.21	-0.21	-0.21	0.75
∝ -	-0.073	0.69	0.7	0.7	<u>~</u> -	-0.017	0.76	0.76	0.76	
S -	0.094	-0.39	0.051	0.069	v -	0.017	-0.49	0.13	0.15	1.00
	Length -	- J9	- Codon+GC	Codon+GC+Cost -		Length -	- J9	Codon+GC -	Codon+GC+Cost -	1.00

Factor

Factor