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

	Pearson correlation coefficient					Spearman correlation coefficient					1.00
∑ -	-0.28	-0.1	0.059	0.054	Σ-	-0.22	-0.16	0.16	0.16		- 1.00
> -	0.18	0.27	-0.042	-0.097	≥ -	0.22	0.37	-0.0047	-0.068		
<b>U</b> -	-0.11	-0.077	0.12	0.13	U -	-0.062	-0.15	0.2	0.19		- 0.75
Δ-	0.16	0.31	-0.12	-0.18	Δ -	0.26	0.35	-0.21	-0.24		
ш -	-0.061	-0.21	0.14	0.18	ш -	-0.081	-0.26	0.11	0.18		- 0.50
<b>LL</b> -	0.094	-0.46	0.45	0.45	ш-	0.035	-0.61	0.61	0.61		- 0.50
エ -	0.022	0.26	-0.032	-0.086	ェ -	0.088	0.34	0.0091	-0.047		
⊻ -	-0.1	-0.82	0.75	0.76	⊻ -	-0.22	-0.83	0.81	0.82		- 0.25
<b>z</b> -	0.11	-0.76	0.77	0.77	<b>z</b> -	-0.018	-0.83	0.83	0.83		
Amino acid Y Q	-0.094	-0.044	0.34	0.33	ο -	-0.097	-0.098	0.4	0.38		- 0.00
Amin.	0.24	-0.54	0.54	0.54	<b>&gt;-</b> -	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
<b>ს</b> -	0.1	0.65	0.64	0.64	<b>ს</b> -	0.23	0.73	0.73	0.73		
۵ -	0.023	0.64	0.64	0.64		0.19	0.77	0.77	0.77		0.50
⊢ -	0.19	0.13	0.067	0.036	<b>-</b> -	0.19	0.15	0.053	0.036		-0.30
> -	-0.11	0.46	-0.035	-0.13	> -	-0.013	0.53	-0.088	-0.17		
	-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>ح</u> -	-0.017	0.76	0.76	0.76		
<b>S</b> -	0.094	-0.39	0.051	0.13	v -	0.017	-0.49	0.13	0.2		1.00
	Length -	- OB Fac	- J9+uopoJ	Codon+GC+Cost -		Length -	- O Fac	- 79+uopoO ctor	Codon+GC+Cost -		1.00