## Bacteria - Correlation of amino acid frequency and proteome factors for genetic code: Mold-protozoan-coelenterate mitochondrial and mycoplasma spiroplasma

Pearson correlation coefficient						Spearman correlation coefficient				1.00
Σ-	-0.28	-0.1	0.17	0.16	∑ -	-0.22	-0.16	0.18	0.17	1.00
> -	0.18	0.27	-0.042	-0.097	≥ -	0.22	0.37	-0.0047	-0.068	
O -	-0.11	-0.077	0.12	0.13	U -	-0.062	-0.15	0.2	0.19	- 0.75
٥ -	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
ш-	0.094	-0.46	0.45	0.45	ш-	0.035	-0.61	0.61	0.61	- 0.50
Ι-	0.022	0.26	-0.032	-0.086	ェ -	0.088	0.34	0.0091	-0.047	
⊻ -	-0.1	-0.82	0.83	0.83	⊻ -	-0.22	-0.83	0.83	0.83	- 0.25
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	0.33	σ-	-0.097	-0.098	0.4	0.38	0.00
Amino acid Y Q	0.24	-0.54	0.54	0.54	<b>&gt;</b> -	0.1	-0.62	0.62	0.62	- 0.00
<b>-</b> -	-0.082	-0.82	0.82	0.82		-0.16	-0.86	0.86	0.86	
<b>V</b> -	0.0085	0.83	0.82	0.82	∢ -	0.098	0.86	0.86	0.86	0.25
ტ -	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	- ۵	0.19	0.77	0.77	0.77	0.50
⊢ -	0.19	0.13	0.067	0.036	⊢ -	0.19	0.15	0.053	0.036	0.50
> -	-0.11	0.46	-0.035	-0.13	> -	-0.013	0.53	-0.088	-0.17	
<b>-</b> -	-0.057	0.15	-0.14	-0.14		-0.053	0.21	-0.21	-0.21	0.75
<b>~</b> -	-0.073	0.69	0.7	0.7	∝ -	-0.017	0.76	0.76	0.76	
S -	0.094	-0.39	0.051	0.13	v -	0.017	-0.49	0.13	0.2	1.00
	Length -	- GC -	- Codon+GC -	Codon+GC+Cost -		Length -	- J9 -	Codon+GC -	Codon+GC+Cost -	1.00

Factor

Factor