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-	-0.23	-0.084	0.16	0.15	Σ-	-0.21	-0.14	0.16	0.15	
> -	0.28	0.26	-0.068	-0.12	≥ -	0.37	0.38	-0.03	-0.091	
<b>U</b> -	-0.038	-0.05	0.1	0.11	U -	0.054	-0.1	0.18	0.17	- 0.75
<b>O</b> -	0.16	0.3	-0.12	-0.18	Δ-	0.27	0.34	-0.23	-0.26	
ш-	-0.069	-0.22	0.14	0.18	ш -	-0.12	-0.28	0.12	0.19	- 0.50
ш-	0.14	-0.44	0.44	0.44	ш-	0.1	-0.58	0.58	0.58	- 0.50
エ -	0.069	0.28	-0.057	-0.11	ェ -	0.14	0.35	-0.011	-0.065	
⊻ -	-0.097	-0.83	0.84	0.84	⊻ -	-0.23	-0.85	0.85	0.85	- 0.25
<b>z</b> -	0.11	-0.76	0.77	0.77	z -	-0.012	-0.83	0.83	0.83	
Amino acid Y Q	-0.05	-0.033	0.32	0.31	ο -	-0.063	-0.095	0.38	0.36	- 0.00
Amine Y	0.25	-0.53	0.54	0.54	<b>&gt;</b> -	0.11	-0.61	0.61	0.61	- 0.00
	-0.1	-0.82	0.82	0.83		-0.17	-0.86	0.86	0.86	
∢ -	0.03	0.83	0.83	0.83	∢ -	0.12	0.87	0.87	0.87	0.25
<b>ს</b> -	0.12	0.66	0.65	0.65	<b>ს</b> -	0.27	0.72	0.72	0.72	
₫ -	0.056	0.63	0.63	0.63	<u>a</u> -	0.22	0.77	0.77	0.77	0.50
⊢ -	0.19	0.13	0.067	0.036	⊢ -	0.18	0.14	0.059	0.043	0.50
> -	-0.095	0.47	-0.019	-0.12	> -	-0.039	0.52	-0.077	-0.15	
<b>-</b> -	-0.047	0.14	-0.14	-0.14		-0.034	0.22	-0.22	-0.22	0.75
∝ -	-0.044	0.73	0.73	0.73	∝ -	0.016	0.8	0.8	0.8	
S -	0.12	-0.38	0.052	0.13	v -	0.059	-0.49	0.13	0.2	1.00
	Length -	- OB	Codon+GC	Codon+GC+Cost -		Length -	- Og Fac	- Ogon+GC	Codon+GC+Cost -	1.00
Factor						Factor				