Bacteria - Correlation of amino acid frequency and proteome factors for genetic code: Candidate division sr1 and gracilibacteria

Pearson correlation coefficient					J	Spearman correlation coefficient					
∑ -	-0.23	-0.084	0.16	0.15	Σ-	-0.21	-0.14	0.16	0.15		1.00
≥ -	0.28	0.26	0.24	0.23	≥ -	0.37	0.38	0.34	0.33		
U -	-0.038	-0.05	0.1	0.11	υ -	0.054	-0.1	0.18	0.17	_	0.75
Δ-	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	エ -	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
z -	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	⊘ -	-0.063	-0.095	0.38	0.36		0.00
Amine Y	0.25	-0.53	0.54	0.54	>-	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
ს -	0.12	0.66	0.65	0.65	ს -	0.27	0.72	0.72	0.72		
₾ -	0.056	0.63	0.63	0.63	₫ -	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		
	-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	ᄯ -	0.016	8.0	0.8	8.0		
S -	0.12	-0.38	0.052	0.13	თ -	0.059	-0.49	0.13	0.2		-1.00
	Length -	- OB Fac	- Ogou+ etor	Codon+GC+Cost -		Length -	- O Fac	-)9+uopo)	Codon+GC+Cost -		1.00