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

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		Pearson correlation coefficient Spearman correlation coefficient							ent		- 1.0
∑ -	-0.28	-0.1	0.17	0.16	∑ -	-0.22	-0.16	0.18	0.17		1.0
≥ -	0.18	0.27	0.25	0.25	≥ -	0.22	0.37	0.35	0.34		
U -	-0.11	-0.077	0.12	0.13	υ -	-0.062	-0.15	0.2	0.19	-	0.
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		•
ш-	0.094	-0.46	0.45	0.45	ш-	0.035	-0.61	0.61	0.61		0.
エ -	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.
z -	0.11	-0.76	0.77	0.77	z -	-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		_
<u>}</u> } ≻ -	0.24	-0.54	0.54	0.54	> -	0.1	-0.62	0.62	0.62	_	0
	-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	_	_
ט -	0.1	0.65	0.64	0.64	ს -	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		
⊢ -	0.19	0.13	0.067	0.036	⊢ -	0.19	0.15	0.053	0.036	_	
> -	-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	_	_
ㄸ -	-0.073	0.69	0.7	0.7	~ -	-0.017	0.76	0.76	0.76		
ω -	0.094	-0.39	0.051	0.13	v -	0.017	-0.49	0.13	0.2		
	Length -	- J9	Codon+GC -	Codon+GC+Cost -		Length -	- 25	Codon+GC -	Codon+GC+Cost -		_

Factor

Factor