Superkingdom	Phylum	GC Content	Type	Functional group	% in proteins
Archaea	Crenarchaeota	Low	disord	Isp	4.01
				Cp	1.23
				Me	0.80
				Pc	1.69
				N.C.	4.52
			order	Isp	62.60
				Cp	73.68
				Me	75.00
				Pc	73.46
				N.C.	70.34
		Medium	disord	Isp	3.65
				Cp	2.01
				Me	0.77
				Pc	1.24
				N.C.	3.56
			order	Isp	64.47
				Cp	74.77
				Me	75.40
				Pc	74.36
				N.C.	71.17
		High	disord	Isp	8.16
				Cp	4.03
				Me	3.44
				Pc	4.63
				N.C.	6.64
			order	Isp	58.53
				Cp	68.58
				Me	68.41
				Pc	64.46
				N.C.	61.20
	Euryarchaeota	Low	disord	Isp	6.96
				Cp	6.56
				Me	2.37
				Pc	2.69
				N.C.	9.16
			order	Isp	56.70
				Cp	63.15
				Me	68.20
				Pc	68.33
				N.C.	60.44
		Medium	disord	Isp	6.94
				Cp	5.64
				Me	2.19
				Pc	4.35
				N.C.	7.24
			order	Isp	58.60
				Cp	63.98
				Me	70.15
				Pc	65.43
				N.C.	58.97
		High	disord	Isp	18.67
				Cp	18.06
				Me	10.07
				Pc	12.06
				N.C.	19.37
			order	Isp	39.47
				Cp	44.77
				Me	53.74
				Pc	52.48
				N.C.	44.52
	Nanoarchaeota	Low	disord	Isp	4.06

Superkingdom	Phylum	GC Content	Type	Functional group	% in proteins
Archaea	Nanoarchaeota	Low	disord	Cp	3.10
				Me	1.02
				Pc	1.68
				N.C.	4.96
			order	Isp	65.28
				Cp	77.24
				Me	76.53
				Pc	74.30
				N.C.	73.98
Bacteria	Aquificae	Medium	disord	Isp	7.89
				Cp	6.41
				Me	2.59
				Pc	3.43
				N.C.	7.55
			order	Isp	55.84
				Cp	64.95
				Me	69.41
				Pc	65.78
				N.C.	59.60
	Bacteroidetes/Chlorobi	Medium	disord	Isp	5.78
				Cp	4.29
				Me	1.43
				Pc	3.55
				N.C.	5.55
			order	Isp	61.35
				Cp	66.40
				Me	69.63
				Pc	68.12
				N.C.	58.34
		High	disord	Isp	21.29
				Cp	21.52
				Me	10.17
				Pc	14.37
				N.C.	23.68
			order	Isp	38.69
				Cp	43.85
				Me	55.02
				Pc	50.48
				N.C.	38.00
	Chlamydiae/Verrucomicrobia	Medium	disord	Isp	4.50
				Cp	7.59
				Me	1.54
				Pc	4.44
				N.C.	15.01
			order	Isp	61.34
				Cp	64.03
				Me	72.28
				Pc	67.45
				N.C.	52.65
	Chloroflexi	Medium	disord	Isp	5.48
				Cp	3.99
				Me	1.25
				Pc	2.89
				N.C.	9.47
			order	Isp	61.81
				Cp	67.02
				Me	72.74
				Pc	67.21
				N.C.	53.19
	Deinococcus-Thermus	High	disord	Isp	15.07
				Cp	12.97

Superkingdom	Phylum	GC Content	Type	Functional group	% in proteins
Bacteria	Deinococcus-Thermus	High	disord	Me	5.91
				Pc	9.09
				N.C.	18.96
			order	Isp	50.92
				Cp	56.42
				Me	65.46
				Pc	58.85
				N.C.	48.36
	Firmicutes	Low	disord	Isp	6.89
				Cp	11.75
				Me	2.69
				Pc	6.39
				N.C.	16.31
			order	Isp	59.19
				Cp	56.78
				Me	67.72
				Pc	63.68
				N.C.	47.96
		Medium	disord	Isp	5.91
				Cp	7.50
				Me	1.81
				Pc	5.01
				N.C.	11.05
			order	Isp	59.67
				Cp	60.68
				Me	69.82
				Pc	65.29
				N.C.	52.04
		High	disord	Isp	10.76
				Cp	10.35
				Me	3.98
				Pc	6.58
				N.C.	15.18
			order	Isp	58.35
				Cp	62.36
				Me	71.63
				Pc	66.31
				N.C.	55.16
	Planctomycetes	Medium	disord	Isp	13.39
				Cp	13.39
				Me	5.77
				Pc	10.50
				N.C.	18.84
			order	Isp	50.40
				Cp	51.92
				Me	59.12
				Pc	54.18
				N.C.	39.89
	Spirochaetes	Low	disord	Isp	6.93
				Cp	8.64
				Me	1.31
				Pc	3.96
				N.C.	7.74
			order	Isp	60.77
				Cp	63.35
				Me	72.41
				Pc	71.39
				N.C.	63.19
		Medium	disord	Isp	7.62
				Cp	6.76
				Me	2.31

Superkingdom	Phylum	GC Content	Type	Functional group	% in proteins
Bacteria	Spirochaetes	Medium	disord	Pc	6.83
				N.C.	7.65
			order	Isp	56.76
				Cp	60.81
				Me	65.92
				Pc	61.01
				N.C.	52.76
	Thermotogae	Medium	disord	Isp	7.26
				Cp	7.12
				Me	1.68
				Pc	2.78
				N.C.	4.36
			order	Isp	58.85
				Cp	64.25
				Me	71.95
				Pc	68.17
				N.C.	63.79