

Concordance factors for branch ID 85

Values are percentages, numbers in brackets are bootstrap 95% CIs

	Ψ_1	Ψ_2	Ψ_3	Ψ_4
gene (n=618)	12.460 (9.709, 14.887)	5.990 (4.207, 7.609)	3.720 (2.427, 5.178)	77.830 (74.757, 81.392)
site (n=159.46)	35.380 (27.593, 42.017)	34.180 (26.339, 41.390)	30.440 (23.830, 37.627)	0.000 (0.000, 0.000)
quartet (n=618)	35.930 (32.201, 39.644)	36.160 (32.362, 40.129)	27.900 (24.434, 31.392)	0.000 (0.000, 0.000)

Concordance or Discordance factor (%)

