

Concordance factors for branch ID 86

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

	Ψ_1	Ψ_2	Ψ_3	Ψ_4
gene (n=531)	46.140 (42.185, 50.471)	4.330 (2.637, 6.215)	4.140 (2.637, 6.026)	45.390 (41.050, 49.534)
site (n=193.02)	44.580 (37.820, 51.808)	30.330 (23.832, 36.784)	25.090 (19.169, 31.085)	0.000 (0.000, 0.000)
quartet (n=530.99)	60.060 (56.122, 64.036)	25.260 (21.658, 29.007)	14.690 (12.053, 17.703)	0.000 (0.000, 0.000)

Concordance or Discordance factor (%)

