Table S1: Table of coefficients for the additional model fitted with uniqueness point and the pre-uniqueness point data. All predictors were centred and scaled before fitting the model.

	Estimate	Std. Error	t value
Intercept	6.290	0.035	181.601
Log phonotactic prob.	-0.019	0.006	-3.167
Trial number	-0.056	0.002	-30.035
Nbhrd density	-0.009	0.010	-0.957
Log moving average response latency	0.195	0.006	31.067
Temporal uniqueness point	0.059	0.004	16.213
N. morph. parses	-0.012	0.003	-3.684
Phonological uniqueness point	-0.057	0.007	-8.218
Log phonotactic prob.*N. morph parses	-0.001	0.004	-0.254

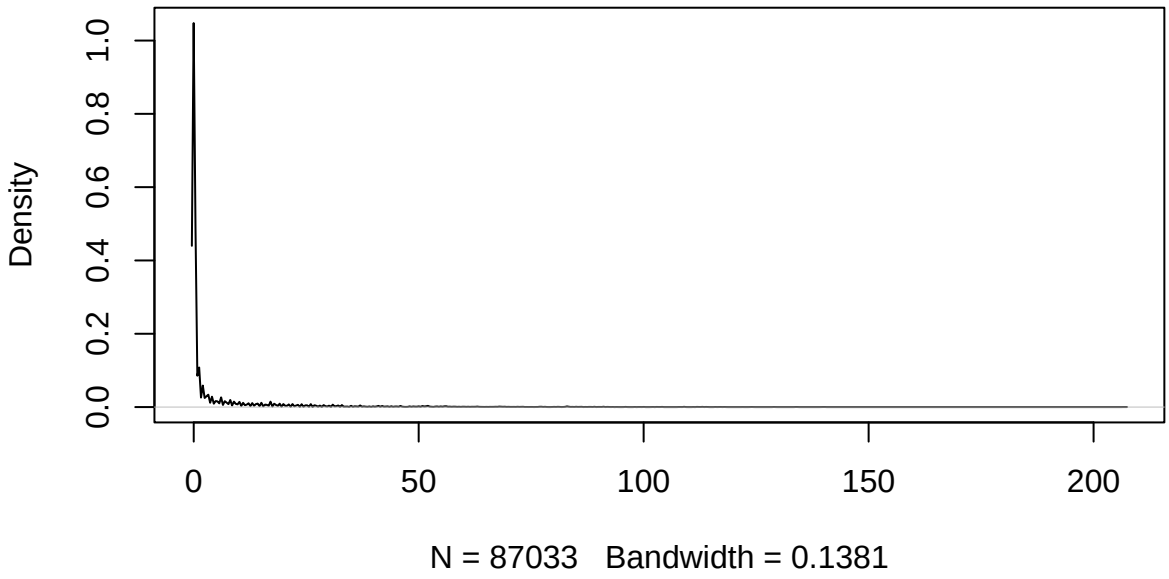

Figure S1: Density plot of the standard calculation of phonological neighbourhood density.

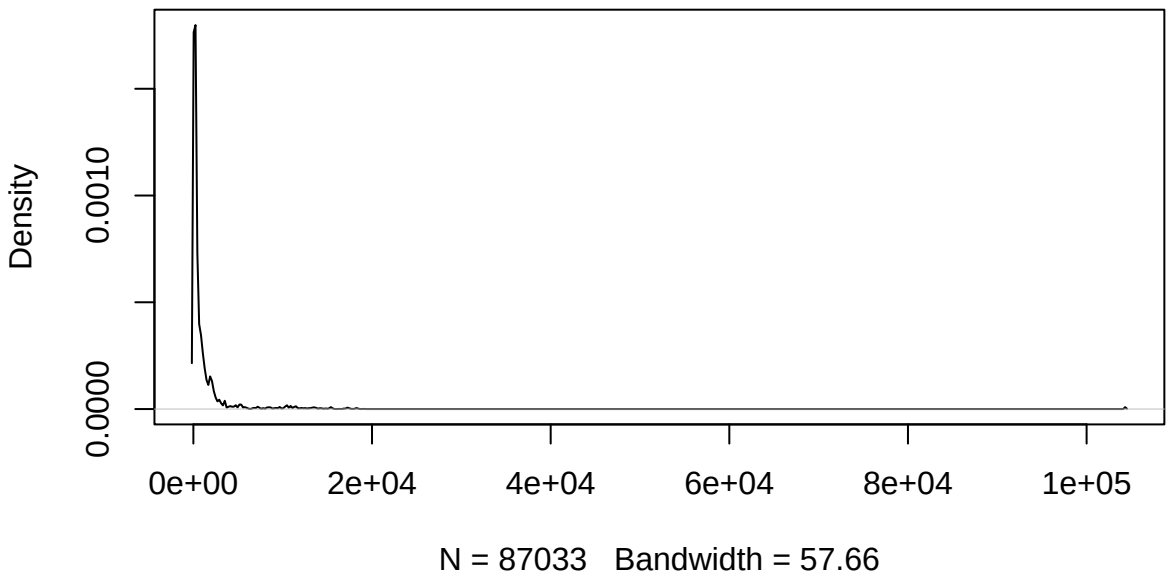

Figure S2: Density plot of the pre-uniqueness point calculation of phonological neighbourhood density.

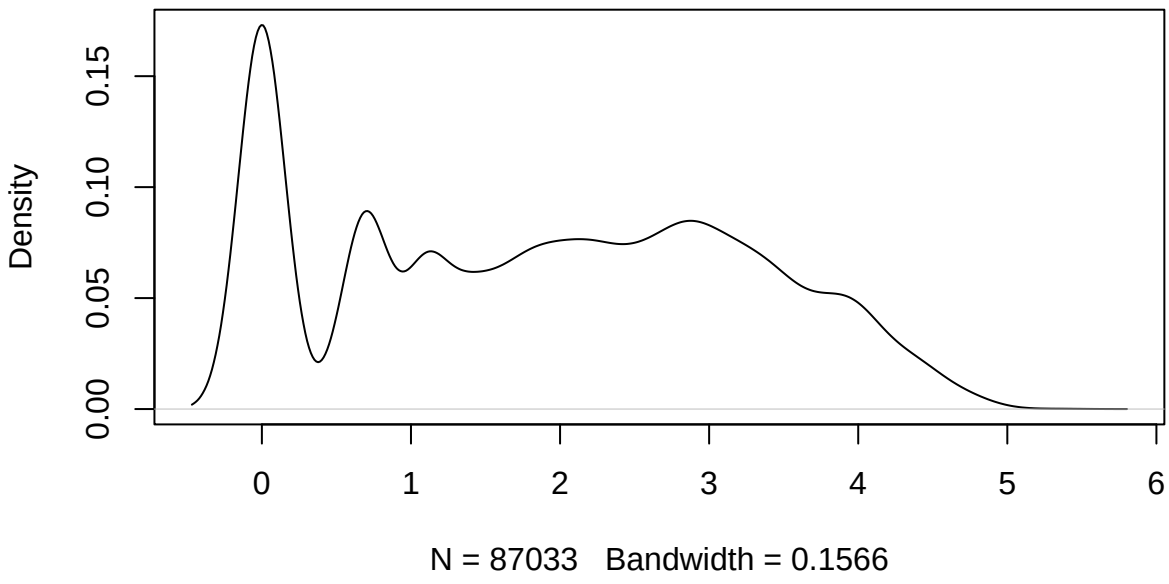

Figure S3: Density plot for the log of the standard calculation of phonological neighbourhood density.

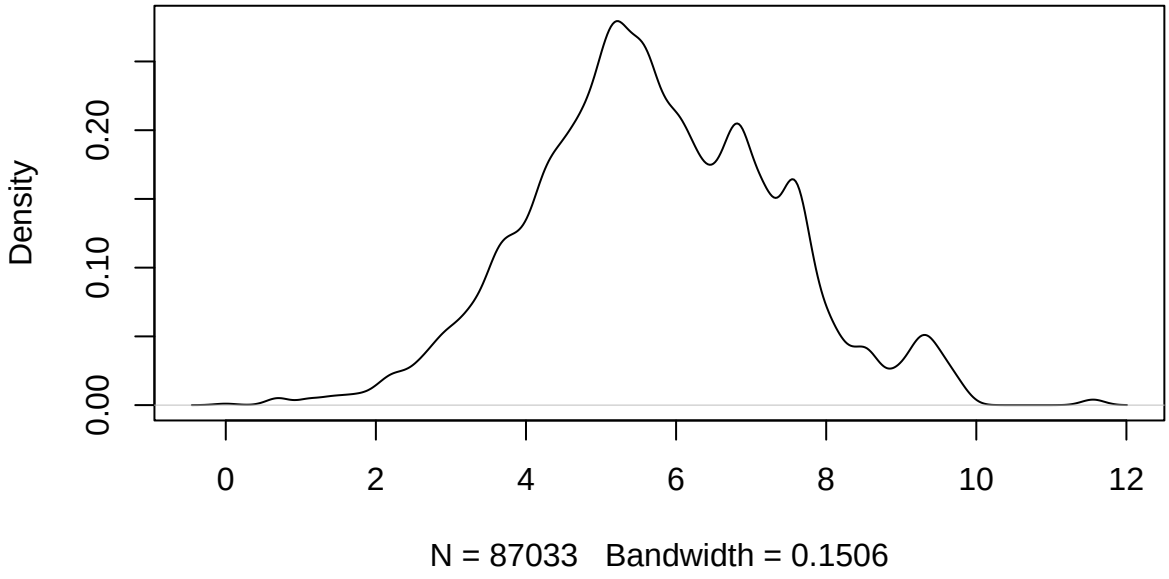

Figure S4: Density plot of the log of the pre-uniqueness point calculation of phonological neighbourhood density.

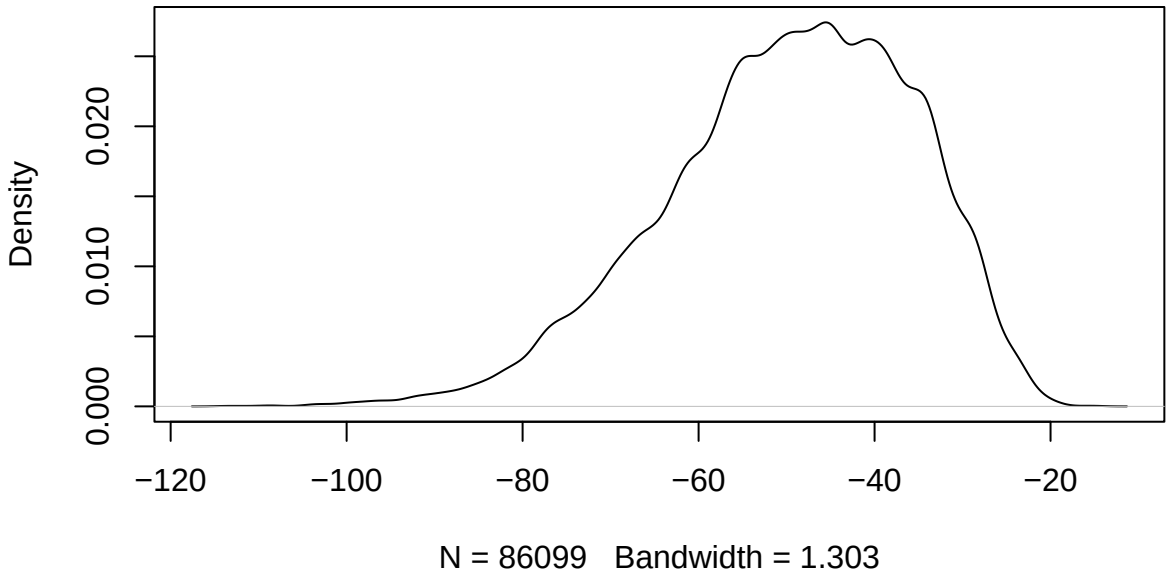

Figure S5: Density plot for standard calculation of log phonotactic probability. Data were additionally subset from the pre-uniqueness point analysis data to remove all 0 values to avoid taking log(0).

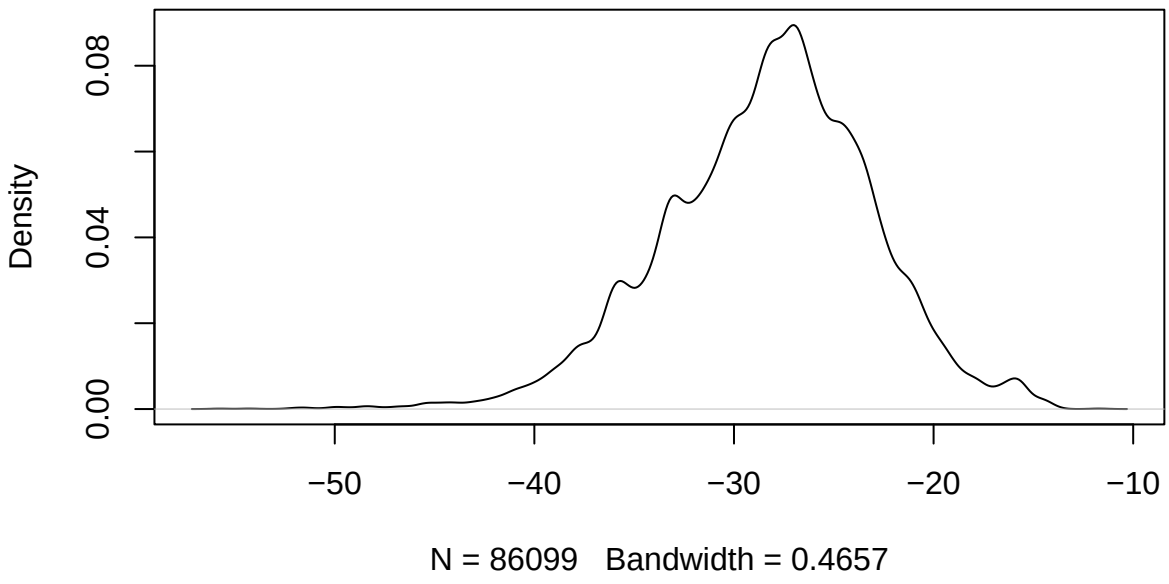

Figure S6: Density plot for pre-uniqueness point calculation of log phonotactic probability.

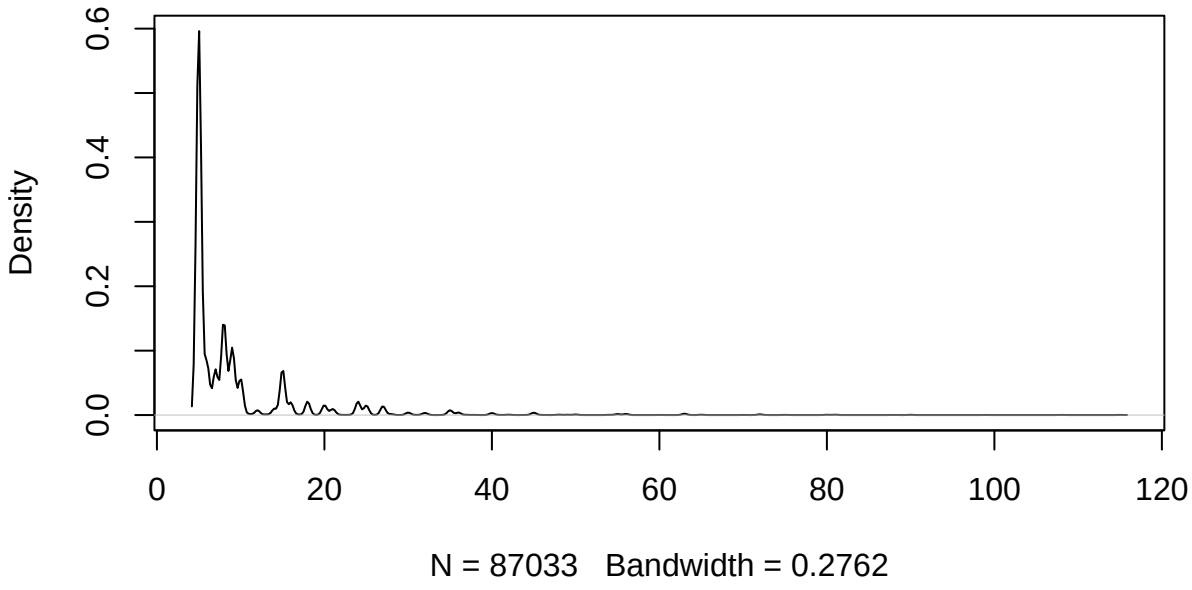

Figure S7: Density plot for standard calculation of number of morphological parses.

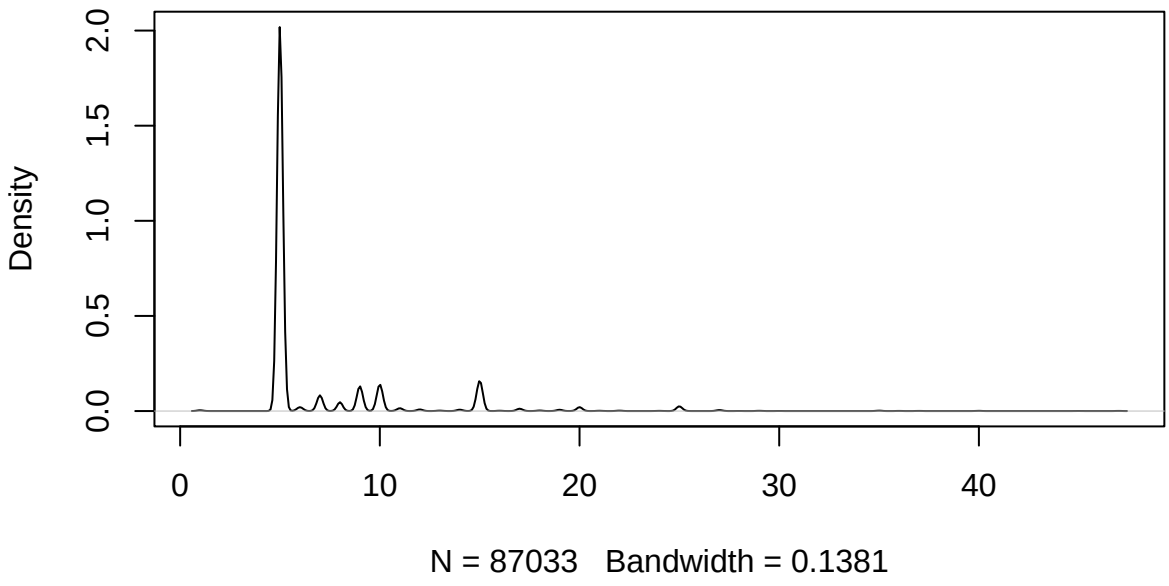

Figure S8: Density plot for pre-uniqueness point calculation of number of morphological parses.