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How network structure shapes languages: disentangling the factors driving variation in communicative agents

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ABSTRACT

Languages show substantial variability between their speakers, but it is currently unclear how the structure of the communicative network contributes to the patterning of this variability. While previous studies have highlighted the role of network structure in language change, the specific aspects of network structure that shape language variability remain largely unknown. To address this gap, we developed a Bayesian agent-based model of language evolution, contrasting between two distinct scenarios: language change and language emergence. By isolating the relative effects of specific global network metrics across thousands of simulations, we show that global characteristics of network structure play a critical role in shaping inter-individual variation in language, while intra-individual variation is relatively unaffected. We effectively challenge the long-held belief that size and density are the main network structural factors influencing language variation, and show that path length and clustering coefficient are the main factors driving inter-individual variation. In particular, we show that variation is more likely to occur in populations where individuals are not well-connected to each other. Additionally, variation is more likely to emerge in populations that are structured in small communities. Our study provides potentially important insights into the theoretical mechanisms underlying language variation.

1. INTRODUCTION

Variation is ubiquitous not only in the natural world, but also in culture. There are intriguing patterns of variation ranging from within to between individuals (in most biological, psychological, cognitive and linguistic characteristics) and groups (most obvious in various aspects of culture). In the language domain, one can think about how two different persons would pronounce the phoneme /r/ in different ways, even if they speak the same language (Sankoff & Blondeau, 2007), or how grammatical constructions vary between English dialects (Trudgill & Chambers, 2017), or even about lexical variation (different use of 'firefly' and 'lightning bug' words in English, for example). Why this variation exists, why it is patterned the

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44 way it is, and what processes govern its dynamics are extremely important and non-trivial
45 scientific questions.

46 Language variation arises from divergences in the evolution of linguistic features over time
47 among different populations. It is customary to distinguish the emergence of an innovation
48 (the so-called “initiation”) from the dynamics of its diffusion in the speaker community
49 (“actuation”) (Croft, 2008; Dediu & Moisik, 2019; Solé & Vives, 2012; Stevens & Harrington,
50 2014; Yu, 2013). Initiation is usually seen as an individual-level process, while actuation is
51 essentially a community-level process involving competing linguistic variants (Blythe & Croft,
52 2012; Fagyal et al., 2010). These processes can take place in two main scenarios: when the
53 language used by the population is being created *de novo* (language *emergence* scenario, as
54 is arguably the case for emergent sign languages such as Al-Sayyid Bedouin Sign Language
55 and Kaka Kolok; Zeshan & de Vos, 2012), and when a language is already in use and
56 undergoing change (language *change* scenario, by far the most frequent and studies case).
57 The selection of variants during the actuation phase can be influenced by many different
58 factors, such as aspects of the environment (e.g., the effect of environment type on the
59 phonemic inventories of languages; Everett et al., 2015; Maddieson & Coupé, 2015), cultural
60 and historical factors (degree of contact with outsiders, or number of L2 speakers; Lupyan &
61 Dale, 2010), or factors related to the structure of the network of linguistic interaction itself
62 (Wray & Grace, 2007; Trudgill, 2011a). Indeed, interactions between individuals in human
63 societies are embedded in social networks with specific characteristics (Beckner et al., 2009;
64 Newman & Park, 2003), which would constrain the transmission of linguistic features
65 (Chambers, 1995). While all human networks tend to share a few features, such as a small
66 average path length, social network structures exhibit a great diversity across populations
67 (Newman, 2003; Nichols, 1992). Our goal in this paper is to disentangle what characteristics
68 in network structures are instrumental to understand how variation unfolds through evolution.

69 It is, however, challenging to carefully disentangle the relative effects of network structure from
70 those of the other factors involved, prompting several researchers to introduce experimental,
71 computer modeling and observational paradigms. The rich tradition of sociolinguistic studies
72 uses fine-grained methods that provide insight into the role of network structure (among many
73 other factors) on language change (Labov, 1972, 2001, 2010; Milroy & Milroy, 1985; Yu, 2013).
74 Following a different approach, Raviv et al. (2019, 2020) tackle this question with an
75 experimental approach involving the learning and transmission of an artificial miniature
76 language within a carefully controlled network structure. Network structure was even put
77 forward to have an influence on cultural evolution (Derex & Boyd, 2016; Migliano et al., 2020).
78 These laboratory experiments emphasize the role of population size and network structure on
79 language variability. However, despite their inherent advantages, laboratory experiments are
80 strongly constrained in terms of population size, network structure and aspects of language
81 considered, limiting the direct extrapolation of these results to understanding the dynamics of
82 actual linguistic communities.

83
84 Artificial multi-agent models do provide ways to address some of these issues, in particular
85 concerning the number of linguistic agents, the structure of their communicative networks, and
86 the usually long time scale necessary for innovations to spread in large populations. Some
87 studies based on the use of such models highlight the role of population structure in shaping
88 convergence (Lee et al., 2015), the formation of categories (Zubek et al., 2017; Gong et al.,
89 2012), the diffusion of innovations (Ke et al., 2008), and linguistic structure (Reali et al., 2014;

90 Lou-Magnuson et al., 2018). Despite using different communication algorithms and
91 addressing various aspects of language, these models share a common approach: they all
92 compare different types of pre-existing network structures (such as fully-connected, star, or
93 small-world networks) and then observe whether language evolves differently in these distinct
94 network structures. For example, Lee et al. (2015) examined populations of learners in fully-
95 connected, linear, Neumann lattice, and bridge network structures and showed that these
96 distinct network typologies result in different patterns of grammar convergence. Departing from
97 these pre-existing network structures, in the field of cultural evolution, Kobayashi et al. (2016)
98 aimed to untangle two key metrics: population size and social connectedness. They propose
99 that social connectedness might exert a stronger influence on the level of culture compared to
100 population size. While most of these models address the dynamics of language change and
101 do not specifically explore language variation, Mudd (2022) and Meir et al. (2012) also
102 emphasize the role of social structure on lexical variation in emergent sign languages. Notably,
103 Mudd (2022) developed computational models grounded in observations from the Kata Kolok
104 community, focusing on various aspects of their social structure, such as connectivity and size.
105 To explain why some sign languages witness more lexical variation than others, she proposed
106 that population size is an important factor influencing lexical variability in these languages.
107 However, while it appears that network structure influences language in both scenarios of
108 emergence and change, it remains unclear whether the impact of network structure differs in
109 these two contexts.

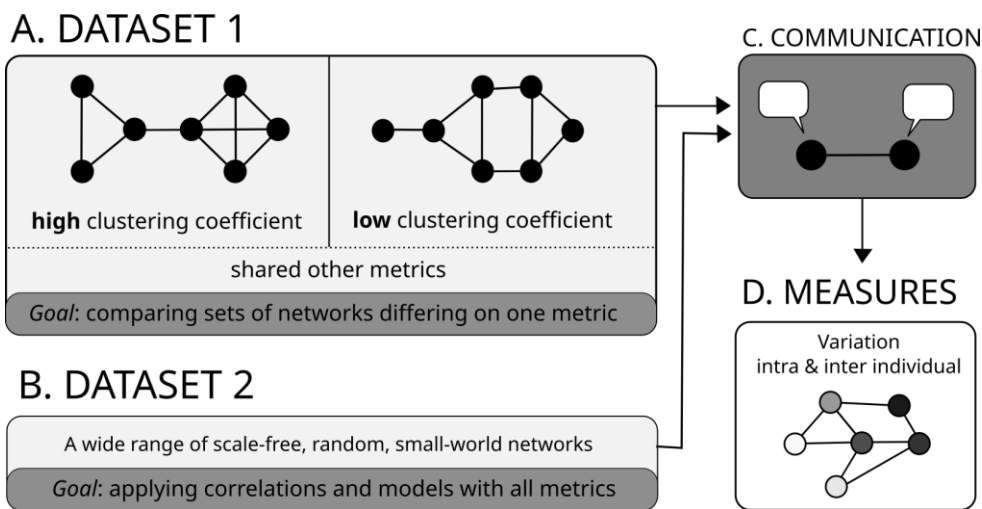
110
111 Thus, a wide range of approaches strongly suggests that network structure likely influences
112 the evolution of many linguistic features. But what exactly is meant by “network structure”?
113 The networks used in such studies are either networks observed in the real world, or are
114 automatically generated with specific properties (such as, for example, being fully connected,
115 scale-free, or small-world). Any network, whether “natural” or “artificial”, can be described
116 using a set of measures (called *metrics*) that summarize various aspects of the network,
117 varying between the relatively global to the quite local (see Illustration 1). As Raviv et al. (2020)
118 observed, it is very difficult to tease apart the contribution of each network characteristic in
119 real-world observational networks. In other words, comparing real-world populations raises
120 the question of understanding which network structural factors are involved in the actuation of
121 linguistic innovations, and in which manner. In contrast, multi-agent models allow a much finer
122 degree of control over these factors. As a consequence, most agent-based modeling studies
123 to date have investigated the separate role of various characteristics of the communicative
124 network. However, this approach has two main limitations: first, when focusing on specific
125 factors in isolation, it is very difficult to compare their relative influence, and to understand the
126 interactions between them. Secondly, working with automatically generated networks has not
127 led to a fine-grained understanding of which network metrics behind these networks actually
128 affect the dynamics of actuation. Indeed, there are intrinsic relationships between various
129 metrics resulting from the type of network being generated. As an example, if communities
130 structured as scale-free networks produce more stable variation than those structured as
131 small-world networks, is this due to their lower centrality, lower clustering coefficient, or any
132 other metric that intrinsically differs between these two types of networks?

133 Addressing these questions makes it essential to disentangle and bring a theoretical basis to
134 the question of what aspects of network structure do, in fact, shape language. This has
135 theoretical implications for understanding how languages adapt to fit their social environments
136 and how network structure shapes language. In turn, it will help understand language

137 differentiation and help explain the great diversity observable in the world’s languages. To
 138 answer these questions, we introduce here a multi-agent model based on the one in
 139 Josserand et al. (2021). The agents use a Bayesian model of language learning in artificially
 140 generated static networks where we control various metrics, and, across thousands of runs,
 141 we effectively isolated the relative influence of several global metrics (average shortest path
 142 length, clustering coefficient, global assortativity, degree distribution, and size) and of local
 143 characteristics of the agents (measures of centrality and clustering coefficients). See
 144 Illustration 1 for an example. We found that these metrics have different contributions to the
 145 evolution of language in this model: while intra-individual variation is not shaped by network
 146 structure, inter-individual variation is strongly affected by the path length and the clustering
 147 coefficient.

148 The paper is structured as follows: in the Data and Methods, we present our Bayesian agent-
 149 based model, along with the networks used, the metrics under consideration, and the overall
 150 procedure of this analysis. In the Results, we investigate whether some of these metrics affect
 151 inter- and intra-individual variability in different types of networks. We close by discussing the
 152 limitations and implications of our findings, and suggest several future directions of study.

153



154

155 *Illustration 1. A. Dataset 1 comprises sets of networks differing on only one metric (here, clustering*
 156 *coefficient, but we applied it to every metric in Table 1). B. Dataset 2 comprises a wide range of different*
 157 *networks: scale-free, small-world, random networks of different sizes and parameters. C. Throughout*
 158 *our study, agents interact using Bayesian communication for an extended period of time. D. Finally, we*
 159 *investigate the language variability within and between agents in the network. To do so, we compare*
 160 *language variability in networks that differ based on global metrics, using dataset 1. We also explored*
 161 *the networks in dataset 2 by computing the correlations between different metrics and measuring their*
 162 *relative contributions to variability in models. More information about the differences between the two*
 163 *datasets will be presented in Data and Methods section.*

164 2. DATA AND METHODS

165 Here we develop a Bayesian model of language evolution (Griffiths & Kalish, 2007) based on
 166 published work (Josserand et al., 2021), where agents exchange linguistic messages as
 167 constrained by a communicative network. Briefly, the network’s nodes represent the agents,

168 which all have an internal representation of language, here reduced to a single feature. The
169 agents that are connected via undirected edges in the network are called “neighbors” and can
170 “talk” to each other. Upon “hearing” their neighbors’ utterances, an agent updates its internal
171 representation of language to take into account the newly “heard” data using Bayes’ rule. We
172 study how the language changes after multiple rounds of such interactions, and, in particular,
173 the manner in which different network structures affect these dynamics in two types of
174 scenarios: language emergence (agents interact without any initial language exposure) and
175 language change (agents interact with an initial language exposure).

176 1.1 The language

177 The language in our model is a multinomial feature that can take one of k exclusive possible
178 values. This simple representation is arguably adequate to approximate numerous aspects of
179 human language, such as lexical choices (using one of several words or expressions to
180 describe a given concept), allophony (different actual sounds for the same phoneme as
181 conditioned by the phonetic environment, socio-linguistic/dialectal choices, or in free
182 variation), or morpho-syntactic structures (different ways of organizing the sentence). Here,
183 we call an *utterance* a token drawn from this multinomial distribution. For example, if the
184 feature refers to the possible allophones of “r” in English, the possible utterances include the
185 alveolar trill [r], the approximant [ɹ], and the retroflex approximant [ɻ], among others. Abstractly,
186 we can represent the feature through the vector of its possible utterances $u = \{u_1, u_2, \dots, u_k\}$.
187 We fixed the number of utterances k to 10 here, but simulations ran with different values of k
188 led to similar conclusions.

189 While we also generalized our feature to a continuous one (which can represent, for example,
190 very fine-grained measurements of vowel height or voice-onset-time (VoT); presented for the
191 Dataset 1 in the **Supplementary Materials** in Part 3.2.2), we decided to focus here on the
192 case of the discrete valued feature. Indeed, using a continuous representation instead of the
193 discrete one does not alter our conclusions, but presenting both would lead to clutter and
194 unnecessary complexity in the presentation of the results.

195 1.2 How do agents represent language?

196 Agents are Bayesian learners (Griffiths & Kalish, 2007). In this framework, there is a fixed
197 “universe” of languages, U , and each possible language h from this universe ($h \in U$) has a
198 certain probability p of being “the real one” from the agent’s “subjective” point of view. Before
199 being exposed to any utterance, each agent has an *a priori* “belief” concerning the probabilities
200 of each possible language, denoted $p(h)$ and known as the *a priori probability distribution* of
201 the languages (or the *prior*). However, each time an agent “hears” an utterance, denoted as
202 d , it updates its subjective distribution of the probabilities of the languages in light of this data
203 using Bayes’ theorem: $p(h|d) = \frac{p(d|h)p(h)}{p(d)}$. Here, $p(d|h)$ is the likelihood of observing data d
204 given that the language is h , while $p(h|d)$ is the updated belief, or *posterior*, i.e. the probability
205 of the language being h after both the *prior* and the observed data d are considered. The
206 denominator $p(d)$ acts as a normalization factor and is not of immediate concern here. While
207 this mechanism is extremely general, we apply it here in a very particular and relatively simple
208 setting.

209 More precisely, an agent’s internal representation of language is given by a vector $\mathbf{p} =$
210 (p_1, p_2, \dots, p_k) , where each component p_i represents the probability of the i^{th} utterance u_i ; as
211 the utterances are mutually exclusive and exhaust the space of possible utterances for this
212 features, $\sum_{i=1}^k p_i = 1$. An agent has an *a priori* distribution that represents its initial belief in
213 the probability of each utterance, which leads naturally to a *multinomial distribution* that
214 represents the likelihood. More precisely, given n such utterances “heard” independently by
215 an agent, the random variable $\mathbf{X} = (X_1, X_2, \dots, X_k)$ distributed as *Multinomial*(n, \mathbf{p}) gives the
216 expected number of times each possible utterance u_i appears. Here, the multinomial
217 distribution models how frequent any one of the k possible utterances should be among the n
218 “heard” independent ones, if their probabilities are given by the vector \mathbf{p} .

219 To compute the posterior distribution after having seen the data, it is possible to use the very
220 computationally expensive Markov Chain Monte Carlo technique. However, in particular
221 cases, one may choose specific conjugate priors, which drastically simplify the application of
222 Bayes’ theorem. The conjugate prior of a multinomial distribution is given by the *Dirichlet*
223 *distribution*, which is defined by a vector of k parameters $\boldsymbol{\alpha} = (\alpha_1, \alpha_2, \dots, \alpha_k) \in R^k$. These
224 parameters α_i are related to the probability p_i of each utterance u_i , so that the vector of
225 probabilities $\mathbf{p} = (p_1, p_2, \dots, p_k)$ is distributed as *Dir*($\boldsymbol{\alpha}$). The prior and posterior distributions of
226 probabilities are modeled by Dirichlet distributions and, thus, characterized by prior (denoted
227 here as $\boldsymbol{\alpha}$) and posterior (denoted here as $\boldsymbol{\alpha}'$) Dirichlet parameters. Using Bayes’ theorem,
228 one can update the prior distribution *Dir*($\boldsymbol{\alpha}$) into the posterior distribution *Dir*($\boldsymbol{\alpha}'$) after seeing a
229 number n of independent utterances (the data, d): it reduces to trivial arithmetic: $\alpha'_i = \alpha_i + X_i$,
230 where, as above, X_i gives the number of times utterance u_i was “heard” in the data d . Thus,
231 this reduces to a simple model where the parameters of the Dirichlet distribution reflect the
232 number of “heard” utterances and the prior belief. It is important to note that, as expected for
233 Bayesian models, both the prior beliefs and the observed data matter. Seeing enough data
234 should be able to overwhelm even very strong *a priori* beliefs and the order in which the
235 utterances in the data arrive does not matter. This generalizes the prior model studied in
236 (Josserand et al., 2021) by allowing more than two possible utterances (i.e., from a Bayesian
237 model with a binary feature using Binomial/Beta conjugate distributions, to a multinomial
238 feature using Multinomial/Dirichlet conjugate distribution).

239 **1.3 The initial language exposure: emergence vs. change**

240 We considered two types of agent populations, differentiated by their “initial language
241 exposure”, which represents the type of scenario (language emergence versus language
242 change). In the populations in a language emergence scenario, the agents are born *without*
243 initial language exposure. Such agents are naïve regarding the language, with all possible
244 utterances having the same probability of occurrence. This is modeled by a Dirichlet
245 distribution with all parameters α_i being equal, $\boldsymbol{\alpha} = (1, 1, \dots, 1)$. It represents the situation
246 where agents are born in a language-less community, but it may as well represent a
247 community with a language where all possible utterances are equally frequent.

248 In contrast, in the language change scenario, agents are born in a population *with* initial
249 language exposure, i.e. born in a community with a pre-existing language that favors a certain
250 utterance at the expense of the others; say, utterance 4, α_4 is 6 times more frequent than the
251 others (for various reasons, including past internally-motivated language change, language
252 contact, or top-down language policies). This situation is represented by a Dirichlet distribution

253 with parameters $\alpha = \{1, 1, 1, 6, 1, \dots, 1\}$. Given how the parameters of the Dirichlet distribution
 254 reflect the number of “heard” utterances, this can be seen as modeling the fact that agents are
 255 expecting, before they engage in communication with other agents, that the utterance u_4 is
 256 much more frequent than all the others. A value of 6 is supposed to represent an initial
 257 language with a very skewed frequency of the utterances, but not too strong (however, we
 258 also investigated the effect of using 21 instead of 6 in the **Supplementary Materials**; see part
 259 3.2.1.8).

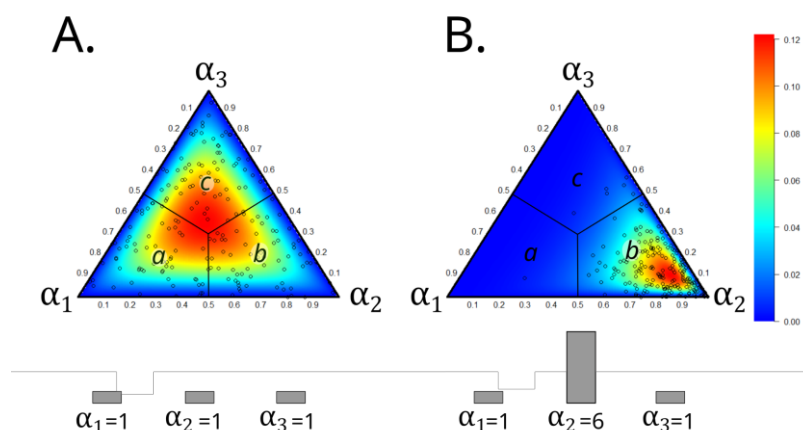
260 1.4 The communication process

261 As described above, the agents are “born” with their initial Dirichlet distribution, which
 262 represents their bias concerning the language feature. The simulation happens in discrete
 263 timesteps (*rounds*), and in each round, the agents “speak” (i.e., produce utterances) and
 264 “hear” (i.e., receive) the utterances produced by their neighbors.

265 To produce utterances, an agent uses the following procedure: first, it normalises the k
 266 parameters of its Dirichlet distribution, α to a probability vector $\mathbf{m} = (m_1, m_2, \dots, m_k)$, where
 267 $m_i = \frac{\alpha_i}{\sum_j \alpha_j}$. Then, it draws a single value $1 \leq j \leq k$ proportional to its probability m_j , such that
 268 each index j can be chosen depending on its probability relative to the other $k-1$ indices in \mathbf{m} .
 269 This is equivalent to the use of random multinomial generator, $j \sim \text{Multinomial}(\mathbf{m})$, as
 270 implemented by, for example, R’s `rmultinom(n, size, prob)`, where `n=1`, `size=1` and
 271 `prob=m`, or Stan’s `multinomial_rng(vector theta, int N)`, where `theta` is the
 272 simplex vector `theta=m` and `N=1`. Finally, this value j represents the utterance that is actually
 273 produced, namely u_j (see Illustration 2). So, we use what is known as a sampler (or ‘SAM’)
 274 strategy (Griffiths & Kalish, 2007; Josserand et al., 2021) in which the agent picks any of the
 275 possible k utterances proportional to its subjective probability (as given by the agent’s current
 276 Dirichlet distribution).

277 Conversely, upon hearing the utterances produced by its neighbors, the agent updates its
 278 current Dirichlet distribution as described above. This update consists of an addition, so that
 279 the Dirichlet hyper-parameters become $\alpha'_i = \alpha_i + X_i$, where $X = \{X_1, X_2, \dots, X_k\}$ represents
 280 the number of times each utterance appears among those that the agent just heard.

281



282

283 *Illustration 2. Representation of the Dirichlet distribution for an agent without initial language exposure*
 284 *(language emergence, panel A) and for an agent with an initial language exposure towards α_2*
 285 *(language change, panel B). For visualization purposes, we plotted here $k=3$ utterances instead of the*
 286 *$k=10$ actually used in the study. The dots represent the random draws from the Dirichlet distribution: if*
 287 *the random draw occurs in the **a** part, the agent will produce an utterance u_1 ; if it occurs in the **b** part, it*
 288 *will produce an utterance u_2 , and it will produce an utterance u_3 when drawn in the **c** region. In the*
 289 *condition without initial language exposure, random draws happen equally probably for all three*
 290 *utterances, but with initial language exposure, they occur mostly in the **b** region, resulting in many more*
 291 *u_2 utterances. The bar plot below each triangle is an alternative way of representing the values of the*
 292 *Dirichlet distribution. The color gradient reflects the density of random draws, with red denoting high*
 293 *density and blue indicating low density.*

294 1.5 The networks: types and metrics

295 The agents form the nodes of a network with a static structure, fixed for the duration of a
 296 simulation, while the edges between the nodes determine the agents that can talk to each
 297 other. Given that we focus on the structure of these networks, our discussion in this section is
 298 framed in terms of nodes and edges, instead of agents and their communicative exchanges.
 299 Here, we use three classes of network topology: random, small-world, and scale-free
 300 networks. The random networks are generated using Erdős & Rényi (1959)'s algorithm, where
 301 each node has a fixed probability of connecting with other nodes in the network. We kept only
 302 networks without any isolated components. The small-world networks are generated using the
 303 Watts-Strogatz algorithm (Watts & Strogatz, 1998): this algorithm starts with a ring of nodes,
 304 where each node is connected to a fixed number N of neighbors on either side, followed by a
 305 rewiring with a fixed probability p . This process leads to the generation of networks with many
 306 real-world properties (Kenett et al., 2018; Kitsak et al., 2010), such as the presence of short
 307 average path lengths (see Milgram (1967)'s "Six degrees of separation"). The scale-free
 308 networks are generated using the preferential attachment algorithm of Barabási et al. (2000),
 309 and exhibit a power-law degree distribution: most nodes have a limited number of neighbors,
 310 while few nodes have very many (Albert, 2005; Albert et al., 1999), resembling real offline and
 311 online social networks. The algorithm gradually adds new nodes to a network such that the
 312 probability p_i that the newly added node is connected to node i is $p_i = \frac{d_i}{\sum_j d_j}$ where d_i is the
 313 degree of node i , and the sum is over all pre-existing nodes j . We use these different classes
 314 of network topology to investigate the effects of specific network metrics (see Table 1).
 315

Name of the global network metric	Short name	Description	Mathematical description	Type of network
Average node degree	Degree	Average number of neighbors for all nodes	$D = \frac{\sum_{i=1}^n k_i}{n}$ where k is the degree of a node i and n is the total number of nodes in the network	Constant for Barabasi-Albert scale-free networks with a fixed size
Network size	Size	Total number of nodes	n	All

Average shortest path length	Path length	On average, how close (defined by the number of intermediate nodes) are two random nodes	The mean of the shortest path length between all pairs of nodes: $l_G = \frac{\sum_{i,j} dist(v_i, v_j)}{n \cdot (n-1)}$, where $dist$ is the shortest path length between two nodes	All, since we removed isolated components in random networks
Global assortativity coefficient	Assortativity	Preference for a network's nodes to attach to others that are similar (here, in terms of nodes's degree)	The mean of the local assortativity based on the node degree, where local assortativity is defined as $\rho = \frac{j(j+1)(\bar{m}_j - \mu_q)}{2M\sigma_q^2}$, where M is the number of edges in the network, j is the excess degree of a particular node and \bar{m}_j is the average excess degree of its neighbors. (The excess degree distribution j is defined as $j(k) = \frac{(k+1)P(k+1)}{D}$). Thus, assortativity can vary between -1 (highly disassortative) to 1 (highly assortative).	All
Average clustering coefficient	Clustering	Measure of the degree to which nodes in a network tend to cluster together	The average local clustering coefficient of all nodes (as in Watts & Strogatz, 1998), where the local clustering is defined as: $C_i = \frac{\lambda_G(v)}{\tau_G(v)}$, where $\lambda_G(v)$ is the number of triangles on $v \in V(G)$, $V(G)$ being the set of all nodes in the undirected graph G , and $\tau_G(v)$ is the number of triangles on $v \in G$. The clustering coefficient varies between 0 (the friends of my friends are never connected) and 1 (the friends of my friends are all connected)	Constant in scale-free networks
Degree distribution: the shape of the power law distribution	Degree distribution	Shape of the degree distribution for power-law distributions: how strongly hubs gather all connections	A power-law distribution is defined as $P(k) \sim k^{-\alpha}$ where α is the exponent of the distribution. The larger the exponent, the rarer the large values are. The exponent is estimated by applying "log" to the degree distribution and looking at the slope of the obtained new graph. Exponents in scale-free networks obtained with Barabasi-Albert algorithm usually present some random fluctuations and vary between 2 and 3.	Only networks exhibiting a power-law distribution

316 In this study, we aim to isolate the effect of the different metrics in order to understand how
317 they may influence language variability. Please note that we focus on these metrics because
318 they capture essential properties of communicative networks that have been claimed, or may
319 arguably contribute to language dynamics within networks. Moreover, some of these metrics
320 are very hard to disentangle and, therefore, their effects might be confounded in most previous
321 studies. To overcome those difficulties, we generated hundreds of thousands of networks, and
322 then selected two sets of networks, each containing 100 networks: for a given metric, we
323 selected the two sets of networks guaranteeing that the averages for the other metrics in the
324 two sets were similar except for the metric under consideration (see Table 2). As significance
325 can be easily attained when it is possible to generate a high number of networks, we ensured
326 that the difference between the two sets was much larger (at least two orders of magnitude)
327 for the metric considered than for the "equalized" metrics. We used scale-free networks to
328 analyze the independent effects of the network size, the path length, the assortativity, and the
329 degree distribution; small-world networks for the clustering coefficient; and random networks
330 for the average degree and path length.

Metric value ¹	Difference between the two sets varying in the network metric (in bold) ²					
	Size	Path length ³	Degree distribution	Assortativity	Clustering	Node degree
<i>Size</i>	(100 / 50) 50	(50 / 50) 0	(50 / 50) 0	(50 / 50) 0	(150 - 150) 0	(50 - 50) 0
<i>Path length</i>	(4.49 / 4.50) 0.01	(4.99 / 3.37) 1.62	(4.09 / 4.10) 0.01	(4.02 / 4.01) 0.01	(1.68 - 1.68) 0	(1.78 / 1.79) 0.01
<i>Degree distribution</i>	(2.85 / 2.85) 0.00	(2.79 / 2.80) 0.01	(3.04 / 2.55) 0.49	(2.75 / 2.75) 0	--	--
<i>Assortativity</i>	(-0.29 / -0.30) 0.01	(-0.31 / 0.31) 0	(-0.35 / -0.35) 0.01	(-0.16 / -0.48) 0.32	(-0.01 / -0.01) 0	(0.02 / 0.01) 0.01
<i>Clustering</i>	(0 / 0) 0	(0 / 0) 0	(0 / 0) 0	(0 / 0) 0	(0.58 - 0.32) 0.26	(0.25 / 0.25) 0.01
<i>Degree</i>	(1.98 / 1.96) 0.02	(1.96 / 1.96) 0	(1.96 / 1.96) 0	(1.96 / 1.96) 0	(48 / 48) 0	(12.69 / 11.98) 0.71

332 These metrics reflect the *global* network structure, but there is *local* heterogeneity between
333 nodes, which is captured by node-level metrics. Using the same model, we investigated the
334 influence of *local* network heterogeneity on language change (Fagyal et al., 2010).
335 Specifically, we extended the results from our previous paper (Josserand et al., 2021). In this
336 earlier work, we found that language adapts more to a group of biased agents—namely, those
337 born with a preference for a specific utterance—when these biased agents have higher
338 centrality. While in the prior study, we defined centrality as betweenness centrality, we
339 complemented these results here by exploring whether various other node-level metrics (local
340 clustering coefficient, node degree, eigenvector centrality, and closeness centrality) might
341 exert different influences. These findings, available only in the **Supplementary Materials** (see
342 part 3.4), not only expand upon but also validate the results presented by Josserand et al.
343 (2021).

344 Measurements

345 Previous studies (see Figure 13 in Josserand et al., 2021) have shown that in a type of
346 Bayesian model of language change similar to the ones implemented here but where the
347 linguistic feature was binary, language stabilized before 1000 rounds in all network types (see
348 our **Supplementary Materials** part 3.1 for visualization). Here, we introduced a conservative

¹ Each row shows the absolute difference between the two sets in the appropriate metric given by the names. The parenthesis indicates the mean value for each set of networks (high / low).

² Each column refers to the two sets of networks varying only in the shown metric.

³ For example, the second column shows the differences between the two sets of networks differing by design in path length: these two sets contain 100 networks each, have the same size, clustering coefficient, average degree, and almost the same degree distribution and assortativity, but (as intended) markedly different path length. These numbers should be interpreted with caution, as the strength of a difference of “1” or “0.01” may vary depending on the metric being analyzed.

349 margin and analyzed the inter- and intra-individual variability after 3000 rounds (when all
350 agents had produced 3000 utterances) in order to make sure that language would have
351 stabilized.

352

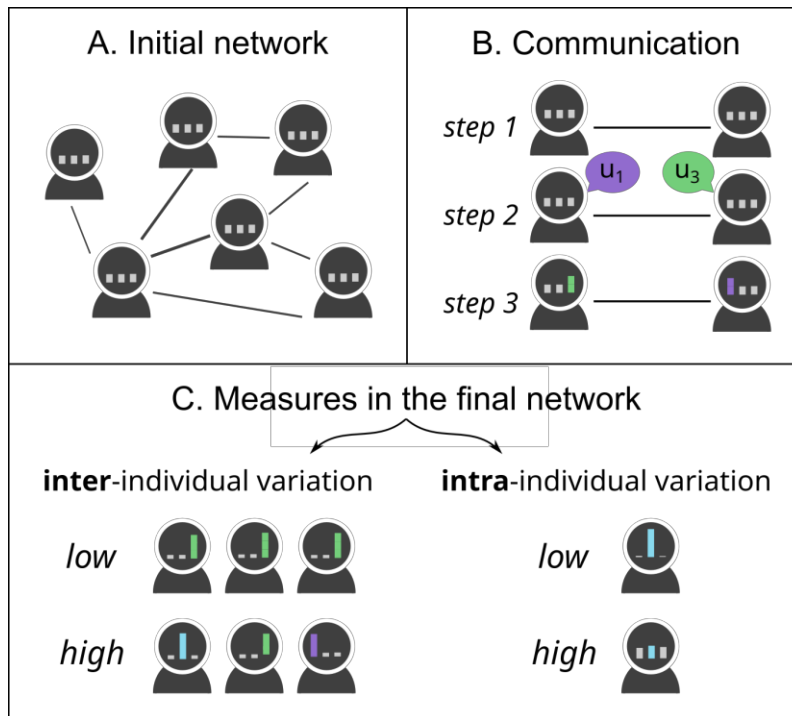
353 First, we define *inter-individual variability* by how different the agents are regarding their
354 internal language representation. For example, if agents tend to use different utterances
355 (some preferentially u_4 while some others preferentially u_3), the inter-individual variability will
356 be high. We estimate inter-individual variation using the Kullback-Leibler divergence, which is
357 a measure of the statistical distance between two probability distributions (as coded in the
358 *entropy* package in R; Hausser & Strimmer, 2021). To do so, we computed for all possible
359 pairs of agents the pairwise Kullback-Leibler divergence between their Dirichlet distributions,
360 and then averaged the resulting pairwise Kullback-Leibler divergences to obtain an average
361 measure of the inter-individual variation in the network.

362

363 Secondly, *intra-individual variability* captures how consistent an agent is in its own language
364 productions. For example, if an agent uses equally likely two utterances (u_2 and u_8 half of the
365 time each, for example), its intra-individual variability will be high. On the contrary, an agent
366 that always uses the same utterance has a low intra-individual variation. We estimate intra-
367 individual variation as the entropy of an agent's Dirichlet distribution (as coded in the *entropy*
368 package, using Jeffreys prior⁴; Hausser & Strimmer, 2021), which is bounded between 0 and
369 $\log_2(k)$, where here $k=10$. The *average* of this intra-individual variation across all agents
370 captures their propensity to consistently use the same utterance (a high average indicates that
371 agents switch between different utterances, while a low average suggests that agents tend to
372 use the same utterance consistently), while its *standard deviation* across all agents captures
373 to what degree there are differences between the agents regarding their intra-individual
374 variation (a high standard deviation suggests that while some agents always use the same
375 utterance, others show a high variability, and a low standard deviation suggests that all agents
376 tend to be equally consistent or not in their language production). See Illustration 3 for a
377 visualization of the entire process.

378

⁴ This ensures that the resulting entropy value is robust and not overly influenced by the specific parameterization of the probability distribution. We also selected other priors, but conclusions were the same.



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Illustration 3. A schematic representation of the multi-agent model. (A) Agents are “born” embedded in a network with a fixed structure and with given characteristics. Each agent has an internal language representation given by the parameters of the Dirichlet distribution (see gray bar plot in Illustration 2). For visualization purposes, we decided to represent $k=3$ utterances here instead of the $k=10$ utterances used in the paper. (B) **Step 1.** Each agent randomly extracts a number from its Dirichlet distribution: here, the probability of picking any of the three utterances is the same. **Step 2.** Each agent produces the chosen utterance (here, u_1 for the agent on the left and u_3 for the one on the right). **Step 3.** Each agent hears the utterance(s) of its neighbor(s) and updates its own Dirichlet distribution accordingly. (C) After each individual talks 3000 times, we measure the inter- and intra-individual variation among the agents in the final network.

391 1.6 The network generation procedure

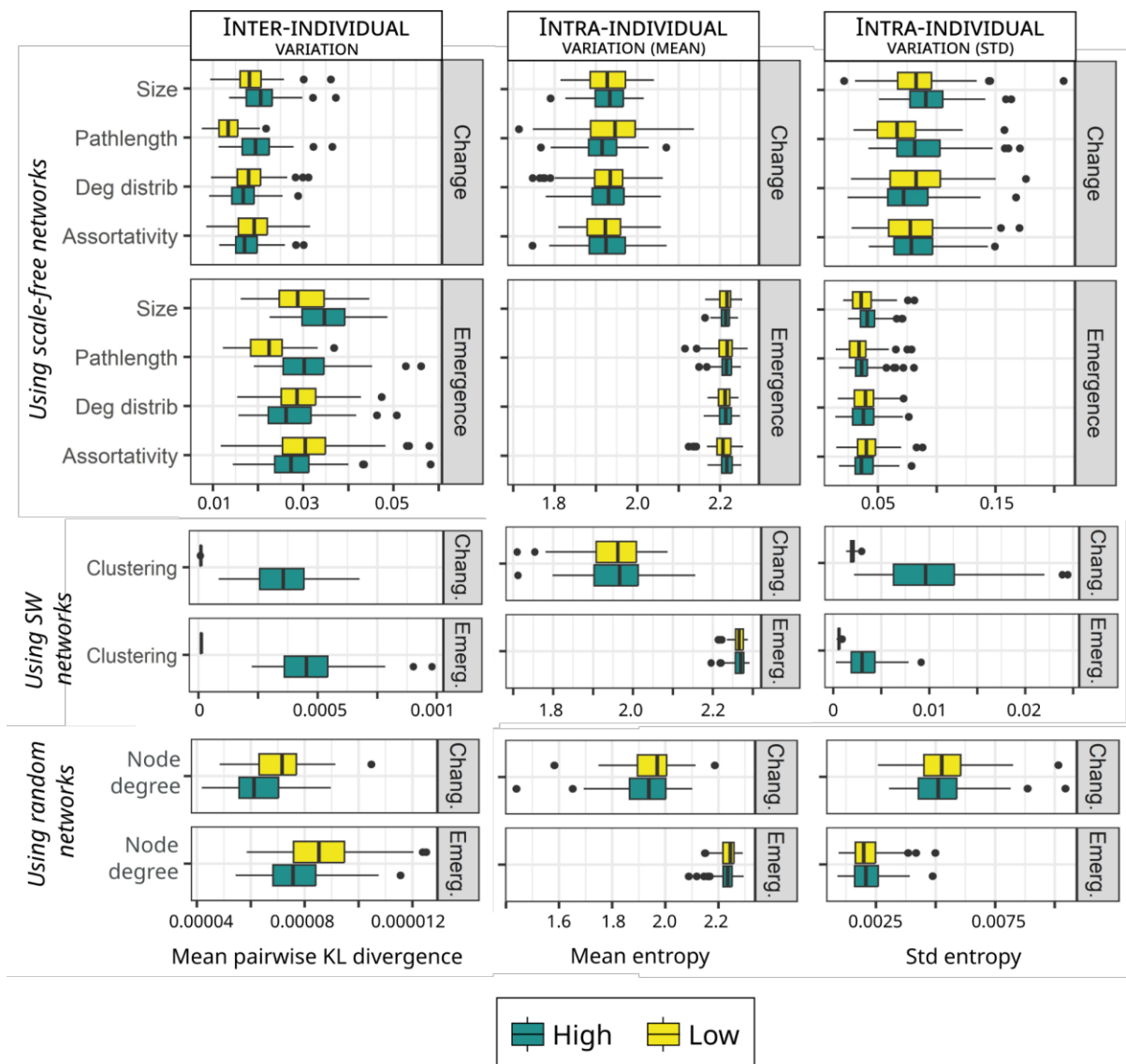
392 We conducted simulations using Netlogo (version 6.2.2; Wilensky, 1999) and generated two
393 datasets. In the first (which we will call *dataset 1*), we systematically compared two sets of
394 networks differing in exactly one metric, as described above. To achieve this, we first
395 generated 100 networks for each combination of parameters which resulted in 80,000 scale-
396 free networks (with varying sizes), 240,000 random networks (with varying connection
397 probability to control for density and pathlength), and 600 small-world networks (with high and
398 low rewiring probability, to examine the influence of clustering). Detailed information about the
399 specific parameters is available in the **Supplementary Materials** (see Part 2.2.1). Following
400 the generation of these networks, and using graphic visualization, we manually selected sets
401 of networks that differed in only one metric. Subsequently, to compare the amount of variation
402 in the two sets of networks and assess whether the difference was meaningful, we used
403 Cohen’s d , a statistical metric that measures the difference between two groups. This metric
404 is computed by dividing the difference between the means of the two groups by the pooled
405 standard deviation of the groups.

406 With dataset 2, we ran a series of simulations with random, scale-free, and small-world
407 networks using a wide range of parameters. For each network type, we generated networks
408 with sizes of 50, 150, and 300. In the case of random networks, we explored connection
409 probabilities ranging from 0.05 to 0.95 in increments of 0.05. For small-world networks, we
410 created networks with varying average numbers of neighbors (4, 8, 16, 48) and different
411 rewiring probabilities (0.1, 0.3, 0.5, 0.7, 0.9). Initially, we examined how the metrics co-varied
412 across a broad range of values using Spearman's and Pearson's correlations. Then, we
413 modeled how these metrics influenced the final variability in the communicative network. We
414 z-scored all the metrics and we predicted inter-individual, mean intra-individual and standard-
415 deviation of intra-individual variation using all our metrics as predictors. To do so, we used
416 three machine learning techniques. First, we measured the predictor importance with *random*
417 *forests* (using package `randomForest`; Liaw & Wiener, 2002) using an accuracy-based index
418 and a gini-based Index. Second, we used *conditional forests* (using `cforest` function in
419 `partykit` package; Hothorn & Zeileis, 2015) to measure the unconditional predictor
420 importance index. Last, we used *Support Vector Machines* (SVM; `fit` function in `rminer`
421 package; Cortez, 2010) for sensitivity analysis. Additionally, for all of them, we measured the
422 goodness of fit using R^2 and RMSE. To prevent overfitting, we trained the model on 80% of
423 the data (stratified by size) and tested the model on the 20% remaining data using 50
424 replications. For easier plotting, we z-scored and then aggregated the results for importance
425 by method (accuracy-based, gini-based, unconditional, and sensitivity), by network type, and
426 by predictor. The values for each method and each predictor, as well as additional details on
427 the exact parameters used in the models, are available in the **Supplementary Materials** (see
428 Part 3.3.4).

429 3. RESULTS

430 3.1 Dataset 1

431 To produce sets of networks that vary only in one metric, we needed to use multiple types of
432 networks. We used scale-free networks to investigate the influence of assortativity, degree
433 distribution, path length, and size. Small-world networks were used to examine the impact of
434 the clustering coefficient, while random networks were used to investigate the effect of the
435 average node degree. The inter- and intra-individual variation (mean and standard-deviation)
436 values are presented in Figure 1, while Figure 2 indicates the effect sizes and significance for
437 each set.
438
439



440
 441 *Figure 1. Measure of inter-individual variation (left column), mean of intra-individual variation (column*
 442 *in the middle), and standard deviation of intra-individual variation (right column). Each row shows the*
 443 *results for one metric: for example, the first row shows the results for 100 networks that have a low size*
 444 *on average (in blue), and 100 networks with a high size (in yellow) (see Table 2), in networks where*
 445 *there is an initial language exposure (language change, upper panel) and networks without initial*
 446 *language exposure (language emergence, lower panel). Depending on the network type used to study*
 447 *the metric of interest, the x axis may vary: the x axis of inter-individual variation shows the mean pairwise*
 448 *Kullback-Leibler divergence, while the two other columns show respectively the mean and the standard*
 449 *deviation of the entropy for all agents, computed using the final Dirichlet distribution for each agent. The*
 450 *quantification of the effect size between the two sets is indicated in Figure 2.*
 451

452 **Clustering coefficient.** Both networks in language emergence (Cohen's $d = 4.2$) and
 453 language change (Cohen's $d = 4.2$) scenarios exhibit a strong influence of clustering
 454 coefficient on inter-individual variability, as shown in Figures 1 and 2. Networks without a
 455 clique structure tend to display lower levels of inter-individual variation. Similar to all other
 456 network structure metrics, the clustering coefficient has no effect on the mean intra-individual
 457 variation. However, higher clustering coefficients lead to more differences among agents in
 458 terms of standard deviation of intra-individual variation (with some agents consistently using
 459 the same utterance and others switching between several different utterances), which is

460 observed in networks with language emergence or change scenarios (Cohen's $d = 2.0$ and
461 Cohen's $d = 2.2$, respectively).

462

463 **Path length.** The path length strongly affects inter-individual variability, in both emergence
464 (Cohen's $d = 1.5$) and change (Cohen's $d = 1.5$) scenarios. Thus, networks where agents are
465 easily reachable by others are likely to show low levels of inter-individual variation. Path length
466 does not affect the mean amount of intra-individual variation, but it does slightly positively
467 affect the standard deviation of intra-individual variation in networks in language change
468 scenario only (Cohen's $d = 0.7$).

469

470 **Size.** The network size also influences language variability, but its effect quickly disappears
471 with higher initial language exposure. Big networks show higher inter-individual variation in
472 networks with language emergence (Cohen's $d = 0.9$) and language change scenarios
473 (Cohen's $d = 0.5$). However, an additional analysis suggests that the stronger the initial
474 language exposure, the less the network size will play a role in shaping inter-individual
475 variation. More specifically, Cohen's d drops to 0.3 when the initial language exposure is
476 strong (namely, if the initial language is skewed with 21 utterances instead of 6; see
477 **Supplementary Materials** part 3.2.1.8). Thus, the size of the population may only affect inter-
478 individual variation in emergent languages. As for intra-individual variation, size does not
479 appear to have any impact on the amount and standard-deviation of intra-individual variation
480 observed.

481

482 **Average node degree.** The average node degree has a discernible but moderate impact on
483 the degree of language variation among individuals, compared to other metrics. In networks
484 with language emergence (Cohen's $d = 0.7$) and language change (Cohen's $d = 0.7$)
485 scenarios, networks with a low average node degree typically exhibit high inter-individual
486 variation. However, the average node degree does not affect any type of intra-individual
487 variation.

488

489 **Assortativity and degree distribution.** Finally, both assortativity and degree distribution
490 have no observable effect on inter-individual nor intra-individual variation. Networks where
491 individuals are preferentially connected to individuals who have a similar degree, and networks
492 with very skewed degree distributions, do not show specific patterns of variation.

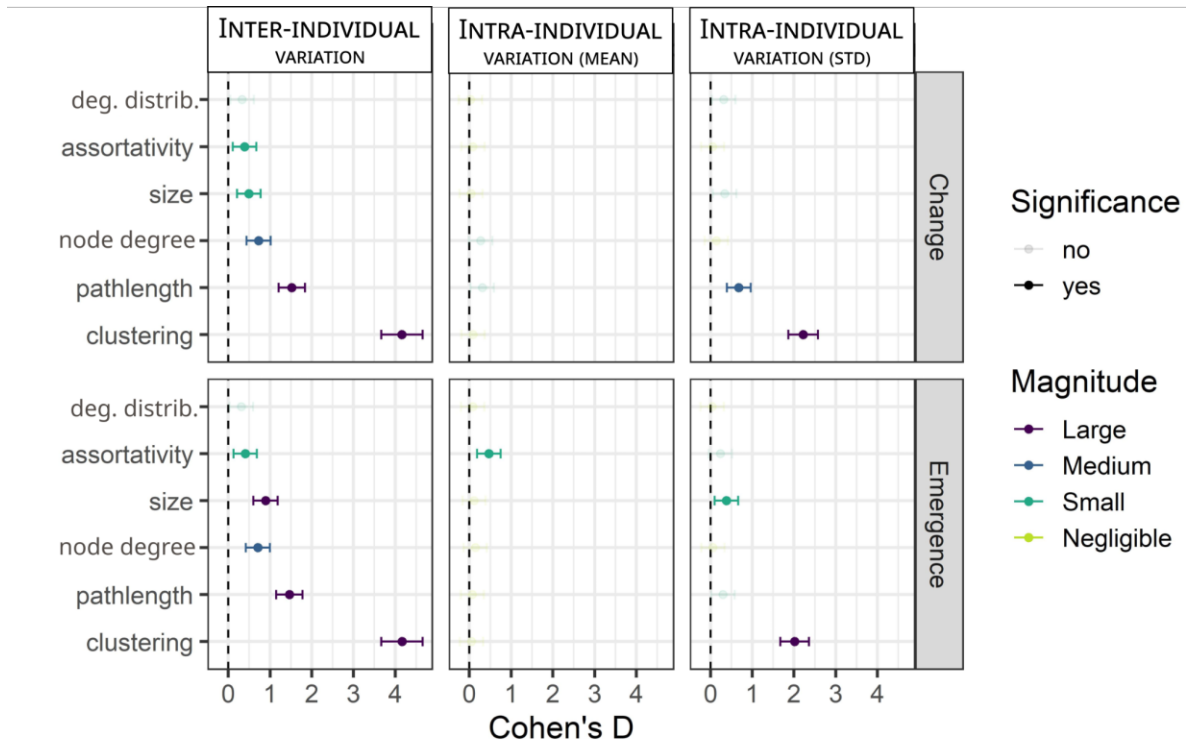
493

494 **Initial language exposure.** The initial language exposure, which reflects whether agents are
495 born in language emergence or language change scenario, affects both inter- and intra-
496 individual variation. Networks in language emergence scenarios exhibit greater inter- and
497 mean intra- individual variation: as expected, agents communicating in a language change
498 scenario exhibit more consistency (lower entropy on average) than agents involved in a
499 language emergence scenario. Notably, the amount of intra-individual variability is solely
500 affected by the initial language exposure. However, the standard-deviation of intra-individual
501 variation is lower in language emergence scenario.

502

503 **Summary.** In summary, our findings indicate that clustering coefficient and path length are
504 the main factors impacting both inter-individual variation and the standard deviation of intra-
505 individual variation. Size and density moderately affect inter-individual variation, specifically in
506 language emergence scenarios. Notably, the mean intra-individual variation remains

507 unaffected by any of our network structure measures. The initial language exposure, reflecting
 508 language change or emergence scenarios, also strongly influences all types of variation.
 509

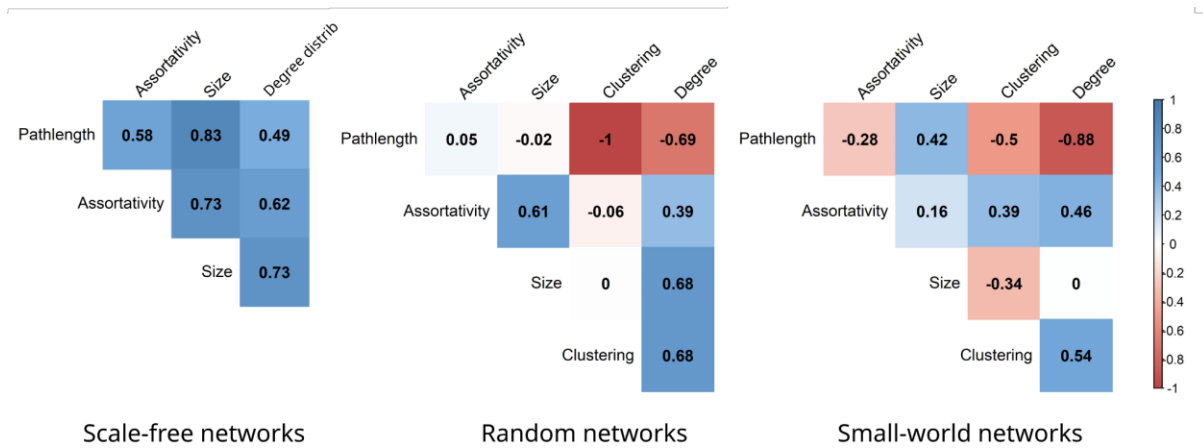


510
 511 *Figure 2. Measure of inter-individual variation (left column), mean intra-individual variation (middle*
 512 *column), and standard deviation of intra-individual variation (right column) in networks with initial*
 513 *language exposure (language change; upper panel) and without initial language exposure (language*
 514 *emergence; lower panel). The Cohen's d index for each predictor indicates the effect size of the*
 515 *difference in variation between the two sets (high and low) for each predictor along with its confidence*
 516 *interval. Gray indicates a non-significant metric, while significant metrics are in shades of color.*

517 3.2 Dataset 2

518
 519 Here, we also use simulations obtained using automatically generated random, small-world,
 520 and scale-free networks (all generated networks were kept without any specific selection
 521 criteria⁵). We analyzed the Spearman's correlations between the metrics (see Figure 3). The
 522 clustering coefficient, the node degree, and the path length are highly correlated with each
 523 other. Thus, the more neighbors a given agent has on average, the more likely the friends of
 524 this agent will be connected to each other, and the lower the average path length will be in the
 525 network. Network size also correlate to the other metrics in scale-free networks: bigger scale-
 526 free networks have higher path length and higher assortativity. These results reflect the
 527 necessity of the sampling procedure used above to study the effect of the different metrics
 528 separately.

⁵ We only excluded random networks that possessed isolated component, in order to be able to compute the path length.



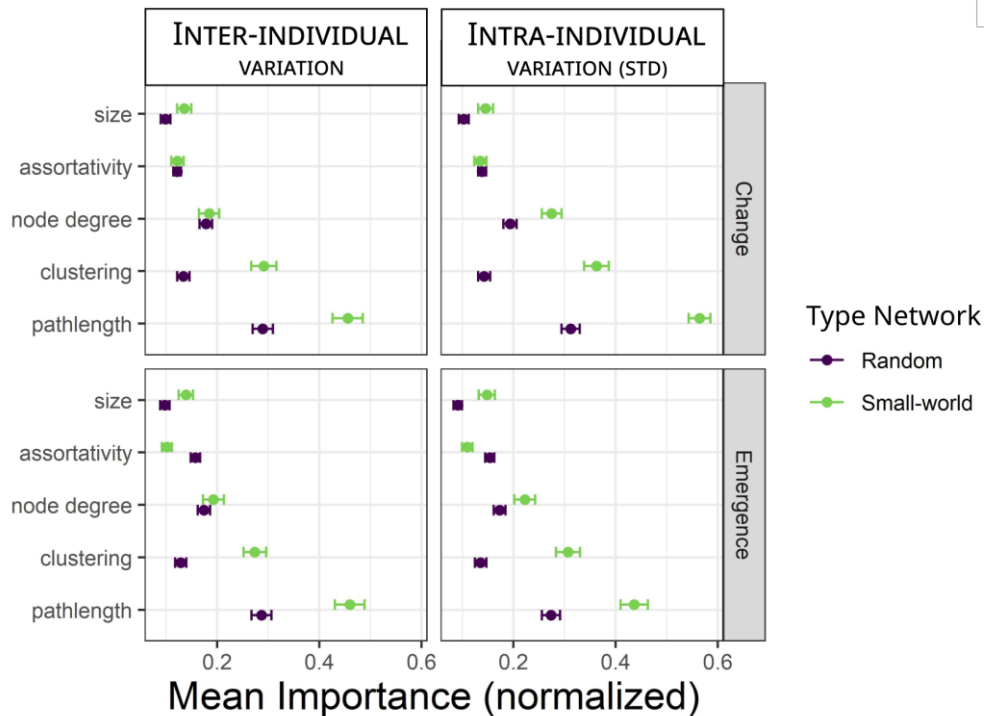
529
 530 *Figure 3. Correlation matrix with the different metrics in scale-free (left), small-world (middle), and*
 531 *random (right) networks.*
 532

533 Using this same set of simulations, we applied several machine learning techniques to predict
 534 variability in different network types using our metrics as predictors (see Figure 4) in both
 535 language emergence and language change scenarios. We observed that the R^2 for the mean
 536 intra-individual variation was notably low, with an average R^2 of 35% for language emergence
 537 and of 2% in language change scenarios (these means were computed for all methods and
 538 all types of networks; please note that there are high differences between the R^2 obtained in
 539 the different methods). Furthermore, our predictors in models using scale-free networks poorly
 540 predicted any types of variations, with R^2 around 37% for inter-individual variation and 4% for
 541 the standard deviation of intra-individual variation. However, in small-world and random
 542 networks, and in both language scenarios, our predictors effectively predicted inter-individual
 543 variation and the standard deviation of intra-individual variation, with R^2 values for these
 544 models around 92% for inter-individual variation and 76% for standard-deviation of intra-
 545 individual variation. Thus, we focus our analysis on these two types of networks and types of
 546 variation. The exact R^2 and RMSE for each method, network type and type of variation are
 547 available in the **Supplementary Materials** (see Part 3.3.4).
 548

549 **Inter-individual variation.** Encouragingly, these results concur with the previous findings
 550 comparing sets of networks. The models consistently indicate that path length and clustering
 551 coefficient are the most reliable predictors for inter-individual variation in both language
 552 change and emergence scenarios. Notably, in this case, path length emerges as a stronger
 553 predictor than the clustering coefficient. Interestingly, the models reveal a substantial impact
 554 of average degree on inter-individual variation, particularly in random networks. The effect
 555 sizes of the other metrics are very small, corroborating the idea that network size and
 556 assortativity are not important for understanding inter-individual variability. There are no big
 557 differences between the language change and language emergence scenarios regarding the
 558 importance of predictors in both random and small-world networks.
 559

560 **Mean intra-individual variation.** Similar to our previous observations, strong differences in
 561 the mean intra-individual variation were identified between language change and language
 562 emergence scenarios. However, our metrics do not contribute significantly to predicting this
 563 type of variation, as evidenced by the relatively low R^2 . This aligns with the findings of Dataset
 564 1.
 565

566 **Standard-deviation of intra-individual variation.** Concerning the standard deviation of intra-
 567 individual variation, the importance of the predictors varies based on the type of network and
 568 the initial language exposure. In both random and small-world networks, path length emerges
 569 as the most crucial predictor. Notably, in small-world networks, it exerts a stronger influence
 570 in language change scenarios compared to language emergence scenarios. In small-world
 571 networks, the clustering coefficient takes the position of the second most important predictor,
 572 followed by node degree. Within random networks, node degree holds the second-highest
 573 importance as a predictor, closely followed by clustering coefficient and assortativity. In both
 574 cases, assortativity and size exhibit the least importance.



575 *Figure 3. Mean importance for inter-individual variation (left column) and standard deviation of intra-*
 576 *individual variation (right column) in language change scenario (upper panel) and language emergence*
 577 *scenario (lower panel). Results for the mean intra-individual variation and for scale-free networks are*
 578 *not presented, as the predictors exhibited poor prediction performance for all types of variation (low R^2).*
 579 *This measure of importance was computed by averaging the following z-scored predictor importance*
 580 *measures: accuracy-based and Gini-based (extracted from random forests models), unconditional*
 581 *(extracted from conditional forests), and sensitivity-based (extracted from support vector machines).*
 582 *The dots represent the mean of these four measures, and the line represents the standard error*
 583 *between these four measures. Results for random networks are shown in purple, while results for small-*
 584 *world networks are displayed in green.*
 585
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 587

588 4. DISCUSSION AND CONCLUSIONS

589 Understanding the patterns of variability within a language provides important insights into the
 590 great diversity observable in the world's languages. While many studies highlight the role of
 591 network structure on language change, no differentiation was made between various network
 592 characteristics in real world-populations or agent-based models, making it impossible to
 593

594 disentangle their relative contributions. In this study, we applied several methods using a
595 Bayesian agent-based model of language evolution to understand how specific characteristics
596 of a network affect language in a language emergence and change scenarios.

597

598 We found that network structure affects inter-individual language variation in both language
599 emergence and change scenarios. Although most earlier studies focused on grammatical
600 complexity and other aspects of language, rather than linguistic variation per se, previous
601 claims supported the idea that the effect of network structure on language was largely
602 attributed to differences in the *size* and *density* of social networks (Dale & Lupyan, 2012;
603 Trudgill, 2011b; Wray & Grace, 2007; Raviv et al., 2019). However, our results suggest that
604 when considering language change, the size and density of a social network do not appear to
605 be the most important factors. Instead, parameters such as path length and the clustering
606 coefficient seem to be more influential in determining language variability. The confusion
607 between size and density on the one hand, and clustering and path length on the other hand,
608 can be easily understood given that these factors are highly correlated: bigger and well-
609 connected populations tend to have shorter path lengths and higher clustering coefficients.
610 Hampering the transmission of information and promoting the existence of distinct
611 communities within a population will favor the emergence of variation in a language. Also, a
612 population growing in size will not necessarily witness more variation, as long as the
613 transmission of information is effective; this is, in particular, the case in populations with hubs,
614 which facilitate the circulation of information. However, interestingly, in the first dataset and in
615 the language emergence scenario only, it appears that the size of a network has an effect:
616 larger populations tend to exhibit more variation. This emphasizes the intrinsic difference
617 between the two scenarios and underscores the need to consider this factor when studying
618 the evolution of language. Quite consistently, network density has a moderate effect in most
619 analyses, but its exact role remains to be determined.

620

621 Additionally, the type of scenario (language change versus emergence) also plays an
622 important role in predicting the amount of inter-individual variation present in a population.
623 Thus, individuals born in a society where there is already a pre-existing language (language
624 change scenario) would be more likely to use homogenous forms of language. In turn, this
625 would explain why emerging sign languages in small remote villages (such as Al Sayyid
626 Bedouin Sign Language sign language in Israel; Jaraisy & Stamp, 2022) usually exhibit more
627 variation than official deaf community sign languages taught at schools, even though these
628 official languages are characterized by sparser and larger populations of signers (Meir et al.,
629 2012; Mudd et al., 2020). We suggest that the pressure for conformity (and more specifically,
630 for using the dominant variant in the population) is the main factor underlying the differences
631 between an established language (language *change* scenario) and an emerging language
632 (language *emergence* scenario). Importantly, the strong influence of the type of scenario
633 suggests that non-structural factors do play a very important role in shaping language
634 variability, and that network structure cannot alone account for all the diversity present in real-
635 world networks. For example, other extrinsic factors such as the role of a shared context (Mudd
636 et al., 2022) and the openness to strangers (Dale & Lupyan, 2012) were not taken into account
637 here, but they might have additional contributions to the effect of network structure.

638

639 On the contrary, intra-individual variability is not shaped by the network structure, but only by
640 the type of scenario of the society in which an individual is born. An individual born in a society
641 with low pressure to use a specific form of language (language *emergence* scenario) will use

642 different types of utterances to refer to the same concept. These findings relate to the study
643 of very recent emergent sign languages, such as Kata Kolok, where high intra-individual
644 variation was observed (Lutzenberger et al., 2021). However, differences between individuals
645 regarding their level of intra-individual variation are affected by the network structure. Here
646 too, a high path length and a high clustering coefficient increase the probability that the
647 population possesses both individuals very confident in their language beliefs and individuals
648 with low confidence in which language form to use. We suggest that both high path length and
649 high clustering coefficients are related to the presence of structural communities (cliques),
650 which then lead to linguistic communities. Individuals bridging these communities are then
651 more likely to possess high intra-individual variation, changing their language according to
652 whom they are speaking to, while individuals inside communities use the language spoken by
653 their group unconditionally.

654
655 Together, these results suggest that differences at the global level of network structure,
656 especially for path length and clustering coefficient, influence language inter-individual
657 variability. They also highlight the weak role of network structure on intra-individual variation.
658 However, one potential limitation of our study is that the degree of contrast between the sets
659 of networks generated based on a single metric may not have been strong enough to capture
660 the full extent of the metric's influence. For instance, the magnitude of the difference between
661 the high and low node degree networks may have been insufficient to fully explain the impact
662 of node degree on language variability. Also, our "flat" Bayesian model contrasts with the
663 human capacity to differentiate and track various language varieties depending on the
664 interlocutors and contexts. This highlights the potential for elaborating more complex
665 simulations (such as hierarchical Bayesian models) in future research, and meanwhile, our
666 conclusions should be extrapolated with caution to real-world language networks. Additionally,
667 the model only accounted for static networks, whereas real-world networks are dynamic and
668 evolve through interactions. While interactions are the driving force behind network structure,
669 further research could explore the mechanisms of preferential attachment to gain a deeper
670 understanding of their impact on language variability.

671

672 5. DATA AVAILABILITY STATEMENT / SUPP MATERIALS

673

674 The datasets and the analysis presented in this study can be found in the online repositories
675 [mathjoss/NetworkStructure_ABM](https://mathjoss.org/NetworkStructure_ABM).

676 6. AUTHOR CONTRIBUTIONS

677

678 BdB, DD, MA-T, and MJ designed the research. MJ performed the research, wrote and ran
679 the simulations. MJ wrote the first draft with substantial contributions from DD and FP. All
680 authors (BdB, DD, MA-T, MJ, and FP) read, contributed to, and approved the manuscript.

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