

# 1 Mean-reverting self-excitation drives evolution: phylogenetic 2 analysis of a literary genre, *waka*, with a neural language model

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## 5 Abstract

6 To elucidate the evolutionary dynamics of culture, we must address fundamental questions such as  
7 whether we can interpolate and extrapolate cultural evolution, whether the time series of cultural  
8 evolution is distinguishable from its reverse, what factors determine the direction of change, and how  
9 the cultural influence of a creative work from the viewpoint of an instant is correlated with that from  
10 the viewpoint of a later instant. To answer these questions, the evolution of classical Japanese poetry,  
11 *waka*, specifically *tanka*, was investigated. Phylogenetic networks were constructed on the basis of the  
12 vector representation obtained using a neural language model. The parent–child relationship in the  
13 phylogenetic networks exhibited significant agreement with a previously established *honkadori* (allusive  
14 variation) phrase-borrowing relationship. The real phylogenetic networks were distinguishable from  
15 the time-reversed and shuffled ones. Two anthologies could be interpolated but not extrapolated. The  
16 number of children of a poem in the phylogenetic networks, the proxy variable of its cultural influence,  
17 evaluated at an instant, was positively correlated with that evaluated later. A poem selected for an  
18 authoritative anthology tended to have 1.1–1.5 times more children than a similar but nonselected  
19 poem, implying the existence of the Matthew effect. A model with mean-reverting self-excitation  
20 replicated these results.

## 21 Introduction

22 A quantitative analysis of cultural evolution is essential in understanding human history because the  
23 history of culture encompasses human history and beyond. Animals, including birds and fish, are  
24 known to transmit culture (Slater, 1986; Boesch and Boesch, 1990; Dugatkin and Godin, 1992). The  
25 oldest stone tools date back 3.3 million years (Harmand et al., 2015), nearly 1 million years before the  
26 emergence of the genus *Homo*, but already show signs of sophistication. Sophisticated stone tools as  
27 well as other creative works are a manifestation of cultural tradition and inherently evolve over time.

28 Naturally, there arise several questions to be addressed to understand the evolutionary dynamics  
29 of culture. Is the culture of a period the intermediate form of those of the preceding and following  
30 periods? Or, are they completely different? Can we predict the future direction of evolution from  
31 the history of the past? In other words, can we interpolate and extrapolate cultural evolution? Can  
32 the time series of cultural evolution be distinguished from its reverse? What factors determine the  
33 direction of cultural evolution? How is the cultural influence of a creative work judged at a time point  
34 correlated with that at a later time point?

35 Some of these questions have been tackled by a considerable amount of studies that investigate  
36 information transmission and replication in the various forms of creative works. Since the seminal  
37 works by Cavalli-Sforza and Feldman (1973a) and Cavalli-Sforza and Feldman (1973b), the evolution

1 of music, scientific literature, social network posts, malware, images, and movies has been studied.  
2 Nakamura and Kaneko (2019) analyzed Western classical music data and found that the frequencies  
3 of dissonant intervals have steadily increased. A study on an English text corpus spanning about 400  
4 years found that the authors in a similar period share the same literary style, which gradually changes  
5 over time (Hughes et al., 2012). The science of science (Fortunato et al., 2018) has elucidated the  
6 dynamics underlying scientific discovery and the scientific community itself: a rare combination of  
7 ideas (Foster et al., 2015), a combination of old and new ideas (Uzzi et al., 2013; Kim et al., 2016;  
8 Wang et al., 2017), and interdisciplinary collaboration (Larivière et al., 2015) result in higher citation  
9 rates; but these features are not advantageous in grant applications (Boudreau et al., 2016; Leahey  
10 and Moody, 2014; Lee and Bozeman, 2005); the number of citations of a paper is boosted by the total  
11 number of previous citations of the authors in the first years after publication (Petersen et al., 2014);  
12 the citation network has a long-tailed distribution (Price, 1965); and the number of citations of a paper  
13 is determined by its fitness, an obsolescence factor, and the number of previous citations (Wang et al.,  
14 2013; Eom and Fortunato, 2011). Scientific papers share a common pattern of a decaying collective  
15 memory with patents, songs, movies, and biographies (Candia et al., 2019). The rapidness of memory  
16 decay changes over time. An analysis of a 500 billion-word corpus revealed that inventions and people  
17 have been forgotten rapidly in recent years (Michel et al., 2011). Similar to the relationship between  
18 scientists and scientific papers, the relationship between users and their posts in social network services  
19 has also been studied (Wei et al., 2013; Weng et al., 2014).

20 One of the most powerful tools to study cultural evolution is phylogenetics. Phylogenetics, originally  
21 a branch of biology, is based on the theory of evolution. The evolutionary, *i.e.*, ancestor–descendant  
22 relationship of organisms estimated in phylogenetics is represented by a phylogenetic tree. However, the  
23 idea of the phylogenetic tree is older than Charles Darwin’s theory of evolution; Friedrich Schlegel’s  
24 language tree and Carl Johan Schlyter’s manuscript phylogeny, or stemma, date back to the early  
25 nineteenth century (Atkinson and Gray, 2005). The method of phylogenetics has successfully been  
26 applied to the analysis of cultural transmission and evolution, forming a field of research called cultural  
27 phylogenetics (Mesoudi, 2011; Straffon, 2016). Cultural phylogenetics has elucidated the origin of a  
28 language family (Gray and Atkinson, 2003), the coevolution of livestock and descent rules (Holden  
29 and Mace, 2003), and the dynamics of political complexity (Currie et al., 2010). These previous works  
30 attempted to distinguish between correlations due to shared ancestry and convergent cultural evolution  
31 (Holden and Mace, 2003). This has been enabled by recent advances in computational methods.

32 Phylogenetic methods have also been applied to the creative works of individuals, such as Paleoin-  
33 dian projectile points (O’Brien et al., 2001) and Turkmen textiles (Tehrani and Collard, 2002). The rise  
34 of specialist forensic need and interest in Internet memes has stimulated techniques to reconstruct the  
35 phylogenetic trees of malware (Goldberg et al., 1998), images (Dias et al., 2010), movies (Dias et al.,  
36 2011), and audio files (Nucci et al., 2013). Extending the notion of classical stemmatics (Marmerola  
37 et al., 2016), Barbrook et al. (1998) employed computerized techniques to reconstruct the stemma of  
38 *The Canterbury Tales*, and Kanojia et al. (2019) used word embeddings to reconstruct the phyloge-  
39 netic tree of a historical Sanskrit text, *Kāśikāvṛtti*. Most of the creative works investigated in these  
40 previous studies have been made by replicating existing ones (as in Internet memes and manuscripts)  
41 or carried an explicit indication of the influence between them as citations in research papers and  
42 reposts on social media. In other words, their phylogenetic networks are easily reconstructed.

43 However, not all kinds of creative works are endowed with such conditions favorable for recon-  
44 structing phylogenetic networks. Some kinds of creative works are not necessarily a direct replication  
45 of existing ones and contain no explicit citations, but nevertheless, are made under the influence of  
46 existing ones. To answer questions regarding the evolutionary dynamics of culture, we must extend the  
47 study of cultural evolution to these kinds of creative works and develop a method to estimate implicit  
48 influence.

49 Given this background, the present study aimed to answer these questions by investigating the  
50 evolution of a literary genre. Literary genres are easier to analyze than other cultural phenomena such  
51 as stone toolmaking, fashion, and rituals because databases of literary genres are publicly available,  
52 and as such, we can take advantage of recent neural language models in analyzing them. The literary

1 genre studied in this paper is *waka*, the most authoritative poetic form in classical Japanese literature.  
2 *Waka* has several advantages in studying evolution dynamics. First, *waka* is a fixed verse with  
3 31 syllables, the brevity of which facilitates quantitative analysis. Using a vector representation of  
4 poems by neural language models, we can estimate the influence among poems and, consequently,  
5 their phylogenetic networks. Second, a comprehensive database of *waka* that contains poems ranging  
6 in date from the eighth to the sixteenth century is available. Third, as is the case with every type  
7 of classical poetry in the world, poems in the form of classical *waka* were written by poets who were  
8 traditionalists and had a thorough knowledge of the great poems of the past. Respect for tradition was  
9 so deeply rooted in the poets' hearts that borrowing words from past great poems (*honkadori*) had  
10 been an established method of poem writing for more than a millennium. Conducting a comparison  
11 with the known *honkadori* relationship is an ideal means to check whether the estimated phylogenetic  
12 network is accurate. Fourth, *waka* is so central to classical Japanese culture that a vast amount of  
13 research is available to shed light on the subject from another angle.

14 Thus, the present paper attempts to estimate the phylogenetic network of *waka* and, thereby,  
15 address the above questions. This paper is organized as follows. The Materials and Methods section  
16 describes the dataset, data preprocessing, and methods of analysis, and also reviews the basic properties  
17 and history of *waka*. The Results section reports an analysis of the phylogenetic networks estimated by  
18 using vector representations generated by BERT (Devlin et al., 2019), one of the most successful neural  
19 language models. First, the estimated phylogenetic network is compared with *honkadori* pointed out  
20 by previous studies. Second, the congruence between language models with different initial parameter  
21 values is quantified. Third, to characterize the time evolution of the real data, the phylogenetic  
22 network is examined by comparing it with those estimated from time-reversed and shuffled data. An  
23 index is shown to be able to distinguish the real data from the time-reversed data. Fourth, to examine  
24 the constancy of the evaluation of the cultural influence of a poem, the numbers of the phylogenetic  
25 children of the poem with the language models trained by using the full dataset and the dataset up to  
26 an anthology are compared. Fifth, the poems in two anthologies are classified to determine whether  
27 interpolation and extrapolation are possible. Sixth, the effect of selecting a poem for an anthology is  
28 measured. More specifically, the increase in the number of phylogenetic children after being selected  
29 for an anthology is observed. Finally, a simple model with mean-reverting self-excitation to reproduce  
30 these results is presented. The Discussion section summarizes and contextualizes the results. First,  
31 the relationship between *honkadori* of *waka* and other literary genres and, thereby, the applicability  
32 of the present results to other genres, are discussed. Second, the possibility of influence from other  
33 literary genres to *waka* is examined. Finally, the limitations of the present study are presented along  
34 with directions for future research.

## 35 **Materials and Methods**

### 36 ***Waka* and its history**

37 *Waka* had been the most authoritative form of Japanese poetry for more than a millennium. Although  
38 it is unclear when the form of *waka* was established, the earliest historically verifiable examples date  
39 back to the early half of the seventh century. The earliest *waka* anthologies were compiled in the  
40 eighth century. In this paper, *waka* refers to *tanka*, the most major poetic form, which consists of  
41 five lines with 5-7-5-7-7 syllables. This form has remained productive to date. The images, allegories,  
42 metaphors, and symbols of *waka* gave birth to *nō*, *haiku*, as well as novels such as *The Tale of Genji*  
43 (Brower and Miner, 1961; Kato et al., 1979; Konishi et al., 1984; Keene, 1999). The authority of *waka*  
44 comes from the fact that it was the most essential communication tool among the society of nobles in  
45 the *Heian* period and an indispensable part of education in the later periods. *Waka* was composed in  
46 the grammar and vocabulary of the early *Heian* period for a millennium (Keene, 1999).

47 The special position occupied by *waka* in classical Japanese high culture is illustrated by the exist-  
48 tence of the Imperial Anthologies (*chokusenshū*). The Imperial Anthologies were the official anthologies

Table 1: List of Imperial Anthologies

ID	Title	Year of publication
1	<i>Kokinshū</i>	905
2	<i>Gosenshū</i>	955
3	<i>Shūishū</i>	1005
4	<i>Goshūishū</i>	1087
5	<i>Kin'yōshū 1</i>	1124
6	<i>Kin'yōshū 2</i>	1125
7	<i>Kin'yōshū 3</i>	1126
8	<i>Shikashū</i>	1151
9	<i>Senzaishū</i>	1187
10	<i>Shinkokinshū</i>	1205
11	<i>Shinchokusenshū</i>	1232
12	<i>Shokugosenshū</i>	1251
13	<i>Shokukokinshū</i>	1265
14	<i>Shokushūishū</i>	1279
15	<i>Shingosenshū</i>	1304
16	<i>Gyokuyōshū</i>	1312
17	<i>Shokusenzaishū</i>	1320
18	<i>Shokugoshūishū</i>	1326
19	<i>Fūgashū</i>	1346
20	<i>Shinsenzaishū</i>	1359
21	<i>Shinshūishū</i>	1364
22	<i>Shingoshūishū</i>	1385
23	<i>Shinshokukokinshū</i>	1439

1 of the imperial court compiled on the order of the emperor or ex-emperor. They were compiled by the  
2 most prominent poets, some of whom were also the most renowned scholars of *waka* at that time. The  
3 Imperial Anthologies were compiled from the tenth to the fifteenth century (Table 1). Poets deeply  
4 revered and intensively studied past Imperial Anthologies and wished their poems to be selected for  
5 future ones. Consequently, this paper focuses on the Imperial Anthologies.

6 As is the case with classical poetry in other regions and periods, *waka* poets were encouraged to  
7 study and imitate great poems of the past. The rhetorical technique of *honkadōri* (allusive variation)  
8 borrows material and phrasing from an older poem or poems (Brower and Miner, 1961; Bialock, 1994).  
9 An example of *honkadōri* is found in a poem by Kiyohara no Fukayabu:

10 Mukashi mishi  
11 haru wa mukashi no  
12 haru nagara  
13 wa ga mi hitotsu no  
14 arazu mo arukana.

15 The original is one of the most famous poems by Ariwara no Narihira:

16 Tsuki ya aranu  
17 haru ya mukashi no  
18 haru naranu  
19 wa ga mi hitotsu wa  
20 moto no mi ni shite.



1 Sharing 14 syllables, these poems are so strikingly similar that Kiyohara no Fukayabu might have  
2 been accused of plagiarism according to present standards. However, *honkadōri* was accepted rhetoric  
3 theorized in *Kindai Shūka* (1209) by Fujiwara no Teika, an anthologist of two Imperial Anthologies.  
4 *Honkadōri* was regarded as a means to enrich and deepen the world behind the poem and represented a  
5 way to show respect to the great poems of the past. This is one of the reasons why the intensive study  
6 of old poems was encouraged in the periods during which the Imperial Anthologies were compiled. To  
7 understand the beauty of *waka*, the audience needs to have deep knowledge of the precedents on which  
8 the poems are based (Konishi et al., 1984). If a poem was a *honkadōri* of an older poem, they could  
9 have easily pointed it out. The widespread practice of *honkadōri* justifies assuming a phylogenetic  
10 network structure among poems. It also allows us to measure the cultural influence of a poem based  
11 on the number of children in the phylogenetic network.

## 12 Dataset

13 The present study used the *waka* database ([https://lapis.nichibun.ac.jp/waka/index\\_era.html](https://lapis.nichibun.ac.jp/waka/index_era.html)) created  
14 by Katsuhiko Seta and maintained by the International Research Center for Japanese Studies. Each  
15 poem in the database is included in an anthology, the date of publication of which ranges from ca. 700  
16 to 1527. This database covers the periods when Japanese classical poetry was prolific and creative.  
17 Some anthologies lack a date of publication and were used as the validation set. Because the database  
18 contains other forms of poetry, poems with more or fewer than five lines were excluded. In addition,  
19 poems with lacunae were excluded. Regarding the data cleansing, line separators were deleted and all  
20 characters were replaced with *hiragana*, the most widely used phonetic lettering system in Japanese.

21 Several variants of poems may be found in an anthology, and some famous poems are included  
22 in multiple anthologies, resulting in multiple entries of a single poem in the database. To determine  
23 the best method to eliminate multiple occurrences of a single poem, the Levenshtein ratio  $l_{ij}$ , which  
24 measures the closeness of two sequences, of poems  $i$  and  $j$  was calculated. Figure 1 shows a histogram  
25 of the maximal Levenshtein ratio of each poem and all other poems, *i.e.*,  $\max_j l_{ij}$ . This histogram  
26 is bimodal, suggesting that a poem should be identified as an existing one if their Levenshtein ratio  
27 exceeds 0.8. Thus, newer poems satisfying this criterion were excluded from the dataset, except in the  
28 analysis shown in Fig. 7. The database contains a total of 206 965 poems in 496 anthologies. The data  
29 cleansing and elimination of the poems identified as existing ones resulted in 146 738 distinct poems,  
30 6343 of which belonged to the validation set.

31 To examine the historical development of *waka*, a total of 24 training sets were used. Training  
32 set 0 contains all poems. Because *Kin'yōshū* has three versions, there are 23 versions of the Imperial  
33 Anthologies (Table 1). Training set  $i$  ( $i = 1, \dots, 23$ ) contains all the anthologies no later than  
34 Imperial Anthology  $i$ . The single validation set was used for all training sets.

35 The database of *honkadōri* was constructed from a modern critical edition of *Shinkokinshū* (Tanaka  
36 and Akase, 1992), in which *honkadōri* reached its highest sophistication. There are 418 poems with  
37 *honkadōri* in *Shinkokinshū*. There are 450 *honkadōri*-original pairs because some poems borrow phrases  
38 from more than one older poem.

## 39 Neural language model and distance metric

40 The vector representation of poems was obtained using BERT (Devlin et al., 2019), a Transformer-  
41 based language model (Vaswani et al., 2017). The language model and data loader were implemented  
42 with HuggingFace Transformers (Wolf et al., 2020) using default parameters unless otherwise stated.  
43 The dimensionality of the feed-forward layers was set to 768. The training sets and validation set were  
44 tokenized by SentencePiece with a token size of 5000 (Kudo and Richardson, 2018). Early stopping  
45 was used to prevent overfitting of the neural language model. If the validation loss had not improved  
46 for the last 10 epochs, the training was stopped, and the parameter values with the smallest loss were  
47 saved. The model was trained four times with different initial parameter values.

1 To measure the similarity between poems, the average vector of the intermediate representation  
2 of all tokens of the last hidden layer was calculated for each poem. The Euclid distance between the  
3 vectors was used as the similarity measure. Because BERT is not a metric learning model, for the  
4 average vectors  $\mathbf{a}$ ,  $\mathbf{b}$ ,  $\mathbf{c}$ , and  $\mathbf{d}$ , the comparability of the distances  $|\mathbf{a} - \mathbf{b}|$  and  $|\mathbf{c} - \mathbf{d}|$  is not necessarily  
5 guaranteed. This means that methods that depend on the axioms of metric space, such as linear  
6 regression and logistic regression, might not be trustworthy. Rather, methods that depend only on a  
7 comparison of the distances from a vector, such as  $|\mathbf{a} - \mathbf{b}|$  and  $|\mathbf{a} - \mathbf{c}|$ , should be used, as these can  
8 be more reliable. Thus, in classification tasks, the  $k$ -nearest neighbor algorithm is used throughout  
9 this paper instead of logistic regression.  $k$  with the greatest validation accuracy was selected from  
10  $k = 1, 3, 5, 7, 9$  using leave-one-out cross-validation.

11 The phylogenetic network was estimated as follows. Each poem is a node in the directed network.  
12 The parent poem of a given poem is the poem closest to it among the poems older than it. Note  
13 that identifying the parent poem of the poem whose vector representation is  $\mathbf{a}$  is done solely by a  
14 comparison of  $|\mathbf{a} - \mathbf{b}|$  and  $|\mathbf{a} - \mathbf{c}|$ . Through this construction, the network comprises a set of directed  
15 trees. As the poems in the oldest anthology have no parent, they become the root nodes. Throughout  
16 this paper, the phylogenetic network estimated by the neural language model trained using training  
17 set  $i$  is referred to as phylogenetic network  $i$ . Phylogenetic network 0 is sometimes simply referred to  
18 as the phylogenetic network. An example of phylogenetic network 0 is visualized in Supplementary  
19 Figure 1.

20 The reasonability of the estimated phylogenetic network was evaluated as follows. In molecular  
21 phylogenetics, species with similar genotypes are placed close to each other. Hence, if a phylogenetic  
22 network is reasonable, each poem and its parent are sufficiently close. This suggests that the distance  
23 between a poem and its parent can be a measure of reasonability. However, as we have seen, the distance  
24 itself is unreliable. In other words, the meaning of the summation of distances such as  $|\mathbf{a} - \mathbf{b}| + |\mathbf{a} - \mathbf{c}|$   
25 remains unclear. Thus, the rank order of the distance is more reliable than the raw distance because it  
26 is calculated based on a comparison of the distances from only  $\mathbf{a}$ . Let us assume that poem  $i$  belongs  
27 to anthology  $a$  and that there are  $n_a$  poems before and  $m_a$  poems after anthology  $a$ . The parent of  
28 poem  $i$  is closest to poem  $i$  among the preceding  $n_a$  poems. Let us define  $r_i = (k_i - 1)/(n_a + m_a - 1)$   
29 if the parent is  $k_i$ -th closest to poem  $i$  among  $n_a + m_a$  poems.  $\bar{r}$  is defined as the average of  $r_i$  over  
30 all non-root poems. The smaller the value of  $\bar{r}$ , the more reasonable the phylogenetic network.

31 Let us note that  $\bar{r}$  can distinguish divergence from and convergence to a poem. Supplementary  
32 Figure 2 shows older and newer poems in lighter and darker colors, respectively. In Supplementary  
33 Figure 2a, poems are diverging from the oldest one. The phylogenetic network shown by the arrows  
34 is constructed by connecting each poem (child) with the closest one among the older poems (parent).  
35 Here, all anthologies are assumed to have only one poem each. The parent of the poem marked by  
36 the asterisk (poem  $*$ ) is the closest to poem  $*$  among all other poems, *i.e.*,  $k_* = 1$  and  $r_* = 0$ . If  
37 a phylogenetic network is constructed from the time-reversed data (Supplementary Figure 2b), the  
38 parent of poem  $*$  is the fifth closest to poem  $*$  among all other poems, *i.e.*,  $k_* = 5$  and  $r_* = 1$ . For  
39 most of the other poems,  $k_i$  and  $r_i$  are greater in the time-reversed phylogenetic network. Thus, poems  
40 diverging from a poem (Supplementary Figure 2a) and poems converging to a poem (Supplementary  
41 Figure 2b) result in a low and a high  $\bar{r}$ , respectively. Note that time reversal does not affect  $\bar{r}$  for  
42 continuously transitioning poems (Supplementary Figure 2c, d). Hence,  $\bar{r}$  can be used as an index of  
43 divergence and convergence.

## 44 Results

### 45 Validity of the estimated phylogenetic networks

46 First, the validity of the estimated phylogenetic networks was examined by comparing it with the  
47 ground truth. Specifically, the estimated child–parent relationship was compared with the original–  
48 *honkadori*-poem relationship described in a critical edition of *Shinkokinshū* (Tanaka and Akase, 1992).

1 Figure 2 shows the relationship between 418 *honkadori* poems in *Shinkokinshū* (child) and their original  
 2 poems (parent). The parent–child relationship in the estimated phylogenetic network is classified into  
 3 seven categories: the original poem of *honkadori* is the parent (parent), an ancestor but not the parent  
 4 (ancestor), a node within 4 hops (close relative), a node more distant than 4 hops but in the same  
 5 connected component (remote relative), in a different connected component (unconnected), found in  
 6 an anthology later than *Shinkokinshū* (anachronism), and not found in the database (not found). If  
 7 a poem is a *honkadori* poem of more than one old poem, the best one was used. This figure shows  
 8 that 6.9% of *honkadori* pairs are captured as the parent and child in phylogenetic network 0 and that  
 9 72.9% of pairs belong to different connected components. This might not seem to be a very accurate  
 10 estimate.

11 However, this is a statistically significant result if it is compared with a hypothetical random  
 12 phylogenetic network (null hypothesis), in which the parents of poems in *Shinkokinshū* are randomly  
 13 drawn from 48 057 poems older than *Shinkokinshū*. Three original poems at most have been identified  
 14 for a *honkadori* poem in *Shinkokinshū*. If one of these poems is the parent in the estimated phylogenetic  
 15 network, the *honkadori* relationship is regarded to be guessed right. Thus, the parent of a poem in the  
 16 random phylogenetic network is identical to one of the original poems of *honkadori* with a probability  
 17 of  $p_h = 3/48\,057$  at most. The number of parents identical to the original,  $n_h$ , obeys the binomial  
 18 distribution  $P(n_h) = p_h^{n_h} (1-p_h)^{n-n_h} n! / \{n_h!(n-n_h)!\}$ , where  $n = 4 \times 418$  because Fig. 2 is the average  
 19 of the four phylogenetic networks. The probability that  $n_k \geq 17$ , *i.e.*, more than 1%, which is much  
 20 lower than 6.9%, of the parents are identical to the original is less than  $p = 5 \times 10^{-32}$ . Hence, the  
 21 phylogenetic network estimated by using the vector representation of a neural language model succeeds  
 22 in identifying at least some of the *honkadori* relationships. This conclusion is robust for change in the  
 23 model parameter values of BERT (Supplementary Figure 3).

## 24 Consistency and reasonability of phylogenetic networks and the arrow of 25 time

26 Second, the cultural influence of poems judged by the models with different initial parameter values  
 27 was quantified to examine the consistency of the estimation. If a poem has a large number of children,  
 28 it is potentially the original poem of a large number of *honkadori* poems, implying that it should be  
 29 judged to be a poem with a great influence on the following poems. Hence, this paper uses the number  
 30 of children of a poem in the estimated phylogenetic network as a proxy variable of its cultural influence.  
 31 However, because, by construction, an earlier poem tends to have a larger number of children than a  
 32 later one, it is not fair to compare the raw number of children. Thus, by standardizing the number  
 33 of children for each anthology, we can quantify the cultural influence of each poem relative to other  
 34 poems in the same anthology. The Spearman correlation coefficients of the standardized number of  
 35 children averaged for all six pairs of the four neural language models are shown in Fig. 3. Most exhibit  
 36 a positive correlation coefficient. Thus, the estimated cultural influence of a poem is consistent among  
 37 the neural language models with different initial parameter values.

38 Third, the reasonability of the structure of the estimated phylogenetic network and the existence of  
 39 the arrow of time were examined. In molecular phylogenetics, closely related species are placed close  
 40 to each other in the phylogenetic tree. Similarly, if a phylogenetic network of *waka* is reasonable, each  
 41 poem and its parent are expected to be sufficiently close. To test this expectation, the phylogenetic  
 42 networks estimated from the dataset in which the order of anthologies is reversed (reversed) and that  
 43 in which the order of anthologies is randomly shuffled (shuffled) were made along with phylogenetic  
 44 network 0 (real). Twenty shuffled networks were made for each neural language model.  $\bar{r}$  measures  
 45 the average distance of the parent–child relationship. Figure 4 shows that the real data exhibit the  
 46 lowest value of  $\bar{r}$ . This means that the phylogenetic network estimated from the real data is more  
 47 reasonable than those estimated from the reversed or shuffled data. The present analysis is consistent  
 48 with and extends the results of Hughes et al. (2012), which showed a gradual change in literary style.  
 49 Particularly, by using  $\bar{r}$ , we can distinguish the real data from the time-reversed data. A low  $\bar{r}$  of the  
 50 real data shows diversification, rather than continuous transition, in *waka*. This result indicates that

1 the arrow of time is present and observable in the evolution of *waka*.

## 2 Cultural influence of poems

3 Fourth, the cultural influence of poems judged from the dataset up to a certain time point was compared  
4 with that judged from the whole dataset. Specifically, the number of children in phylogenetic network  
5  $i$  was compared with that in phylogenetic network 0. In other words, the congruence of the cultural  
6 influence of a poem measured by a language model trained with a limited corpus of poems and that  
7 with the whole corpus in the database is examined. It is likened to asking poets in the *Shinkokinshū*  
8 (1205) era “Which do you think are most influential among *Kokinshū* (905) poems?” and comparing  
9 their answers with ours. The standardized number of children of poems up to Imperial Anthology  
10  $i$  in phylogenetic networks  $i$  and 0 was positively correlated in most cases (Fig. 5a). Moreover, the  
11 standardized number of children until Imperial Anthology  $i$  in phylogenetic network  $i$  was positively  
12 correlated with the number of children after Imperial Anthology  $i$  in phylogenetic network 0 (Fig. 5b).  
13 Therefore, the cultural influence of a poem from the viewpoint of a certain time point is correlated  
14 with the cultural influence of the poem from the viewpoint of a later time point. Analogically, the  
15 influence of *Kokinshū* poems evaluated by poets in the *Shinkokinshū* era is positively correlated with  
16 our estimation of the influence of *Kokinshū* poems on poems later than *Shinkokinshū*. However, the  
17 correlation was not so strong presumably because of the collective memory decay of a poem (Candia  
18 et al., 2019).

## 19 Interpolation and extrapolation

20 Fifth, to investigate whether a culture in a period is an intermediate form of those in the preceding and  
21 following periods and whether the past history allows us to predict future culture, the interpolation and  
22 extrapolation of the Imperial Anthologies were examined using classification. The  $k$ -nearest neighbor  
23 algorithm discriminating the first Imperial Anthology *Kokinshū* (class label 0) and the last Imperial  
24 Anthology *Shinshokukokinshū* (class label 1) was applied to all Imperial Anthologies. Figure 6a shows  
25 the average of the class label predicted by the model for all Imperial Anthologies. The validation  
26 accuracy of leave-one-out cross-validation is used as the average class label of the labeled anthologies.  
27 This figure shows that the average class label of the anthologies in between exhibit intermediate values,  
28 that is, this  $k$ -nearest neighbor classifier can interpolate the anthologies between the two anthologies.  
29 However, the  $k$ -nearest neighbor model discriminating *Kokinshū* and *Shinkokinshū* exhibits no signs of  
30 extrapolation (Fig. 6b). The anthologies following *Shinkokinshū* are no more *Shinkokinshū*-like than  
31 *Shinkokinshū*. Taken together, these results indicate the presence of detectable and gradual, albeit  
32 unpredictable, change over time.

## 33 Effect of being selected for an Imperial Anthology

34 Sixth, the effect of being selected for an Imperial Anthology was examined. The Imperial Anthologies  
35 were so authoritative that being selected for them is expected to increase the number of children.  
36 This expectation was tested by comparing a pair of poems, the first of which is contained in Imperial  
37 Anthology  $i$ , referred to as  $x$ , and the second of which is not contained in Imperial Anthology  $i$  but  
38 regarded to be similar to the first, referred to as  $y$ . All poems that are included in Imperial Anthology  
39  $i$  but first appear in an earlier anthology were classified as  $x$ . For each  $x$ ,  $y$  was sampled under the  
40 condition that it appears first in the same anthology as  $x$  and that it gives birth to the same number  
41 of children as  $x$  in phylogenetic network  $i$  in the period preceding Imperial Anthology  $i$ . The numbers  
42 of children after Imperial Anthology  $i$  in phylogenetic network 0 were compared for the pairs of  $x$  and  
43  $y$  (Fig. 7).

44 For most Imperial Anthologies, the average number of children of  $x$  after the Imperial Anthology  
45 tended to be greater than that of  $y$ , meaning that the effect of being selected is positive. In most  
46 cases, a poem selected for an Imperial Anthology gains 1.1–1.5 times more children than a poem that

1 is not. However, this effect is inconclusive for some anthologies, particularly for the later ones. This  
 2 may be the result of a decline in quality of, or loss of interest in, these Imperial Anthologies (Keene,  
 3 1999). The positive effect of most Imperial Anthologies can be interpreted in two ways. The first is  
 4 that entry into the Imperial Anthologies boosted its fame and increased its number of children. This  
 5 is a form of the Matthew effect (Merton, 1968). The second is that  $x$  is closer to the taste of a later  
 6 period than  $y$ , and thereby had more children in later periods. At any rate, these results indicate that  
 7 the number of children is largely affected by chance.

8 The difference among the three versions of *Kin'yōshū* (ID 5–7) is worth noticing (Supplementary  
 9 Figure 4). Of these, being selected for the first and third versions (ID 5 and 7) does not seem to affect  
 10 the number of children. In other words, it exhibits a weaker effect than the second version (ID 6).  
 11 This is consistent with the fact that the second version was the most circulated. Particularly, the third  
 12 version had been forgotten until the nineteenth century (Keene, 1999). The difference among them  
 13 supports the existence of the Matthew effect.

## 14 Model

15 Both the chance factor’s role in the number of children and the impossibility of future prediction  
 16 suggest that randomness is a key feature of the evolution of *waka*. This leads us to a model that  
 17 qualitatively replicates these results. This model assumes that a poem is generated in the vicinity of  
 18 existing poems. Specifically, poem 0 is generated on  $\mathbf{x}_0 = \mathbf{0}$ , where  $\mathbf{x}_i$  is a  $d$ -dimensional vector. Poem  
 19  $t \geq 1$  is randomly drawn from the Gaussian mixture

$$p(\mathbf{x}_t) = \sum_{s=0}^{t-1} \frac{k^{t-1-s}(1-k)}{1-k^t} \frac{1}{(2\pi)^{d/2}} \exp\left(-\frac{|\mathbf{x}_t - \alpha\mathbf{x}_s|^2}{2}\right),$$

20 where  $k$  is the decay constant and  $\alpha$  is a positive constant less than one.  $\alpha$  makes this process a mean-  
 21 reverting self-excitatory process. A poem facilitates the generation of another poem in the vicinity  
 22 of itself and the origin. This is a self-exciting stochastic dynamical model (Golosovsky and Solomon,  
 23 2012). Because the influence of a poem decays at rate  $k$ , the poems as a whole can exhibit a long-term  
 24 drift. If a poem is generated in the close vicinity of another poem by chance, the number of children  
 25 of the latter can be boosted.

26 Figure 8 shows the results of the model with 24 000 poems and the following parameter values:  
 27  $d = 100$ ,  $k = \exp(-1/5000)$ , and  $\alpha = 0.6$ . To obtain poems in the steady state, a total of 74 000 poems  
 28 were generated. The first 50 000 poems were discarded, and the last 24 000 poems were divided into 24  
 29 anthologies containing 1000 poems each. The  $k$ -nearest neighbor algorithm with the first anthology as  
 30 class 0 and the 12th anthology as class 1 shows a steady increase up to the 11th anthology, but plateaus  
 31 thereafter (Fig. 8a). This is consistent with the possibility of interpolation and the impossibility of  
 32 extrapolation. The phylogenetic network made in the same way as *waka* has a lower  $\bar{r}$  than those made  
 33 from the time-reversed or shuffled anthologies (Fig. 8b). This is because poems diverging from and  
 34 converging to a point exhibit a low and high  $\bar{r}$ , respectively. The standardized number of children before  
 35 anthology  $i$  is positively correlated with that after anthology  $i$  (Fig. 8c). Mean reversion is essential  
 36 because the model without mean reversion ( $\alpha = 1$ ) exhibits a much weaker correlation (Supplementary  
 37 Figure 5). Thus, the model succeeded in replicating the results qualitatively.

## 38 Discussion

39 The present study attempted to elucidate the evolutionary dynamics of culture in estimating the  
 40 phylogenetic network of *waka*. The results are summarized as follows. First, the phylogenetic network  
 41 reflects a significant part of the *honkadōri* relationship. Second, the vector representation obtained  
 42 using BERT gives reproducible results. Third, the estimated phylogenetic network is distinguishable  
 43 from the phylogenetic network constructed from the time-reversed data. That is, the arrow of time

1 is observable in a literary genre. Fourth, the cultural influence of a poem, which is measured by the  
2 number of children at a certain time point, is correlated with that at a later time point. Fifth, we can  
3 successfully perform the interpolation, but not the extrapolation, of the poetic style. In other words,  
4 we cannot predict a style in the future. Sixth, the number of phylogenetic children increases after  
5 being selected for an anthology. Last, a mean-reverting self-excitation model replicates these results.  
6 If not complete, these results are at least partial answers to the questions raised at the beginning of  
7 the paper.

8 It is quite natural to ask whether these results are universal to other literary genres and creative  
9 works. Although *honkadori* is a rhetorical technique characteristic of *waka*, borrowing and imitating  
10 phrases from old great literary works are prevalent in a diverse range of classical literary genres. Kato  
11 et al. (1979) pointed out that the *xikūn* style in the early Sung dynasty and “Waste Land” by T. S. Eliot  
12 can be seen as a parallel to *honkadori*. Gahan (1987) presented a detailed analysis of a tragedy from  
13 the Silver Age of Latin literature and illustrated the abundance of *imitatio* and *aemulatio*, that is, the  
14 technique of borrowing phrases and ideas. Hence, the technique of borrowing is universal to classical  
15 literary genres. An analysis similar to the present study can also shed new light on these genres.

16 To analyze other creative works, the method of estimating the phylogenetic structure in this paper  
17 should be extended. The present study has assumed that a poem is a child of another poem and ignored  
18 exogenous factors. This assumption is justifiable because *waka* had been the most authoritative genre  
19 in Japanese literature and, consequently, influence from *waka* to other genres in Japanese literature  
20 exceeds *vice versa*. However, because the influence of Chinese literature is indisputable (Konishi et al.,  
21 1984), this should be taken into account in future analyses. In addition, although the present study  
22 has assumed that there is only one parent poem for a given poem, a poem can be a *honkadori* poem  
23 of multiple poems. Thus, phylogenetic networks allowing multiple parents should be examined in the  
24 future.

25 The effect of being selected for an Imperial Anthology is greater in earlier Imperial Anthologies  
26 but diminished in later ones. There are two possible explanations for this result. First, the Imperial  
27 Anthologies after *Shinkokinshū* were regarded to be of low quality and thus less intensively studied  
28 (Keene, 1999). Second, poems in older anthologies tend to have a larger number of children than  
29 those in newer ones. Thus, poems in the later Imperial Anthologies tend to have a smaller number of  
30 children, deteriorating the signal-to-noise ratio.

31 The present paper has proposed a model that replicates the results qualitatively. In this model,  
32 a poem is an event in a self-exciting point process. Although the timing of poem generation was not  
33 formulated in the present model, the spatiotemporal Hawkes process might be hopeful. Identifying a  
34 poem with an individual in population genetics, we can regard this model to be closely related to the  
35 neutral theories, which stress the importance of the interaction of selection and drift (Kimura et al.,  
36 1968; Ohta, 2002; Akashi et al., 2012). Testing whether the word frequency obeys the distribution  
37 predicted from the neutral model will be of interest (Bentley and Shennan, 2003; Bentley et al., 2004).  
38 Although there may sometimes be fixed directionality in evolution, the results of the present study  
39 indicate that the cultural evolution of *waka* is approximated by mean-reverting self-excitation.

40 However, this study has some limitations. First, the training set may not have been sufficiently  
41 large. Including other literary genres in the corpus might improve the performance. In particular, the  
42 poems in *The Tale of Genji* had a substantial influence on the history of classical Japanese literature.  
43 Taking the influence from and to *The Tale of Genji* into account could enable us to create a more  
44 holistic picture of the historical development of Japanese literature. However, other literary genres,  
45 such as novels, lack dating more often than *waka*. In fact, many of the poems that were excluded  
46 from the training sets and included in the validation set because of missing dating were from novels.  
47 Thus, this may be difficult to implement. Second, using the chronology of the *waka* anthologies as  
48 the chronology of the poems in them may not be appropriate in some cases. This is because an  
49 anthology can contain a poem by a poet from an older generation. Thus, the order of poem writing  
50 and publication may be reversed. Incorporating information from the poem description (*kotobagaki*),  
51 author, and volume name in the anthology might also improve the results. Third, as stated above, the  
52 influence of Chinese literature was not taken into consideration in the present analysis.

1 There are several possible future directions. First, the present paper has left a detailed analysis of  
2 the specificities of each anthology for future work. What is and is not influential, what determines the  
3 strength of influence, and what are the long-term trends in the strength of influence remain questions  
4 that need to be addressed. Second, examining whether the phylogenetic network reflects the schools  
5 of *waka*, such as *Nijō*, *Kyōgoku*, and *Reizei*, would also be of interest. Third, a method to measure  
6 the speed of evolution needs to be developed. This was difficult to measure in the present analysis,  
7 which utilized an intermediate vector representation of a neural language model. If a large number  
8 of nearly identical poems are in the training set, they might take a large volume in the vector space.  
9 Conversely, if we introduce Chinese poetry into the training set, the volume covered by *waka* would  
10 be compressed. This means that the distance between successive poems can be affected by both the  
11 speed of evolution and the relative abundance of similar poems. Thus, the development of a measure  
12 of evolution speed that is insensitive to the relative abundance is needed. Whether the dynamics  
13 follow biased cultural transmission (Henrich, 2001) would also be of interest. Because the styles  
14 of English authors are similar among contemporaries and differ from preceding generations (Hughes  
15 et al., 2012), conformity bias in a generation and anti-conformity bias between generations could also be  
16 observed in *waka*. In addition, we might be able to measure conformity in a school and anti-conformity  
17 between schools. Fourth, applying this method to other literary genres and creative works would be  
18 of interest. Specifically, whether similar results can be obtained for longer works such as novels should  
19 be examined. Other forms of creative works, such as music (Nakamura and Kaneko, 2019), fine arts  
20 (Cetinic et al., 2019; Sandoval et al., 2019), and Internet memes, should also be subjects of the present  
21 method. Self-supervised representation learning, such as SimCLR (Chen et al., 2020) for images, gives  
22 a vector representation of input without labeled data. This can be used in the same way as the vector  
23 representations produced by language models. Fifth, the present study may provide hints about how  
24 to train generative artificial intelligence (AI) with output of another generative AI. This procedure  
25 has been reported to deteriorate the quality of output (Alemohammad et al., 2023; Shumailov et al.,  
26 2024). If this is the result of an unbounded random walk of the generated data, it can be mitigated  
27 by mean reversion, which led to the persistent influence of poems in the present model. Similarly to  
28 the model’s mean reversion, selecting the output that is closer to the mean of existing creative works  
29 could prevent deterioration. Furthermore, in multimodal generative AI, natural photos and sounds  
30 could not only serve as the “mean” but also prevent model collapse (Shumailov et al., 2024).

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10 38–45

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## 13 **Competing interests**

14 The author declares no competing interests.

## 15 **Data availability**

16 All data used in this paper can be found at [https://lapis.nichibun.ac.jp/waka/index\\_era.html](https://lapis.nichibun.ac.jp/waka/index_era.html). The  
17 program files for data preprocessing, model training, phylogenetic network reconstruction, and figure  
18 generation can be found at <https://github.com/tanaka-takuma-lab/>.

## 19 **Ethical approval**

20 Not applicable.

## 21 **Informed consent**

22 Not applicable.

## 629 **Author contributions**

630 Not applicable.

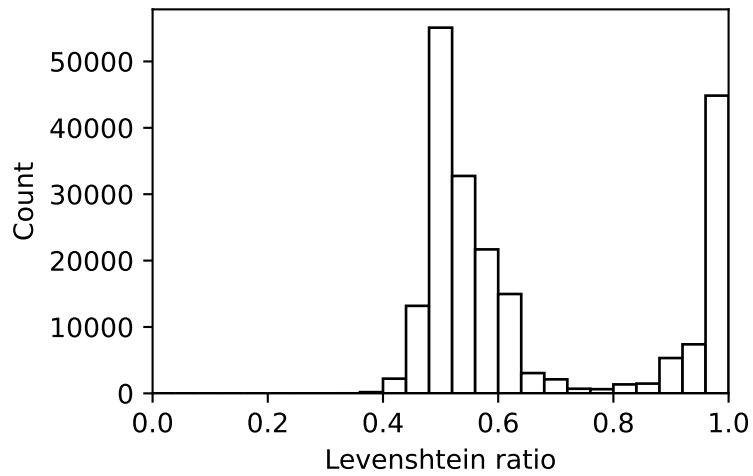


Figure 1: Histogram of the maximal Levenshtein ratio,  $\max_j l_{ij}$ , for all  $i$ .

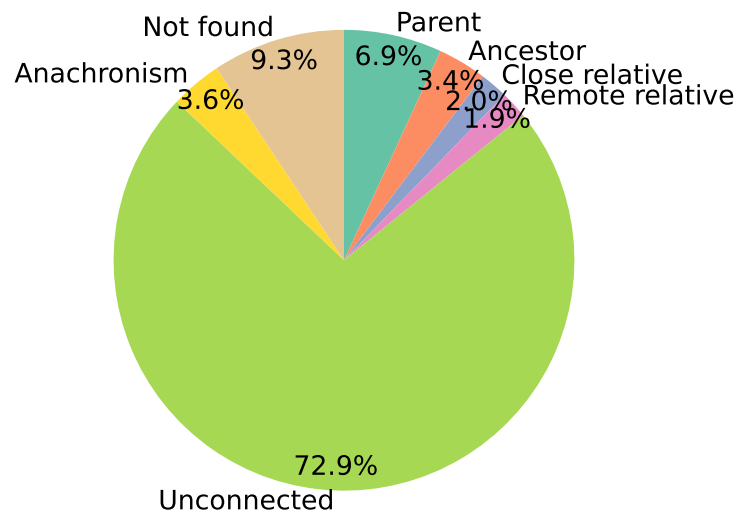


Figure 2: Classification of *honkadori* relationships in the phylogenetic network. The average for four neural language models is shown.

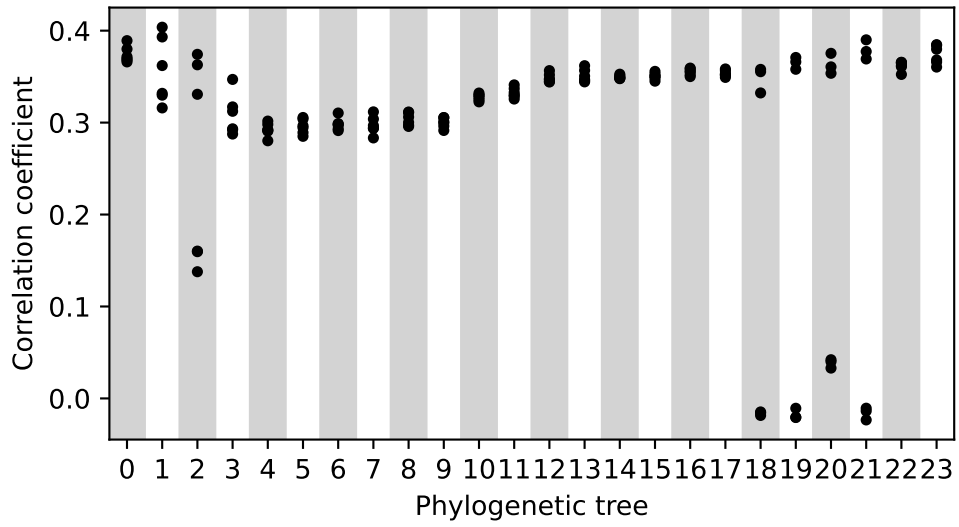


Figure 3: Correlation coefficients of the standardized numbers of children in phylogenetic network  $i$  between neural language models with different initial conditions.

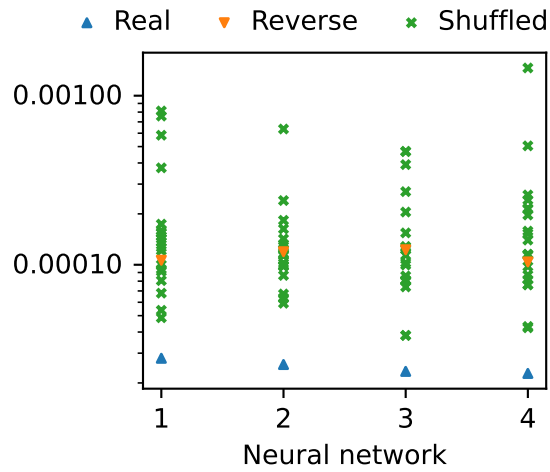


Figure 4: The  $\bar{r}$  values for four neural language models with different initial conditions.

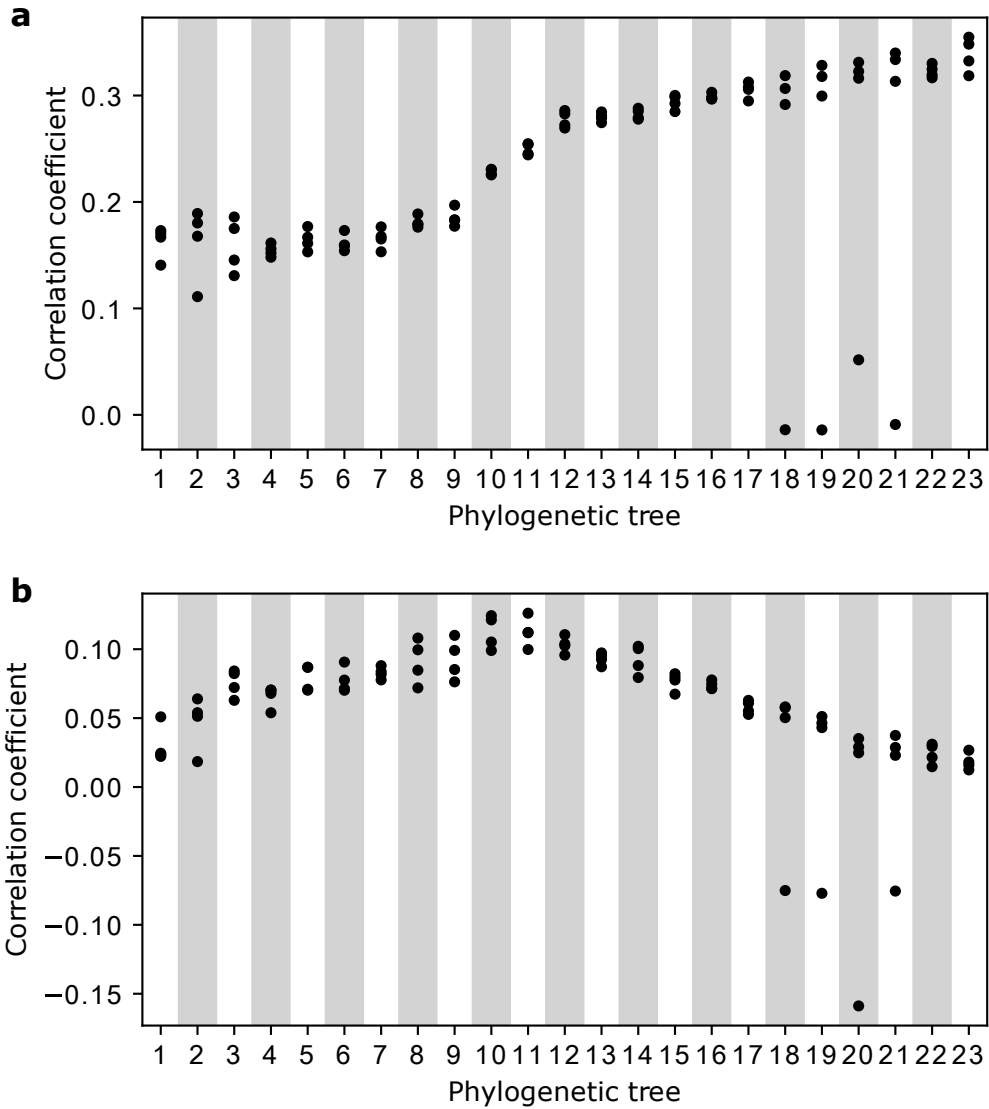


Figure 5: Correlation coefficients of the standardized numbers of (a) children before Imperial Anthology  $i$  in phylogenetic network  $i$  and phylogenetic network 0 and (b) children before Imperial Anthology  $i$  in phylogenetic network  $i$  and children after Imperial Anthology  $i$  in phylogenetic network 0. The panel labels indicate  $i$ .

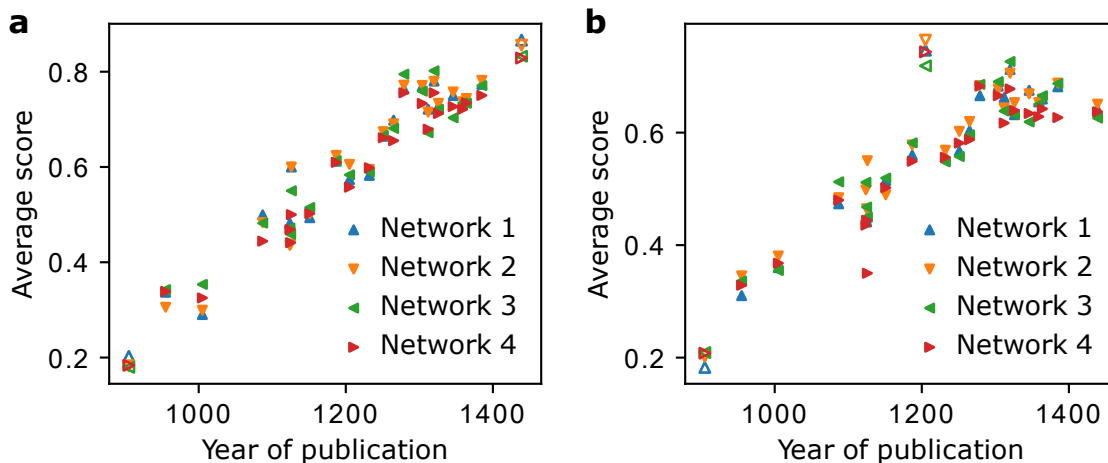


Figure 6: The  $k$ -nearest neighbor classification discriminating (a) the first (*Kokinshū*) and last (*Shinshokukokinshū*) Imperial Anthologies, and (b) the first and tenth (*Shinkokinshū*) Imperial Anthologies. The results for the anthologies used as the training set of the  $k$ -nearest neighbor classification are indicated by open triangles.

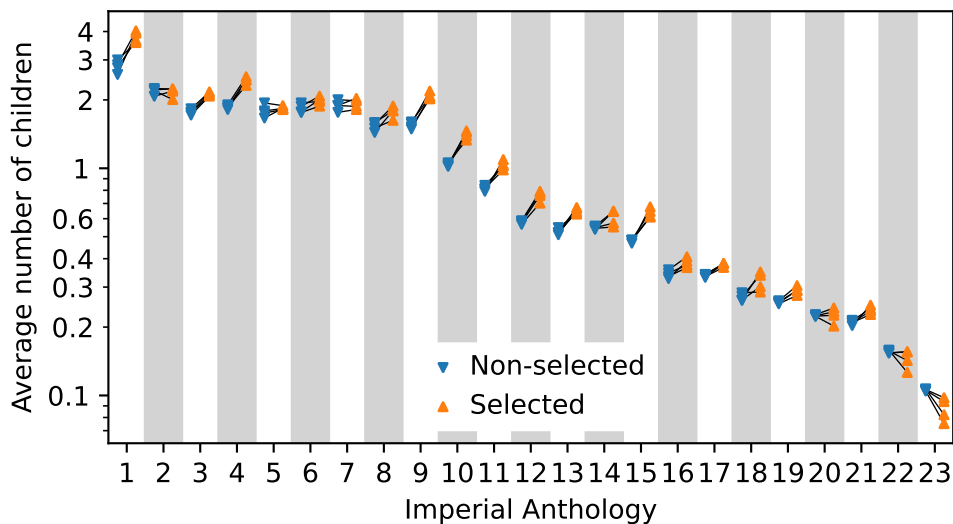


Figure 7: Effects of being selected for an Imperial Anthology. The average numbers of children of poems selected and not selected for Imperial Anthology  $i$  after this anthology are compared in panel  $i$ . The phylogenetic networks estimated from the neural language models with the same initial conditions are connected by lines.

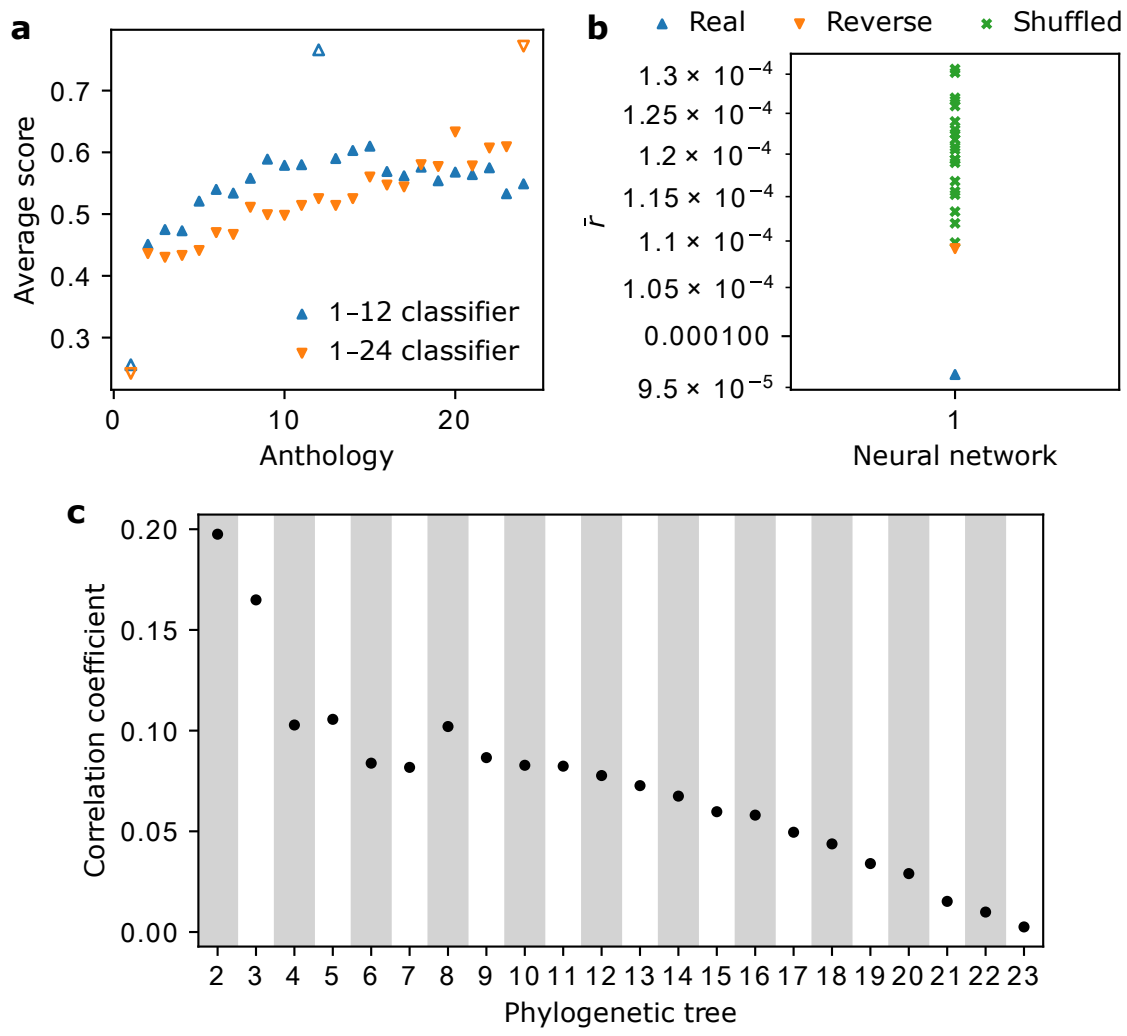
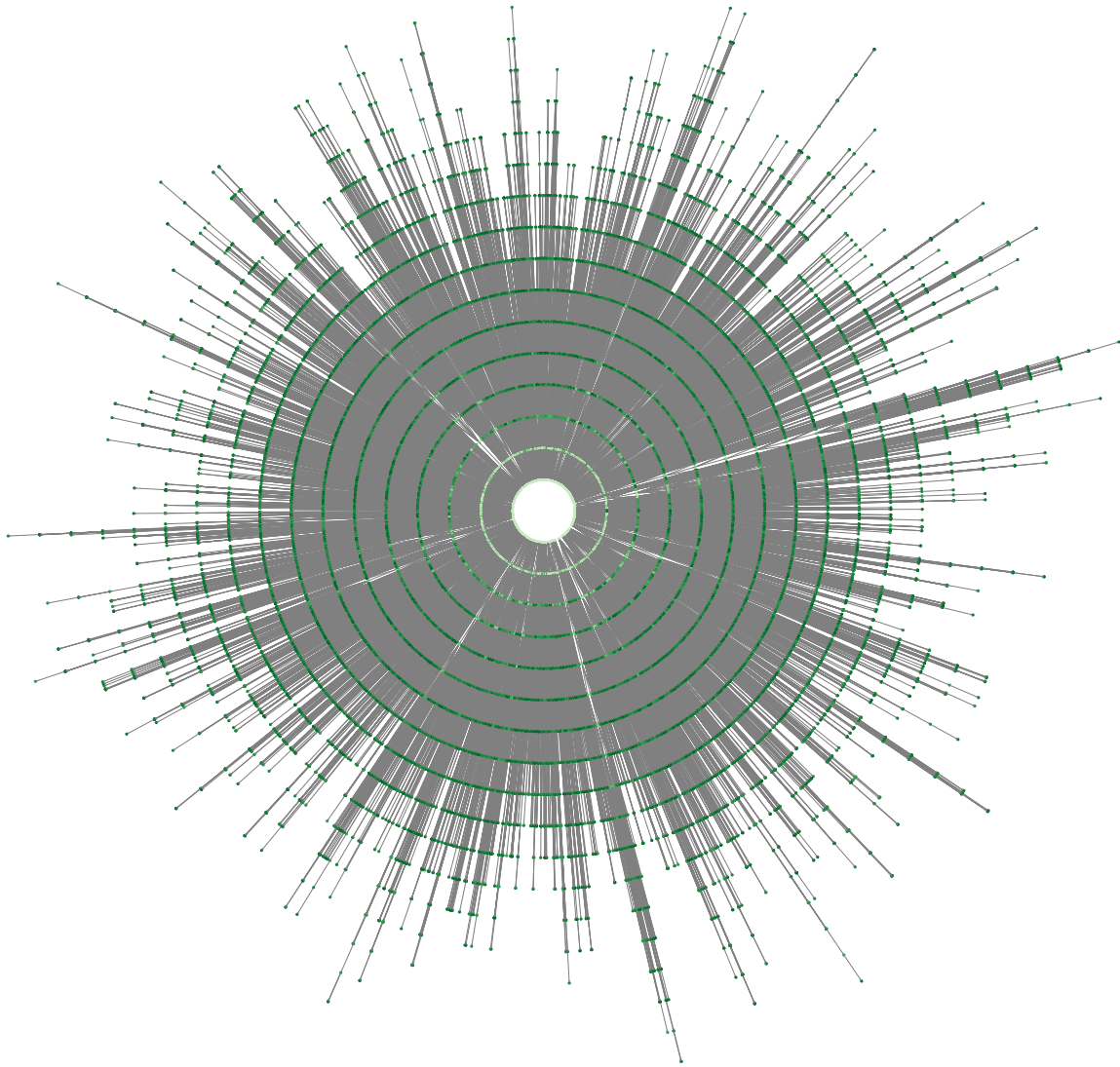
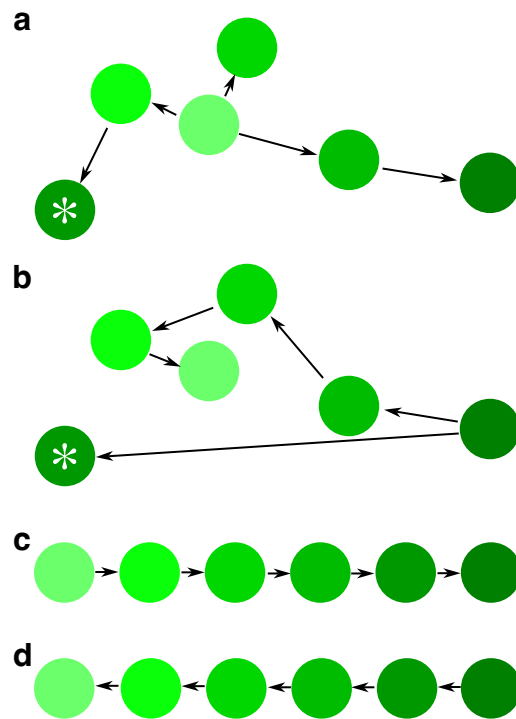


Figure 8: Model results. (a) Performance of  $k$ -nearest neighbor classifiers discriminating anthologies 1 and 12 and anthologies 1 and 24. (b) Values of  $\bar{r}$  for the real, time-reversed, and shuffled data. (c) Correlation coefficient between the standardized numbers of children before and after anthology  $i$ .

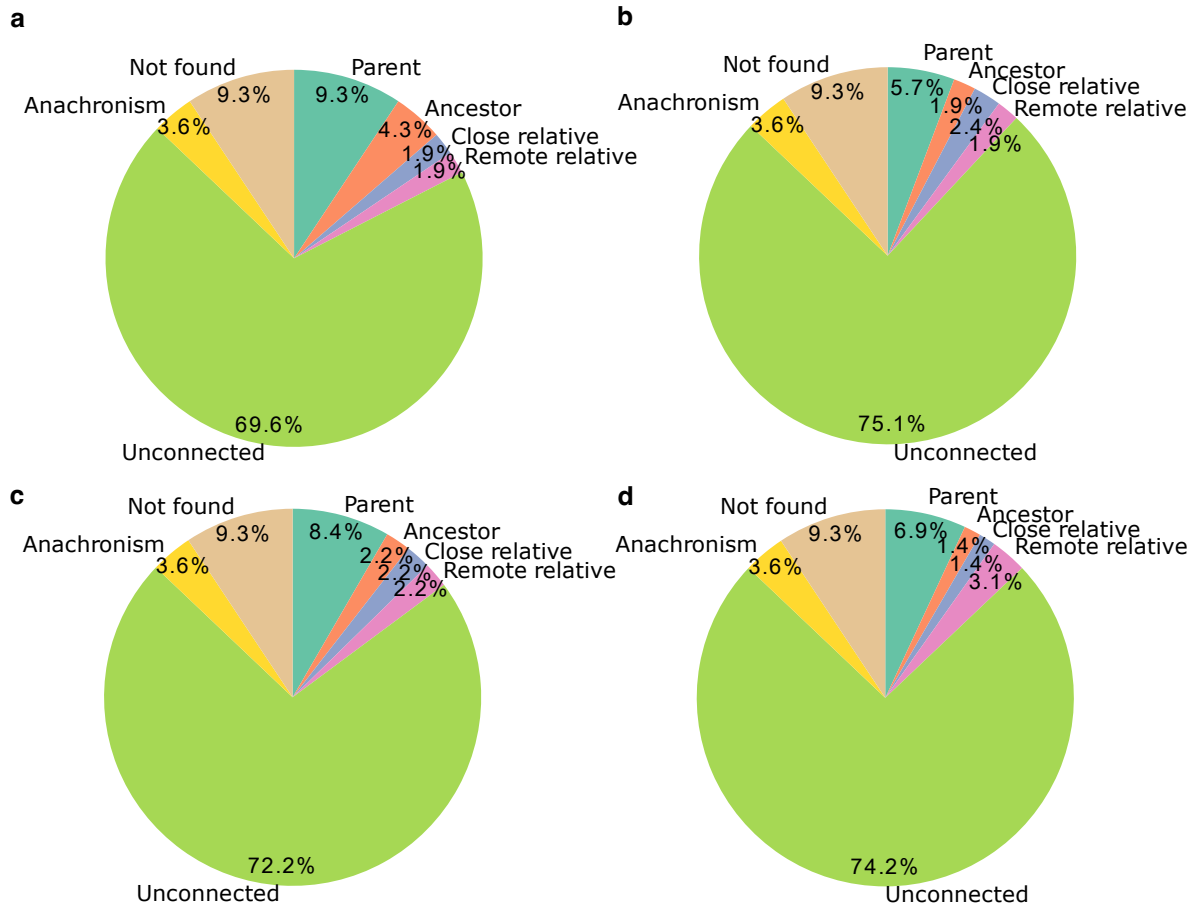




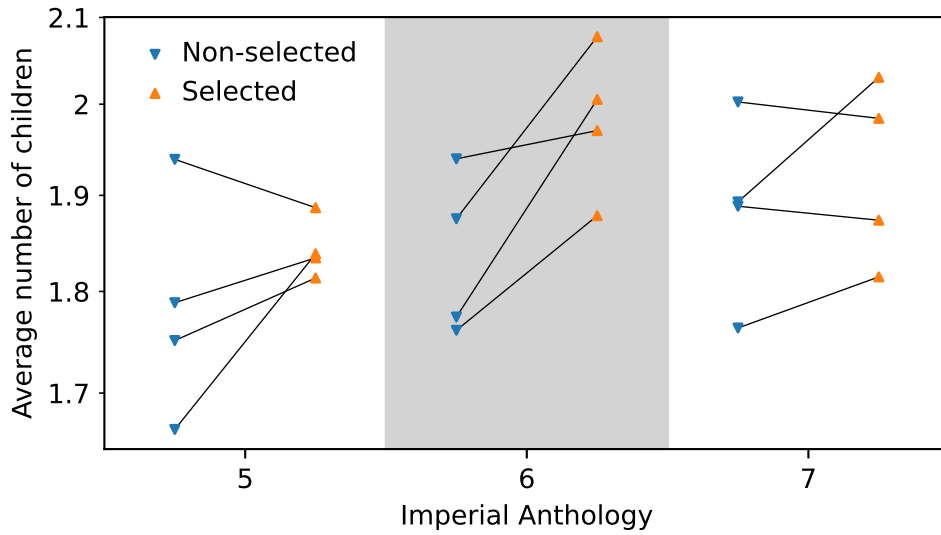
Supplementary Figure 1: A reconstructed phylogenetic network. Poems and parent–child relationships are indicated by colored disks and arrows, respectively. Light and dark green indicate earlier and later poems, respectively. Each of the concentric circles contains one generation of poems, the disks on the innermost circle being the poems in the oldest anthology, *i.e.*, the first generation poems.



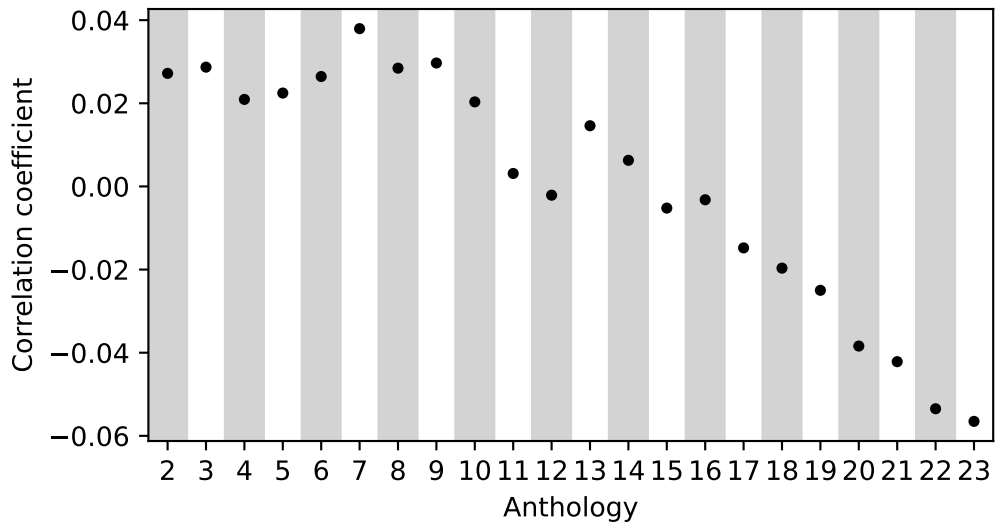
Supplementary Figure 2: Phylogenetic networks diverging from a poem (a), converging to a poem (b), and transitioning continuously (c, d). *b* and *d* are the time-reversed networks of *a* and *c*, respectively.



Supplementary Figure 3: *Honkadori* relationships in the phylogenetic network generated with (a) a token size of 3000, (b) a token size of 10000, (c) the dimensionality of the intermediate layers of 256, and (d) the dimensionality of the intermediate layers of 1024. The same model parameter values were used unless otherwise stated.



Supplementary Figure 4: Effects of being selected for *Kin'yōshū* 1, 2, and 3.



Supplementary Figure 5: Correlation coefficient between the standardized numbers of children before and after anthology  $i$  in the model with  $\alpha = 1$ .