A mathematical model for universal semantics

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Abstract—We characterize the meaning of words with language-independent numerical fingerprints, through a mathematical analysis of recurring patterns in texts. Approximating texts by Markov processes on a long-range time scale, we are able to extract topics, discover synonyms, and sketch semantic fields from a particular document of moderate length, without consulting external knowledge-base or thesaurus. Our Markov semantic model allows us to represent each topical concept by a low-dimensional vector, interpretable as algebraic invariants in succinct statistical operations on the document, targeting local environments of individual words. These language-independent semantic representations enable a robot reader to both understand short texts in a given language (automated question-answering) and match medium-length texts across different languages (automated word translation). Our semantic fingerprints quantify local meaning of words in 14 representative languages across 5 major language families, suggesting a universal and cost-effective mechanism by which human languages are processed at the semantic level. Our protocols and source codes are publicly available on https://github.com/yajun-zhou/linguae-naturalis-principia-mathematica

Index Terms—recurring patterns in texts, semantic model, recurrence time, hitting time, word translation, question answering

1 Introduction

A QUANTITATIVE model for the meaning of words not only helps us understand how we transmit information and absorb knowledge, but also provides foothold for algorithms in machine processing of natural language texts. Ideally, a universal mechanism of semantics should be based on numerical characteristics of human languages, transcending concrete written and spoken forms of verbal messages. In this work, we demonstrate, in both theory and practice, that the time structure of recurring language patterns is a good candidate for such a universal semantic mechanism. Through statistical analysis of recurrence times and hitting times, we numerically characterize connectivity and association of individual concepts, thereby devising language-independent semantic fingerprints (LISF).

Concretely speaking, we define semantics through algebraic invariants of a stochastic text model that approximately governs the empirical hopping rates on a web of word patterns. Such a stochastic model explains the distribution of recurrence times and outputs recurrence eigenvalues as semantic fingerprints. Statistics of recurrence times allow machines to tell non-topical words from topical ones. A comparison of hitting and recurrence times further generates quantitative fingerprints for topics, enabling machines to overcome language barriers in translation tasks and perform associative reasoning in comprehension tasks, like humans.

Akin to the physical world, there is a hierarchy of length scales in languages. On short scales such as syllables, words, and phrases, human languages do not exhibit a universal pattern related to semantics. Except for a few onomatopoeias, the sounds of words do not affect their meaning [1]. Neither do morphological parameters [2] (say,

singular/plural, present/past) or syntactic rôles [3] (say, subject/object, active/passive). In short, there are no universal semantic mechanisms at the phonological, lexical or syntactical levels [4]. Grammatical "rules and principles" [2], [3], however typologically diverse, play no definitive rôle in determining the inherent meaning of a word.

Motivated by the observations above, we will build our quantitative semantic model on long-range and language-independent textual features. Specifically, we will measure the lengths of text fragments flanked by *word patterns* of interest (Fig. 1). Here, a word pattern is a collection of *content words* that are identical up to morphological parameters and syntactic rôles. A content word signifies definitive concepts (like *apple*, *eat*, *red*), instead of serving purely grammatical or logical functions (like *but*, *of*, *the*). Fragment length statistics will tell us how tightly/loosely one concept is connected to another. This in turn, will provide us with quantitative criteria for inclusion/exclusion of different concepts within the same (computationally constructed) semantic field. Such statistical semantic mining will then pave the way for machine comprehension and machine translation.

2 METHODOLOGY

We quantify the time structure of an individual word pattern W_i through the statistics of its recurrence times τ_{ii} . We characterize the dynamic impact of a word pattern W_i on another word pattern W_j by the statistics of their hitting times τ_{ij} . In what follows, we will describe the statistical analyses of τ_{ii} and τ_{ij} , on which we build a language-independent Markov model for semantics.

2.1 Recurrence times and topicality

Assuming uniform reading speed,¹ we measure the recurrence times τ_{ii} for a word pattern W_i through n_{ii} samples

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^{1.} On the scale of words (rather than phonemes), this assumption works fine in most languages that are written alphabetically. However, this working hypothesis does not extend to Japanese texts, which interlace Japanese syllabograms (lasting one mora per written unit) with Chinese ideograms (lasting one or more morae per written unit).

 $W_i := \text{happ(ier|ily|iness|y)} \equiv \{\text{happier, happily, happiness, happy}\}, \quad W_j := \text{marr(iage|ied|y)} \equiv \{\text{marriage, married, marry}\}$ $\dots \text{ lorem ipsum happy dolor sit amet, happy, consectetur adipiscing unhappy elit, happiness} \xrightarrow{L_{ii}} \text{ happiness} \xrightarrow{L_{ii}} \text{$

Fig. 1. Counting long-range transitions between word patterns. A transition from W_i to W_j counts towards long-range statistics, if the *underlined* text fragment in between contains no occurrences of W_i , and lasts strictly longer than the longest word in $W_i \cup W_j$. For each long-range transition, the effective fragment length L_{ij} discounts the length of the longest word in $W_i \cup W_j$.

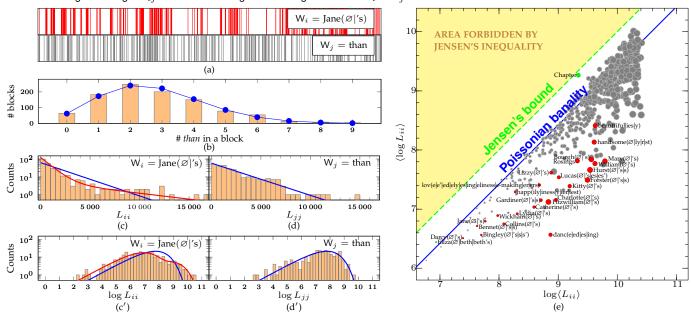


Fig. 2. Statistical analysis of recurrence times and topicality. (a) Barcode representations (adapted from [5, Fig. 2]) for the coverage of $W_i = \text{Jane}(\varnothing|\text{'s})$ (291 occurrences) and $W_j = \text{than}$ (282 occurrences) in the whole text of *Pride and Prejudice*. Horizontal axis scales linearly with respect to the text length measured in the number of constituting letters, spaces and punctuation marks. (b) Counts of the word *than* within a consecutive block of 1217 words (spanning about 1% of the entire text), drawn from 1000 randomly chosen blocks, fitted to a Poisson distribution with mean 2.776 (*blue* curve). (c) Histogram of effective fragment length L_{ii} (see Fig. 1 for its definition) for the topical pattern $W_i = \text{Jane}(\varnothing|\text{'s})$, fitted to an exponential distribution (*blue* line in the semi-log plot) and a weighted mixture of two exponential distributions $c_1k_1e^{-k_1t} + c_2k_2e^{-k_2t}$ (*red* curve, with $c_1 : c_2 \approx 1 : 3$, $k_1 : k_2 \approx 1 : 7$). (d) Histogram of L_{jj} for the function word $W_j = \text{than}$, fitted to an exponential distribution (*blue* line in the semi-log plot). All the parameter estimators in panels b-d are based on maximum likelihood. (c')-(d') Reinterpretations of panels c-d, with logarithmic binning on the horizontal axes, to give fuller coverage of the dynamic ranges for the statistics. (e) Recurrence statistics for word patterns in Jane Austen's *Pride and Prejudice*, where (\cdots) denotes averages over n_{ij} samples of long-range transitions. Data points in *gray*, *green* and *red* have radii $\frac{1}{4\sqrt{n_{ii}}}$. Labels for proper names and some literary motifs are attached next to the corresponding colored dots. Jensen's bound (*green dashed line*) has unit slope and zero intercept. Exponentially distributed recurrence statistics reside on the line of Poissonian banality (*blue line*), with unit slope and negative intercept. *Red* (resp. *green*) dots mark significant downward (resp. upward) departure from the *blue line*.

of the effective fragment lengths L_{ii} (Figs. 1, 2a). Here, while counting as in Fig. 1, we ignore contacts between short-range neighbors, which may involve language-dependent redundancies.²

2.1.1 Recurrence of non-topical patterns

In a memoryless (hence banal) Poisson process (Fig. 2b), recurrence times are exponentially distributed (Fig. 2d,d'). The same is also true for word recurrence in a randomly reshuffled text [5]. If we have n_{ii} independent samples of exponentially distributed random variables L_{ii} , then the statistic $\delta_i := \log \langle L_{ii} \rangle - \langle \log L_{ii} \rangle - \gamma_0 + \frac{1}{2n_{ii}}$ satisfies an inequality

$$|\delta_i| < \frac{2}{\sqrt{n_{ii}}} \sqrt{\frac{\pi^2}{6} - 1 - \frac{1}{2n_{ii}}} \tag{1}$$

2. For example, a German phrase *liebe Studentinnen und Studenten* with short-range recurrence is the gender-inclusive equivalent of the English expression *dear students*. Some Austronesian languages (such as Malay and Hawaiian) use reduplication for plurality or emphasis.

with probability 95% (see Theorem 1 in Appendix A for a two-sigma rule). Here, $\gamma_0 := \lim_{n \to \infty} \left(-\log n + \sum_{m=1}^n \frac{1}{m} \right)$ is the Euler–Mascheroni constant.

As a working definition, we consider a word pattern W_i non-topical if its n_{ii} counts of effective fragment lengths L_{ii} are exponentially distributed $\mathbb{P}(L_{ii} > t) \sim e^{-kt}$, within 95% margins of error [that is, satisfying (1) above].

2.1.2 Recurrence of topical patterns

In contrast, we consider a word pattern W_i topical if its diagonal statistics n_{ii}, L_{ii} constitute significant departure from the Poissonian line $\langle \log L_{ii} \rangle - \log \langle L_{ii} \rangle + \gamma_0 = 0$ (Fig. 2e, blue line), violating the bound in (1).

Notably, most data points for topics (colored dots on Fig. 2e) in Jane Austen's *Pride and Prejudice* mark systematic downward departures from the Poissonian line. This suggests that the topical recurrence times $\tau = L_{ii}$ follow

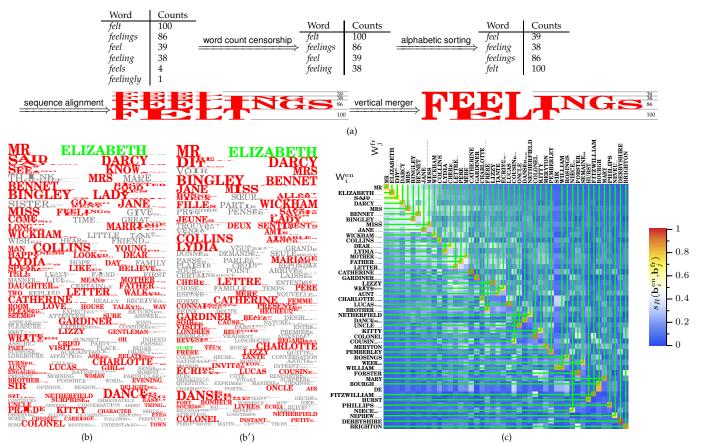


Fig. 3. Automated topic extraction and raw alignment across bilingual corpora. (a) Schematic diagram illustrating our graphical representation of morphologically related words (identified by supervised algorithms in *Supplementary Materials*) in a word pattern. To avoid unprintably small characters, rarely occurring forms (less than 5% of the total sum of all the words ranked above) are ignored in graphical display. To enhance the visibility of word stems, we print shared letters only once, and compress other letters vertically, with heights proportional to their corresponding word counts. (b) Word patterns W_i in Jane Austen's *Pride and Prejudice*, sorted by descending n_{ii} , with font size proportional to the square root of $e^{-\langle \log L_{ii} \rangle}$ (a better indicator of reader's impression than the number of recurrences $n_{ii} \propto e^{-\log(L_{ii})}$). Topical (that is, significantly non-Poissonian) patterns painted in *red* (resp. *green*) reside below (resp. above) the critical line of Poissonian banality (*blue line* in Fig. 2e), where the deviations exceed the error margin prescribed in (1) of §2.1. (b') A similar service on a French version of *Pride and Prejudice* (tr. Valentine Leconte & Charlotte Pressoir). (c) A low-cost and low-yield word translation, based on chapter-wise word counts $b_i^{\rm en}$ and $b_j^{\rm fr}$. Ružička similarities $s_R(b_i^{\rm en}, b_j^{\rm fr})$ between selected topics (sorted by descending $n_{ii} \geq 20$) in English and French versions of *Pride and Prejudice*. Rows and columns with maximal $s_R(b_i^{\rm en}, b_j^{\rm fr})$ less than 0.7 are not shown. Correct matchings are indicated by green cross-hairs.

weighted mixtures of exponential distributions (Fig. 2c,c'):

$$\mathbb{P}(\tau > t) \sim \sum_{m} c_m e^{-k_m t},\tag{2}$$

(where $c_m, k_m > 0$, and $\sum_m c_m = 1$), which impose an inequality constraint on the recurrence time $\tau = L_{ii}$:

$$\langle \log L_{ii} \rangle - \log \langle L_{ii} \rangle + \gamma_0$$

$$= \sum_{m} c_m \log \frac{1}{k_m} - \log \sum_{m} \frac{c_m}{k_m} \le 0.$$
 (3)

2.1.3 Raw alignment of topical patterns

If a word pattern W_i qualifies as a topic by our definition (Fig. 3b,b'), then the signals in its coarse-grained timecourse (say, a vector $\mathbf{b}_i = (b_{i,1}, \dots, b_{i,61})$ representing word counts in each chapter of *Pride and Prejudice*) are not overwhelmed by Poisson noise.

This vectorization scheme, together with the Ružička similarity [6]

$$s_R(\mathbf{b}_i^{\mathrm{A}}, \mathbf{b}_j^{\mathrm{B}}) := \frac{\|\mathbf{b}_i^{\mathrm{A}} \wedge \mathbf{b}_j^{\mathrm{B}}\|_1}{\|\mathbf{b}_i^{\mathrm{A}} \vee \mathbf{b}_j^{\mathrm{B}}\|_1}$$
(4)

between two vectors with non-negative entries, allow us to align some topics found in parallel versions of the same document, in languages A and B (Fig. 3c). Here, in the definition of the Ružička similarity, \land (resp. \lor) denotes componentwise minimum (resp. maximum) of vectors; $\|\mathbf{b}\|_1$ sums over all the components in \mathbf{b} .

2.2 Markov text model

2.2.1 Transition probabilities via pattern analysis

The diagonal statistics n_{ii} , L_{ii} (Fig. 1) have enabled us to extract topics automatically through recurrence time analysis (Figs. 2e and 3b,b'). The off-diagonal statistics n_{ij} , L_{ij} (Fig. 1) will allow us to determine how strongly one word pattern W_i binds to another word pattern W_j , through hitting time analysis. In an empirical Markov matrix $\mathbf{P} = (p_{ij})$, the long-range transition rate p_{ij} is estimated by

$$p_{ij} := \frac{n_{ij}e^{-\langle \log L_{ij} \rangle}}{\sum_{k=1}^{N} n_{ik}e^{-\langle \log L_{ik} \rangle}},$$
(5)

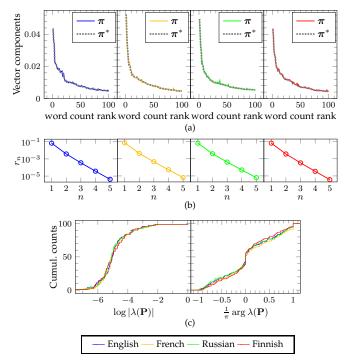


Fig. 4. Quantitative properties of Markov text model. (a) Dominant eigenvector π of a 100×100 Markov matrix \mathbf{P} , computed from one of the four versions of Pride and Prejudice, in comparison with π^* , the list of normalized frequencies for top 100 word patterns. (b) Precipitous decays of $r_n := \frac{1}{2} \sum_{1 \le i,j \le 100} \left| \pi_i p_{ij}^{(n)} - \pi_j p_{ji}^{(n)} \right|$ from the initial value $r_1 \approx 0.07$, for matrix powers $\mathbf{P}^n = (p_{ij}^{(n)})_{1 \le i,j \le 100}$ constructed from four versions of Pride and Prejudice. (In contrast, one has $r_1 \approx 0.33$ for a random 100×100 Markov matrix.) Such quick relaxations support our working hypothesis about detailed balance $\pi_i^* p_{ij}^* = \pi_j^* p_{ji}^*$. (c) Distributions of eigenvalues λ of empirical Markov matrices \mathbf{P} , with nearly language-independent modulus $|\lambda(\mathbf{P})|$ and phase-angle $\arg \lambda(\mathbf{P})$.

where n_{ij} counts the number of long-range transitions from W_i to W_j , and L_{ij} is a statistic that measures the effective fragment lengths of such transitions (Fig. 1).

2.2.2 Equilibrium state and detailed balance

Numerically, we find that our empirical Markov matrix $\mathbf{P}=(p_{ij})$ defined in (5) is a fair approximation to an $ergodic^3$ matrix $\mathbf{P}^*=(p_{ij}^*)$, which in turn, governs the stochastic hoppings between content word patterns during text generation.

Each ergodic Markov matrix $\mathbf{P}^* = (p_{ij}^*)_{1 \leq i,j \leq N}$ possesses a unique equilibrium state $\pi^* = (\pi_i^*)_{1 \leq i \leq N}$. The equilibrium state π^* represents a probability distribution (that is, $\pi_i^* \geq 0$ for $1 \leq i \leq N$ and $\sum_{i=1}^N \pi_i^* = 1$) that satisfies $\pi^*\mathbf{P}^* = \pi^*$ (that is, $\sum_{1 \leq i \leq N} \pi_i^* p_{ij}^* = \pi_j^*$ for $1 \leq j \leq N$). In our numerical experiments, the dominant eigenvector π (satisfying $\pi \mathbf{P} = \pi$) consistently reproduces word frequency statistics that are proportional to the ideal equilibrium state π^* (Fig. 4a).

Furthermore, through numerical experimentation, we find that our empirical Markov matrix $\mathbf{P} \approx \mathbf{P}^*$ approximately honors the detailed balance condition $\pi_i^* p_{ij}^* = \pi_j^* p_{ji}^*$ for $1 \leq i,j \leq N$. The approximation $\pi_i p_{ij}^{(n)} \approx \pi_j p_{ji}^{(n)}$

3. If a Markov chain is ergodic, then there is a strictly positive probability to transition from any Markov state (that is, any individual word pattern in our model) to any other state, after finitely many steps.

becomes closer as we go to higher iterates $\mathbf{P}^n = (p_{ij}^{(n)})$, where n is a small positive integer (Fig. 4b).

On an ergodic Markov chain with detailed balance, one can show that recurrence times are distributed as weighted mixtures of exponential decays (see Theorem 3 in Appendix B.3), thus offering a theoretical explanation for (2).

2.2.3 Spectral invariance under translation

The spectrum $\sigma(\mathbf{P})$ (collection of eigenvalues) is approximately invariant against translations of texts (Fig. 4c), which can be explained by a matrix equation

$$\mathbf{P}_{\mathbf{A}}\mathbf{T}_{\mathbf{A}\to\mathbf{B}} = \mathbf{T}_{\mathbf{A}\to\mathbf{B}}\mathbf{P}_{\mathbf{B}}.\tag{6}$$

Here, both sides of the identity above quantify the transition probabilities from words in language A to words in language B, from the impressions of Alice and Bob, two monolingual readers in a thought experiment. On the left-hand side, Alice first processes the input in her native language A by a Markov matrix \mathbf{P}_{A} , and then translates into language B, using a dictionary matrix $\mathbf{T}_{A \to B}$; on the right-hand side, Bob needs to first translate the input into language B, using the same dictionary $\mathbf{T}_{A \to B}$, before brainstorming in his own native language, using \mathbf{P}_{B} . Putatively, the matrix equation holds because semantic content is shared by native speakers of different languages. In the ideal scenario where translation is lossless (with invertible $\mathbf{T}_{A \to B}$), the Markov matrices \mathbf{P}_{A} and \mathbf{P}_{B} are indeed linked to each other by a similarity transformation that leaves their spectrum intact.

2.3 Localized Markov matrices and semantic cliques

2.3.1 Semantic contexts for recurrent topics

Specializing spectral invariance to individual topical patterns, we will be able to generate semantic fingerprints through a list of topic-specific and language-independent eigenvalues. Here, we will be particularly interested in recurrence eigenvalues of individual topical patterns, which correspond to multiple decay rates in the weighted mixtures of exponential distributions.

Unlike the single exponential decays associated to non-topical recurrence patterns, the multiple exponential decay modes will enable our robot reader to easily discern one topic from another. In general, it is numerically challenging to recover multiple exponential decay modes from a limited amount of recurrence time measurements [7]. However, in text processing, we can circumvent such difficulties by off-diagonal statistics n_{ij} and L_{ij} that provide semantic contexts for individual topical patterns.

To quantitatively define the semantic content of a topical pattern W_i , we specify a local, directed, and weighted graph, corresponding to a localized Markov transition matrix $\mathbf{P}^{[i]}$.

2.3.2 Localized Markov contexts of topical patterns

To localize, we need to remove edges between two vertices W_i and W_j , when the hitting times L_{ij} and L_{ji} are "long enough" relative to what one could naïvely expect from recurrence time statistics n_{ij} , n_{ji} and L_{ii} , L_{jj} . Here, for naïve expectation, we approximate the probability

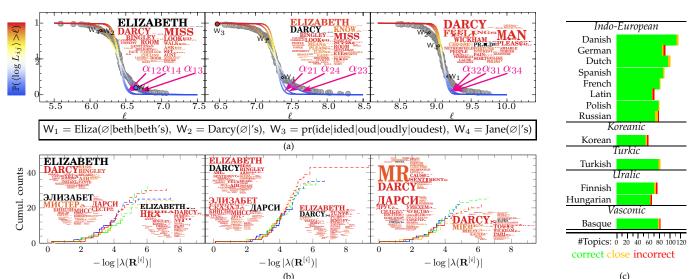


Fig. 5. Semantic cliques and their applications to word translation. (a) Empirical distributions of $\langle \log L_{ij} \rangle$ in *Pride and Prejudice*, as *gray* and *colored* dots with radii $\frac{1}{4\sqrt{n_{ij}}}$, compared to Gaussian model $\alpha_{ij}(\ell)$ (colored curves parametrized by (7) and (8)). The numerical samplings of W $_j$'s exhaust all the textual patterns available in the novel, including topical word patterns, non-topical word patterns and function words. Only those textual patterns with over 40 occurrences are displayed as data points. *Inset* of each frame shows the semantic clique \mathscr{S}_i surrounding topic W $_i$ (painted in black), color-coded by the $\alpha_{ij}(\langle \log L_{ij} \rangle)$ score. The areas of the bounding boxes for individual word patterns are proportional to the components of $\pi^{[i]}$ (the equilibrium state of $\mathbf{P}^{[i]}$). (b) Distributions for the magnitudes of eigenvalues (LISF) in the recurrence matrices $\mathbf{R}^{[i]}$, for three concepts from four versions of *Pride and Prejudice*. The color encoding for languages follows Fig. 4. The largest $\lfloor e^{\eta_i} \rfloor$ magnitudes of eigenvalues are displayed as *solid* lines, while the remaining terms are shown in *dashed* lines. *Inset* of each frame shows the semantic clique \mathscr{S}_i , counterclockwise from top-left, in French, Russian and Finnish. (c) Yields from bipartite matching of LISF (see Fig. 6 for English-French) for topical words between the English original of *Pride and Prejudice* and its translations into 13 languages out of 5 language families.

 $\mathbb{P}(\langle \log L_{ij} \rangle > \ell)$ by a Gaussian model $\alpha_{ij}(\ell)$ (colored curves in Fig. 5a)

$$\mathbb{P}(\langle \log L_{ij} \rangle > \ell) \approx \alpha_{ij}(\ell) := \sqrt{\frac{n_{ij}}{2\pi\beta_i}} \int_{\ell}^{\infty} e^{-\frac{n_{ij}(x-\ell_i)^2}{2\beta_i}} dx,$$
(7)

whose mean and variance are deducible from n_{ij} and L_{ii} (see Theorem 4 in Appendix B.4):

$$\ell_i := \frac{\langle L_{ii} \log L_{ii} \rangle}{\langle L_{ii} \rangle} - 1, \quad \beta_i := \frac{\langle L_{ii} (\ell_i - \log L_{ii})^2 \rangle}{\langle L_{ii} \rangle}. \quad (8)$$

The parameters in the Gaussian model are justified by the relation between hitting and recurrence times [8] on an ergodic Markov chain with detailed balance, and become asymptotically exact if distinct word patterns are statistically independent (such as α_{13} , α_{24} , α_{31} , α_{34} in Fig. 5a). Here, statistical independence justifies additivity of variances, hence the $\sqrt{n_{ij}}$ factor in (7); sums of independent samples of $\log L_{ij}$ become asymptotically Gaussian, thanks to the central limit theorem. Failing that, the actual ranking of $\langle \log L_{ij} \rangle$ may deviate from the Gaussian model prediction in (7), such as the intimately related pairs of words *Elizabeth/Darcy*, *Elizabeth/Jane*, *Darcy/Elizabeth*, *Darcy/pride* and *pride/Darcy*.

2.3.3 Markov criteria for semantic cliques

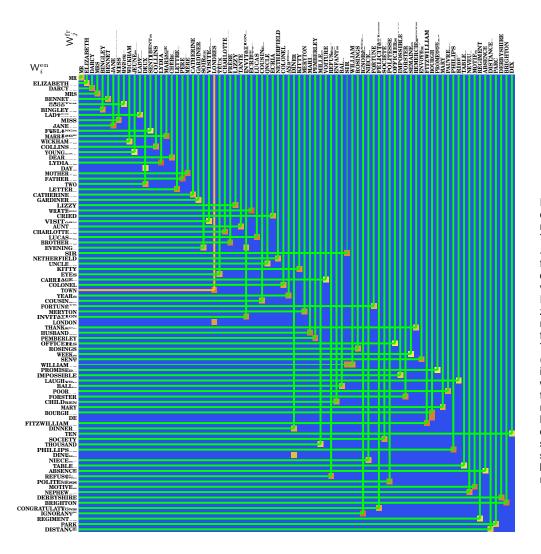
Empirically, we find that higher $\alpha_{ij}(\ell)$ scores point to closer affinities between word patterns (Fig. 5a), attributable to kinship (*Elizabeth*, *Jane*), courtship (*Darcy*, *Elizabeth*), disposition (*Darcy*, *pride*) and so on. Our robot reader automatically detects such affinities, without references other than the novel itself. Therefore, we can use the $\alpha_{ij}(\ell)$ scores as guides to numerical approximations of semantic fields, hereafter referred to as *semantic cliques*.

We invite a topical pattern W_j to the semantic clique \mathscr{S}_i (Figs. 5a and b, insets) surrounding W_i , if $\min\{\alpha_{ij}(\langle \log L_{ij}\rangle), \alpha_{ji}(\langle \log L_{ji}\rangle)\} > \alpha_*$ for a standard Gaussian threshold $\alpha_* := \frac{1}{\sqrt{2\pi}} \int_{-\infty}^1 e^{-x^2/2} \mathrm{d}\,x \approx 0.8413$. This operation emulates the brainstorming procedure of a human reader, who associates one word with another only when they stay much closer than two randomly picked words, according to his/her impression.

Indeed, by numerical brainstorming from W_i , our semantic cliques \mathcal{S}_i (Figs. 5a and b, insets) inform us about their center word W_i , through several types of semantic relations, including, but not limited to

- Synonyms (pride and vanity in English, orgeuil and fierté in French, etc.);
- Temperaments (Elizabeth, a delightful girl, often laughs, corresponding to French verbs sourire and rire);
- Co-references (e.g. Darcy as a personification of pride);
- Causalities (such as *pride* based on *fortune*).

On a local graph with vertices $\mathscr{S}_i = \{W_{i_1} = W_i, W_{i_2}, \ldots, W_{i_{N_i}}\}$, we specify the connectivity of each directed edge by a localized Markov matrix $\mathbf{P}^{[i]} = (p_{jk}^{[i]})_{1 \leq j,k \leq N_i}$. This localized Markov matrix is the row-wise normalization of an $N_i \times N_i$ subblock of \mathbf{P} with the same set of vertices as \mathscr{S}_i . Resetting the entries $p_{1k}^{[i]}$ and $p_{j1}^{[i]}$ as zero, one arrives at the localized recurrence matrix $\mathbf{R}^{[i]}$. We call $\mathbf{R}^{[i]}$ a recurrence matrix, because one can use it to compute the distribution for recurrence times to the Markov state W_i in \mathscr{S}_i . As we will see soon in the applications below, the eigenvalues of $\mathbf{R}^{[i]}$, when properly arranged, become language-independent semantic fingerprints.



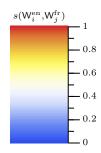


Fig. 6. Automated alignments of vectorized topics via bipartite matching of semantic similarities. The similarities semantic $s(W_i^{en}, W_i^{fr})$ computed are topics for selected (sorted by descending $n_{ii} \geq$ 20) in two versions of Pride and Prejudice. Rows and columns filled with zeros are not shown. Cross-hairs meet at optimal nodes that solve the bipartite matching problem. The thickness of each horizontal vertical) cross-hair inversely proportional to the rowwise (resp. column-wise) ranking of the similarity score for the optimal node. Green (resp. amber) crosshair indicates an exact (resp. a close but non-exact) match. At the same confidence level (0.7) for similarities, this experiment has better recall than Fig. 3c, without much cost of precision.

3 Applications

3.1 Automated word translations from bilingual documents

Experimentally, we resolve the connectivity of an individual pattern W_i through the recurrence spectrum $\sigma(\mathbf{R}^{[i]})$ (Fig. 5b). The dominant eigenvalues of $\mathbf{R}^{[i]}$ are concept-specific while remaining nearly language-independent (a localized version of the invariance in Fig. 4c). Such empirical evidence motivates us to define the language-independent semantic fingerprint (LISF) of a word pattern W_i by a descending list for the magnitudes of eigenvalues

$$\mathbf{v}_i = (|\lambda_1(\mathbf{R}^{[i]})|, |\lambda_2(\mathbf{R}^{[i]})|, \dots), \tag{9}$$

computable from its semantic clique \mathscr{S}_i . We zero-pad this vector from the $(\lfloor e^{\eta_i} \rfloor + 1)$ st component onwards, where η_i is the Kolmogorov–Sinai entropy production rate of the Markov matrix $\mathbf{P}^{[i]}$, measured in nats per word.⁴

4. The entropy production rate $\eta(\mathbf{P}) := -\sum_{i,j} \pi_i p_{ij} \log p_{ij}$ [9, (4.27)] of a Markov matrix \mathbf{P} represents the weighted average (assigning probability mass π_i to the ith Markov state) of Boltzmann's partition entropies $-\sum_j p_{ij} \log p_{ij}$ [10, §8.2]. We have $\eta(\mathbf{P}) \leq \log N$ for an $N \times N$ Markov matrix \mathbf{P} with strictly positive entries [10, Theorem 14.1].

Via bipartite matching (Fig. 6) of word vectors \mathbf{v}_i across languages, our algorithm translates words from parallel texts at very high precision (Fig. 5c), being competitive with state-of-the-art algorithms for bilingual word mapping [11], [12].

Unlike the vector \mathbf{b}_i (Fig. 3c) that captures only chapter-scale features of W_i , the semantic fingerprint \mathbf{v}_i in (9) characterizes the kinetic behavior of W_i on all the long-range time scales.

Given a topical pattern W_i^A in language A, its semantic fingerprint \mathbf{v}_i^A (a descending list of recurrence eigenvalues, as in Fig. 5b) allows us to numerically locate a semantically close pattern in a parallel text written in another language B, in two steps:

(1) Divide the document into K chapters, and define the semantic similarity function as $s(\mathsf{W}_i^{\mathrm{A}}, \mathsf{W}_j^{\mathrm{B}}) := s_R(\mathbf{v}_i^{\mathrm{A}}, \mathbf{v}_j^{\mathrm{B}})$ if

$$s_R(\mathbf{b}_i^{\mathrm{A}}, \mathbf{b}_j^{\mathrm{B}}) \ge \max \left\{ 1 - 0.07\sqrt{K}, 1 - \sqrt{\frac{\|\mathbf{b}_i^{\mathrm{A}} \wedge \mathbf{b}_j^{\mathrm{B}}\|_0}{\|\mathbf{b}_i^{\mathrm{A}} \vee \mathbf{b}_j^{\mathrm{B}}\|_1}} \right\}$$
(10)

(which is a ballpark screening more robust than Fig. 3c, with $\|\mathbf{b}\|_0$ counting the number of non-zero components in b) and $s_R(\mathbf{v}_i^{\mathrm{A}},\mathbf{v}_j^{\mathrm{B}}) \geq 0.7$; $s(\mathsf{W}_i^{\mathrm{A}},\mathsf{W}_j^{\mathrm{B}}) := 0$ otherwise.

WikiQA-Q26: How did Anne Frank die? Reference: "Anne Frank" (Wikipedia)

FRANK
ANNE TYPHUS
SURVIVED CENTRERSTORED
GRAMSTURIFARIL ****INICIALIST
GRAMSTURIFARIL STREED
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(1) Anne Frank and her sister, Margot, were eventually transferred to the Bergen-Belsen concentration camp, where they died of typhus in March 1945.

(2) Annells "Anne" Marie Frank (, ?, ; 12 June 1929early March 1945) is one of the most discussed Jewish victims of the Holocaust.

(3) Otto Frank, the only survivor of the family, retuned to Amsterdam after the war to find that Anne's diary had been saved, and his efforts led to its publication in 1947.

(3) Approximation of the Review Inguilation in consolation 1947, the family went into hiding in the hidden rooms of Anne's father, Ouo Frank's, office building.







Fig. 7. Applications of semantic cliques to question-answering. (a) A construction of semantic clique $\mathcal{Q} \cup \mathcal{Q}'$ (based on $\mathcal{Q} = \{Anne, Frank, die\}$) weighted by the PageRank equilibrium state $\widetilde{\pi}$ and subsequent question-answering. Top 5 candidate answers, with punctuation and spacing as given by WikiQA, are shown with font sizes proportional to the entropy production score in (11). Here, the top-scoring sentence with *highlighted* background is the same as the official answer chosen by the WikiQA team. Like a human reader, our algorithm automatically detects the place "Bergen-Belsen concentration camp", cause "typhus", and year "1945" of Anne Frank's death. (b) Evaluations of our model (LISF and LISF*) on the WikiQA data set, in comparison with established algorithms.

(2) Solve a bipartite matching problem (Fig. 6) that maximizes $\sum_{i,j} s(\mathsf{W}_i^{\mathrm{A}}, \mathsf{W}_j^{\mathrm{B}})$, using the Hungarian Method [13] attributed to Jacobi–Kőnig–Egerváry–Kuhn [14].

3.2 Machine-assisted text comprehension on WikiQA data set

By automatically discovering related words through numerical brainstorming (Figs. 5a and b, insets), our semantic cliques \mathscr{S}_i are useful in text comprehension and question answering. We can expand a set of question words $\mathscr{Q} = \{W_{q_1}, \ldots, W_{q_K}\}$ into $\mathscr{Q} \cup \mathscr{Q}' = \bigcup_{k=1}^K \mathscr{S}_{q_k}$, by bringing together the semantic cliques \mathscr{S}_{q_k} generated from a reference text by each and every question word W_{q_k} .

As before, we construct a localized Markov matrix $\mathbf{P} = (p_{ij})_{1 \leq i,j \leq N}$ on this subset of word patterns $\mathcal{Q} \cup \mathcal{Q}'$. We further use the Brin–Page damping [15] to derive an ergodic Markov matrix $\widetilde{\mathbf{P}} = (\widetilde{p}_{ij})_{1 \leq i,j \leq N}$, where $\widetilde{p}_{ij} = 0.85p_{ij} + \frac{0.15}{N}$.

By analogy to the behavior of internet surfing [15], [16], we model the process of associative reasoning [17] as a navigation through the nodes $\mathcal{Q} \cup \mathcal{Q}'$ according to $\widetilde{\mathbf{P}}$, which quantifies the click-through rate from one idea to another. The PageRank recursion [16] ensures a unique equilibrium state $\widetilde{\pi}$ attached to $\widetilde{\mathbf{P}}$. If our question Q and a candidate answer A contain, respectively, words from W_{Q_1} , ..., $W_{Q_m} \in \mathcal{Q}$ and W_{A_1} , ..., $W_{A_n} \in \mathcal{Q} \cup \mathcal{Q}'$ (counting multiplicities, but excluding function words and patterns with fewer than 3 occurrences in the reference document), then we assign the following entropy production score

$$\mathscr{F}[Q,A] := -\sum_{i=1}^{m} \sum_{j=1}^{n} \widetilde{\pi}_{Q_i} \widetilde{p}_{Q_i A_j} \log \widetilde{p}_{Q_i A_j}$$
 (11)

to this question-answer pair.⁵

A sample work flow is shown in Fig. 7a, to illustrate how our rudimentary question-answering machine handles a query. To answer a question, we use a single Wikipedia page (without infoboxes and other structural data) as the only reference document and training source. Like a typical human reader of Wikipedia, our numerical associative reasoning generates a weighted set of nodes $\mathcal{Q} \cup \mathcal{Q}'$ (presented graphically as a thought bubble in Fig. 7a), without the help of external stimuli or knowledge feed. Here, the relative

5. One may compare the score $\mathscr{F}[Q,A]$ to the Kolmogorov–Sinai entropy production rate [9,(4.27)] $\eta(\mathbf{P})=-\sum_{i=1}^{N}\sum_{j=1}^{N}\pi_{i}p_{ij}\log p_{ij}$ of a Markov matrix $\mathbf{P}=(p_{ij})_{1\leq i,j\leq N}$. The score $\mathscr{F}[Q,A]$ is modeled after Boltzmann's partition entropies, weighted by words in the question, and sifted by topics in the answer. Such a weighting and sifting method is analogous to the definition of scattering cross-sections in particle physics.

weights in the nodes of $\mathcal{Q} \cup \mathcal{Q}'$ are computed from the equilibrium state $\tilde{\pi}$ of $\tilde{\mathbf{P}}$, via the PageRank algorithm.

We then test our semantic model (LISF in Fig. 7b) on all the 1242 questions in the WikiQA data set, each of which is accompanied by at least one correct answer located in a designated Wikipedia page. Our algorithm's performance is roughly on par with LCLR and CNN benchmarks [18], improving upon the baseline by significant margin. This is perhaps remarkable, considering the relatively scant data at our disposal. Unlike the LCLR approach, our numerical discovery of synonyms does not draw on the WordNet database [19] or pre-existent corpora of question-answer pairs. Unlike the CNN method, we do not need pre-trained word2vec embeddings [20] as semantic input.

Moreover, our algorithm (LISF* in Fig. 7b) performs slightly better on a subset of 990 questions that do not require quantitative cues (*How big? How long? How many? How old? What became of? What happened to? What year?* and so on). This indicates that, with a Markov chain description of two-body interactions between topics, our structural model fits associative reasoning better than rule-based reasoning [17], while imitating human behavior in the presence of limited data. To enhance the reasoning capabilities of our algorithm, it is perhaps appropriate to apply a Markov random field [21, §4.1.3] to graphs of word patterns, to capture many-body interactions among different topics.

4 Conclusion

In our current work, we define semantics through algebraic invariants that are concept-specific and language-independent. To construct such invariants, we develop a stochastic model that assigns a semantic fingerprint (list of recurrence eigenvalues) to each concept via its long-range contexts. Consistently using a single Markov framework, we are able to extract topics (Figs. 2e, 3b,b'), translate topics (Figs. 3c, 4c, 5b,c, 6) and understand topics (Figs. 5a,b, 7a,b), through statistical mining of short and medium-length texts. In view of these three successful applications, we are probably close to a complete set of semantic invariants, after demystifying the long-range behavior of human languages.

Notably, our algorithms apply to documents of moderate lengths, similar to the experience of human readers. This contrasts with data-hungry algorithms in machine learning [18], [22], which utilize high-dimensional numerical representations of words and phrases [11], [12], [20], [23] from large corpora. Our semantic mechanism exhibits universality on long-range linguistic scales. This adds to our quantitative understanding of diversity on shorter-range linguistic scales, such as phonology [24], [25], [26], morphology [27], [28], [29], [30] and syntax [3], [30], [31], [32], [33].

Thanks to the independence between semantics and syntax [3], our current model conveniently ignores the non-Markovian syntactic structures which are essential to fluent speech. In the near future, we hope to extend our framework further, to incorporate both Markovian and non-Markovian features across different ranges. *The Mathematical Principles of Natural Languages*, as we envision, must and will combine the statistical analysis of a Markov model with linguistic properties on shorter time scales that convey morphological [27], [28], [29], [30] and syntactical [3], [30], [31], [32], [33] information.

APPENDIX A POISSONIAN BANALITY AND NON-TOPICALITY

Here, we first present a proof of our statistical criterion for Poissonian banality (which we identify with non-topicality of word patterns), as stated in (1).

Theorem 1 (Sums and products of exponentially distributed random variables). Let X_1, \ldots, X_N be independent random variables, each obeying an exponential distribution with mean 1. The probability distribution of

$$Y_N := \log\left(\frac{1}{N}\sum_{i=1}^{N}X_i\right) - \frac{1}{N}\sum_{i=1}^{N}\log X_i$$
 (12)

is asymptotic to a Gaussian law

$$\mathbb{P}\left(\frac{Y_N - (\gamma_0 - \frac{1}{2N})}{\frac{1}{\sqrt{N}}\sqrt{\frac{\pi^2}{6} - 1 - \frac{1}{2N}}} < a\right) \sim \int_{-\infty}^a e^{-x^2/2} \frac{\mathrm{d}\,x}{\sqrt{2\pi}} \tag{13}$$

Proof: By definition, we have a moment-generating function

$$\mathbb{E}e^{-tY_{N}} = \int_{(0,+\infty)^{N}} \frac{\prod_{i=1}^{N} x_{i}^{t/N}}{\left(\frac{1}{N} \sum_{i=1}^{N} x_{i}\right)^{t}} e^{-\sum_{i=1}^{N} x_{i}} \, \mathrm{d} \, x_{1} \cdots \, \mathrm{d} \, x_{N}$$

$$= 2^{N} \int_{(0,+\infty)^{N}} \frac{\prod_{i=1}^{N} \xi_{i}^{2t/N+1}}{\left(\frac{1}{N} \sum_{i=1}^{N} \xi_{i}^{2}\right)^{t}} e^{-\sum_{i=1}^{N} \xi_{i}^{2}} \, \mathrm{d} \, \xi_{1} \cdots \, \mathrm{d} \, \xi_{N}. \tag{14}$$

To evaluate the last multiple integral, we use the spherical coordinates $\xi_1 = r\cos\theta_1$, $\xi_2 = r\sin\theta_1\cos\theta_2$, $\xi_3 = r\sin\theta_1\sin\theta_2\cos\theta_3$, ..., $\xi_N = r\prod_{j=1}^{N-1}\sin\theta_j$ (where r>0, and $0<\theta_j<\pi/2$ for all $j\in\{1,\ldots,N-1\}$), with volume element

$$d\xi_1 \cdots d\xi_N = r^{N-1} dr \prod_{j=1}^{N-1} \sin^{N-1-j} \theta_j d\theta_j.$$
 (15)

The result reads

$$\mathbb{E}e^{-tY_N} = N^t \Gamma(N) \prod_{j=1}^{N-1} \frac{\Gamma\left(\frac{N+t}{N}\right) \Gamma\left(\frac{(N-j)(N+t)}{N}\right)}{\Gamma\left(\frac{(N-j+1)(N+t)}{N}\right)}$$
$$= \frac{N^t \Gamma(N)}{\Gamma(N+t)} \left[\Gamma\left(1 + \frac{t}{N}\right)\right]^N, \tag{16}$$

where $\Gamma(s) := \int_0^\infty x^{s-1} e^{-x} dx$ is Euler's gamma function.

Consequently, the proof of (13) builds on a cumulant expansion of (16), that is, development of $\log \mathbb{E} e^{-tY_N}$ up to $O(t^2)$ terms.

If we have N samples of recurrence times from a Poisson process, then the statistic Y_N satisfies the inequality in (1) with probability $\int_{-2}^2 e^{-x^2/2} \frac{\mathrm{d}\,x}{\sqrt{2\pi}} \approx 0.95$, in view of the theorem above.

APPENDIX B

RECURRENCE TIMES ON AN ERGODIC MARKOV CHAIN WITH DETAILED BALANCE

B.1 Background in probability theory

Based on numerical evidence (Fig. 4), we postulate that at the *discourse level* (the longest time scale in Friederici's [4] neurobiological hierarchy), the production of natural language texts can be caricatured by the stochastic transitions on a stationary and ergodic Markov chain $\mathcal{M}=(\mathcal{S},\mathbf{P}).^6$ Here, the state space $\mathcal{S}=\{W_1,\ldots,W_N\}$ runs over finitely many word patterns occurring in the text, which in turn is representable as a discrete-time stochastic process $(X(0),X(1),\ldots,X(n),\ldots)$; the transition matrix $\mathbf{P}=(p_{ij})$ describes the conditional hopping probability on the web of word patterns: $p_{ij}=\mathbb{P}(X(n+1)=W_j|X(n)=W_i)$, for $n\in\mathbb{Z}_{\geq 0}$. Depending on context, we also model a document by a localized Markov chain $\mathcal{M}'=(\mathcal{S}',\mathbf{P}')$, where certain word patterns W_i are removed from the state space \mathcal{S} to form a proper subset $\mathcal{S}'\subsetneq\mathcal{S}$.

In a more formal setting, our notation $\mathscr{M} = (\mathscr{S}, \mathbf{P})$ for the Markov chain should be expanded into $\mathscr{M} = (\Omega, \mathscr{F}, \{\mathbb{P}^{\mathsf{W}} : \mathsf{W} \in \mathscr{S}\}, \{X(t) : t \in \mathbb{Z}_{\geq 0}\}, \{\widehat{\theta}_n : n \in \mathbb{Z}_{\geq 0}\})$, whose components are explained below.

- The sample space Ω consists of all stochastic trajectories $\omega = (X(0), X(1), \ldots, X(t), X(t+1), \ldots) = (X(t))_{t \in \mathbb{Z}_{\geq 0}}$ on the state space \mathscr{S} , namely, all possible texts that can be analyzed by our particular model.
- Each member in the field of events $\mathscr{F}=2^{\Omega}$ (the totality of all subsets in the sample space Ω) is a set containing zero or more stochastic processes that can be regarded as caricatures of text productions.
- The family of probability measures $\{\mathbb{P}^{\mathbb{W}}: \mathbb{W} \in \mathscr{S}\}$ are related to the transition matrix $\mathbf{P} = (p_{ij})_{1 \leq i,j \leq N}$ by the identity $p_{ij}^{(t)} = \mathbb{P}^{\mathbb{W}_i}(X(t) = \mathbb{W}_j) = \mathbb{P}(X(t) = \mathbb{W}_j | X(0) = \mathbb{W}_i), \forall \mathbb{W}_i, \mathbf{W}_j \in \mathscr{S}, \forall t \in \mathbb{Z}_{>0}.$
- The shift operator $\widehat{\theta}_n: \Omega \longrightarrow \Omega$ acts on an arbitrary trajectory $(X(0),X(1),\ldots,X(t),X(t+1),\ldots)=\omega\in\Omega$ in the following manner: $\widehat{\theta}_n\circ\omega=(X(n),X(n+1),\ldots,X(t+n),X(t+n+1),\ldots), \forall t,n\in\mathbb{Z}_{\geq 0}.$

For discussions of the stopping times (a class of random variables on Markov chains) as well as the Markov property, it is convenient to further introduce the notation \mathscr{F}_n for $n \in \mathbb{Z}_{\geq 0}$. Here, \mathscr{F}_n is the smallest σ -algebra containing all the events in the form of $(X(0) = W_{i_0}, X(1) = W_{i_1}, \ldots, X(n) = W_{i_n})$. The family of σ -algebras $\{\mathscr{F}_n : \mathcal{F}_n : \mathcal{F}$

6. To reduce notational burden, we will not use superscripted asterisks to mark Markov matrices, beyond this point.

 $n \in \mathbb{Z}_{\geq 0}$ forms a filtration: $\mathscr{F}_n \subseteq \mathscr{F}_m$ if $n \leq m$. For any $n \in \mathbb{Z}_{\geq 0}$, $B \in \mathscr{F}$, $W \in \mathscr{S}$, we have the following relation concerning conditional expectations:

$$\mathbb{E}^{\mathsf{W}}(\mathbf{1}_{B} \circ \widehat{\theta}_{n} | \mathscr{F}_{n}) = \mathbb{E}^{X(n)}(\mathbf{1}_{B}) := \mathbb{P}^{X(n)}(B), \tag{17}$$

(for every time-step $n \in \mathbb{Z}_{\geq 0}$ and Markov state $W \in \mathscr{S}$) which is merely a reformulation of the Markov condition $\mathbb{P}^W(X(n+1) = W_{i_{n+1}}|X(n) = W_{i_n}, X(n-1) = W_{i_{n-1}}, \ldots, X(0) = W_{i_0}) = \mathbb{P}^{W_{i_n}}(X(1) = W_{i_{n+1}}) = \mathbb{P}^W(X(n+1) = W_{i_{n+1}}|X(n) = W_{i_n})$. In other words, for any $A \in \mathscr{F}_n$, $B \in \mathscr{F}$, $W \in \mathscr{S}$, we have the following statement of the *Markov property*:

$$\mathbb{E}^{\mathsf{W}}(\mathbf{1}_{B} \circ \widehat{\theta}_{n}; A) = \mathbb{E}^{\mathsf{W}}(\mathbb{E}^{X(n)}(\mathbf{1}_{B}); A). \tag{17'}$$

In both (17) and (17'), one can replace the indicator function $\mathbf{1}_B$ for event $B \in \mathscr{F}$ by any random variable with finite expectation.

B.2 Hitting time and return time on a Markov chain

We need to first precisely define the probability distributions for the hitting time and return time (the latter also known as "recurrence time") on our Markov chain $\mathcal{M} = (\mathcal{S}, \mathbf{P})$.

For each state $W_i \in \mathscr{S}$ and trajectory $\omega = (X(t))_{t \in \mathbb{Z}_{\geq 0}} \in \Omega$, we define

$$\tau_i(\omega) := \inf\{n \in \mathbb{Z}_{>0} : X(n) = \mathsf{W}_i\},\tag{18}$$

then $\tau_i:\Omega\longrightarrow\mathbb{Z}_{>0}\cup\{+\infty\}$ is a stopping time. Suppose that our pattern of interest corresponds to a subset of states $\mathscr{W}\subseteq\mathscr{S}$ on the web of words. We define another stopping time $\tau_{\mathscr{W}}:\Omega\longrightarrow\mathbb{Z}_{>0}\cup\{+\infty\}$ as

$$\tau_{\mathscr{W}}(\omega) := \inf\{n \in \mathbb{Z}_{>0} : X(n) \in \mathscr{W}\}
= \inf\{\tau_i(\omega) : \mathsf{W}_i \in \mathscr{W}\}.$$
(19)

Clearly, $\tau_{\mathscr{W}}(\omega)$ is equal to the first time when a forward stepwise search lands on the set of interest \mathscr{W} . Recalling the invariant measure $\pi=(\pi_1,\ldots,\pi_N)$, we define the cumulative distribution function (CDF) for the hitting time to the set of patterns \mathscr{W} as

$$H_{\mathscr{W}}(t) := \sum_{\mathsf{W}_i \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_i}(\tau_{\mathscr{W}} < t) \pi_i, \quad t \in \mathbb{Z}_{>0}.$$
 (20)

Similarly, the CDF for the return time to the set of patterns \mathcal{W} is defined as

$$R_{\mathscr{W}}(t) := \sum_{\mathsf{W}_i \in \mathscr{W}} \mathbb{P}^{\mathsf{W}_i}(\tau_{\mathscr{W}} < t) \frac{\pi_i}{\sum_{\mathsf{W}_j \in \mathscr{W}} \pi_j}, \quad t \in \mathbb{Z}_{>0.} \quad (21)$$

In the next theorem, we present an identity that connects hitting and return times of Markov states, which in turn, is a discrete analog of the Haydn–Lacroix–Vaient relation for continuous-time ergodic dynamical systems [8].

Theorem 2 (Relation between hitting time and return time distributions). For a stationary and ergodic Markov chain $\mathcal{M} = (\mathcal{S}, \mathbf{P})$, and a subset of states $\mathcal{W} \subseteq \mathcal{S}$, we have the following identity regarding the probability distribution of hitting and return times to \mathcal{W} :

$$H_{\mathcal{W}}(1) = R_{\mathcal{W}}(1) = 0; \ \frac{H_{\mathcal{W}}(t+1)}{\sum_{W_j \in \mathcal{W}} \pi_j} = \sum_{n=1}^t [1 - R_{\mathcal{W}}(n)]$$

for all $t \in \mathbb{Z}_{>0}$.

Proof: Clearly, our task is equivalent to the verification of the following formula:

$$\sum_{\mathsf{W}_i \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_i} (\tau_{\mathscr{W}} \le t) \pi_i = \sum_{n=1}^t \sum_{\mathsf{W}_j \in \mathscr{W}} [1 - \mathbb{P}^{\mathsf{W}_j} (\tau_{\mathscr{W}} < t)] \pi_j$$
(22')

for all $t \in \mathbb{Z}_{>0}$.

For t = 1, the left-hand side of (22') can be computed as follows:

$$\sum_{\mathsf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_{i}}(\tau_{\mathscr{W}} = 1)\pi_{i} = \sum_{\mathsf{W}_{i} \in \mathscr{S}} \sum_{\mathsf{W}_{j} \in \mathscr{W}} \mathbb{P}^{\mathsf{W}_{i}}(X(1) = \mathsf{W}_{j})\pi_{i}$$

$$= \sum_{\mathsf{W}_{i} \in \mathscr{S}} \sum_{\mathsf{W}_{j} \in \mathscr{W}} \pi_{i}p_{ij} = \sum_{\mathsf{W}_{j} \in \mathscr{W}} \pi_{j}.$$
(23)

This is equal to the right-hand side of (22'), because $\mathbb{P}^{\mathsf{W}_j}(\tau_{\mathscr{W}}<1)=0.$

For $t \in \mathbb{Z}_{>0}$, we can compute

$$\sum_{\mathbf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathbf{W}_{i}} (\tau_{\mathscr{W}} \leq t + 1) \pi_{i}$$

$$= \sum_{\mathbf{W}_{i} \in \mathscr{S}} \mathbb{E}^{\mathbf{W}_{i}} ((\mathbf{1}_{(\tau_{\mathscr{W}} \leq t)} \circ \widehat{\theta}_{1}) (\mathbf{1}_{(X(1) \notin \mathscr{W})}) + \mathbf{1}_{(X(1) \in \mathscr{W})}) \pi_{i}$$

$$= \sum_{\mathbf{W}_{i} \in \mathscr{S}} [\mathbb{E}^{\mathbf{W}_{i}} (\mathbb{E}^{X(1)} (\mathbf{1}_{(\tau_{\mathscr{W}} \leq t)}; X(1) \notin \mathscr{W}))$$

$$+ \mathbb{E}^{\mathbf{W}_{i}} (\mathbf{1}_{(X(1) \in \mathscr{W})}) \pi_{i}$$
(24)

from the Markov property (17'). Here, we have $\sum_{W_i \in \mathscr{S}} \mathbb{E}^{W_i} (\mathbf{1}_{(X(1) \in \mathscr{W})}) \pi_i = \sum_{W_i \in \mathscr{S}} \mathbb{P}^{W_i} (\tau_{\mathscr{W}} = 1) \pi_i = \sum_{W_j \in \mathscr{W}} \pi_j$ according to (23), and

$$\sum_{\mathbf{W}_{i} \in \mathscr{S}} \mathbb{E}^{\mathbf{W}_{i}} (\mathbb{E}^{X(1)} (\mathbf{1}_{(\tau_{\mathscr{W}} \leq t)}; X(1) \notin \mathscr{W})) \pi_{i}$$

$$= \sum_{\mathbf{W}_{i} \in \mathscr{S}} \sum_{\mathbf{W}_{k} \in \mathscr{S} \setminus \mathscr{W}} \mathbb{P}^{\mathbf{W}_{i}} (X(1) = \mathbf{W}_{k}) \mathbb{E}^{\mathbf{W}_{k}} (\mathbf{1}_{(\tau_{\mathscr{W}} \leq t)}) \pi_{i}$$

$$= \sum_{\mathbf{W}_{k} \in \mathscr{S} \setminus \mathscr{W}} \mathbb{E}^{\mathbf{W}_{k}} (\mathbf{1}_{(\tau_{\mathscr{W}} \leq t)}) \pi_{k}$$

$$= \sum_{\mathbf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathbf{W}_{i}} (\tau_{\mathscr{W}} \leq t) \pi_{i} - \sum_{\mathbf{W}_{i} \in \mathscr{W}} \mathbb{P}^{\mathbf{W}_{j}} (\tau_{\mathscr{W}} \leq t) \pi_{j}. \tag{25}$$

Therefore, the recursion

$$\sum_{\mathsf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_{i}} (\tau_{\mathscr{W}} \leq t + 1) \pi_{i}$$

$$= \sum_{\mathsf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_{i}} (\tau_{\mathscr{W}} \leq t) \pi_{i} + \sum_{\mathsf{W}_{i} \in \mathscr{W}} [1 - \mathbb{P}^{\mathsf{W}_{j}} (\tau_{\mathscr{W}} \leq t)] \pi_{j} \quad (26)$$

for all $t \in \mathbb{Z}_{>0}$ allows us to build (22') inductively on the t = 1 case (23).

B.3 Positive definite return time distributions on a Markov chain with detailed balance

Thanks to Theorem 2, our analysis of a return time distribution can be built on the analysis of the corresponding hitting time, the latter of which is usually easier to compute (in theoretical analysis).

For a stationary and ergodic Markov chain $(\mathscr{S}, \mathbf{P} = (p_{ij})_{1 \leq i,j \leq N})$, and a pattern of interest $\mathscr{W} \subsetneq \mathscr{S}$, one can

use the Markov property to show that the hitting time distribution satisfies

$$\sum_{\mathsf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_{i}}(\tau_{\mathscr{W}} > 1)\pi_{i} = \sum_{\mathsf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_{i}}(X(1) \notin \mathscr{W})\pi_{i}$$

$$= \sum_{\mathsf{W}_{i} \in \mathscr{S}} \sum_{\mathsf{W}_{m} \in \mathscr{S} \setminus \mathscr{W}} \pi_{i}p_{im} = \sum_{\mathsf{W}_{m} \in \mathscr{S} \setminus \mathscr{W}} \pi_{m}$$
(27)

and

$$\sum_{\mathbf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathbf{W}_{i}} (\tau_{\mathscr{W}} > t) \pi_{i}$$

$$= \sum_{\mathbf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathbf{W}_{i}} (X(n) \notin \mathscr{W}, \forall n \in \mathbb{Z} \cap [1, t]) \pi_{i}$$

$$= \sum_{\mathbf{W}_{i} \in \mathscr{S}} \sum_{\mathbf{W}_{m_{n}} \in \mathscr{S} \setminus \mathscr{W}, n \in \mathbb{Z} \cap [1, t]} \pi_{i} p_{im_{1}} \prod_{n \in \mathbb{Z} \cap [1, t-1]} p_{m_{n}m_{n+1}}$$

$$= \sum_{\mathbf{W}_{m_{n}} \in \mathscr{S} \setminus \mathscr{W}, n \in \mathbb{Z} \cap [1, t]} \pi_{m_{1}} \prod_{n \in \mathbb{Z} \cap [1, t-1]} p_{m_{n}m_{n+1}}, \qquad (28)$$

for all $t \in \mathbb{Z}_{>1}$.

Consider the abelian semigroup $\Sigma = \mathbb{Z}_{\geq 0}$. A semicharacter [34, p. 92, Definition 2.1] $\rho: \mathbb{Z}_{\geq 0} \longrightarrow \mathbb{C}$ must assume the following form: $\rho(0) = 1$; $\rho(s) = [\rho(1)]^s$ for $s \in \mathbb{Z}_{>0}$. Since the spectral radius of our recurrence matrix is strictly less than 1, we will only concern ourselves with $\rho_0(s) := \mathbf{1}_{\{0\}}(s)$ and $\rho_\lambda(s) := \lambda^s, s \in \mathbb{Z}_{>0}$ for $\lambda \in (-1,0) \cup (0,1)$.

According to the harmonic analysis on semigroups, a bounded function $f:\mathbb{Z}_{\geq 0}\longrightarrow \mathbb{C}$ can be represented as a weighted mixture of semicharacters $f(s)=\int_{-1<\lambda<1}\rho_{\lambda}(s)\,\mathrm{d}\,\mu(\lambda)$ if and only if it is positive definite [34, p. 93, Theorem 2.5]: the inequality

$$\sum_{r=1}^{m} \sum_{s=1}^{m} c_r \overline{c_s} f(t_r + t_s) \ge 0 \tag{29}$$

holds for arbitrary positive integers $m \in \mathbb{Z}_{>0}$, and arbitrarily chosen m-dimensional "vectors" $(t_1, \ldots, t_m) \in \Sigma^m$, $(c_1, \ldots, c_m) \in \mathbb{C}^m$.

In the following theorem, we check the positive definiteness criterion (29) for the hitting time distribution $1-H_{\mathscr{W}}(s+2), s\in\mathbb{Z}_{\geq 0}$ on a Markov chain satisfying the detailed balance condition. This would imply that the return time distribution $1-R_{\mathscr{W}}(s+2), s\in\mathbb{Z}_{\geq 0}$ is also a weighted mixture of semicharacters $\rho_{\lambda}(s)$ for $-1<\lambda<1$, according to the finite difference relation (22) in Theorem 2.

Theorem 3 (Positive definite return time distributions). Suppose that a non-void pattern $\mathscr{W} \subsetneq \mathscr{S}$ on a stationary and ergodic Markov chain $\mathscr{M} = (\Omega, \mathscr{F}, \{\mathbb{P}^{\mathsf{W}} : \mathsf{W} \in \mathscr{S}\}, \{X(t) : t \in \mathbb{Z}_{\geq 0}\}, \{\widehat{\theta}_n : n \in \mathbb{Z}_{\geq 0}\})$ satisfies $\pi_i p_{ij} = \pi_j p_{ji}$ for all $\mathsf{W}_i, \mathsf{W}_j \in \mathscr{S} \setminus \mathscr{W}$. Then, the cumulative distribution for the return time to \mathscr{W} has the following integral representation

$$\frac{1 - R_{\mathscr{W}}(s+2)}{1 - R_{\mathscr{W}}(2)} = \frac{\sum_{\mathsf{W}_i \in \mathscr{W}} \mathbb{P}^{\mathsf{W}_i} (\tau_{\mathscr{W}} > s+1) \pi_i}{\sum_{\mathsf{W}_i \in \mathscr{W}} \mathbb{P}^{\mathsf{W}_i} (\tau_{\mathscr{W}} > 1) \pi_i}$$

$$= \int \rho_{\lambda}(s) \, d \mathscr{P}_{\mathscr{W}}(\lambda), \quad s \in \mathbb{Z}_{\geq 0}, \quad (30)$$

for a certain probability measure $\mathscr{P}_{\mathscr{W}}(\lambda)$ supported on $\lambda \in (-1,1).$

Proof: By (28), we have

$$\sum_{\mathbf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathbf{W}_{i}} (\tau_{\mathscr{W}} > r + s + 1) \pi_{i}$$

$$= \sum_{\mathbf{W}_{m_{n}} \in \mathscr{S} \setminus \mathscr{W}, n \in \mathbb{Z} \cap [1, r + s + 1]} \pi_{m_{1}} \prod_{n \in \mathbb{Z} \cap [1, r + s]} p_{m_{n} m_{n+1}}$$

$$= \sum_{\mathbf{W}_{m_{n}} \in \mathscr{S} \setminus \mathscr{W}, n \in \mathbb{Z} \cap [1, r + s + 1]} \pi_{m_{1}} \times \prod_{n \in \mathbb{Z} \cap [1, r]} p_{m_{n} m_{n+1}} \prod_{n \in \mathbb{Z} \cap [r + 1, r + s]} p_{m_{n} m_{n+1}}, \quad (31)$$

for $r,s \in \mathbb{Z}_{>0}$. We then apply the identity $\pi_i p_{ij} = \pi_j p_{ji}, \forall \mathsf{W}_i, \mathsf{W}_j \in \mathscr{S} \setminus \mathscr{W}$ to the product over $n \in \mathbb{Z} \cap [1,r]$, which brings us

$$\sum_{\mathsf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_{i}} (\tau_{\mathscr{W}} > r + s + 1) \pi_{i}$$

$$= \sum_{\mathsf{W}_{m_{n}} \in \mathscr{S} \smallsetminus \mathscr{W}, n \in \mathbb{Z} \cap [1, r + s + 1]} \pi_{m_{1}} \times \prod_{n \in \mathbb{Z} \cap [1, r]} \frac{\pi_{m_{n+1}}}{\pi_{m_{n}}} p_{m_{n+1} m_{n}} \prod_{n \in \mathbb{Z} \cap [r+1, r + s]} p_{m_{n} m_{n+1}}$$

$$= \sum_{\mathsf{W}_{m_{r+1}} \in \mathscr{S} \smallsetminus \mathscr{W}} \mathbb{P}^{\mathsf{W}_{m_{r+1}}} (\tau_{\mathscr{W}} > r) \mathbb{P}^{\mathsf{W}_{m_{r+1}}} (\tau_{\mathscr{W}} > s) \pi_{m_{r+1}}$$
(32)

for $r, s \in \mathbb{Z}_{>0}$. In other words, we have verified

$$\sum_{\mathsf{W}_{i} \in \mathscr{S}} \mathbb{P}^{\mathsf{W}_{i}} (\tau_{\mathscr{W}} > r + s + 1) \pi_{i}$$

$$= \sum_{\mathsf{W}_{i} \in \mathscr{S} \setminus \mathscr{W}} \mathbb{E}^{\mathsf{W}_{i}} (\mathbf{1}_{(\tau_{\mathscr{W}} > r)}) \mathbb{E}^{\mathsf{W}_{i}} (\mathbf{1}_{(\tau_{\mathscr{W}} > s)}) \pi_{i}$$
(33)

for $r, s \in \mathbb{Z}_{>0}$. Using the Markov property, one can readily generalize the identity above to $r, s \in \mathbb{Z}_{\geq 0}$. Thus, we have

$$\sum_{r=1}^{m} \sum_{s=1}^{m} \sum_{\mathsf{W}_{i} \in \mathscr{S}} c_{r} \overline{c_{s}} \mathbb{P}^{\mathsf{W}_{i}} (\tau_{\mathscr{W}} > t_{r} + t_{s} + 1) \pi_{i}$$

$$= \sum_{\mathsf{W}_{i} \in \mathscr{S} \setminus \mathscr{W}} \left| \mathbb{E}^{\mathsf{W}_{i}} \left(\sum_{r=1}^{m} c_{r} \mathbf{1}_{(\tau_{\mathscr{W}} > t_{r})} \right) \right|^{2} \pi_{i} \geq 0 \qquad (34)$$

for $t_r, t_s \in \mathbb{Z}_{\geq 0}$. By the positive definiteness criterion in (29), we see that $1 - H_{\mathscr{W}}(s+2), s \in \mathbb{Z}_{\geq 0}$ is a weighted mixture of semicharacters. In view of the recursion identity relating hitting and return time distributions (26), we have confirmed the statement in (30).

With bin sizes far larger than 2 time units on the Markov chain, our L_{ii} is nearly a continuous random variable and all the period-2 oscillatory decays $\rho_{\lambda}(s)$ for $-1 < \lambda < 0$ behave just like noise terms in the histogram. Then, Theorem 3 tells us that the probability distribution for L_{ii} can be approximated by a weighted mixture⁷ of exponential decays $e^{-kL_{ii}}$, in the continuum limit.

7. We note that there exist exceptions to stationarity and ergodicity in realistic documents. A novel may involve the birth and/or death of leading/supporting characters. An academic treatise may place particular emphasis on certain concepts in specific chapters. In such scenarios, the overall recurrence kinetics may still follow the generic law given in (2), as a consequence of both detailed balance (in locally applied Markov models) and a heterogeneous mixture of several stationary and ergodic Markov models that patch together as a whole.

B.4 A statistical criterion for numerical independence between different word patterns

If we define \widetilde{L}_{ij} as the effective length of a text fragment free from word pattern W_j that is simultaneously flanked by W_i to the left and by W_j to the right, then $\sum_{W_i \in \mathscr{S}} \pi_i \mathbb{P}(\widetilde{L}_{ij} > t) \sim \sum_{W_i \in \mathscr{S}} \pi_i \mathbb{P}^{W_i}(\tau_j > t)$ represents the hitting time distribution to the Markov state W_j . According to Theorem 2, the hitting time distribution can be obtained by integrating the return time distribution $\mathbb{P}(\widetilde{L}_{jj} > t) \sim \mathbb{P}^{W_j}(\tau_j > t) \sim \sum_n c_n e^{-k_n t}$. Here, the positive weights $c_n > 0$ are normalized to one: $\sum_n c_n = 1$. In other words, for fixed W_j , we can predict some statistical properties of \widetilde{L}_{ij} by analyzing the recurrence statistic \widetilde{L}_{jj} , as described in the theorem below.

Theorem 4 (Mean and variance for the logarithm of hitting time). Using the continuous time approximation, we have the following relations for a fixed W_j :

$$\mathbb{E}\log\widetilde{L}_{ij} = \frac{\langle \widetilde{L}_{jj}\log\widetilde{L}_{jj}\rangle}{\langle \widetilde{L}_{jj}\rangle} - 1, \qquad (35)$$

$$\mathbb{E}(\log \widetilde{L}_{ij} - \mathbb{E}\log \widetilde{L}_{ij})^2 = \frac{\langle \widetilde{L}_{jj}(\ell_* - \log \widetilde{L}_{jj})^2 \rangle}{\langle \widetilde{L}_{ij} \rangle}, \quad (36)$$

where ℓ_* equals the right-hand side of (35), and the expectation \mathbb{E} denotes average over all the states W_i , weighted by π_i .

Proof: In the continuous time approximation, we can assume that the probability density function for \widetilde{L}_{jj} is $\sum_n c_n k_n e^{-k_n t}$, (where $c_n, k_n > 0$) so that the probability density function for \widetilde{L}_{ij} (weighted average over all the states W_i) is

$$\frac{\sum_{n} c_n e^{-k_n t}}{\sum_{n} c_n / k_n} = \frac{\sum_{n} c_n e^{-k_n t}}{\langle \widetilde{L}_{jj} \rangle}.$$
 (37)

In view of (37), we can compute the left-hand side of (35) as

$$\int_0^\infty \sum_n c_n e^{-k_n t} \log t \frac{\mathrm{d} t}{\langle \widetilde{L}_{jj} \rangle} = -\sum_n \frac{c_n (\gamma_0 + \log k_n)}{k_n \langle \widetilde{L}_{jj} \rangle}.$$
(38)

Meanwhile, we may evaluate the right-hand side of (35) as follows:

$$\int_{0}^{\infty} \sum_{n} c_{n} k_{n} e^{-k_{n}t} t \log t \frac{\mathrm{d} t}{\langle \widetilde{L}_{jj} \rangle} - 1$$

$$= -\sum_{n} \frac{c_{n} (\gamma_{0} - 1 + \log k_{n})}{k_{n} \langle \widetilde{L}_{jj} \rangle} - 1. \tag{39}$$

Thus, (35) is confirmed.

In a similar vein, we can argue that the left-hand side of (36) equals

$$\int_0^\infty \sum_n c_n e^{-k_n t} \log^2 t \frac{\mathrm{d} t}{\langle \widetilde{L}_{jj} \rangle} - \ell_*^2$$

$$= \sum_n \frac{c_n [6 \log k_n (2\gamma_0 + \log k_n) + 6\gamma_0^2 + \pi^2]}{6k_n \langle \widetilde{L}_{jj} \rangle} - \ell_*^2, \quad (40)$$

while the right-hand side of (36) amounts to

$$\int_0^\infty \sum_n c_n k_n e^{-k_n t} t (\ell_* - \log t)^2 \frac{\mathrm{d} t}{\langle \widetilde{L}_{jj} \rangle}$$

$$= \int_{0}^{\infty} \sum_{n} c_{n} k_{n} e^{-k_{n}t} t \log^{2} t \frac{\mathrm{d} t}{\langle \widetilde{L}_{jj} \rangle} - 2\ell_{*} - \ell_{*}^{2}$$

$$= \sum_{n} \frac{c_{n} [6 \log k_{n} (2\gamma_{0} - 2 + \log k_{n}) + 6\gamma_{0} (\gamma_{0} - 2) + \pi^{2}]}{6k_{n} \langle \widetilde{L}_{jj} \rangle}$$

$$+ \sum_{n} \frac{2c_{n} (\gamma_{0} + \log k_{n})}{k_{n} \langle \widetilde{L}_{jj} \rangle} - \ell_{*}^{2}. \tag{41}$$

This completes the proof of (36).

The detailed balance condition $\pi_i p_{ij} = \pi_j p_{ji}$ means that a Markov chain is reversible, and the reverse chain has the same transition matrix **P** [35, §4.7, p. 203].

By the reversibility of our Markov chain, we may extend the results from the theorem above to a dual situation, namely, the statistical properties for L_{ij} , the effective length of a text fragment free from word W_i that is simultaneously flanked by W_i to the left and by W_j to the right (as considered in Fig. 1).

Trading L_{ij} (resp. L_{jj}) in (35)–(36) for L_{ij} (resp. L_{ii}), we recover (8): for a given i, we have used (35) as an estimate for $\langle \log L_{ij} \rangle$, and have used $1/n_{ij}$ times (36) as an estimate for the variance of $\langle \log L_{ij} \rangle$, which is taken over n_{ij} independent samples.

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