

# The Geometry of Syntax

Matilde Marcolli

TGSI: Topological and Geometrical Structures of Information  
CIRM Luminy  
August 28–September 1, 2017

## Talk based on some recent and ongoing work

- 1 Matilde Marcolli, Robert Berwick, Kevin Shu, *Phylogenetics of Indo-European Language families via an Algebro-Geometric Analysis of their Syntactic Structures*, in preparation;
- 2 Andrew Ortegaray, Matilde Marcolli, *A Heat Kernel analysis of the space of Syntactic Parameters*, in preparation;
- 3 Alexander Port, Matilde Marcolli, *Persistent topology and syntactic parameters of Indo-European languages*, in preparation.
- 4 Matilde Marcolli, *Syntactic Parameters and a Coding Theory Perspective on Entropy and Complexity of Language Families*, Entropy 2016, 18(4), 110
- 5 Kevin Shu, Matilde Marcolli, *Syntactic Structures and Code Parameters*, Math. Comput. Sci. 11 (2017), no. 1, 79–90.

## and some previous work

- 1 Alexander Port, Iulia Gheorghita, Daniel Guth, John M. Clark, Crystal Liang, Shival Dasu, Matilde Marcolli, *Persistent Topology of Syntax*, arXiv:1507.05134
- 2 Karthik Siva, Jim Tao, Matilde Marcolli, *Spin Glass Models of Syntax and Language Evolution*, arXiv:1508.00504
- 3 Jeong Joon Park, Ronnel Boettcher, Andrew Zhao, Alex Mun, Kevin Yuh, Vibhor Kumar, Matilde Marcolli, *Prevalence and recoverability of syntactic parameters in sparse distributed memories*, arXiv:1510.06342
- 4 Kevin Shu, Sharjeel Aziz, Vy-Luan Huynh, David Warrick, Matilde Marcolli, *Syntactic Phylogenetic Trees*, arXiv:1607.02791

## General Question: Language and Machines

- Natural Language Processing has made enormous progress in problems like automated translation
- **but** can we use computational (mathematical) techniques to better understand how the human brain processes language?
- some of the main questions:
  - Language acquisition (poverty of the stimulus): how does the learning brain converge to *one* grammar?
  - How is language (in particular syntax) stored in the brain?
  - How do languages change and evolve in time? quantitative, predictive modeling?
- **Plan:** approach these questions from a mathematical perspective, using tools from geometry and theoretical physics
- focus on the “large scale structure” of language: **syntax**

## Syntax and Syntactic Parameters

- one of the key ideas of modern Generative Linguistics:

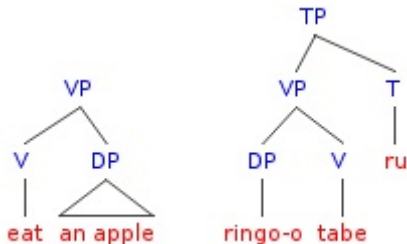
### Principles and Parameters (Chomsky, 1981)

- *principles*: general rules of grammar
- *parameters*: **binary variables** (on/off switches) that distinguish languages in terms of syntactic structures
- this idea is very appealing for a mathematician: at the level of syntax a language can be described by a set of **coordinates** given by binary variables
- however, surprisingly no mathematical model of Principles and Parameters formulation of Linguistics has been developed so far

## What are the binary variables?

- Example of parameter: **head-directionality**  
(head-initial versus head-final)

English is head-initial, Japanese is head-final



VP= verb phrase, TP= tense phrase, DP= determiner phrase

- Other examples of parameters:
  - *Subject-side*
  - *Pro-drop*
  - *Null-subject*

## Main Problems

- there is **no complete classification** of syntactic parameters
- there are hundreds of such binary syntactic variables, but not all of them are “true” syntactic parameters (conflations of deep/surface structure)
- **Interdependencies** between different syntactic parameters are poorly understood: what is a good independent set of variables, a good set of coordinates?
- syntactic parameters are **dynamical**: they change historically over the course of language change and evolution
- collecting **reliable data** is hard! (there are thousands of world languages and analyzing them at the level of syntax is much more difficult for linguists than collecting lexical data; few ancient languages have enough written texts)

## Databases of syntactic structures of world languages

- ① Syntactic Structures of World Languages (SSWL)  
<http://sswl.railsplayground.net/>
  - ② TerraLing <http://www.terraling.com/>
  - ③ World Atlas of Language Structures (WALS)  
<http://wals.info/>
  - ④ another set of data from Longobardi–Guardiano, *Lingua* 119 (2009) 1679-1706
  - ⑤ more complete set of data by Giuseppe Longobardi, 2016
- **Data Analysis** of syntax of world languages with various mathematical tools (persistent topology, etc.)



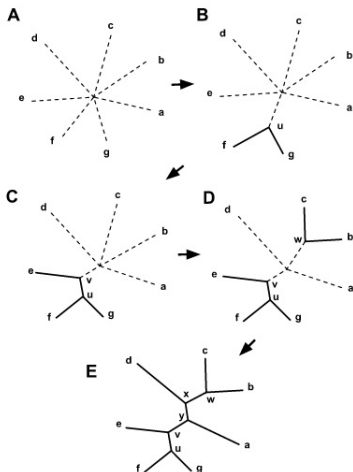
## Problems of SSWL data

- Very **non-uniformly mapped** across the languages of the database: some are 100% mapped, while for some only very few of the 116 parameters are mapped
- Linguists criticize the **choice of binary variable** (not all of them should count as “true” parameters)
- the data of Longobardi–Guardiano and the more recent data of Longobardi are more reliable: 83 parameters and 62 languages (mostly Indo-European), more completely mapped, “true” parameters
- linguistic question: can languages that are far away in terms of historical linguistics end up being close in terms of syntactic parameters?

## Phylogenetic Algebraic Geometry of Languages (ongoing work with R.Berwick, K.Shu)

- Linguistics has studied in depth how languages change over time (Philology, Historical Linguistics)
- Usually via lexical and morphological analysis
- **Goal**: understand the historical relatedness of different languages, subdivisions into families and sub-families, phylogenetic trees of language families
- Historical Linguistics techniques work best for language families where enough ancient languages are known (Indo-European and very few other families)
- Can one reconstruct phylogenetic trees **computationally** using only information on the modern languages?
- **controversial results** about the Indo-European tree based on *lexical data*: Swadesh lists of lexical items compared on the existence of cognate words (many problems: synonyms, loan words, false positives)

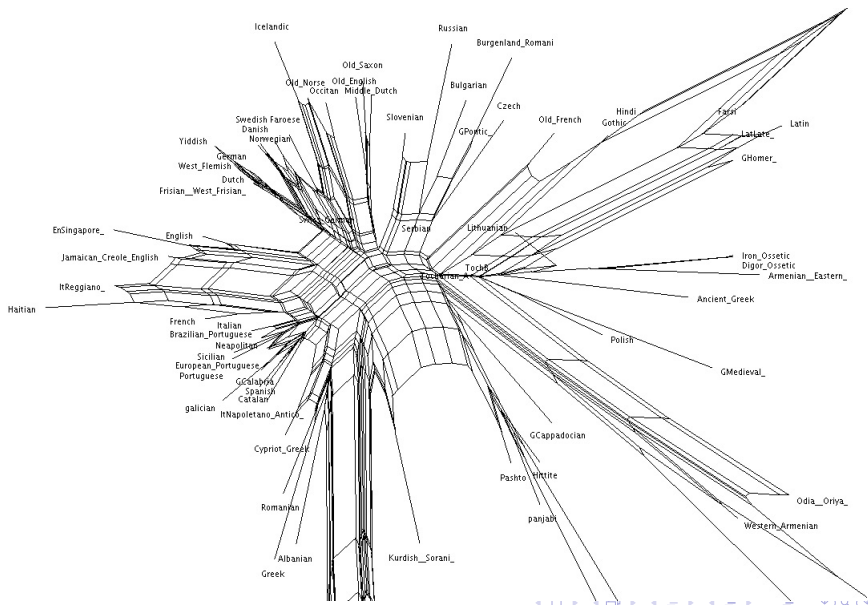
- Some phylogenetic tree reconstructions using syntactic parameters by Longobardi–Guardiano using their parameter data
- Hamming distance between binary string of parameter values + neighborhood joining method



## Expect problems: SSWL data and phylogenetic reconstructions

- known problems related to the use of Hamming metric for phylogenetic reconstruction
  - SSWL problems mentioned above (especially non-uniform mapping)
  - dependence among parameters (not independent random variables)
  - syntactic proximity of some unrelated languages
- **Phylogeny Programs** for trees and networks
    - PHYLIP
    - Splittree 4
    - Network 5

# Checking on the Indo-European tree where good Historical-Linguistics



## Indeed Problems

- misplacement of languages within the correct family subtree
  - placement of languages in the wrong subfamily tree
  - proximity of languages from unrelated families (all SSWL)
  - incorrect position of the ancient languages
- different approach: subdivide into subfamilies (some a priori knowledge from morpholexical linguistic data) and use **Phylogenetic Algebraic Geometry** (Sturmfels, Pachter, et al.) for statistical inference of phylogenetic reconstruction

## General Idea of Phylogenetic Algebraic Geometry

- Markov process on a binary rooted tree (Jukes-Cantor model)
- probability distribution at the root  $(\pi, 1 - \pi)$   
(frequency of 0/1 for parameters at root vertex) and transition matrices along edges  $M^e$  bistochastic

$$M^e = \begin{pmatrix} 1 - p_e & p_e \\ p_e & 1 - p_e \end{pmatrix}$$

- observed distribution at the  $n$  leaves polynomial function

$$p_{i_1, \dots, i_n} = \Phi(\pi, M^e) = \sum_{w_v \in \{0,1\}} \pi_{w_v} \prod_e M^e_{w_{s(e)}, w_{t(e)}}$$

with sum over “histories” consistent with data at leaves

- polynomial map that assigns

$$\Phi : \mathbb{C}^{4n-5} \rightarrow \mathbb{C}^{2^n}, \quad \Phi(\pi, M^e) = p_{i_1, \dots, i_n}$$

defines an *algebraic variety*

$$V_T = \overline{\Phi(\mathbb{C}^{4n-5})} \subset \mathbb{C}^{2^n}$$

- (Allman–Rhodes theorem) ideal  $\mathcal{I}_T$  defining  $V_T$  generated by all  $3 \times 3$  minors of all *edge flattenings* of tensor  $P = (p_{i_1, \dots, i_n})$ :  $2^r \times 2^{n-r}$ -matrix  $Flat_{e,T}(P)$

$$Flat_{e,T}(P)(u, v) = P(u_1, \dots, u_r, v_1, \dots, v_{n-r})$$

where edge  $e$  removal separates boundary distribution into  $2^r$  variable and  $2^{n-r}$  variables



## Procedure

- set of languages  $\mathcal{L} = \{\ell_1, \dots, \ell_n\}$  (selected subfamily)
- set of SSWL (or Longobardi) syntactic parameters mapped for all languages in the set:  $\pi_i, i = 1, \dots, N$
- gives vectors  $\pi_i = (\pi_i(\ell_j)) \in \mathbb{F}_2^n$
- compute frequencies

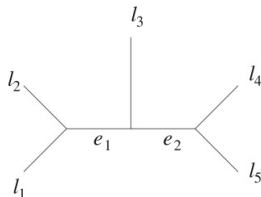
$$P = \{p_{i_1, \dots, i_n} = \frac{N_{i_1, \dots, i_n}}{N}\}$$

with  $N_{i_1, \dots, i_n}$  = number of occurrences of binary string  $(i_1, \dots, i_n) \in \mathbb{F}_2^n$  among the  $\{\pi_i\}_{i=1}^N$

- Given a *candidate tree*  $T$ , compute all  $3 \times 3$  minors of each flattening matrix  $Flat_{e,T}(P)$ , for each edge
- evaluate  $\phi_T(P)$  minimum absolute value of these minors
- smallest  $\phi_T(P)$  selects best among candidate trees
- **likelihood function**: distance of  $P$  from  $V_T$  via singular values of flattening matrices

## Simple examples

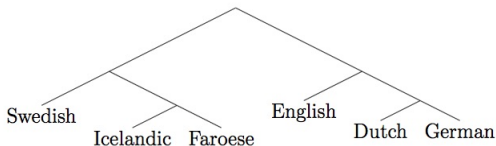
- PHYLIP and Splittree 4 misplace the position of Portuguese among the Latin languages, but phylogenetic invariants identify the correct tree ( $\ell_1$  = French,  $\ell_2$  = Italian,  $\ell_3$  = Latin,  $\ell_4$  = Spanish,  $\ell_5$  = Portuguese)



$$\text{Flat}_{e_1}(P) = \begin{pmatrix} p_{00000} & p_{00001} & p_{00010} & p_{00011} & p_{00100} & p_{00101} & p_{00110} & p_{00111} \\ p_{01000} & p_{01001} & p_{01010} & p_{01011} & p_{01100} & p_{01101} & p_{01110} & p_{01111} \\ p_{10000} & p_{10001} & p_{10010} & p_{10011} & p_{10100} & p_{10101} & p_{10110} & p_{10111} \\ p_{11000} & p_{11001} & p_{11010} & p_{11011} & p_{11100} & p_{11101} & p_{11110} & p_{11111} \end{pmatrix}$$

$$\text{Flat}_{e_2}(P) = \begin{pmatrix} p_{00000} & p_{00001} & p_{00010} & p_{00011} \\ p_{00100} & p_{00101} & p_{00110} & p_{00111} \\ p_{01000} & p_{01001} & p_{01010} & p_{01011} \\ p_{01100} & p_{01101} & p_{01110} & p_{01111} \\ p_{10000} & p_{10001} & p_{10010} & p_{10011} \\ p_{10100} & p_{10101} & p_{10110} & p_{10111} \\ p_{11000} & p_{11001} & p_{11010} & p_{11011} \\ p_{11100} & p_{11101} & p_{11110} & p_{11111} \end{pmatrix}$$

- PHYLIP and Splittree 4 misplace the relative position of sub-branches of the Germanic languages, but phylogenetic invariants identify the correct tree (similar computation)

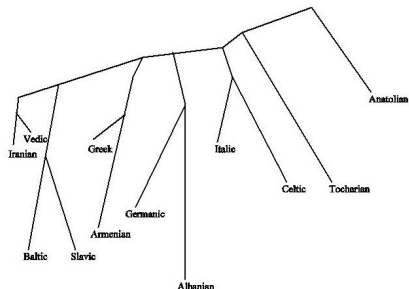
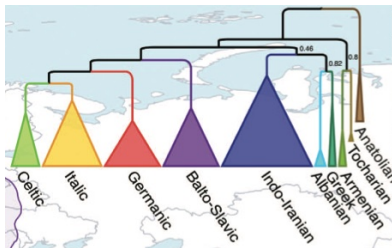


with correct subdivision into North Germanic and West Germanic sub-branches

**Conclusion:** work with smaller subfamilies, then paste together subtrees; use PHYLIP to generate candidate subtrees and phylogenetic algebraic geometry to select among them

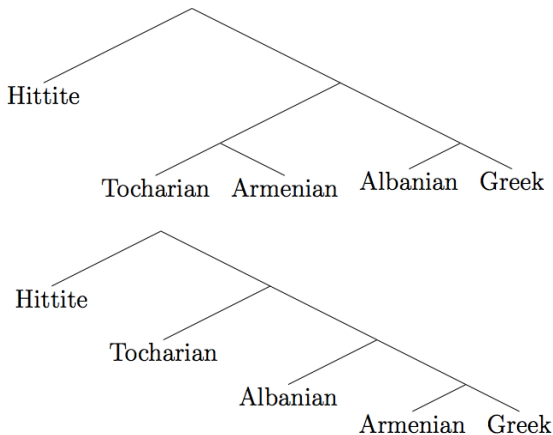
**Main Question:** can one use this method to obtain new results on the “Indo-European controversy”?

- What is the controversy? Early branches of the tree of Indo-European languages
  - The relative positions of the Greco-Armenian subtrees;
  - The position of Albanian in the tree;
  - The relative positions of these languages with respect to the Anatolian-Tocharian subtrees.
- Controversial claims by Gray and Atkinson (Nature, 2003); disputed via morphological analysis (Ringe, Warnow, Taylor, 2002)
- A. Perelysvaig, M.W. Lewis, *The Indo-European controversy: facts and fallacies in Historical Linguistics*, Cambridge University Press, 2015.



The Atkinson–Gray early Indo-European tree and the Ringe–Warnow–Taylor tree

Focus on this part of the tree:



Can detect the difference from syntactic parameters?

## Using Phylogenetic Algebraic Geometry of Syntactic Parameters?

- **Problem:** SSWL data for Hittite, Tocharian, Albanian, Armenian, and Greek have a small number of parameters that is completely mapped for all these languages (and these parameters largely agree); Hittite and Tocharian not mapped in Longobardi data.
- the SSWL data appear to favor the Atkinson–Gray tree, *but the data is too problematic to be trusted!* ...need better syntactic data on these languages (especially Hittite and Tocharian that are poorly mapped in all available databases)



**Coding Theory** to study how syntactic structures differ across the landscape of human languages

- Kevin Shu, Matilde Marcolli, *Syntactic Structures and Code Parameters*, Math. Comput. Sci. 11 (2017), no. 1, 79–90.
- Matilde Marcolli, *Syntactic Parameters and a Coding Theory Perspective on Entropy and Complexity of Language Families*, Entropy 2016, 18(4), 110

- select a group of languages  $\mathcal{L} = \{\ell_1, \dots, \ell_N\}$
- with the binary strings of  $n$  syntactic parameters form a code  $\mathcal{C}(\mathcal{L}) \subset \mathbb{F}_2^n$
- compute code parameters  $(R(\mathcal{C}), \delta(\mathcal{C}))$  code rate and relative minimum distance
- analyze position of  $(R, \delta)$  in space of code parameters
- get information about “syntactic complexity” of  $\mathcal{L}$



code parameters  $\mathcal{C} \subset \mathbb{F}_2^n$

- **transmission rate** (encoding)

$$R(\mathcal{C}) = \frac{k}{n}, \quad k = \log_2(\#\mathcal{C}) = \log_2(N)$$

for  $q$ -ary codes in  $\mathbb{F}_q^n$  take  $k = \log_q(N)$

- **relative minimum distance** (decoding)

$$\delta(\mathcal{C}) = \frac{d}{n}, \quad d = \min_{\ell_1 \neq \ell_2} d_H(\ell_1, \ell_2)$$

Hamming distance of binary strings of  $\ell_1$  and  $\ell_2$

- error correcting codes: optimize for maximal  $R$  and  $\delta$  but constraints that make them inversely correlated
- **bounds** in the space of code parameters  $(R, \delta)$

## Bounds on code parameters

- **Gilbert-Varshamov curve** (q-ary codes)

$$R = 1 - H_q(\delta), \quad H_q(\delta) = \delta \log_q(q-1) - \delta \log_q \delta - (1-\delta) \log_q(1-\delta)$$

q-ary Shannon entropy: asymptotic behavior of volumes of Hamming balls for large  $n$

- The Gilbert-Varshamov curve represents the typical behavior of large random codes (Shannon Random Code Ensemble)
- **Plotkin curve**  $R = 1 - \delta/q$ : asymptotically codes below Plotkin curve  $R \leq 1 - \delta/q$

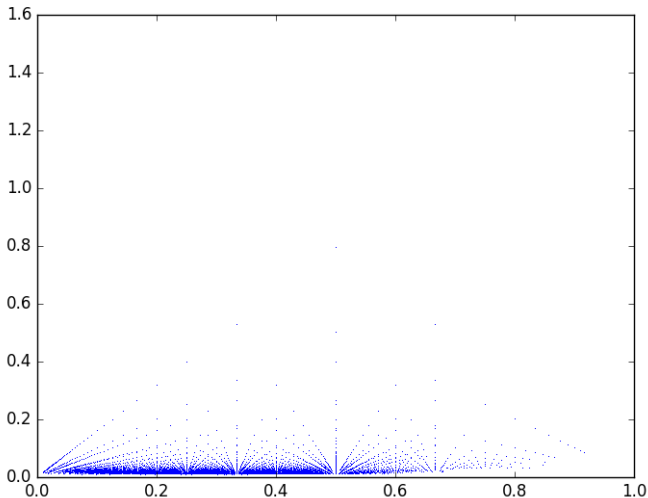
- more significant **asymptotic bound** (Manin '82) between Gilbert-Varshamov and Plotkin curve

$$1 - H_q(\delta) \leq \alpha_q(\delta) \leq 1 - \delta/q$$

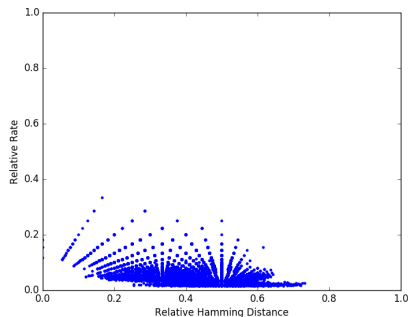
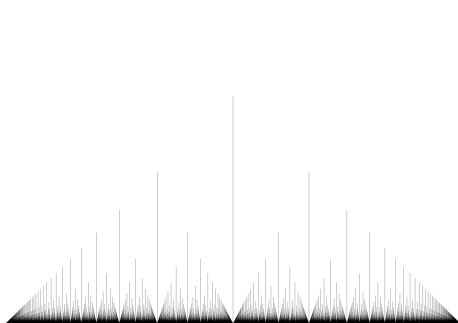
separates a region with dense code points with infinite multiplicities (below) and one with isolated code points with finite multiplicity (good codes above): difficult to find examples

- asymptotic bound not explicitly computable (related to Kolmogorov complexity of codes, Manin–Marcolli)
- difficult to construct codes above the asymptotic bound: examples from algebro-geometric codes from curves (but only for  $q \geq 49$  otherwise entirely below the GV curve)

- look at the distribution of code parameters for small sets of languages (pairs or triples) and SSWL data

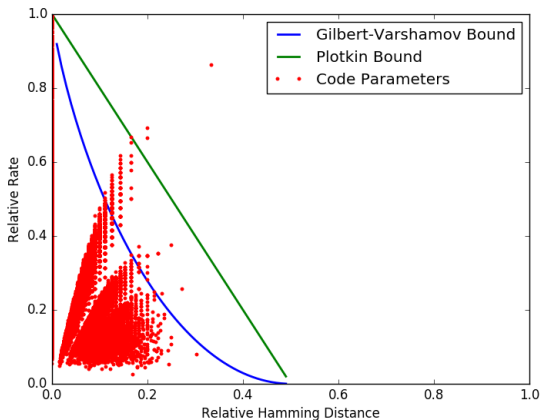


- in lower region of code parameter space a superposition of two Thomae functions ( $f(x) = 1/q$  for  $x = p/q$  coprime, zero on irrationals)



and behaves like the case of random codes with fixed  $k = \log_2(N)$

- more interesting what happens in the upper regions of the code parameter space
- take larger sets of randomly selected languages and syntactic parameters in the SSWL database



codes better than algebraic-geometric above GV, asymptotic, and Plotkin

## Spin Glass model of Language Evolution

- Karthik Siva, Jim Tao, Matilde Marcolli, *Spin Glass Models of Syntax and Language Evolution*, arXiv:1508.00504, to appear in Linguistic Analysis
- syntactic parameters are dynamical: change over time, effects of *interaction between languages* (Ancient Greek switched SOV to SVO from Homeric to Classical; Sanskrit also switched by influence of Dravidian languages; also Old English to Middle English)
- physicist viewpoint: binary variables (up/down spins) that flip by effect of interactions: **Spin Glass Model**

## Building a Spin Glass Model

- **graph**: vertices = languages, edges = language interaction (strength proportional to bilingual population)
- over each vertex a set of spin variables (syntactic parameters)
- if all syntactic parameters independent: uncoupled Ising models (low temperature: converge to more prevalent up/down state in initial configuration; high temperature fluctuations around zero magnetization state)
- role of **temperature**: fluctuations in bilingual users between different structures (“code-switching” in Linguistics)
- **Interactions between parameters!** .... *coupled* Ising models
- Hamiltonian modeling *entailment relations* in Longobardi–Guardiano data (case where one state of a parameter can make another parameter undefined)



- variables:  $S_{\ell,p_1} = \exp(\pi i X_{\ell,p_1}) \in \{\pm 1\}$ ,  $S_{\ell,p_2} \in \{\pm 1, 0\}$  and  $Y_{\ell,p_2} = |S_{\ell,p_2}| \in \{0, 1\}$
- Hamiltonian  $H = H_E + H_V$

$$H_E = H_{p_1} + H_{p_2} = - \sum_{\ell, \ell' \in \text{languages}} J_{\ell\ell'} \left( \delta_{S_{\ell,p_1}, S_{\ell',p_1}} + \delta_{S_{\ell,p_2}, S_{\ell',p_2}} \right)$$

$$H_V = \sum_{\ell} H_{V,\ell} = \sum_{\ell} J_{\ell} \delta_{X_{\ell,p_1}, Y_{\ell,p_2}}$$

$J_{\ell} > 0$  anti-ferromagnetic

- two parameters: *temperature* as before and coupling *energy of entailment*
- if freeze  $p_1$  and evolution for  $p_2$ : Potts model with external magnetic field

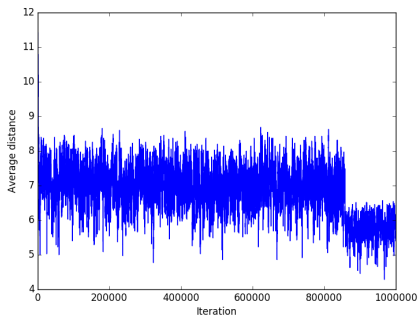
- **Metropolis–Hastings dynamics** (some binary some ternary variables)

$$\pi_A(s \rightarrow s \pm 1 \pmod{3}) = \begin{cases} 1 & \text{if } \Delta_H \leq 0 \\ \exp(-\beta \Delta_H) & \text{if } \Delta_H > 0. \end{cases}$$

$$\Delta_H := \min\{H(s + 1 \pmod{3}), H(s - 1 \pmod{3})\} - H(s)$$

- obtain interesting dynamics in the case of a small number of languages and parameters with strong entailment relations
- **Problem:** when consider more realistic models (28 languages and 63 parameters of Longobardi–Guardiano with all the entailment relations) *very slow convergence* of the Metropolis–Hastings dynamics, even for low temperature
- how to get better information on the dynamics? consider set of languages as codes and an *induced dynamics in the space of code parameters*

- Spin Glass Model dynamics for a set of languages  $\mathcal{L}$  induces dynamics on codes  $\mathcal{C}(\mathcal{L})$  and on code parameters  $(R, \delta)$
- without entailment (independent parameters) fixed  $R$  and  $\delta$  flows towards zero (spoiling code)
- with entailment parameters dynamics can improve code making  $\delta$  larger ( $R$  fixed)
- in some cases can see better the dynamics on code parameter than with average magnetization of spin glass model



**The Manifold of Syntax?** looking for relations between parameters  
(ongoing work with Andrew Ortegaray)

- Geometric methods of dimensional reduction: *Belkin–Niyogi heat kernel method*
- M. Belkin, P. Niyogi, *Laplacian eigenmaps for dimensionality reduction and data representation*, Neural Comput. 15 (6) (2003) 1373–1396.
- *Problem*: low dimensional representations of data sampled from a probability distribution on a manifold
- *Want* more efficient methods than Principal Component Analysis
- *Main Idea*: build a graph with neighborhood information, use Laplacian of graph, obtain low dimensional representation that maintains the local neighborhood information using eigenfunctions of the Laplacian

- **setting**: data points  $x_1, \dots, x_k \in \mathcal{M} \subset \mathbb{R}^\ell$  on a manifold; find points  $y_1, \dots, y_k$  in a low dimensional  $\mathbb{R}^m$  ( $m \ll \ell$ ) that *represent* the data points  $x_i$
- Step 1 (a): **adjacency graph ( $\epsilon$ -neighborhood)**: an edge  $e_{ij}$  between  $x_i$  and  $x_j$  if  $\|x_i - x_j\|_{\mathbb{R}^\ell} < \epsilon$
- Step 1 (b): **adjacency graph ( $n$  nearest neighborhood)**: edge  $e_{ij}$  between  $x_i$  and  $x_j$  if  $x_i$  is among the  $n$  nearest neighbors of  $x_j$  or viceversa
- Step 2: **weights on edges: heat kernel**

$$W_{ij} = \exp\left(-\frac{\|x_i - x_j\|^2}{t}\right)$$

if edge  $e_{ij}$  and  $W_{ij} = 0$  otherwise; heat kernel parameter  $t > 0$

- Step 3: **Eigenfunctions** for connected graph (or on each component)

$$L\psi = \lambda D\psi$$

diagonal matrix of weights  $D_{ii} = \sum_j W_{ji}$ ; Laplacian  $L = D - W$  with  $W = (W_{ij})$ ; eigenvalues  $0 = \lambda_0 \leq \lambda_1 \leq \dots \leq \lambda_{k-1}$  and  $\psi_j$  eigenfunctions

$$\psi_i : \{1, \dots, k\} \rightarrow \mathbb{R}$$

defined on set of vertices of graph

- Step 4: **Mapping by Laplace eigenfunctions**

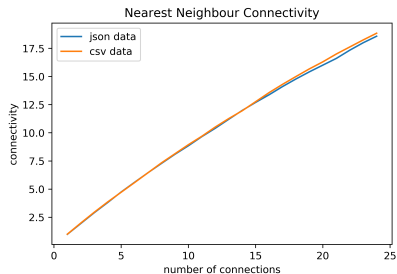
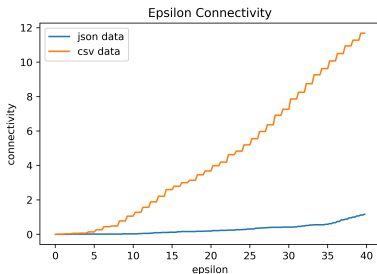
$$\mathbb{R}^\ell \supset \mathcal{M} \ni x_i \mapsto (\psi_1(i), \dots, \psi_m(i)) \in \mathbb{R}^m$$

map by first  $m$  eigenfunctions

- Belkin–Niyogi: *optimality* of embedding by Laplace eigenfunctions

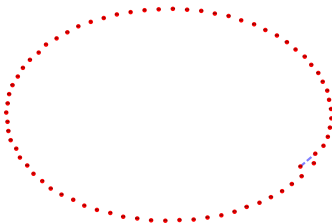
## Heat Kernel analysis of Syntactic Parameters

- Connectivity in  $\epsilon$ -neighborhood and nearest-neighbor (difference between SSWL data (json) and Longobardi data (csv))

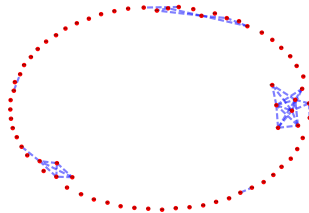


## Graphs with $\epsilon$ -neighborhood Longobardi data

Epsilon-Neighbourhood,epsilon = 1.000000



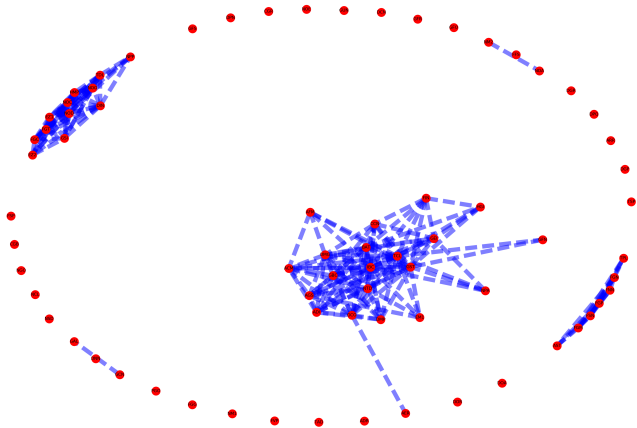
Epsilon-Neighbourhood,epsilon = 8.000000





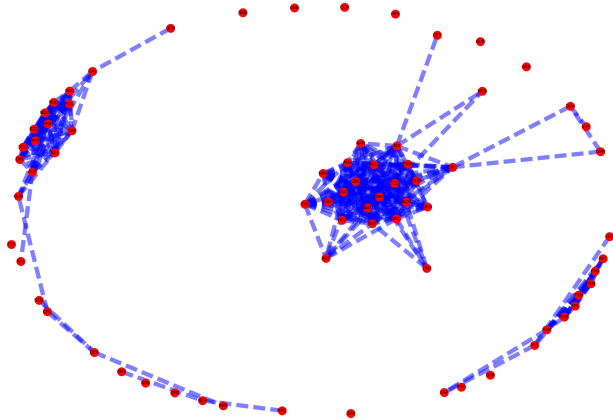
## Graphs with $\epsilon$ -neighborhood Longobardi data

Epsilon-Neighbourhood, epsilon = 15.000000



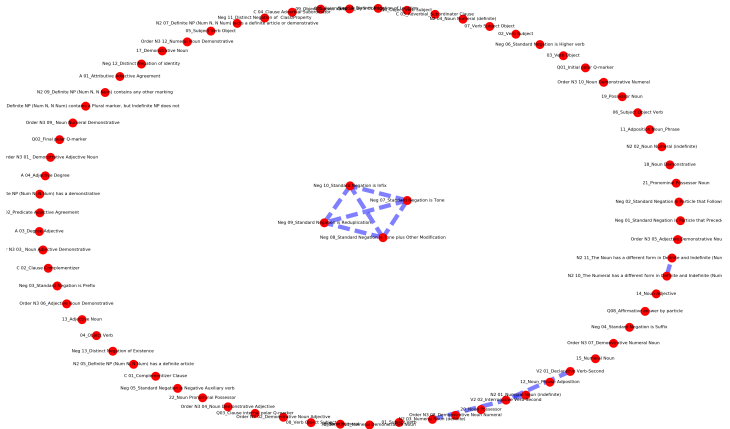
## Graphs with $\epsilon$ -neighborhood Longobardi data

Epsilon-Neighbourhood, epsilon = 22.000000



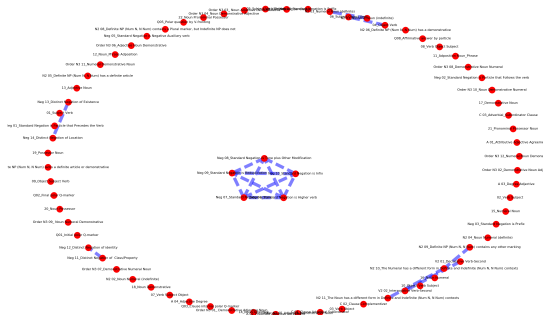
## Graphs with $\epsilon$ -neighborhood SSWL data

Epsilon-Neighbourhood,epsilon =15.000000



# Graphs with $\epsilon$ -neighborhood SSWL data

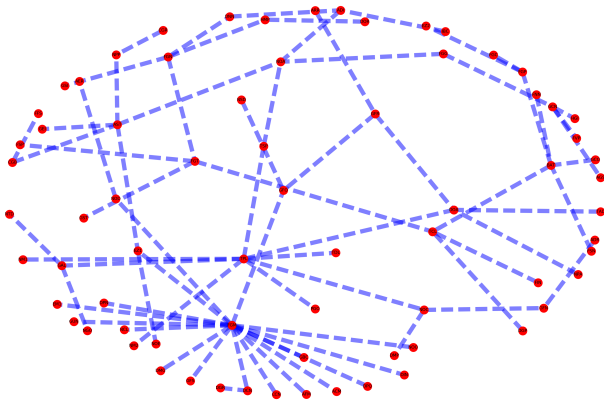
Epsilon-Neighbourhood,  $\epsilon = 22.000000$



The  $\epsilon$ -neighborhood construction is better suited to gain connectivity information in the Longobardi data: the SSWL data remain highly disconnected (only small local structures)

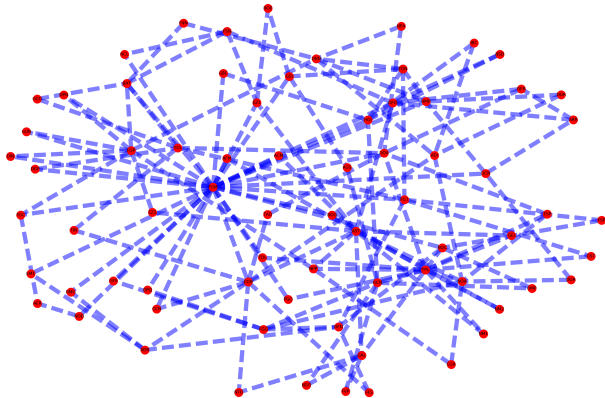
## Graphs with $n$ -neighborhood Longobardi data

Nearest 1 Connections



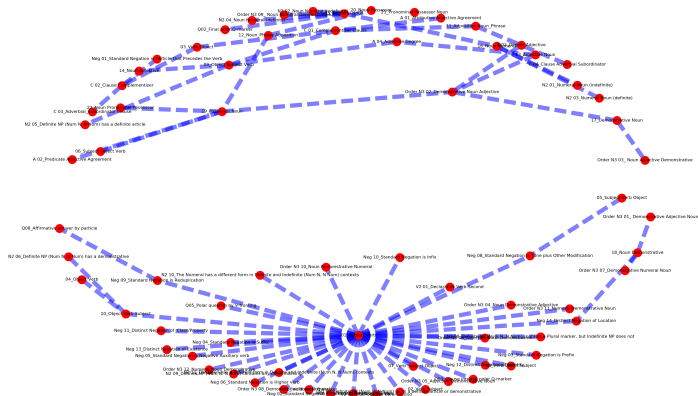
## Graphs with $n$ -neighborhood Longobardi data

Nearest 2 Connections



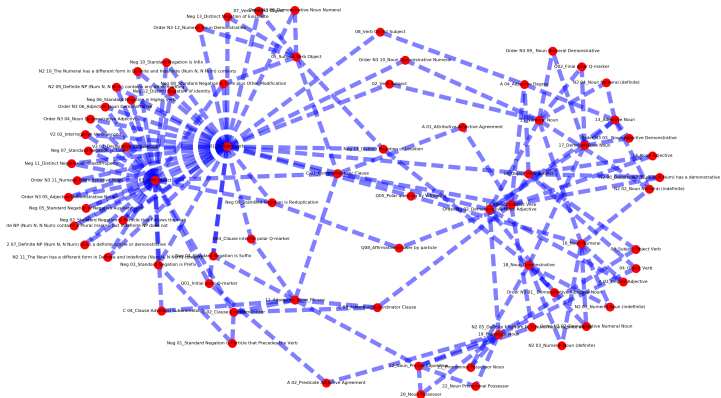
# Graphs with $n$ -neighborhood SSWL data

## Nearest 1 Connections



## Graphs with $n$ -neighborhood SSWL data

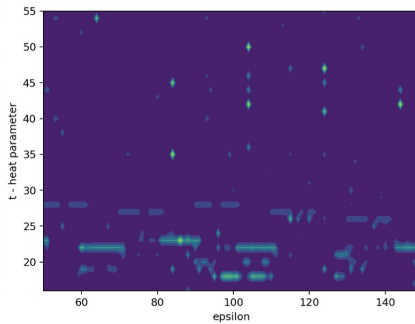
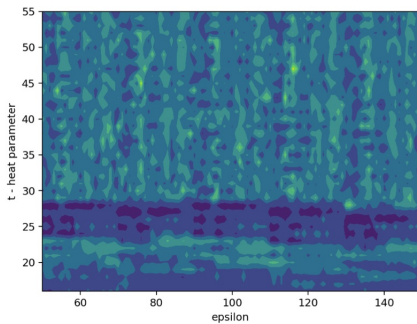
### Nearest 2 Connections





## Regions of $\epsilon$ - $t$ space

- Graphs depend on  $\epsilon$ -neighborhood and on  $t$ -heat kernel variable
- explore  $\epsilon$ - $t$  space: determine regions where high variance in distribution of each parameter under the heat kernel mapping
- high variance in a parameter suggests it is highly independent (similar to PCA method)
- contour plots of variance; plots of number of outliers produced in set of coordinates for a given parameter



## Further Questions

- an in depth linguistic analysis of the meaning of these clustering structures is still needed (ongoing work)
- comparison of the heat kernel technique with other dimensional reduction techniques (PCA etc.)
- more detailed discussion of different regions of the  $\epsilon$ - $t$  space in the heat kernel model (specific parameters with high independence measure)
- manifold  $\mathcal{M}$  reconstruction? Belkin-Niyogi results

## Topological Structures of Syntactic Parameters

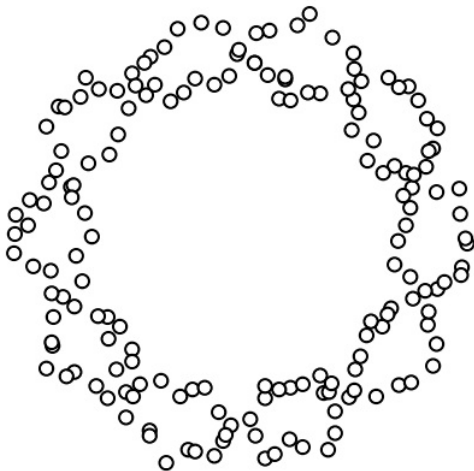
persistent homology (ongoing work with Alex Port)

- previous work computing persistent homology of SSWL data

Alexander Port, Iulia Gheorghita, Daniel Guth, John M. Clark, Crystal Liang, Shival Dasu, Matilde Marcolli, *Persistent Topology of Syntax*, arXiv:1507.05134

- ongoing work: persistent homology in the new Longobardi data
- main questions:
  - persistent generators of  $H_0$  and phylogenetic trees?
  - meaning of persistent generators of  $H_1$ ?

## Persistent Topology of Data Sets

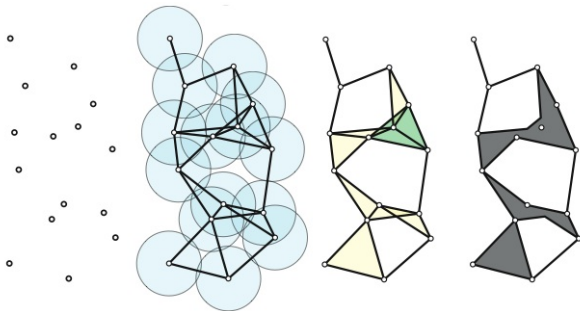


how data cluster around topological shapes at different scales

## Vietoris–Rips complexes

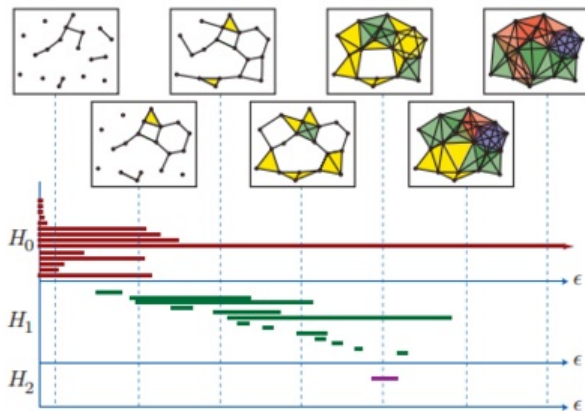
- set  $X = \{x_\alpha\}$  of points in Euclidean space  $\mathbb{E}^N$ , distance  $d(x, y) = \|x - y\| = (\sum_{j=1}^N (x_j - y_j)^2)^{1/2}$
- Vietoris-Rips complex  $R(X, \epsilon)$  of scale  $\epsilon$  over field  $\mathbb{K}$ :

$R_n(X, \epsilon)$  is  $\mathbb{K}$ -vector space spanned by all unordered  $(n + 1)$ -tuples of points  $\{x_{\alpha_0}, x_{\alpha_1}, \dots, x_{\alpha_n}\}$  in  $X$  where all pairs have distances  $d(x_{\alpha_i}, x_{\alpha_j}) \leq \epsilon$



(image by Jeff Erickson)

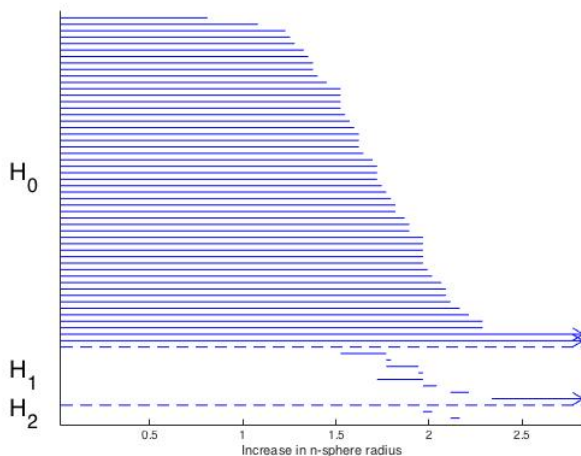
- inclusion maps  $R(X, \epsilon_1) \hookrightarrow R(X, \epsilon_2)$  for  $\epsilon_1 < \epsilon_2$  induce maps in homology by functoriality  $H_n(X, \epsilon_1) \rightarrow H_n(X, \epsilon_2)$



(image by forty.to)

barcode diagrams: births and deaths of persistent generators

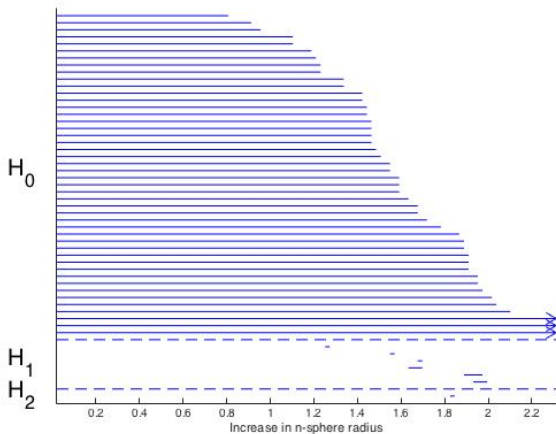
## Persistent Topology of Indo-European Languages (SSWL data)



- Two persistent generators of  $H_0$  (Indo-Iranian, European)
- One persistent generator of  $H_1$

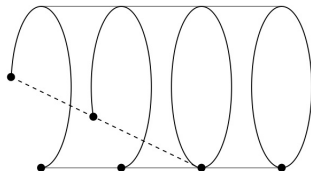
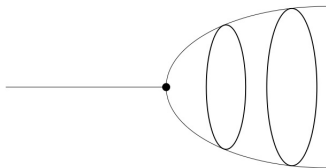


## Persistent Topology of Niger–Congo Languages (SSWL data)



- Three persistent components of  $H_0$  (Mande, Atlantic-Congo, Kordofanian)
- No persistent  $H_1$

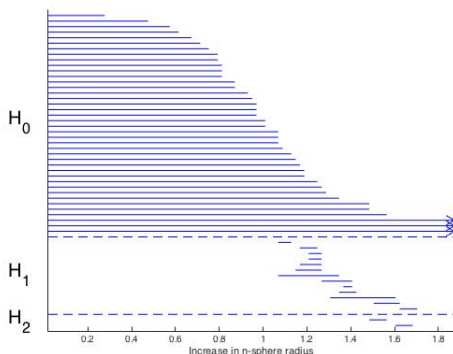
## Sources of Persistent $H_1$



- “Hopf bifurcation” type phenomenon
- two different branches of a tree closing up in a loop

two different types of phenomena of historical linguistic development within a language family

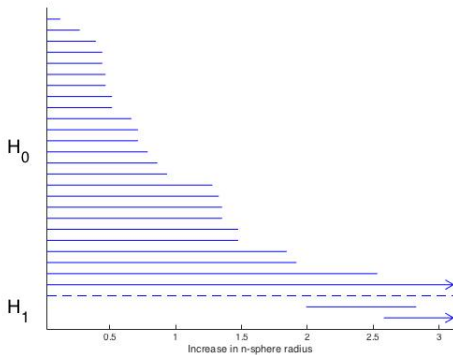
## What is the Indo-European $H_1$ ?



Persistent topology with Hellenic (and Indo-Iranic) branch removed

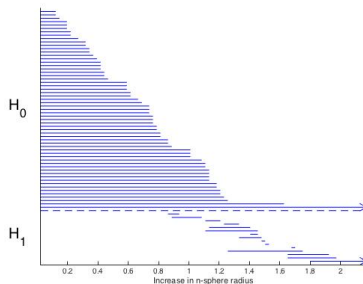
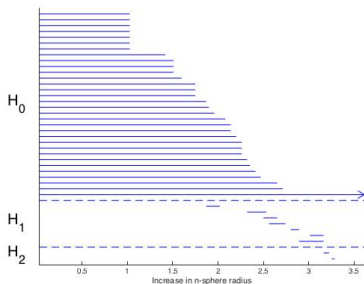
- related to influences (at the syntactic level) of the Hellenic branch on languages in other branches (like some Slavic languages)
- consistent with some previous independent linguistic observations by Longobardi
- what about the new Longobardi data analyzed topologically?

## Topological Analysis of New Syntactic Data (ongoing with Alex Port)



Longobardi data (2016): persistent generator of  $H_1$  still present

## Evidence of further structures at different scales



Linguistic interpretation: behavior of  $H_0$  versus phylogenetic trees;  
interpretation of generators of  $H_1$ ?

## Longer Term Goals

- import a set of different mathematical techniques (phylogenetic algebraic geometry, persistent topology, coding theory, statistical mechanics, geometric models of associative memory) in order to *study natural languages as dynamical objects*

- create mathematical and computational models of

- ① how languages are acquired?
- ② how languages are stored in the brain?
- ③ how languages change and evolve dynamically in time?

*for human languages viewed at the level of their syntactic structures*