# Detecting Path Anomalies in Sequential Data on Networks

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#### This Talk

Motivation: Understanding mechanisms behind sequential data on networks

#### Today:

Motivate the study of path anomalies

Introduce de Bruijn graph representation of sequential data

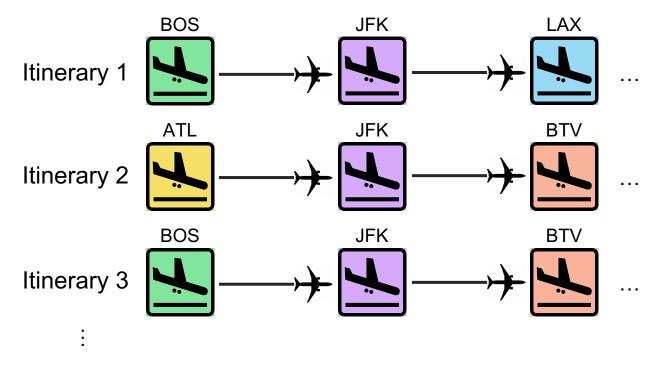
Develop tractable **null model** to measure deviation of path data from expectation

Validate null model in synthetic data + compare with naïve baseline method

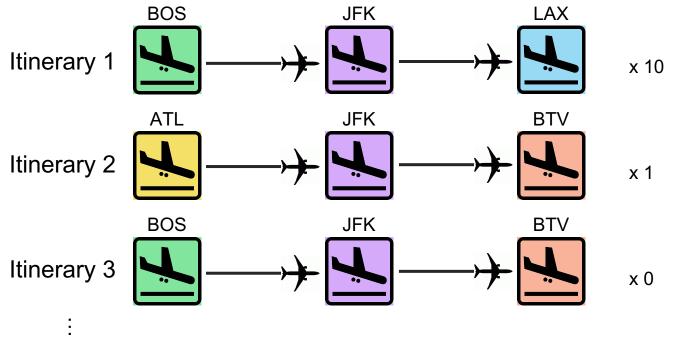
**Application** of methodology to a real system



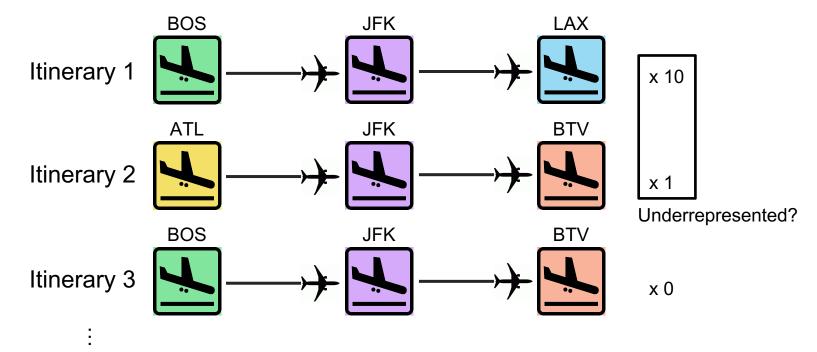




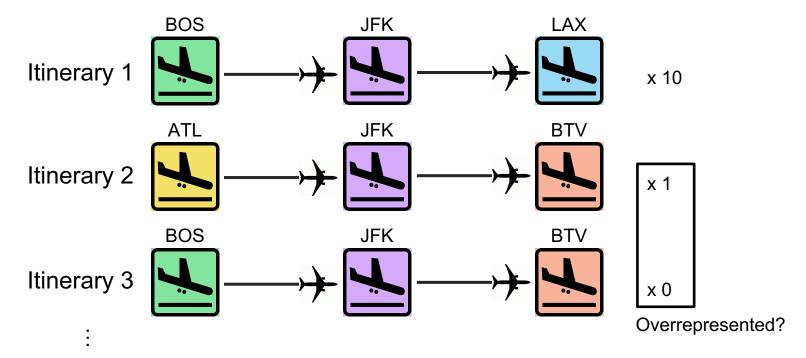








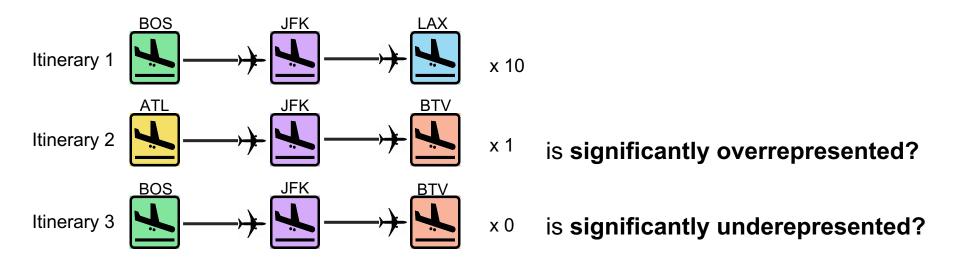






#### **Research Question**

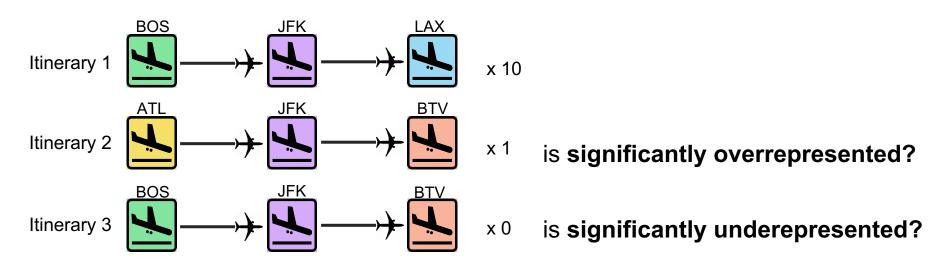
Given this pathway dataset, can we determine whether...





#### **Research Question**

Given this pathway dataset, can we determine whether...





In other words: Which **paths** are **anomalous**?

### Problem: Path anomaly detection

For a given pathway dataset S, graph G, and integer k, identify paths of length k through G whose observed frequencies in S deviate significantly from random expectation in a (k-1)-order model of paths through G.



# Problem: Path anomaly detection

For a given pathway dataset S, graph G, and integer k, identify paths of length k through G whose observed frequencies in S deviate significantly from random expectation in a (k-1)-order model of paths through G.

When k=2, this corresponds to comparing a random walk with a single step of memory to a memoryless (Markovian) random walk on G.



### Toy Example

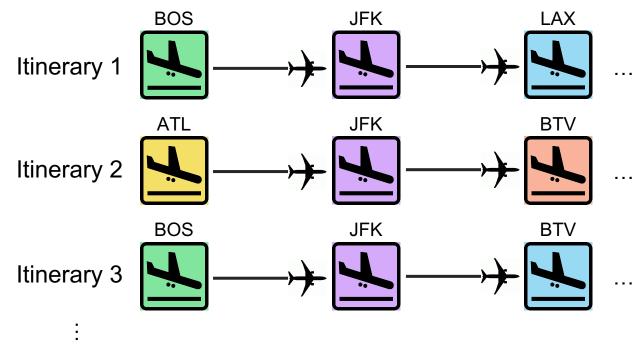
#### Three Goals:

- 1. Introduce de Bruijn graphs as representations of sequential data
- 2. Show how path anomalies emerge in a simple setting
- 3. Show how path anomalies can be detected through a random walk simulation approach

(Spoiler: Simulation approach is infeasible for real world datasets!)

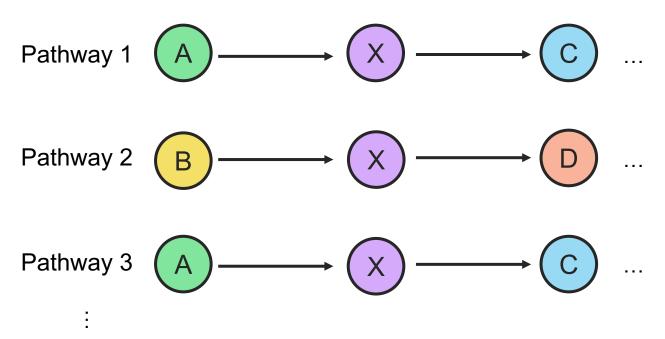


# Toy Example





## Toy Example



# Toy Example: Data

$$S = \left\{ \begin{array}{cccc} p_1 & A & X & C & \dots \\ & p_2 & B & X & D & \dots \\ & & & & & & & & & & & & \\ & p_3 & A & X & C & \dots & & & \\ & \vdots & & & & & & & & \\ & p_N & & & & & & & & \\ \end{array} \right.$$

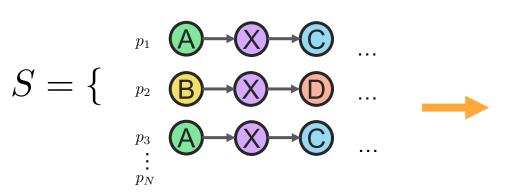


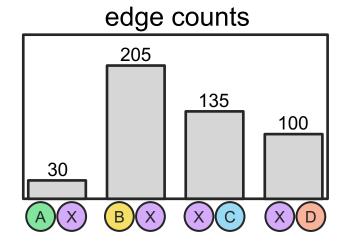
## Toy Example: Data

$$p_1$$
  $A$   $X$   $C$  ...  $S$  =  $\{$   $p_2$   $B$   $X$   $D$  ...  $p_3$   $A$   $X$   $C$  ...  $p_N$ 

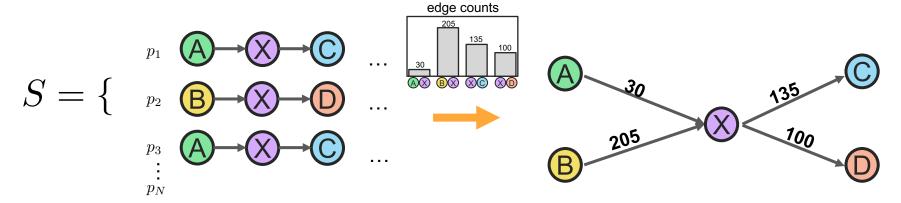


# Toy Example: Data to (first-order) graph

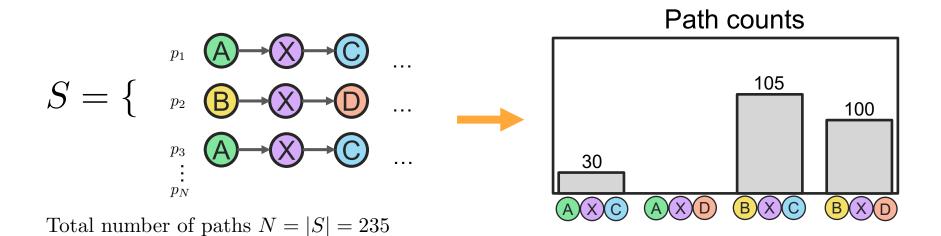




# Toy Example: Data to (first-order) graph

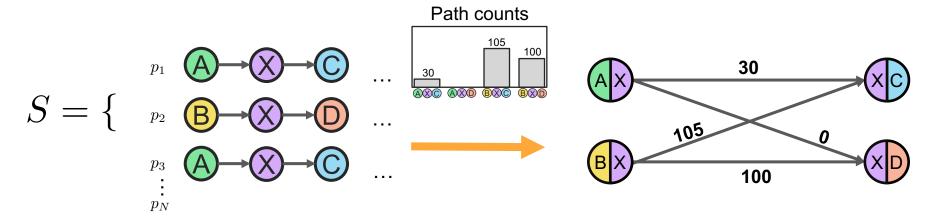


# Toy Example: Data to 2<sup>nd</sup> order de Bruijn graph





# Toy Example: Data to 2<sup>nd</sup> order de Bruijn graph



# Toy Example: Path Anomalies via Simulations

For a given pathway dataset S, graph G, and integer k, identify paths of length k through G whose observed frequencies in S deviate significantly from random expectation in a (k-1)-order model of paths through G.

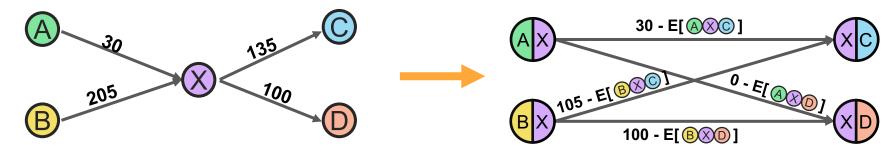


# Toy Example: Path Anomalies via Simulations

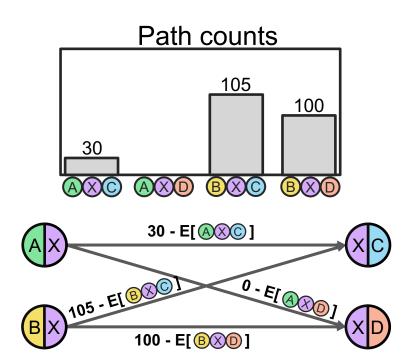
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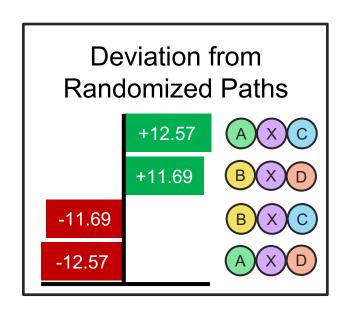
Simulate many random walk datasets

Compute expected frequency of each pathway and subtract from observed value



#### Toy Example: Path Anomalies via Simulations







# Challenges



## Path Anomaly Detection: Challenges

Detecting path anomalies via simulations → computationally intensive

Result is expected value, no concrete notion of significance

Alternative: detect path anomalies analytically by developing a tractable null model



### Null Model: Challenges

Traditional null models (e.g. configuration model) cannot be applied directly



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Edges between higher-order nodes can not be randomized by stub-matching





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Edges between higher-order nodes can not be randomized by stub-matching



Need to randomize *edge weight distribution* in de Bruijn graph models, since connectivity structure is fixed by 1<sup>st</sup>-order topology

#### **HYPA: Efficient Detection of Path Anomalies**



#### Generalized Hypergeometric Ensembles: Statistical Hypothesis Testing in Complex Networks

Giona Casiraghi,<sup>1,\*</sup> Vahan Nanumyan,<sup>1,†</sup> Ingo Scholtes,<sup>1,2,‡</sup> and Frank Schweitzer<sup>1,§</sup> <sup>1</sup>ETH Zürich, Chair of System Design, Weinbergstrasse 56/58, 8092 Zürich, Switzerland <sup>2</sup>AIFB. Karlsruhe Institute of Technology, Karlsruhe, Germany (Dated: 5th August 2016)

Statistical ensembles of networks, i.e., probability spaces of all networks that are consistent with given aggregate statistics, have become instrumental in the analysis of complex networks. Their numerical and analytical study provides the foundation for the inference of topological patterns, the definition of network-analytic measures, as well as for model selection and statistical hypothesis testing. Contributing to the foundation of these data analysis techniques, in this Letter we introduce generalized hypergeometric ensembles, a broad class of analytically tractable statistical ensembles of finite, directed and weighted networks. This framework can be interpreted as a generalization of the classical configuration model, which is commonly used to randomly generate networks with a given degree sequence or distribution. Our generalization rests on the introduction of dyadic link propensities, which capture the degree-corrected tendencies of pairs of nodes to form edges between each other. Studying empirical and synthetic data, we show that our approach provides broad perspectives for model selection and statistical hypothesis testing in data on complex networks.

PACS numbers: 89.75.Hc, 02.50.Sk, 89.75.Kd

#### Generalised hypergeometric ensembles of random graphs: the configuration model as an urn problem

Giona Casiraghi\* Vahan Nanumyan<sup>†</sup>

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#### Abstract

We introduce a broad class of random graph models: the generalised hypergeometric ensemble (GHypEG). This class enables to solve some long standing problems in random graph theory. First, GHypEG provides an elegant and compact formulation of the well-known configuration model in terms of an urn problem. Second, GHypEG allows to incorporate arbitrary tendencies to connect different vertex pairs. Third, we present the closed-form expressions of the associated probability distribution ensures the analytical tractability of our formulation. This is in stark contrast with the previous state-of-the-art, which is to implement the configuration model by means of computationally expensive procedures.

Generalization of the configuration model to weighted, directed networks.



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Fixes the *expected* weight of every node, rather than the *exact* degree sequence.

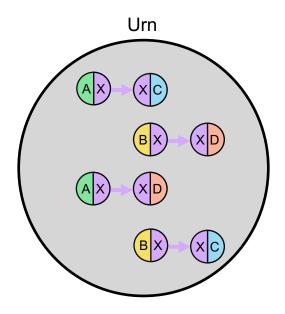


Generalization of the configuration model to weighted, directed networks.

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#### **Urn Problem Intuition:**

 Each pair of nodes that can possibly connect is assigned a color

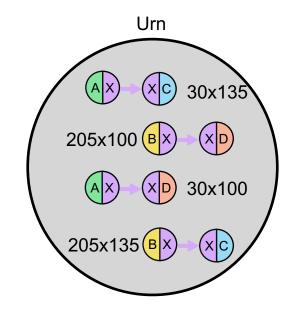


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- Each pair of nodes that can possibly connect is assigned a color
- $\circ$  Add  $K_{ij}$  balls, where  $K_{ij}=k_i^{
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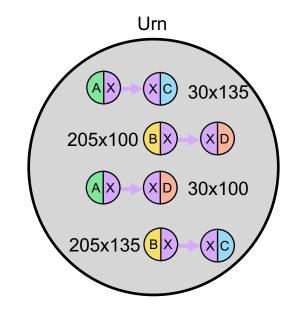


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  m out} k_j^{
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- Oraw m edges to sample a network from the urn, where  $m = \sum_{ij} W_{ij}$





# Hypergeometric Ensemble

$$m = \sum_{ij} W_{ij}$$
$$K_{ij} = k_i^{\text{out}} k_j^{\text{in}}$$

$$Pr(X_{vw} = f(v, w)) \propto {\binom{K_{vw}}{f(v, w)}} {\binom{\sum_{ij} K_{ij} - K_{vw}}{m - f(v, w)}}$$



## Hypergeometric Ensemble

$$m = \sum_{ij} W_{ij}$$
$$K_{ij} = k_i^{\text{out}} k_j^{\text{in}}$$

$$Pr(X_{vw} = f(v, w)) \propto {K_{vw} \choose f(v, w)} {\sum_{ij} K_{ij} - K_{vw} \choose m - f(v, w)}$$

Probability of observing frequency f(v, w) given the entire weighted network structure.



# Hypergeometric Ensemble

$$m = \sum_{ij} W_{ij}$$
$$K_{ij} = k_i^{\text{out}} k_i^{\text{in}}$$

$$Pr(X_{vw} = f(v, w)) \propto {\binom{K_{vw}}{f(v, w)}} {\binom{\sum_{ij} K_{ij} - K_{vw}}{m - f(v, w)}}$$

Number of ways to pick f(v,w) multiedges from  $K_{vw}$  possible.



## Hypergeometric Ensemble

$$m = \sum_{ij} W_{ij}$$
$$K_{ij} = k_i^{\text{out}} k_i^{\text{in}}$$

$$Pr(X_{vw} = f(v, w)) \propto {\binom{K_{vw}}{f(v, w)}} {\binom{\sum_{ij} K_{ij} - K_{vw}}{m - f(v, w)}}$$

Number of ways to pick everything else.

## Putting it all together: HYPA scores

$$HYPA^{(k)}(\overrightarrow{v}, \overrightarrow{w}) := \Pr(X_{\overrightarrow{v}\overrightarrow{w}} \le f(\overrightarrow{v}, \overrightarrow{w}))$$

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If "close" to **0**, then the pathway is **underrepresented**.

If "close" to 1, then pathway is **overrepresented**.

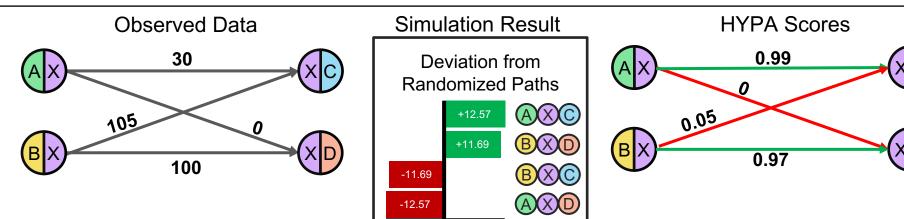


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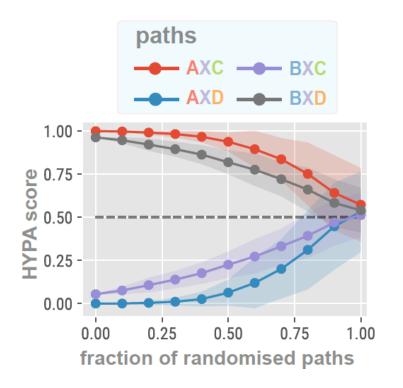




### Validation

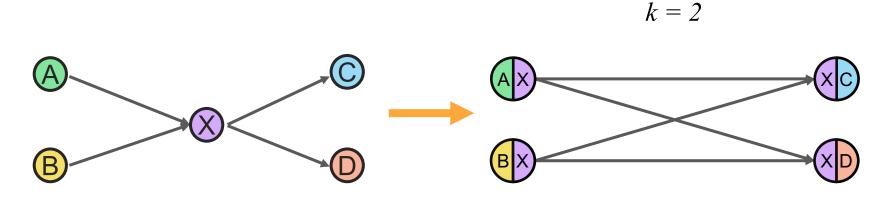


### Noise via Path Randomization

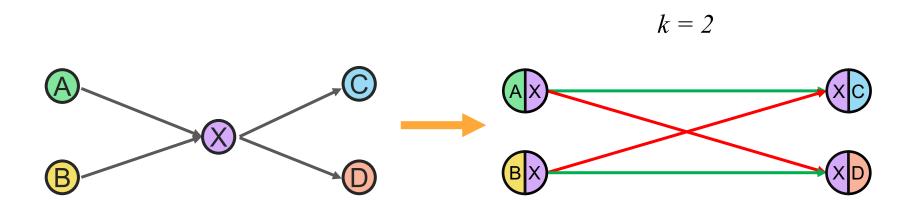




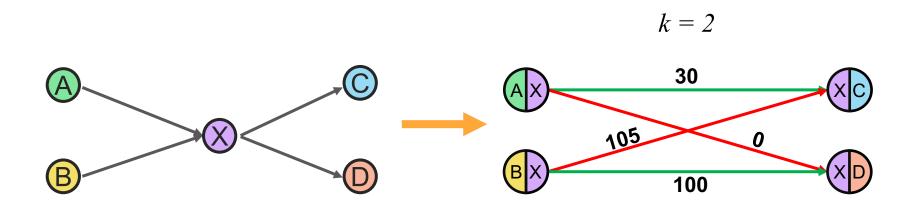
Start with an arbitrary first order topology, then construct the *kth*-order de Bruijn graph



Randomly choose some edges to label over-represented

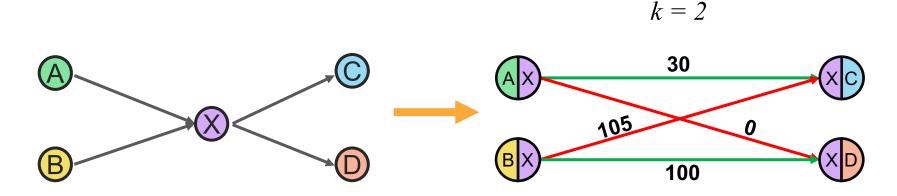


Assign heterogeneous weights based on label





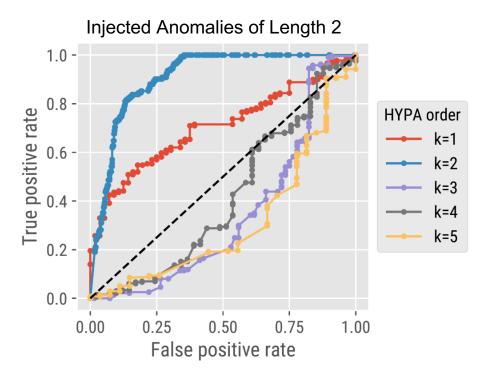
Generate paths via random walks on this model, then evaluate ability of HYPA to detect injected anomalies (binary classifier).



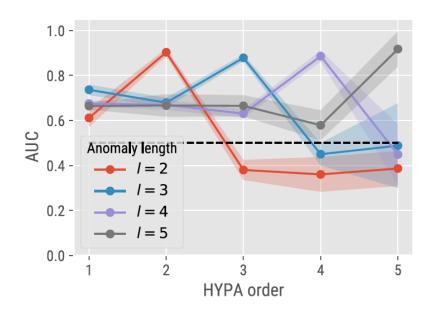
## Synthetic Anomalies: ROC Example



### Synthetic Anomalies: ROC Example



### Synthetic Anomalies: AUC Results





## Application to Flight Data



### **Airlines**

5% sample of all US domestic flights in 2018

	Topology		Sequences			
Data	Nodes	Edges	Total	Unique	l <sup>max</sup>	$\langle l \rangle$
<b>Flights</b>	382	6933	185871	88539	10	2.48



### **Airlines**

### Hypotheses:

1. Return flights should be over-represented, since people most often travel round trip.



## Airlines: Return trips are over-represented

α	Return	Non-return
0.05	0.915	0.340
0.01	0.851	0.130
0.001	0.760	0.023
0.0001	0.688	0.004
0.00001	0.628	0.001

Fraction of over-represented return/non-return flights for various discrimination thresholds.



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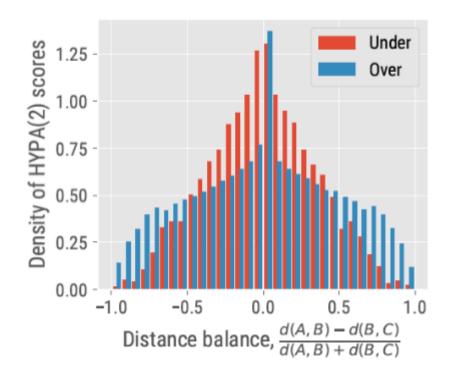


### Airlines

### Hypotheses:

- 1. Return flights should be over-represented, since people most often travel round trip.
- 2. Over-represented non-return flights are due to regional/national hubs, since people need to fly from small airports  $\rightarrow$  regional hub  $\rightarrow$  large airport.

### Airlines: Trip Balance



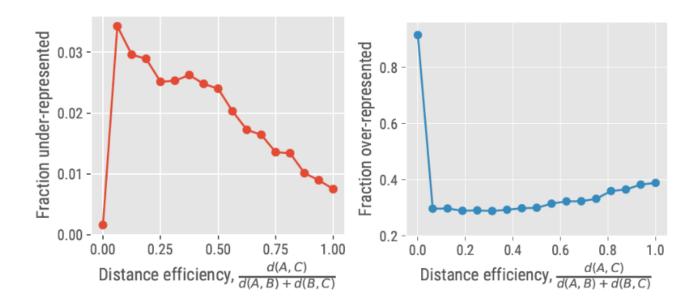
### **Airlines**

### Hypotheses:

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- 2. Over-represented non-return flights are due to regional/national hubs, since people need to fly from small airports  $\rightarrow$  regional hub  $\rightarrow$  large airport.
- 3. "Efficient" paths are more likely to be over-represented.



### Airlines: Efficiency



### Thanks!

### Tim LaRock

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https://arxiv.org/abs/1905.10580

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Scholtes, Ingo. "When is a network a network?: Multi-order graphical model selection in pathways and temporal networks." *Proceedings of the 23rd ACM SIGKDD International Conference on Knowledge Discovery and Data Mining.* ACM, 2017.

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R. TransStat. Origin and destination survey database. http://www.transtats.bts.gov/Tables.asp?DB\_ID=125, 2018.

### **Detecting Path Anomalies in Time Series Data on Networks**

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https://arxiv.org/abs/1905.10580



### Definition: kth-order de Bruijn Graph

For a given graph G = (V, E) and positive integer k we define a k-th order De Bruijn graph of paths in G as a graph  $G^k = (V^k, E^k)$ , where (i) each node  $\vec{v} := v_0 v_1 . \vec{.} . v_{k-1} \in V^k$  is a path of length k-1 in G, and (ii)  $(\vec{v}, \vec{w}) \in E^k$  iff  $v_{i+1} = w_i$  for  $i = 0, \ldots, k-2$ .



### Pseudocode

**Algorithm 1** Compute HYPA(S, k): Compute kth order HYPA scores for sequence dataset S.

**Input:** S (sequences), k (desired order)

**Output:** HYPA $^{(k)}$  score for all k-th order paths

1:  $G^k \leftarrow \text{DeBruijnGraph}(S, k) \# \text{Construct } k \text{th order graph}$ 

2:  $\Xi \leftarrow \text{fitXi}(G^k, \text{tolerance}) \# \text{Optimization (Algorithm 2 in Appendix A.1)}$ 

- 3: for  $(\overrightarrow{v}, \overrightarrow{w}) \in G^k$  do
- 4:  $\text{HYPA}^{(k)}(\overrightarrow{v}, \overrightarrow{w}) \leftarrow \Pr(x_{vw} \leq (\overrightarrow{v}, \overrightarrow{w}) \mid m, \Xi)$ # Compute CDF
- 5: **return**  $HYPA^{(k)}$



## Naïve Baseline Comparison

Frequency-Based Anomaly Detection (FBAD)

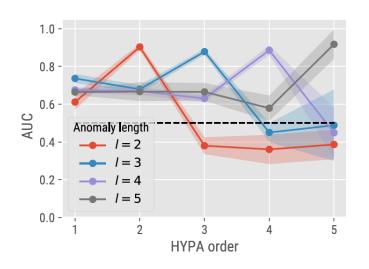
Compute mean,  $\mu$ , and standard deviation,  $\sigma$ , of kth order edge weights

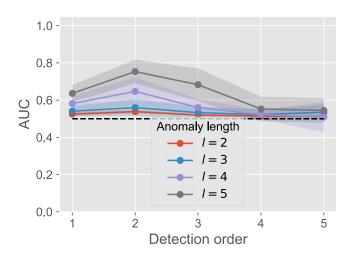
Given scaling factor  $\alpha$ , label edges as

- Overrepresented if frequency is larger than  $\mu + \sigma \alpha$
- Underrepresented if frequency is smaller than  $\mu$   $\sigma\alpha$



## Synthetic Anomalies



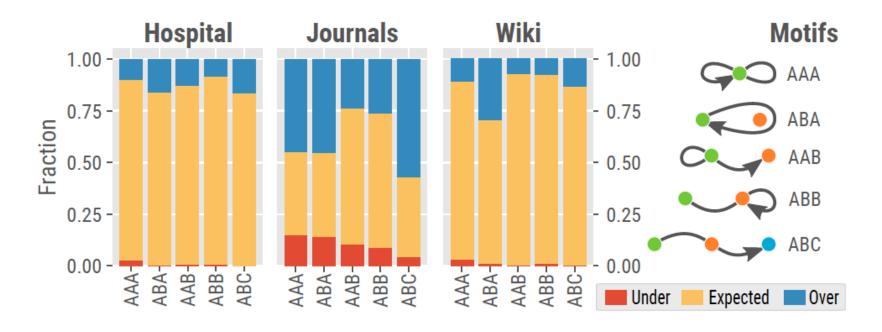


### Real Data

	Topology		Sequences			
Data	Nodes	Edges	Total	Unique	$l^{\max}$	$\langle l \rangle$
Tube	268	646	4295731	67015	35	6.75
<b>Flights</b>	382	6933	185871	88539	10	2.48
Journals	283	1743	480496	309565	35	14.8
Hospital	75	1138	28422	2561	5	1.19
Wiki	100	1598	29682	7431	21	1.64



## **Exploring Motifs**





### Case Study: London Tube

#### Data:

- Origin → destination statistics between London Tube stations
  - (origin, destination, #observations)
- Shortest paths between stations
  - Assume people follow shortest paths



### London Tube

#### Hypothesis:

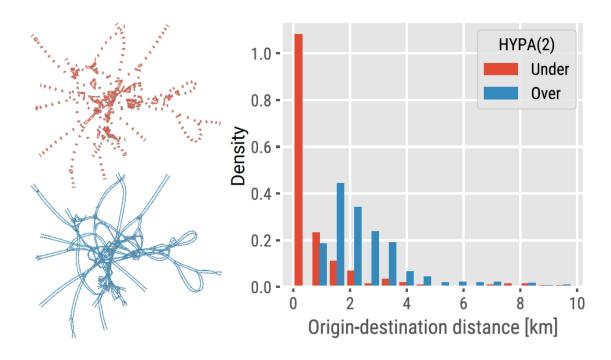
- People typically use public transportation to travel large geographic distances
- Overrepresented pathways should cover larger distances

#### Test:

- Measure distance between every station
- For 2nd order transitions A-B-C, compute distance between nodes A and C
- Analyze distributions of distance in over vs. under represented transitions
  - Expect to see distribution shifted towards higher values for over-represented transitions



### London Tube





### London Tube

$\mathbf{HYPA}^{(k)}$	k = 2	k = 3	k = 4	k = 5	k = 6
Under [km]	0.00	2.38	3.29	4.60	5.43
Over [km]	2.20	2.93	3.79	5.21	5.63
<i>p</i> -value	$< 10^{-170}$	$< 10^{-7}$	$< 10^{-4}$	0.006	0.08

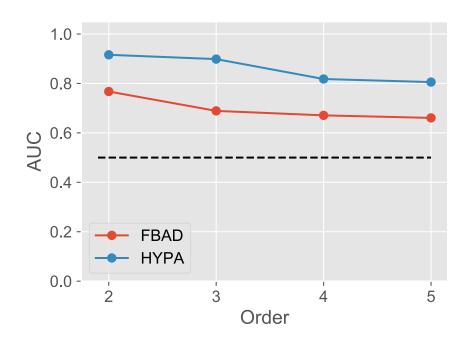
Median distance between source and destination nodes in under/over represented transitions.

### Constructing Ground Truth

Construct ground truth based on the method discussed earlier:

- Randomize path data using k-1<sup>st</sup> order random walks
- Compute kth-order path statistics
- Repeat m times, noting the frequency of each path
- Estimate multinomial distribution and its CDF from these statistics
- If CDF(path) > threshold, label over-represented

### Tube Data - Ground Truth





### Computational complexity

$$O(N+|V|^2\lambda_1^k)$$



## Scalability

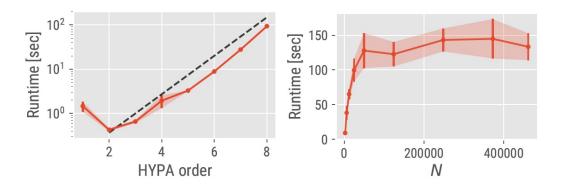


Figure 8: Empirical scalability of HYPA. Left: Required time to detect path anomalies of length k for the Tube data. Right: Runtime in Flights data for detection order k=1 and varying data size N randomly sampled from the data. All data points correspond to the mean of ten repeated measurements, with the standard deviations shown as bars.