

# Modeling Hypergraph Data with Diversity

Xianshi Yu

Department of Computer Science  
University of Wisconsin-Madison

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# Overview of my research

## Network Data Analysis

## Signal Processing   EHR Data

- Network community detection
- Collaborative filtering using social networks
- Hypergraph data analysis

# Hypergraph data—examples

Hypergraph data characterize ‘multi-actor’ relations

## Collaborations

paper 1-- Kulesza, A. & Taskar, B. (2012)

paper 2-- Brunel, V.-E., Moitra, A., Rigollet, P., & Urschel, J. (2017)

...

paper 8-- Gartrell, M., Paquet, U., & Koenigstein, N. (2017)

## Medical codes in electronic health records (EHR)

patient visit 1-- J44.9, J45.9, B44.9, O60, L50.5

patient visit 2-- J45.9, J46, O60, L50.5

...

patient visit 7-- Z11.52, Z20.822, Z86.16, U07.1, J12.82

## Shopping orders

order 1-- scissors, pencil, cheese, spinach

order 2-- tape, tissues, lemons

...

order 9-- pork, vitamins, pan, lock, brush

## Ingredients in cooking recipes

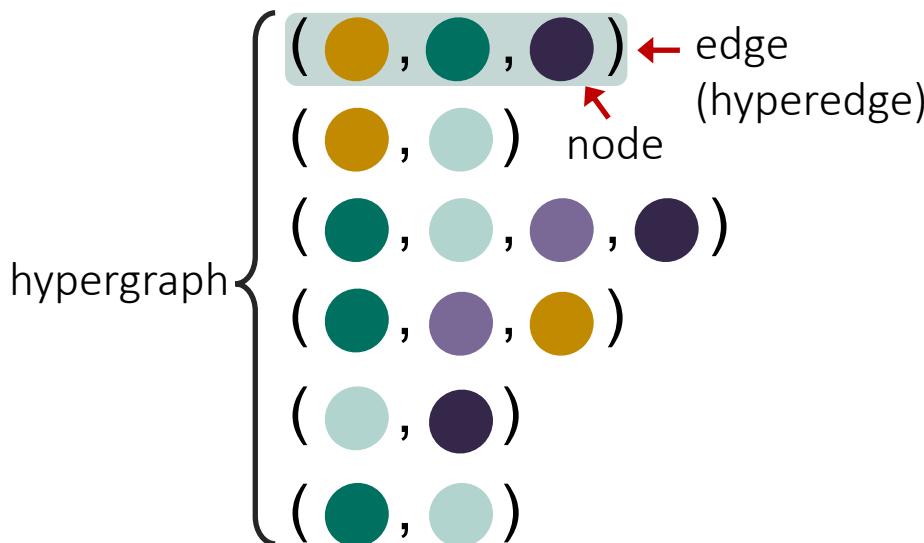
recipe 1--

recipe 2--

...

recipe 8--

# Hypergraph data—concepts



A hypergraph is a collection of sets.

## Electronic health records (EHR)

J44.9, J45.9, B44.9, O60, L50.5 ← edge = patient visit  
 J45.9, J46, O60, L50.5  
 ...  
 Z11.52, Z20.822, Z86.16, U07.1, J12.82

## Cooking recipes

, , , , ← edge = recipe  
, , ,   
 ...  
, , , , ,

# Hypergraph data – what can we learn from it?

Row = edge = recipe

Node = ingredient

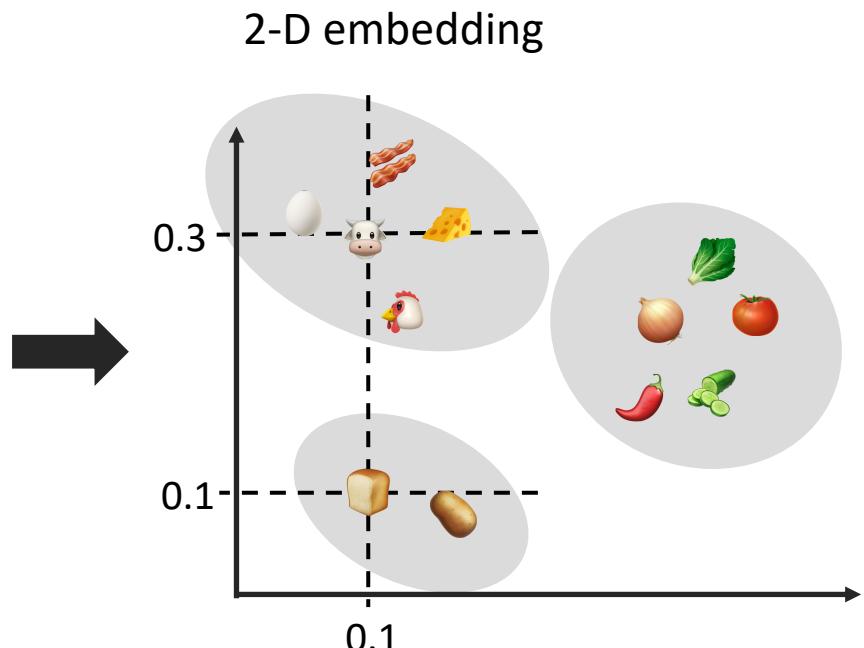
recipe 1 (bread, lettuce, bacon, cheese, tomato)

recipe 2 (tomato, potato, cow)

recipe 3 (cucumber, peanut, chicken, chili, onion)

recipe 4 (chili, chicken, garlic)

recipe 5 (tomato, egg)



- Node embedding, e.g.,  $\text{cow}=(0.1, 0.3)$ ,  $\text{bread}=(0.1, 0.1)$  enables clustering

# Hypergraph data – what can we learn from it?

Row = edge = patient visit

Node = medical code

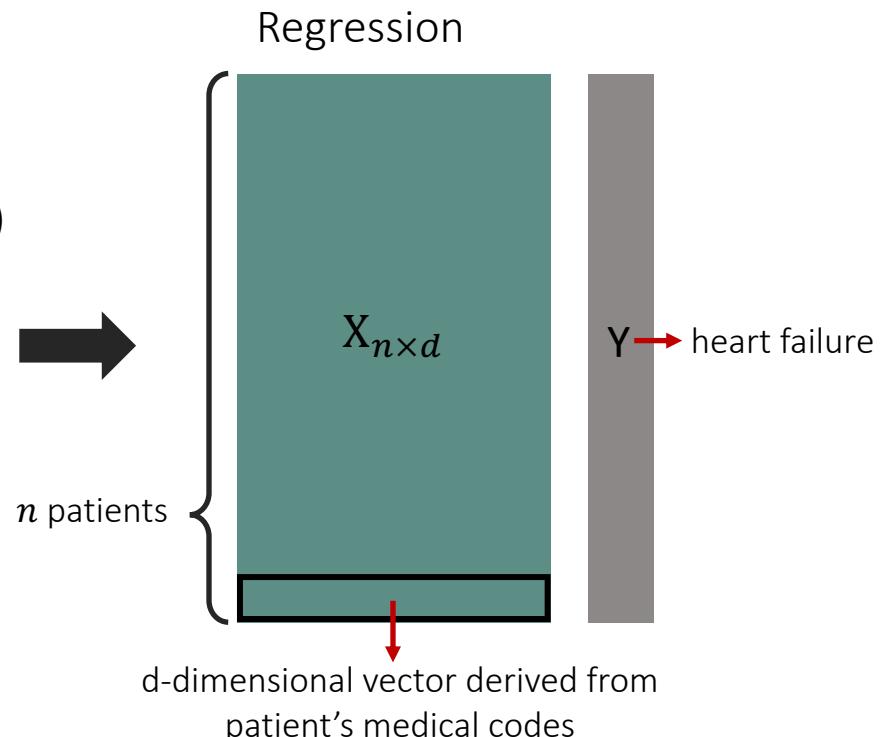
visit 1 (J44.9, J45.9, B44.9, O60, L50.5)

visit 2 (J45.9, J46, O60, L50.5)

visit 3 (Z11.52, Z20.822)

visit 4 (Z86.16, U07.1, J12.82)

visit 5 (O60, L50.5)



- Node embedding  
enables clustering, regression, can preserve privacy
- Edge prediction, e.g., (🐮, 🌶, 🧅, ?)

# Gap in literature on hypergraph modeling

In practice, hypergraph data are often projected to a network (i.e., pairwise relations), causing **information loss**. Direct modeling of hypergraph data is a relatively open area.

	e.g., to allow recipes to have different numbers of ingredients	e.g., to allow the same set of medical codes be observed in multiple patient visits	Model characteristic
	Edges with varying cardinality	Edge multiplicity	
Ghoshdastidar and Dukkipati (2014, 2015, 2017b); Chien et al. (2018); Kim et al. (2018); Ke et al. (2019)			Clustering of nodes
Lyu et al. (2021); Yuan and Qu (2021)			<b>Latent space model</b>
Stasi et al. (2014)	✓		$\beta$ -model
Zhang and McCullagh (2015)	✓		Hereditary hypergraph
Lunagómez et al. (2017)	✓		Random geometric graph
Ghoshdastidar and Dukkipati (2017a)	✓		Clustering of nodes
Turnbull et al. (2019)	✓		<b>Latent space model</b>
Zhen and Wang (2021)	✓		Clustering and <b>latent position</b> of nodes
Chodrow et al. (2021)	✓	✓	Clustering of nodes
Ng and Murphy (2021)	✓	✓	Clustering of hyperedges
<b>Yu and Zhu (2023 +)</b>	✓	✓	<b>Latent space model</b>

# Gap in literature on hypergraph modeling

In practice, hypergraph data are often projected to a network (i.e., pairwise relations), causing **information loss**. Direct modeling of hypergraph data is a relatively open area.

## Up next: the proposed model

- ✓ General types of hypergraphs
  - edges can have varying numbers of nodes
  - a given edge can appear more than once
- ✓ Node embedding
- ✓ Edge prediction

# Model — notation

## Notation

$\mathcal{V} = \{1, 2, \dots, n_v\}$  is the set of all nodes.

Each observed edge is a subset of  $\mathcal{V}$ .

$e_1, \dots, e_{n_e}$  denote all edges.

Thus, there are  $n_v$  nodes and  $n_e$  edges.

$$\mathcal{V} = \{ \textcolor{orange}{1}, \textcolor{teal}{2}, \textcolor{purple}{3}, \textcolor{lightblue}{4}, \textcolor{violet}{5} \}$$

$$e_1 = (\textcolor{orange}{1}, \textcolor{teal}{2}, \textcolor{purple}{3})$$

$$e_2 = (\textcolor{orange}{1}, \textcolor{lightblue}{4})$$

$$e_3 = (\textcolor{teal}{2}, \textcolor{lightblue}{4}, \textcolor{purple}{3}, \textcolor{purple}{5})$$

$$e_4 = (\textcolor{teal}{2}, \textcolor{purple}{3}, \textcolor{orange}{1})$$

$$e_5 = (\textcolor{lightblue}{4}, \textcolor{purple}{3})$$

$$e_6 = (\textcolor{teal}{2}, \textcolor{lightblue}{4})$$

$$n_v = 5, n_e = 6$$

# Model — motivations from real-world observations

## Diversity within each edge

- Nodes in an edge often complement each other,  
*e.g., different expertise in a collaboration.*
- Diversity also appears when selections are made to prevent redundancy,  
*e.g., products of various categories in a shopping order.*

## Heterogeneous node popularity

- Different nodes appear with very different frequencies.

# Model – how to encourage diversity and heterogeneity?

## Diversity

- Each node is represented by a vector of latent features
  - An edge contains multiple nodes
  - $\text{Prob}[\text{observing an edge}]$  is large when the corresponding set of vectors have different “directions”



## Heterogeneous popularity

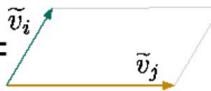
- Each node is associated with a popularity parameter
  - $\text{Prob}[\text{observing an edge}]$  is large when nodes in the edge have large popularity parameters

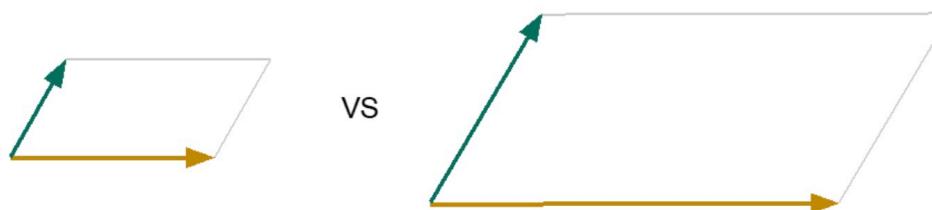


# Model — how to encourage diversity and heterogeneity?

$$\tilde{v}_i = \left( \underbrace{v_{i1}, \dots, v_{id}}_{\text{latent feature vector}}, \underbrace{0, \dots, 0, a_i, 0, \dots 0}_{\text{popularity parameter}} \right) \text{ for node } i$$

How to use  $\tilde{v}_i$ ?

- Let  $E$  denote a random edge.  $P(E = \{i, j\}) \propto \text{area}^2(\tilde{v}_i, \tilde{v}_j) = \frac{\tilde{v}_i}{\tilde{v}_j}$
- $P(E = \{i, j\})$  is large when  $\tilde{v}_i, \tilde{v}_j$  are long
- 



# Model — how to encourage diversity and heterogeneity?

latent feature vector      popularity parameter  
 $\tilde{v}_i = (\overbrace{v_{i1}, \dots, v_{id}}^{\text{latent feature vector}}, \underbrace{0, \dots, 0, a_i, 0, \dots 0}_{\text{popularity parameter}})$  for node  $i$

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- $P(E = \{i, j\})$  is large when  $\tilde{v}_i, \tilde{v}_j$  are long and have separated directions, i.e., as close to orthogonal as possible.



vs



# Model – how to encourage diversity and heterogeneity?

$$\tilde{v}_i = (\underbrace{v_{i1}, \dots, v_{id}}_{\text{latent feature vector}}, \underbrace{0, \dots, 0, a_i, 0, \dots 0}_{\text{popularity parameter}}) \text{ for node } i$$

live on a sphere      pairwise orthogonal

How to use  $\tilde{v}_i$ ?

- Let  $E$  denote a random edge.  $P(E = \{i, j\}) \propto \text{area}^2(\tilde{v}_i, \tilde{v}_j) = \frac{1}{2} \|\tilde{v}_i - \tilde{v}_j\|_2^2$   
 $P(E = \{i, j\})$  is large when  $\tilde{v}_i, \tilde{v}_j$  are long and have separate directions, i.e., as close to orthogonal as possible.
- $\|\tilde{v}_i\|_2$  is constant across  $i$ ,  $a_i > 0$  is the  $i$ th entry of  $(0, \dots, 0, a_i, 0, \dots 0)_{1 \times n_v}$
- The lengths are driven by the popularity parameters  $a_i, a_j$
- The separation is mainly driven by the ‘diversity’ of the feature vectors  $v_i, v_j$

$$P(E = \{i, j, k\}) \propto \text{volume}^2(\tilde{v}_i, \tilde{v}_j, \tilde{v}_k) = \frac{1}{6} \|\tilde{v}_i - \tilde{v}_j\|_2^2 \|\tilde{v}_i - \tilde{v}_k\|_2^2 \|\tilde{v}_j - \tilde{v}_k\|_2^2$$


# Model — nice properties

The random edge  $E$  can be any subset  $e$  of  $\mathcal{V}$ , e.g.,  $e = \{1,3\}, e = \{1,3,5,7\}$ .

$$P(E = e) = \frac{\text{volume}^2(\tilde{v}_i | i \in e)}{\sum_{e' \subset \mathcal{V}} \text{volume}^2(\tilde{v}_i | i \in e')}$$

Observed edges  $e_1, \dots, e_{n_e}$  are i.i.d realizations of  $P$ .

## Nice properties

- Explicit formulas for marginal probability & conditional probability
  - $P(i \in E)$ , e.g.,  $P(\text{●} \in E)$
  - $P(e \subset E)$ , e.g.,  $P((\text{●}, \text{●}, \text{●}) \subset E)$
  - $P(E = e' | e \subset E)$ , e.g.,  $P(E = (\text{●}, \text{●}, \text{●}) | (\text{●}) \subset E)$
- Easy-to-apply sampling algorithm
- Distribution of  $|E|$  has an explicit characterization

# Method

## Model fitting

- Maximum likelihood estimation (MLE)

$$\arg \max_{\substack{v_i \in \mathbb{R}^d, a_i > 0, \|v_i\|_2 \\ \text{is constant across } i}} -\log \underbrace{\sum_{e' \subset \mathcal{V}} \text{volume}^2(\tilde{v}_i \mid i \in e')}_{\text{determinant of a matrix defined by } v_i, a_i \text{ for all } i} + \frac{1}{n_e} \sum_{\ell=1}^{n_e} \underbrace{\log \text{volume}^2(\tilde{v}_i \mid i \in e_\ell)}_{\text{determinant of a matrix defined by } v_i, a_i \text{ for } i \text{ in } e_\ell}$$

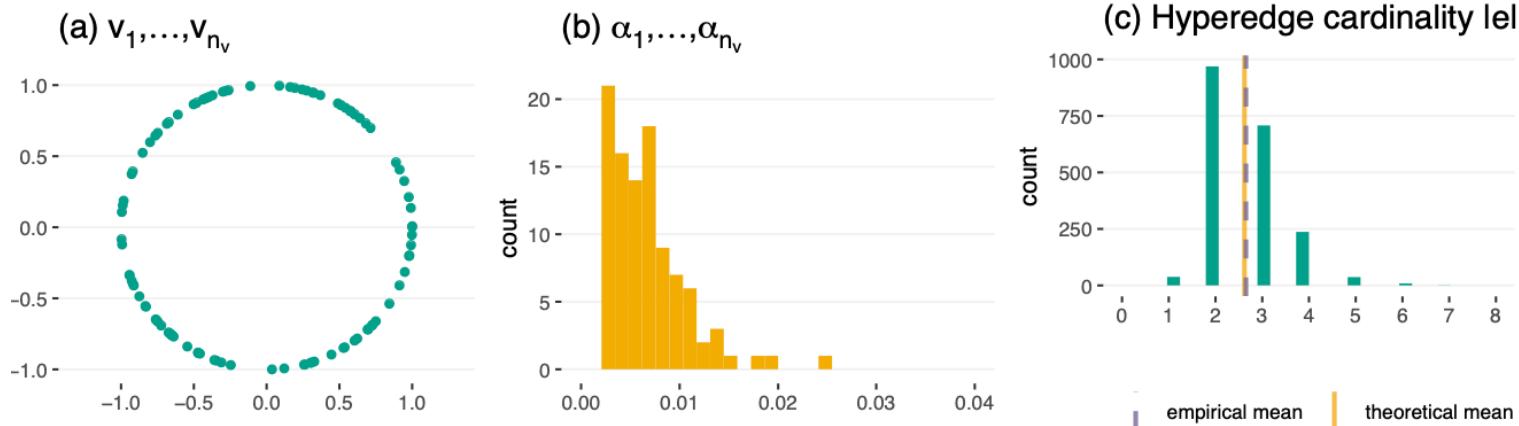
- Gradient descent algorithm

- Accelerated projected gradient methods for nonconvex programming <sup>†</sup>
- Mini batch gradient descent, i.e., using a small sample of edges in each iteration, and Adam adaptive learning rate

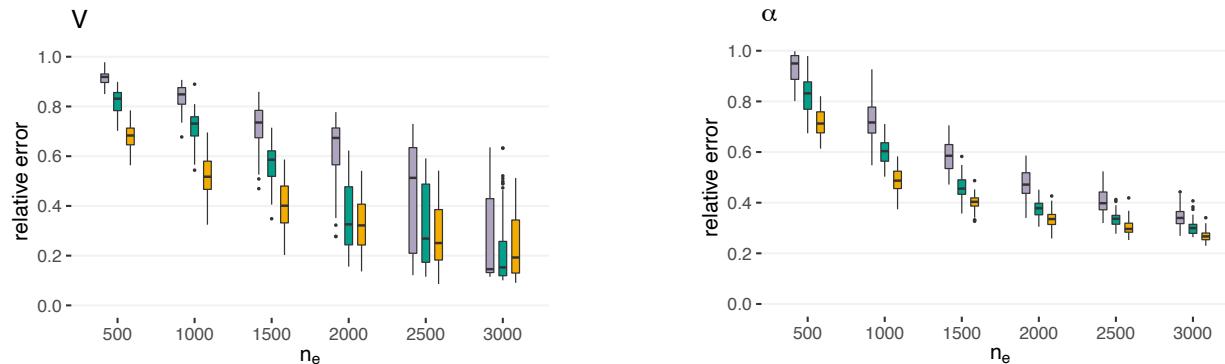
<sup>†</sup> Li & Lin (2015) NeurIPS

# Simulations (consistency)

Model setting:  $n_v=100, d = 2, n_e = 2000$



Results of 50 repeated simulations:  $n_v=100, d = 2, 3, 4$



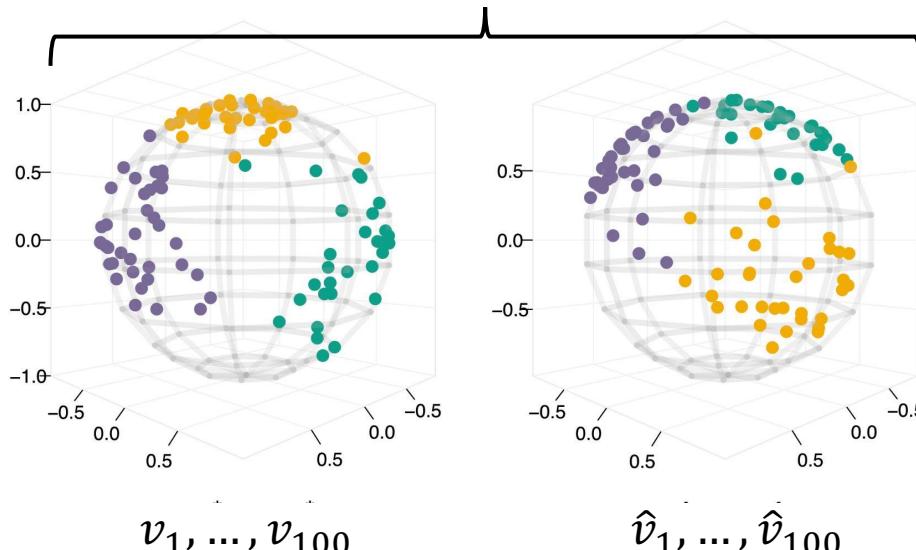
$$\dagger \quad \alpha_i = a_i^2, \alpha = (\alpha_1, \dots, \alpha_{n_v}); V_{\cdot i} = \frac{v_i}{\|v_i\|_2}.$$

# Simulation (clustering performance)

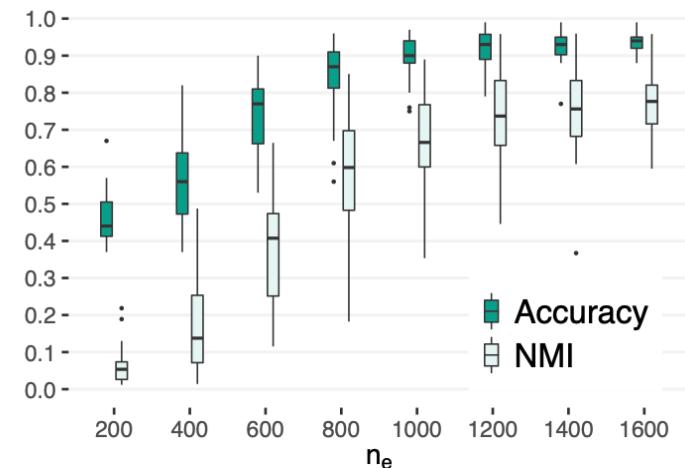
## Evaluating clustering performance

- $n_v = 100$  nodes are assigned to three even clusters.
- $v_i$ 's are on the unit sphere in  $\mathbb{R}^3$  and are generated by three von Mises–Fisher distributions.

color = true cluster label;  $n_e = 2000$



Clustering results



Clustering result as  $n_e$  varies

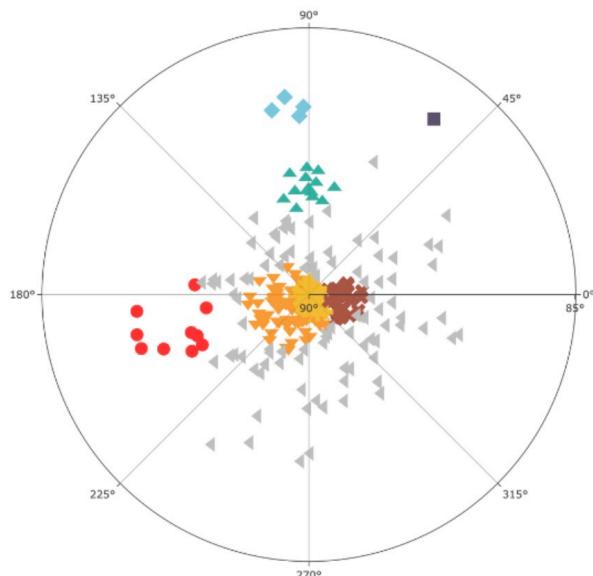
\* NMI = normalized mutual information

## Numerical results — recipe data

Recipes on Yummly.com

$n_e = 2673$  recipes involving  $n_v = 906$  ingredients; fit a model with  $d = 3$

Estimated latent vectors  $\hat{v}_i$  (embeddings)



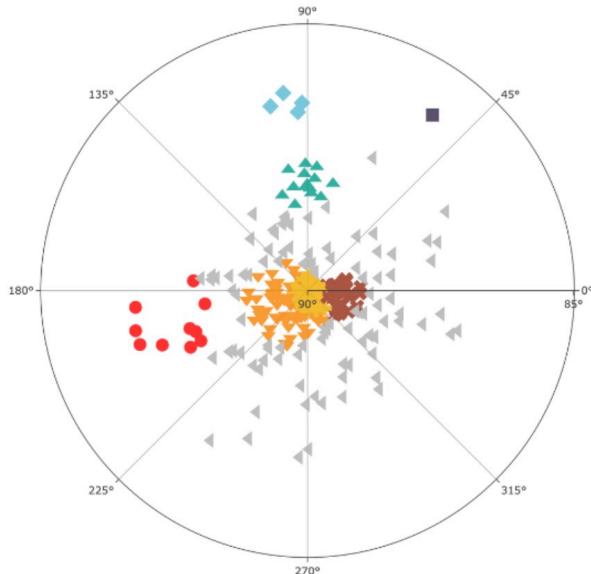
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Applications of the fitted model

- Clustering ingredients using embedding  $\hat{v}_i$ 's, which lie on a sphere in  $\mathbb{R}^3$



Clustering method: fit a mixture of von Mises–Fisher model (i.e., analogy to Gaussian on a sphere) on embeddings of ingredients that appear in 10+ recipes (298 ingredients in total)

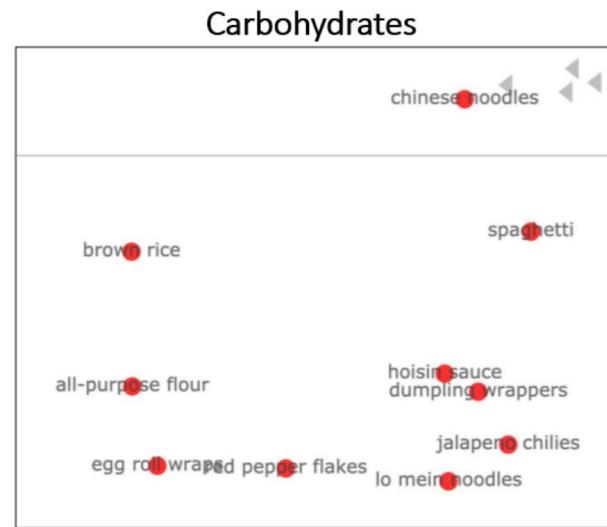
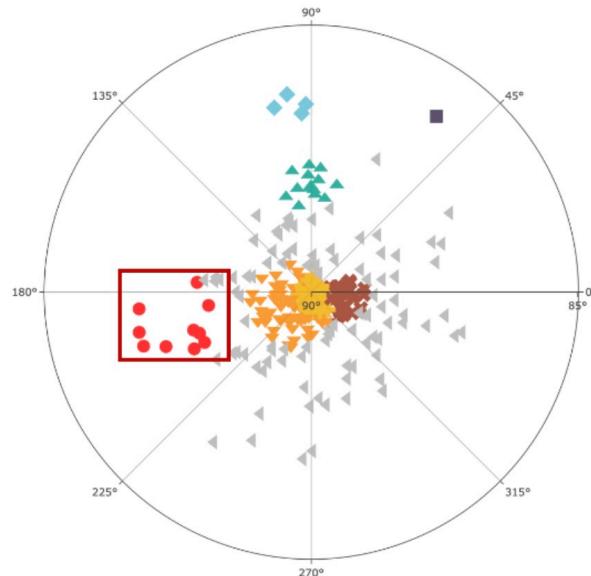
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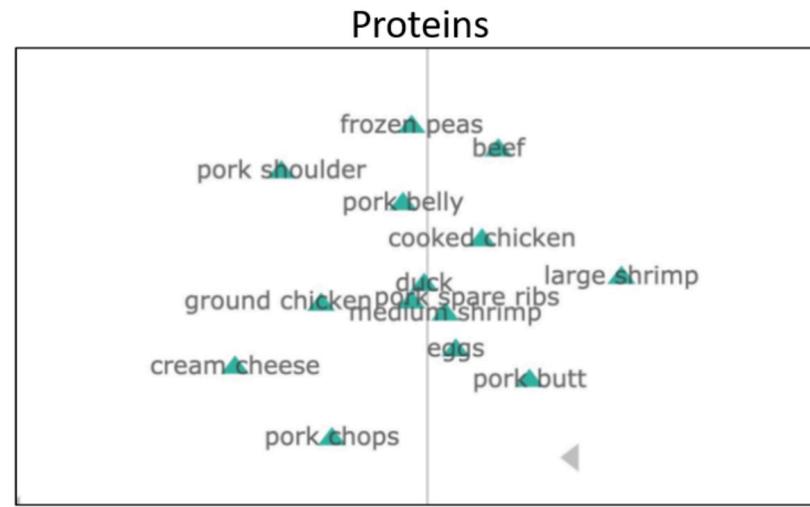
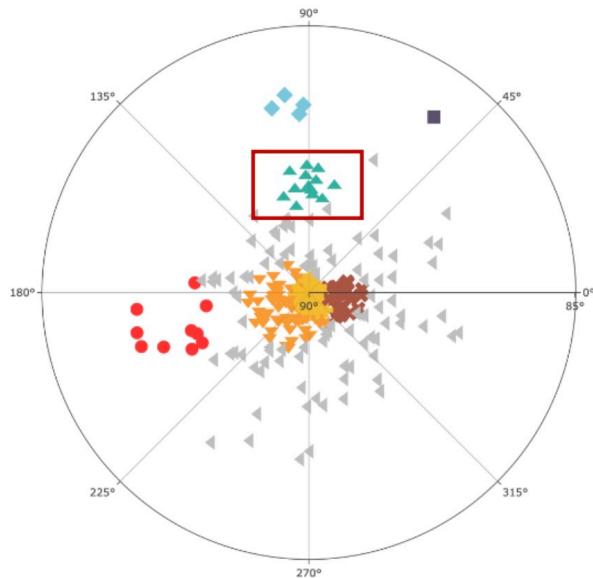
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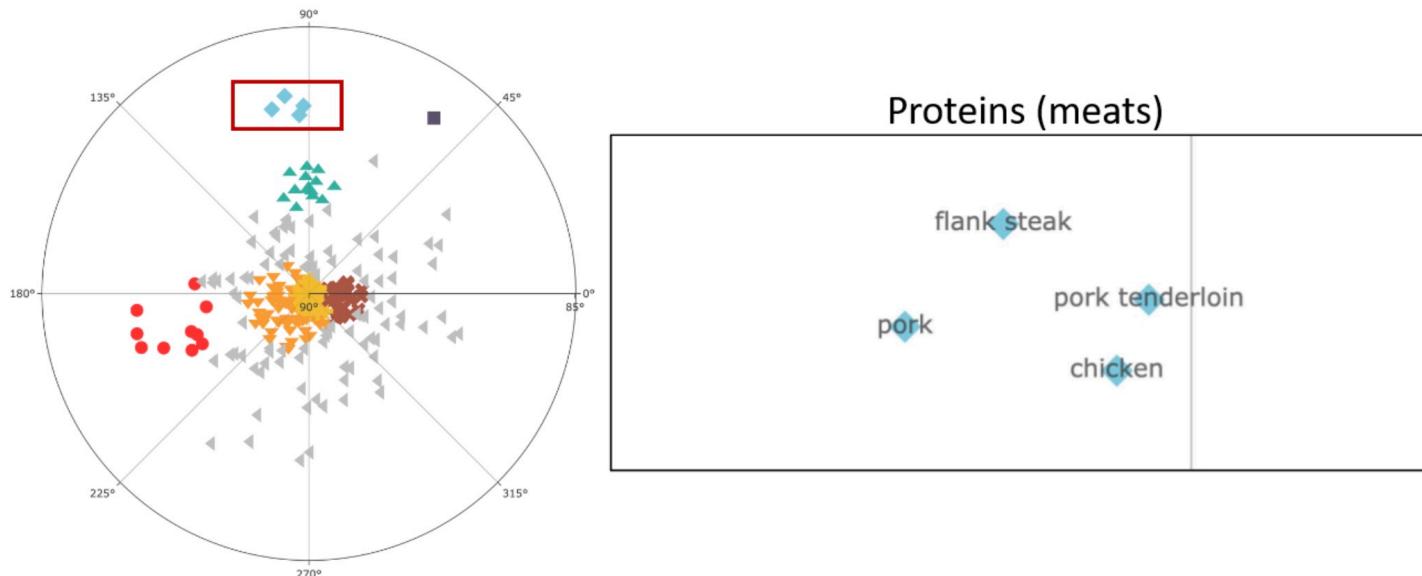
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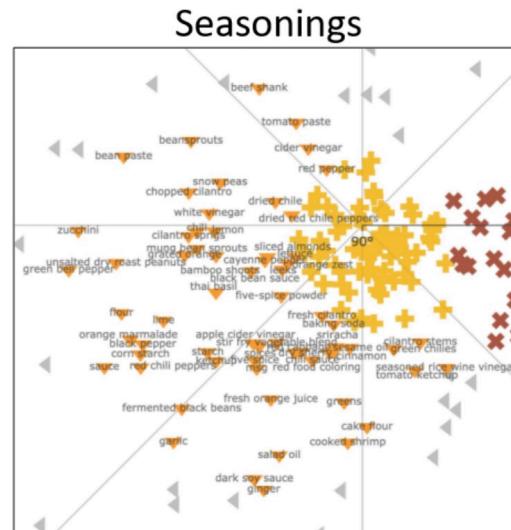
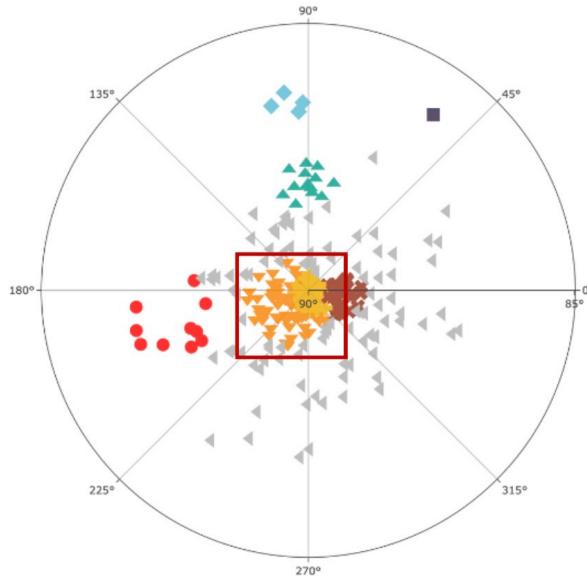
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# Numerical results — recipe data

## Applications of the fitted model

- Completing recipes
- e.g., (, , , , , ?)

```
> selected=c("pork belly", "green bell pepper", "cooking oil", "soy sauce", "sugar")
> recommend_one_ingredient(L_hat, selected, ingredients)
```

Knowing that a recipe has (PORK BELLY, GREEN BELL PEPPER, COOKING OIL, SOY SAUCE, SUGAR), the one additional ingredient that is most likely to be in this recipe is GARLIC.

$$\arg \max_{i \notin e} \hat{P}(i \in E | e \subset E)$$

$\hat{P}$  is parameterized by  
 $\hat{\nu}_i$  and  $\hat{a}_i$  for all  $i$

The probability that ingredient  $i$  is in a recipe, given that ingredients in the selected set  $e$  are in the recipe

# Theory — consistency

## Proposition (Identifiability)

If  $n_v > 2d$ , then, given any fixed model,  $a = (a_1, \dots, a_{n_v})$  is identifiable and  $v_1, \dots, v_{n_v}$  are identifiable up to a shared rotation and individual sign flips (i.e., multiplication with  $\pm 1$ ).

## Theorem 1 (Consistency)

If  $n_v > 2d$  and  $\{v_1, \dots, v_{n_v}\}$  span  $\mathbb{R}^d$ , then, as  $n_e \rightarrow \infty$ , with proper rotation and sign flips of  $\hat{v}_i$

$$\sum_{i=1}^{n_v} \|\hat{v}_i - v_i\|_2 \xrightarrow{p} 0,$$

$$\|\hat{a} - a\|_2 \xrightarrow{p} 0.$$

# Theory — asymptotic distribution

## Parameterize the model using matrix $L$

- The proposed model is a special determinantal point process (DPP)
- A DPP is defined using a matrix  $L$  ( $L$  can be any semi-definite matrix)
- \* For the proposed model,  $L = (v_i^T v_j)_{i,j=1}^{n_v} + \text{diag}(a_1^2, \dots, a_{n_v}^2)$

### Theorem 2 (Asymptotic normality)

If  $n_v > 2d$ ,  $\{v_1, \dots, v_{n_v}\}$  span  $\mathbb{R}^d$  and can not be divided into multiple groups that are mutually orthogonal, then, as  $n_e \rightarrow \infty$ , with proper sign flips of  $\hat{v}_i$ ,  $\|\hat{L} - L\|_F \rightarrow 0$  in probability and

$$\sqrt{n_e} \cdot \text{vec}(\hat{L} - L) \xrightarrow{\text{dist}} N(\mathbf{0}, \Sigma).$$

Here  $\Sigma$  is a matrix that we have derived which is defined by  $v_i$ ,  $a_i$ ,  $i \in \mathcal{V}$ .

# Theory — asymptotic result (challenges)

- This asymptotic result is one regarding **constrained** M-estimation, since the **parameter space** of  $L$  is **special**.
- The theoretical development requires non-trivial analysis of the local geometry of this parameter space, where we applied **recent results in variational geometry**.
- This is the **first** asymptotic result, to our knowledge, for **structured determinantal point processes**.

# Take home messages

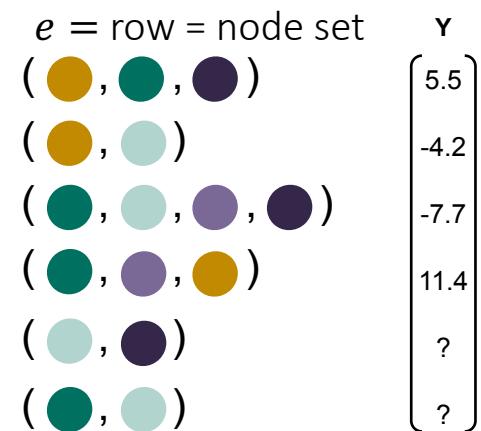
- The proposed model is the **first** hypergraph model that
  - considers **diversity**
  - enables **embedding** while allowing **edges** to have different **numbers of nodes** and to **appear more than once** in data
- The model can be applied for
  - node **embedding**
  - node **clustering**
  - **edge prediction**
- We have established the **consistency** and **asymptotic normality** of the estimates of model parameters.

[Ornes, Stephen. "How Big Data Carried Graph Theory Into New Dimensions." Quanta Magazine \(Aug 2021\).](#)

# Planned future work on hypergraph data

## Develop regression model on node set

- Consider  $y = f(e)$  for  $e \subset \mathcal{V}$  and estimate  $f$
- e.g.,  $\hat{f}$  predicts quality of collaboration for a given team, future medical needs given current medical codes

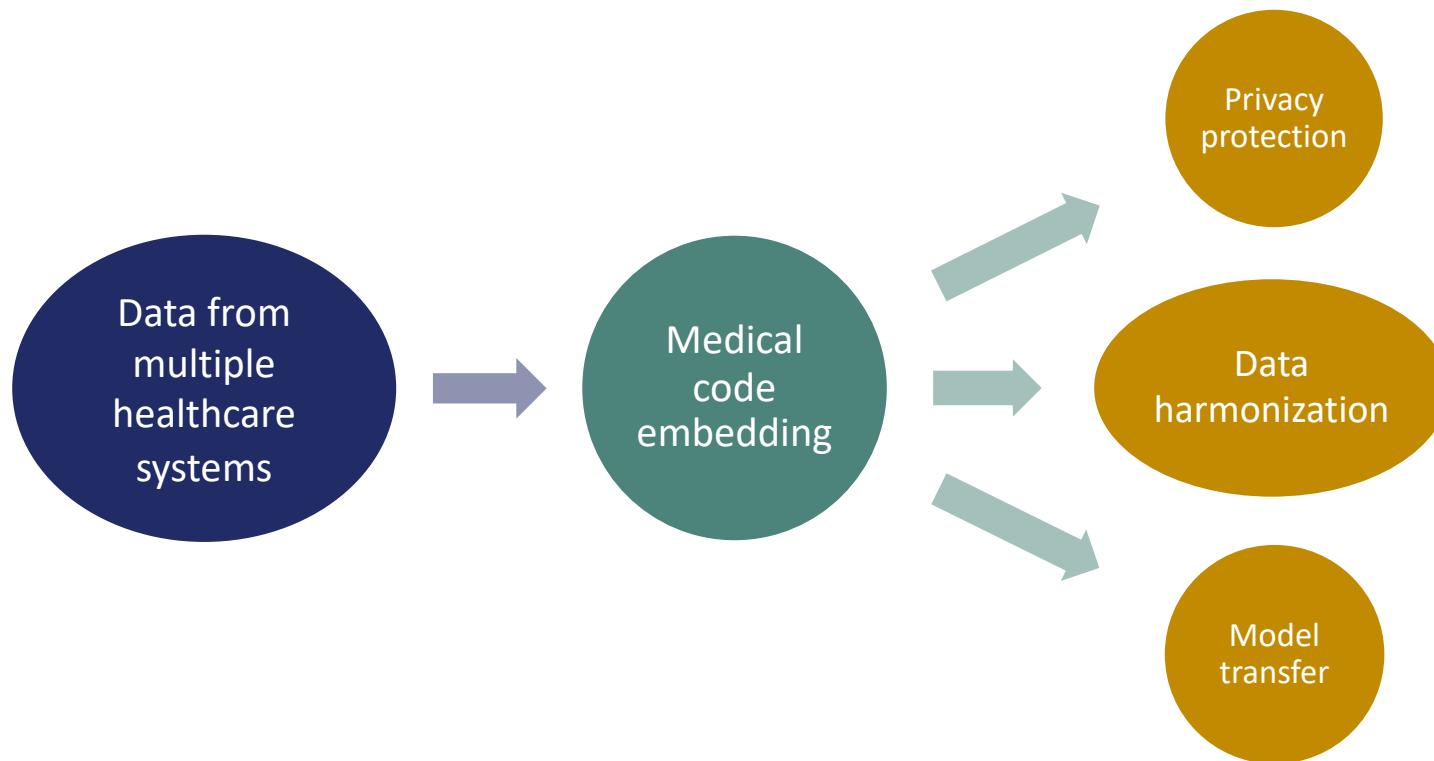


## Extend the hypergraph model to incorporate

- observed covariates of nodes
- more complex mechanism beside diversity and popularity
- information on nodes' different roles within edges

# Planned future work on hypergraph data

## Application to distributed EHR data network



Thank you

# References

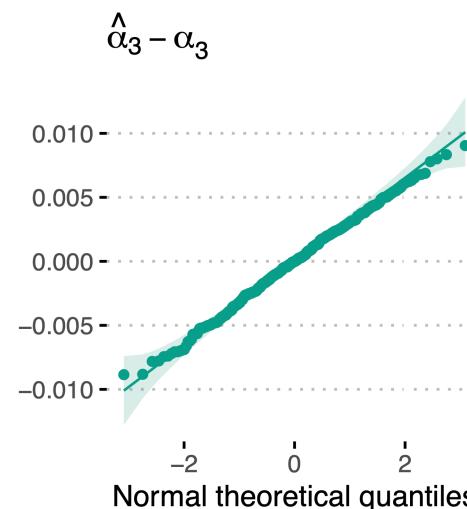
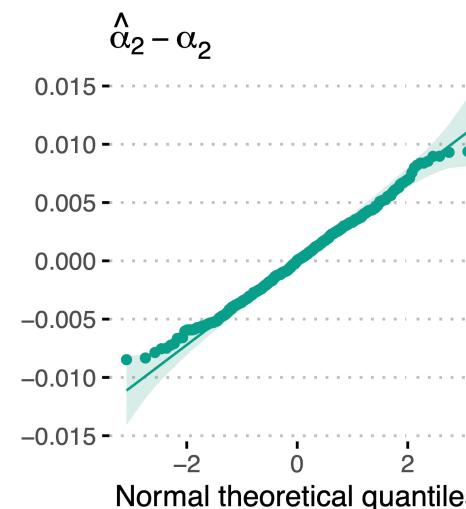
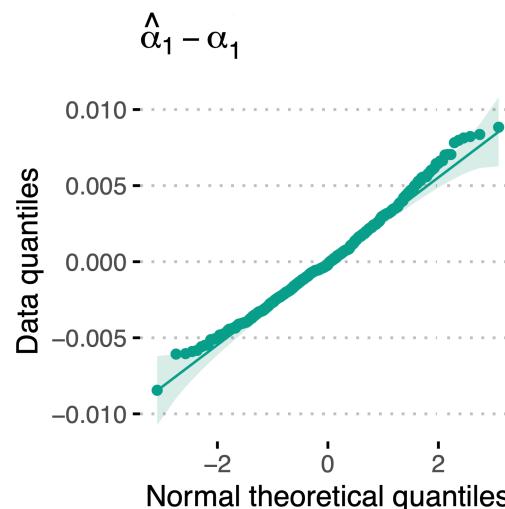
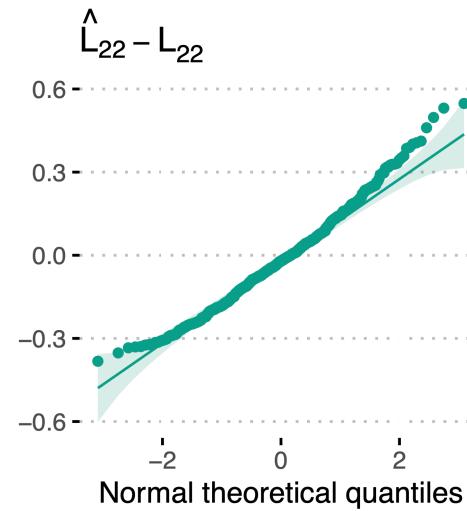
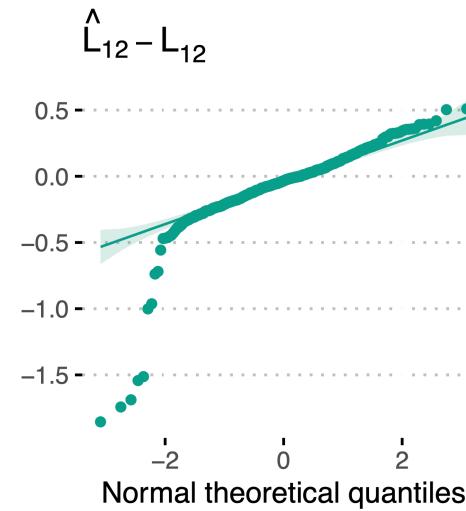
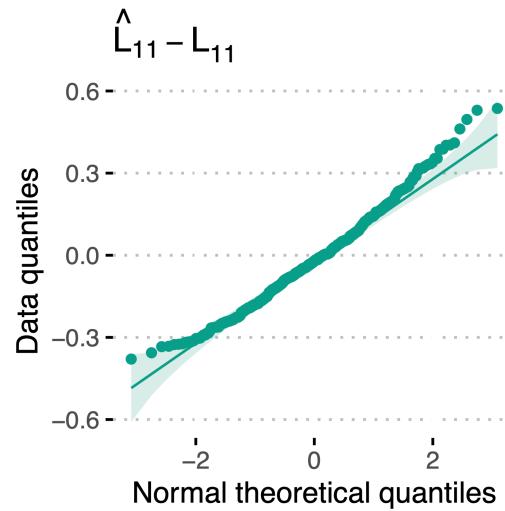
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# Appendix – simulations (asymptotic normality)

Results of 500 repeated simulations:  $n_v=100, d = 2, n_e = 3000$



# Appendix – simulations (asymptotic normality)

Results of 500 repeated simulations:  $n_v=100, d = 2, n_e = 3000$

