

# Non-Gaussian Factor Model for Interpretable Ordinal Time Series Analysis

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# Research overview (Scott)

Develop computationally efficient, adaptive methodology for analyzing large, complex time series data and producing optimal summary measures.

## Complexity:

- Nonstationarity
- Covariate-dependent
- High-dimensionality
- Replicated data
- Spatio-temporal data

## Themes:

- Frequency and time domain
- Nonparametric
- Learning tasks
- Bayesian computation
- Theoretical support

**Transdisciplinary Collaboration:** Sleep research, kinesiology, neuroscience, climate science.

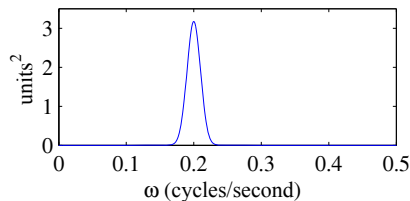
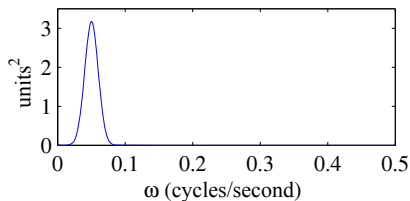
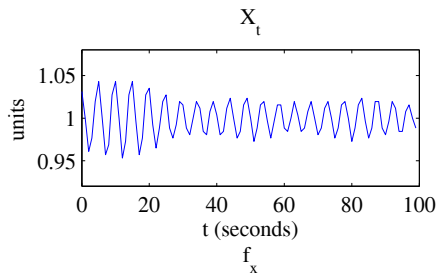
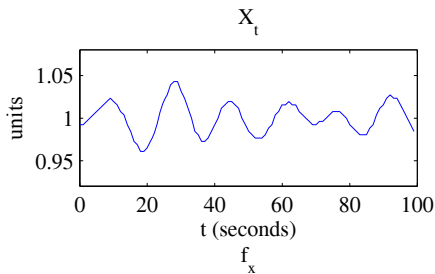
# Frequency domain approach

- Consider a **stationary** univariate time series  $X_t$ .
- Cramér Representation:

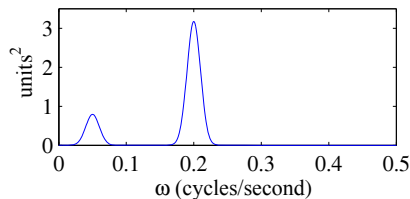
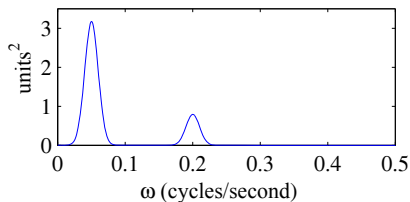
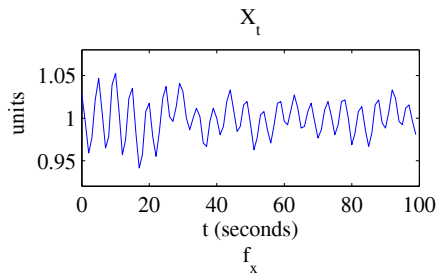
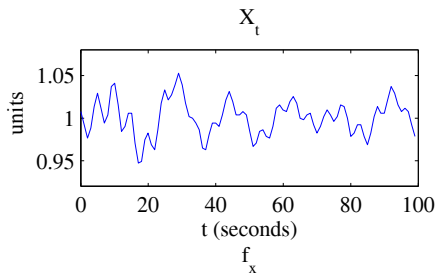
$$X_t = \int_{-1/2}^{1/2} A(\omega) \exp(2\pi i \omega t) dZ(\omega).$$

- **Power spectrum**:  $f(\omega) = |A(\omega)|^2$ .
- Type of **frequency ANOVA**:  $\text{var}(X_t) = \int_{-1/2}^{1/2} f(\omega) d\omega$ .
- Interpretation: **variability** of  $X_t$  attributable to periodic signals at **frequency**  $\omega \in \mathbb{R}$ .
- Can be extended to accommodate changes across time and/or covariate.

# Two simulated examples



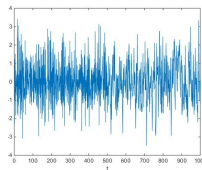
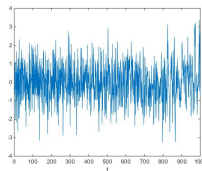
# Two more simulated examples



# Adaptive Bayesian spectral analysis

**Goal:** Develop **flexible** methodology for adaptively estimating complex frequency domain **structure and dependencies**.

- Capture **smooth and abrupt** changes over different dimensions.
- **Adaptive partitioning** via reversible-jump Markov chain Monte Carlo.
- General framework for modeling **nonstationary and covariate-dependent** power spectra\*.

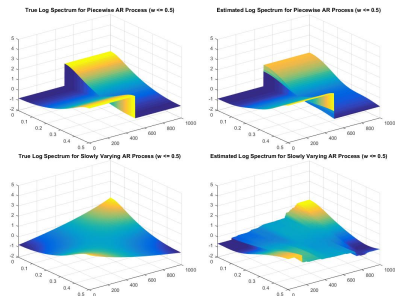


\*Scott A. Bruce, Martica H. Hall, Daniel Buysse, and Robert T. Krafty (2018), Conditional adaptive Bayesian spectral analysis of nonstationary biomedical time series. *Biometrics*, 74:260-269.

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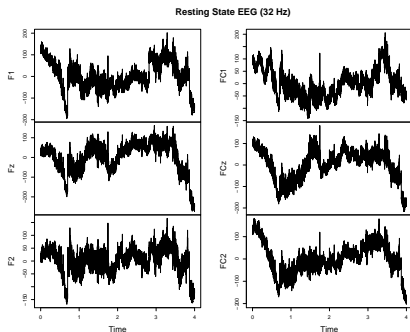


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**What's new:** Rebecca Lee developing hierarchical extension of this work.

# Adaptive frequency band learning

**Goal:** Develop a **quantitative** framework for identifying frequency bands that **optimally** preserve characteristics of power spectra.



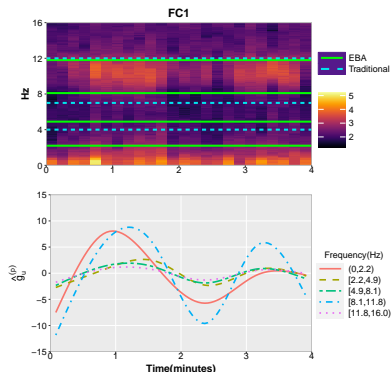
- Traditional frequency bands used in practice are often **subjectively** determined.
- Offer **consistent** estimation of frequency partition points\*.
- Preserve **important** features:
  - time-varying dynamics
  - variability across replicates
  - discriminative power
  - cluster structure
  - local extrema

\*Scott A. Bruce, Cheng Yong Tang, Martica H. Hall, and Robert T. Krafty (2020) Empirical Frequency Band Analysis of Nonstationary Time Series, *Journal of the American Statistical Association*, 115(532):1933-1945.



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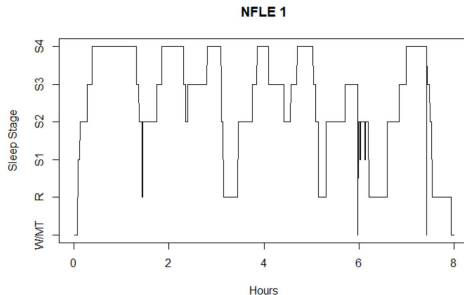
\*Scott A. Bruce, Cheng Yong Tang, Martica H. Hall, and Robert T. Krafty (2020) Empirical Frequency Band Analysis of Nonstationary Time Series, *Journal of the American Statistical Association*, 115(532):1933-1945.

**What's new:** Connor Brubaker developing extension for multiple time series.

# Categorical time series learning

**Goal:** Develop frequency-domain framework for supervised and unsupervised learning for **categorical** time series data.

- Motivation: studying **sleep stage** time series for different sleep disorders.
- Idea: use **spectral envelope** for feature-based learning.
- Offer **consistent** classification using frequency domain structure\*.



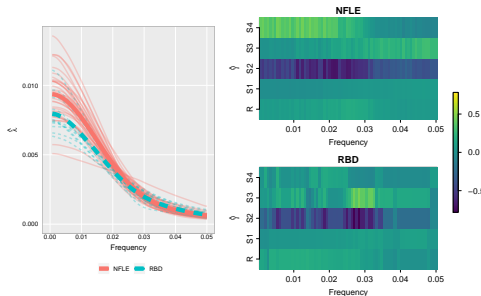
**NFLE:** Nocturnal Frontal Lobe Epilepsy, **RBD:** REM Behavior Disorder

\*Zeda Li, Scott A. Bruce, and Tian Cai. (2022) "Classification of Categorical Time Series Using the Spectral Envelope and Optimal Scalings". Journal of Machine Learning Research. 23(299):1-31.  
<http://jmlr.org/papers/v23/21-0369.html>

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**What's new:** Developing extension for nonstationary time series (interested students may contact me).

# Identifying Granger Causal Structures and Stationarity Regions via Deep Learning

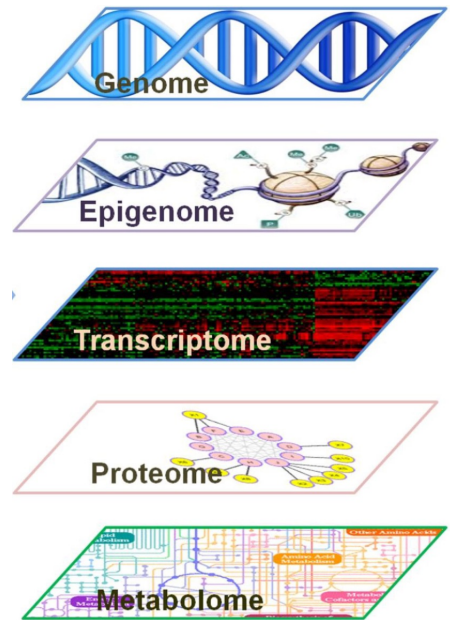
More to come...

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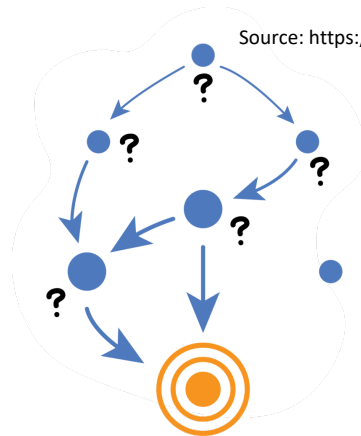
More to come...

**What's new:** Eric Gao testing feasibility and developing architecture.

# Research Overview



# Research Interests



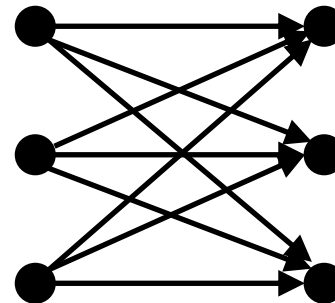
Source: <https://www.inguo.io/>

## Methods

Causal Discovery

Clustering and Bayesian Nonparametrics

Graphical Models

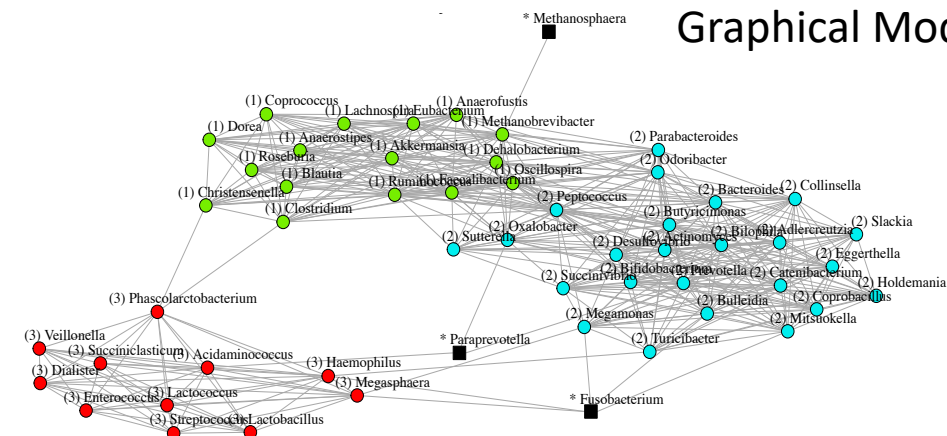


## Applications

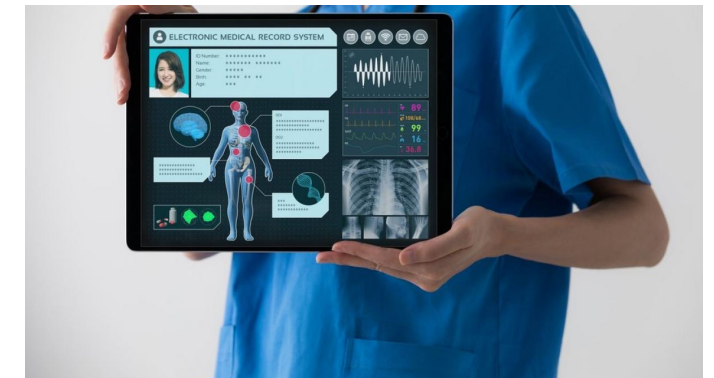
Single-Cell Multi-Omics

Microbiome Multi-Omics

Digital Health



Source: aamc.org



# Causal Discovery



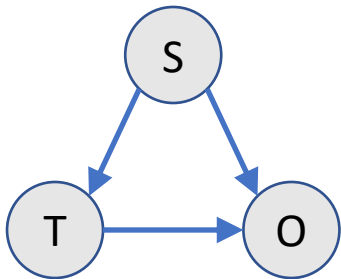
# Simpson's Paradox

- Kidney stone treatment

	Treatment A	Treatment B
	78%(273/350)	83%(289/350)

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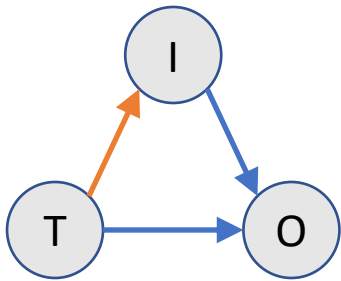
- Kidney stone treatment



	Treatment A	Treatment B
Small Stones	93%(81/87)	87%(234/270)
Large Stones	73%(192/263)	69%(55/80)
Total	78%(273/350)	83%(289/350)

# Simpson's Paradox

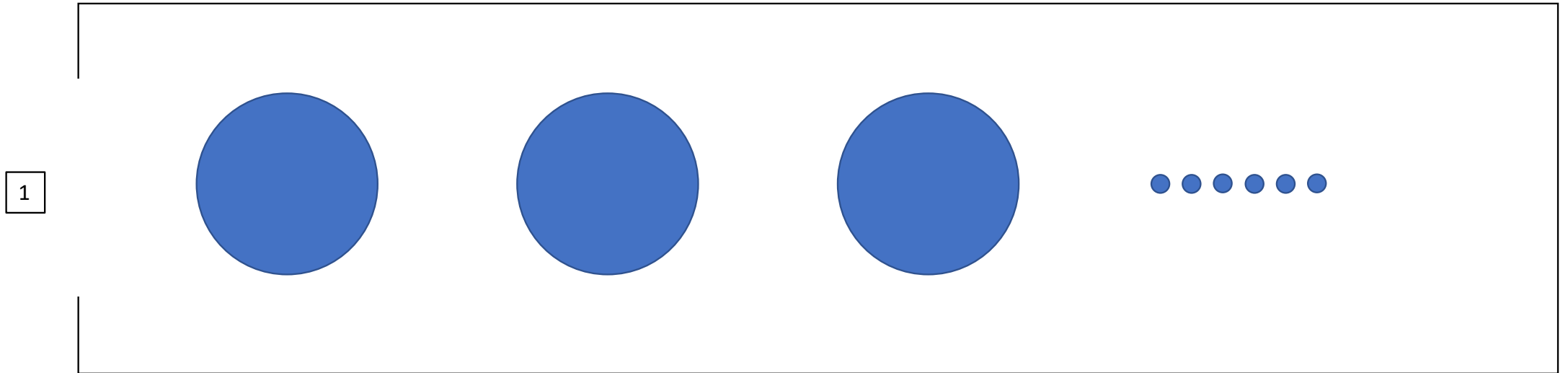
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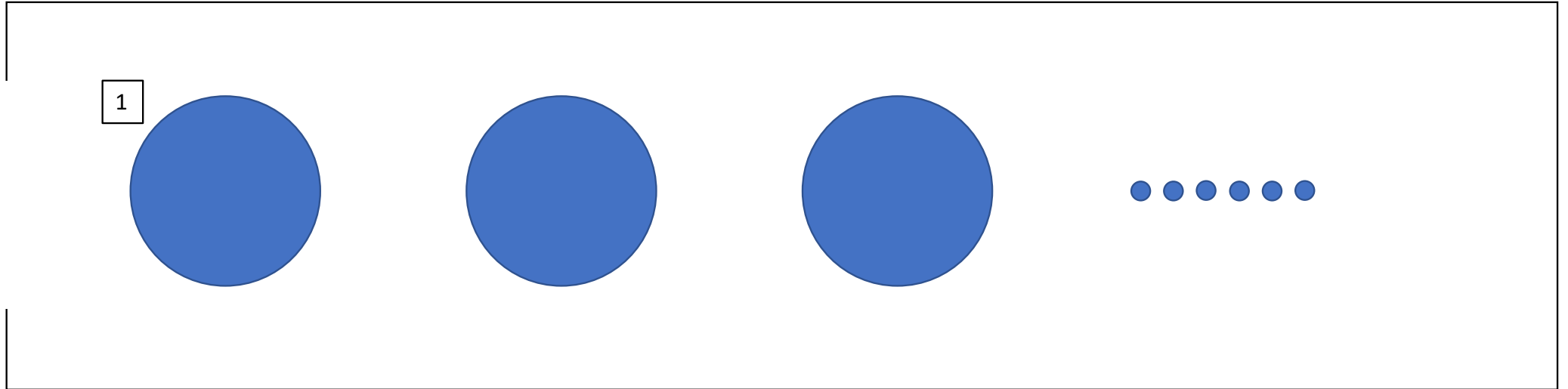
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Non-Infection	93%(81/87)	87%(234/270)
Infection	73%(192/263)	69%(55/80)
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# Bayesian Nonparametrics

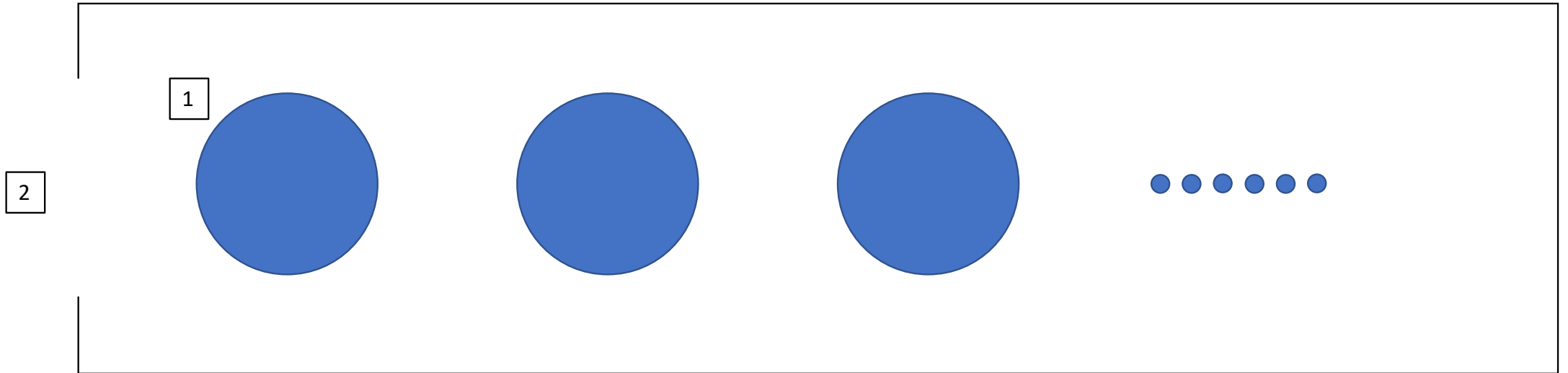
# Chinese Restaurant Process



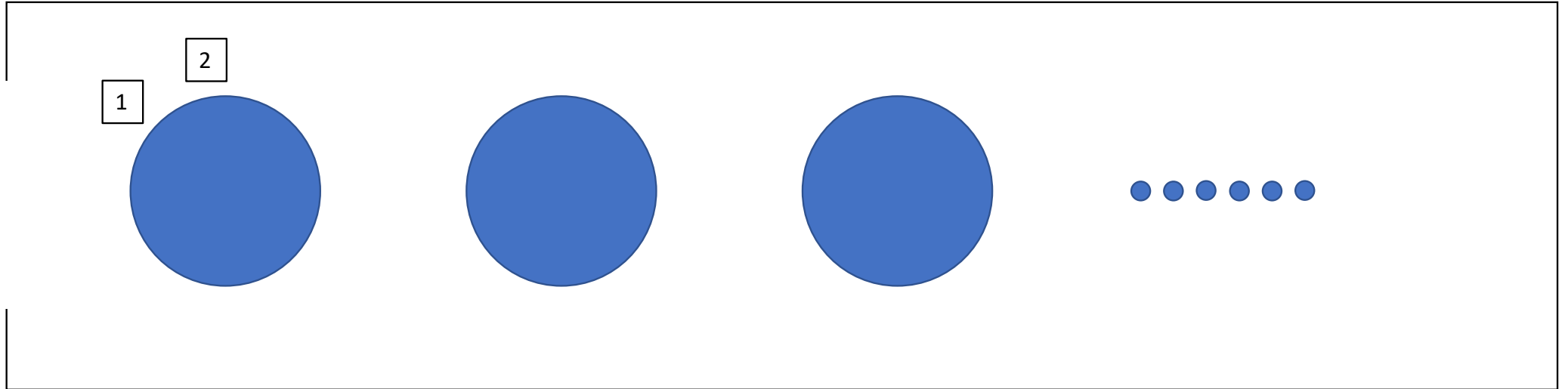
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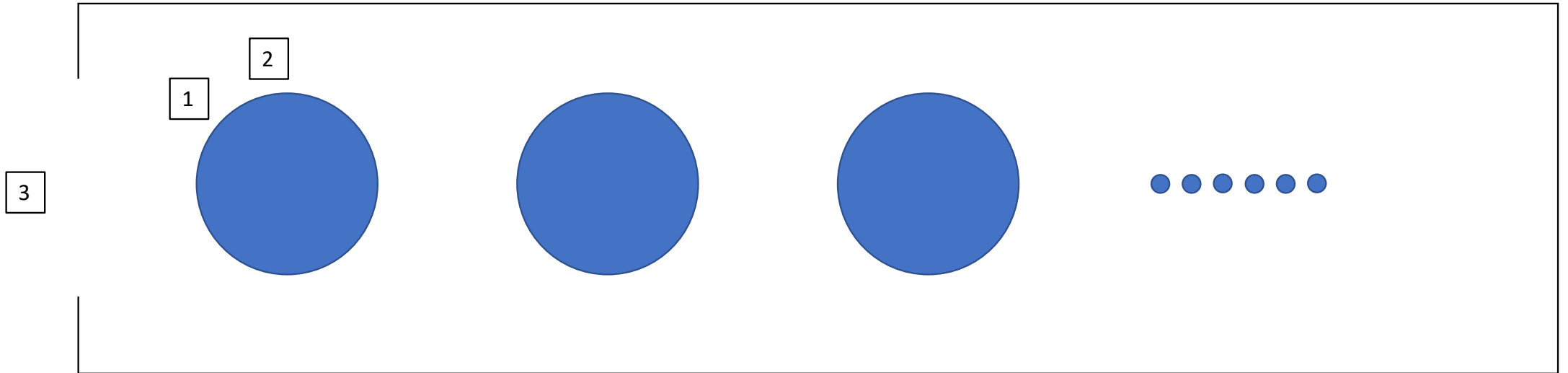


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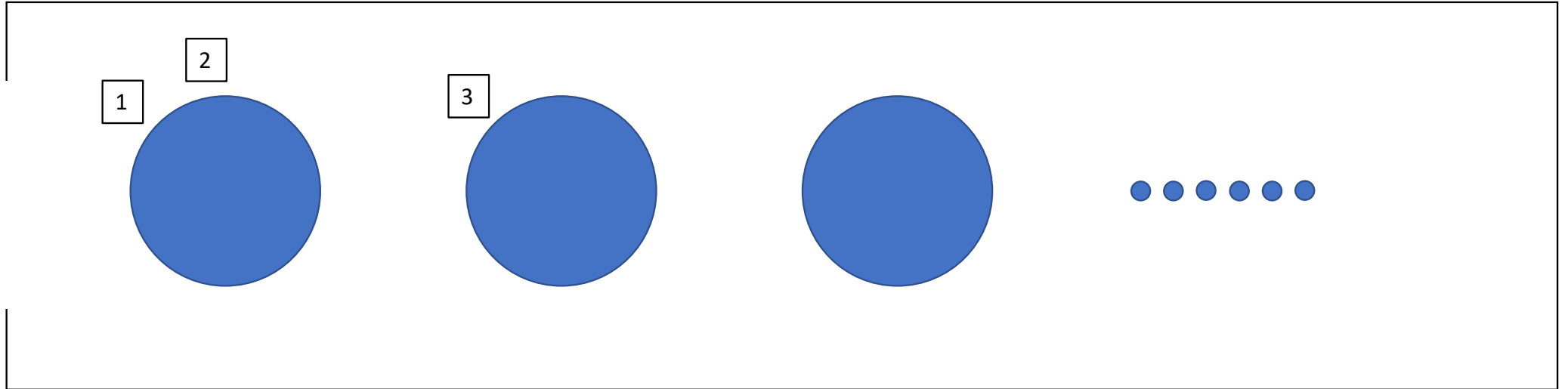




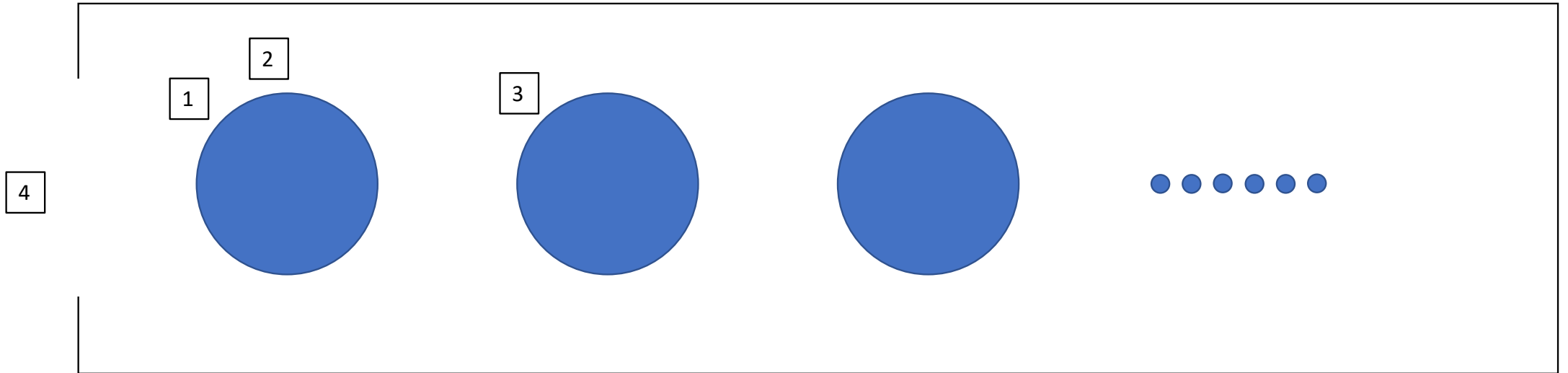
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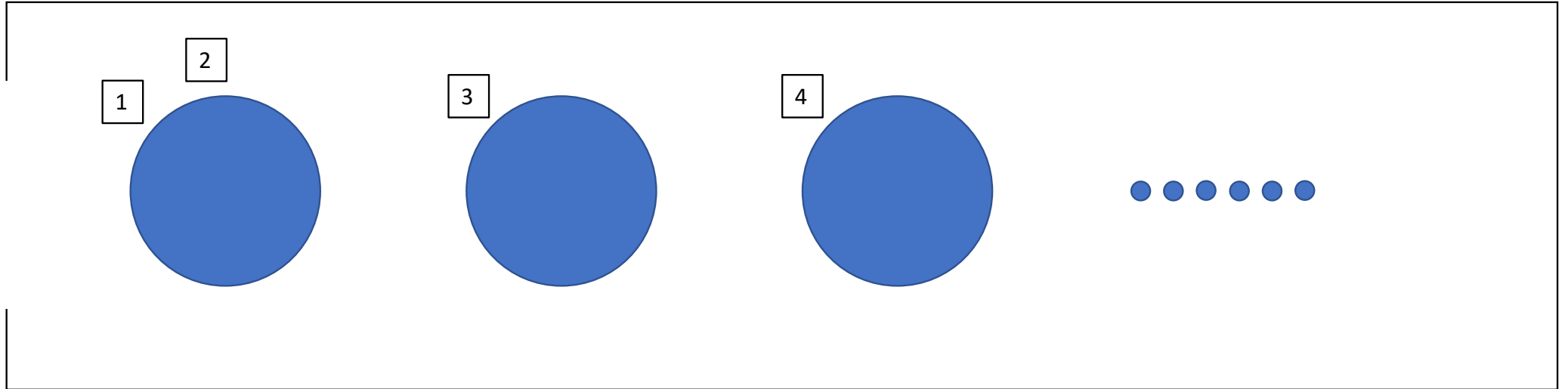
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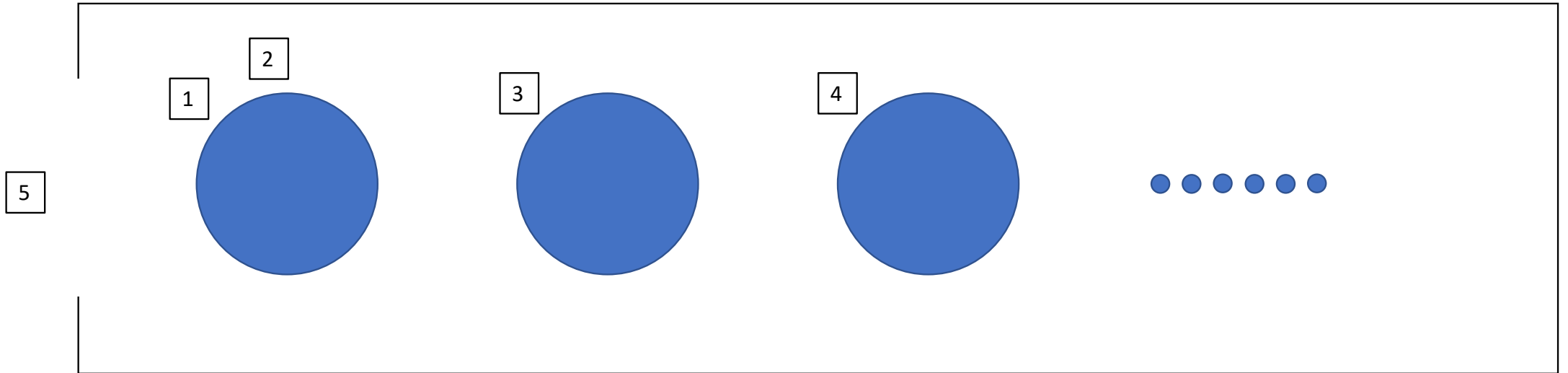
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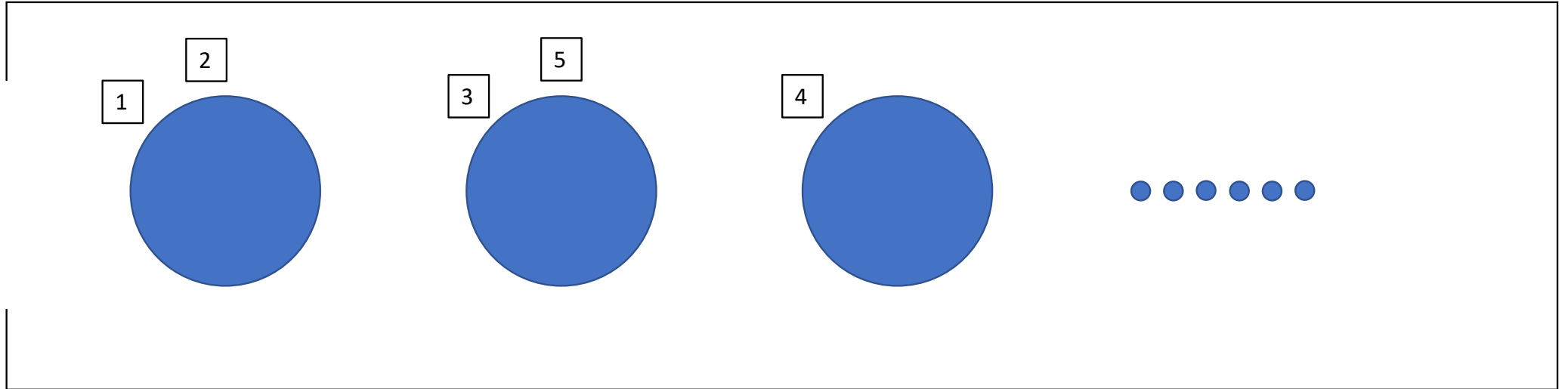
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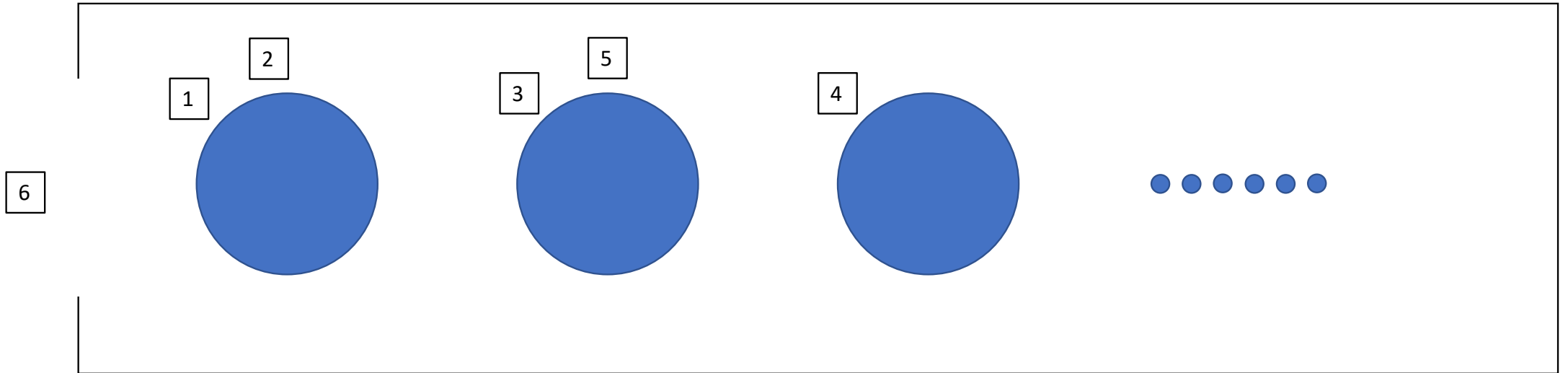
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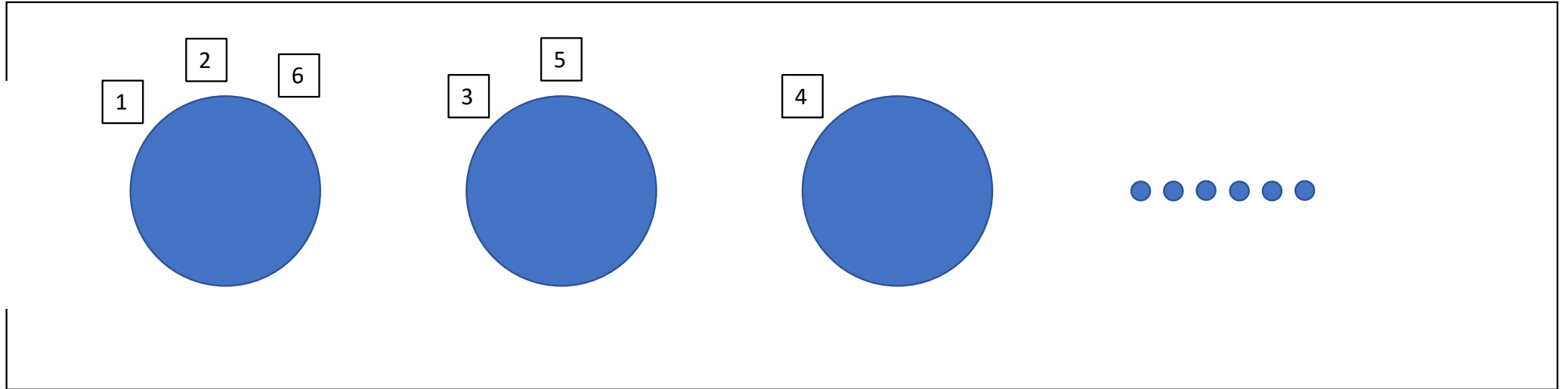
# Chinese Restaurant Process



# Chinese Restaurant Process



# Chinese Restaurant Process





# Indian Buffet Process



# Indian Buffet Process



1

# Indian Buffet Process



1

2

# Indian Buffet Process



1

2



2



# Indian Buffet Process



1



2



2

3

# Indian Buffet Process



1

2



2



3





# Indian Buffet Process



1



2



3



2

4

# Indian Buffet Process



1

2

4



2

4



3

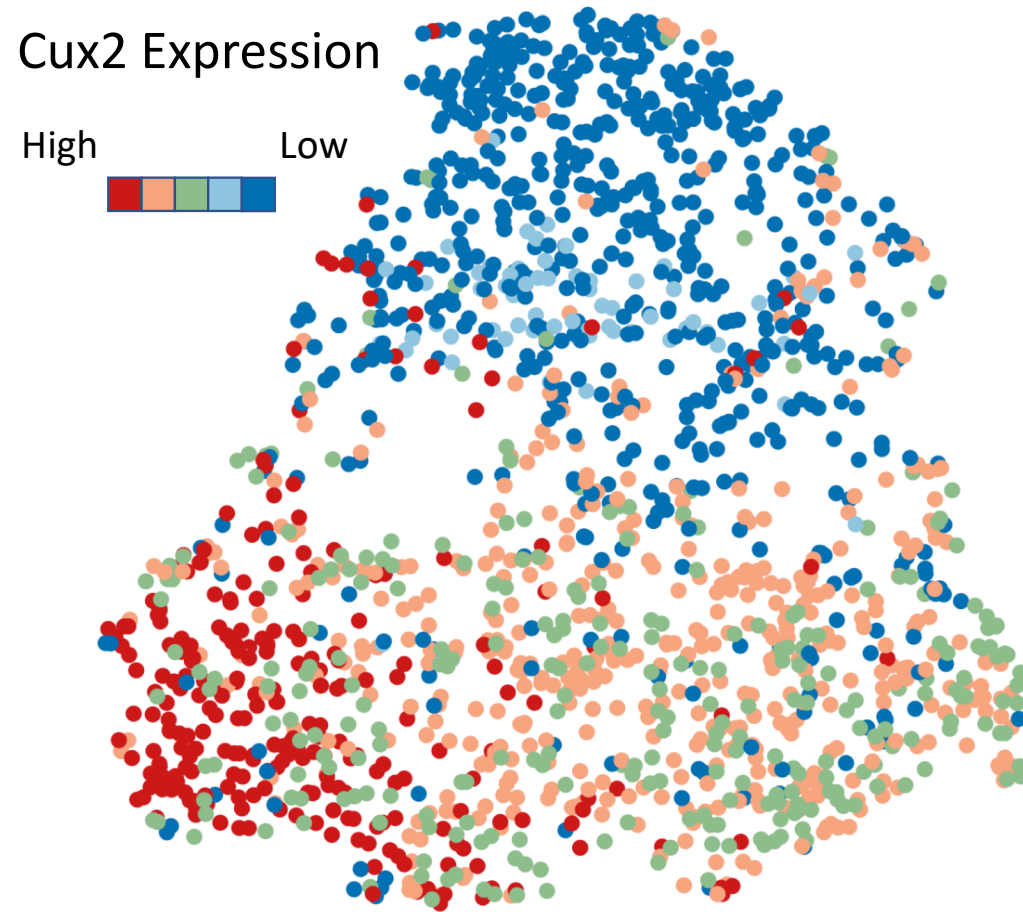
4





# Single-Cell Multi-Omics

# scRNA-seq



## Features:

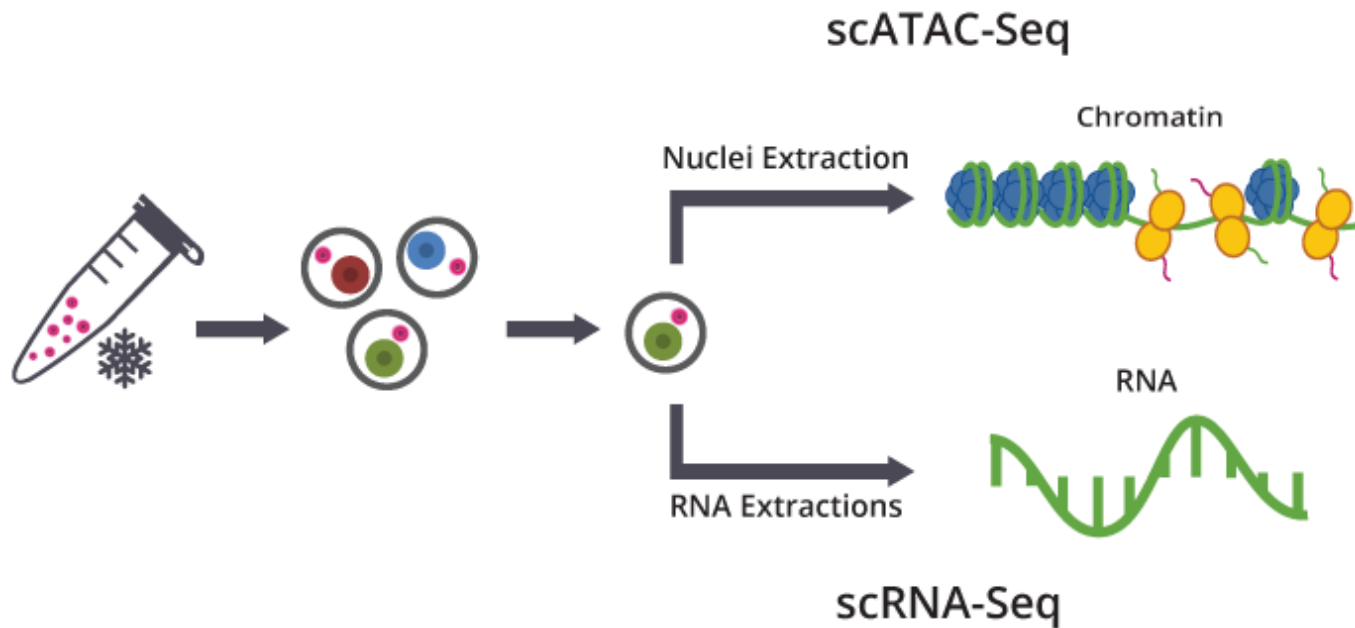
- Zero-inflated counts
- Cells may be spatially correlated (spatial transcriptomics)
- A tissue region may be non-convex

## Problems:

- Find cell types – clustering
- Estimate gene regulatory networks – causal discovery and graphical models

# scRNA-seq + scATAC-seq

Source: <https://www.genewiz.com/Public/Services/Next-Generation-Sequencing/Single-Cell-ATAC-Seq/>



## Features:

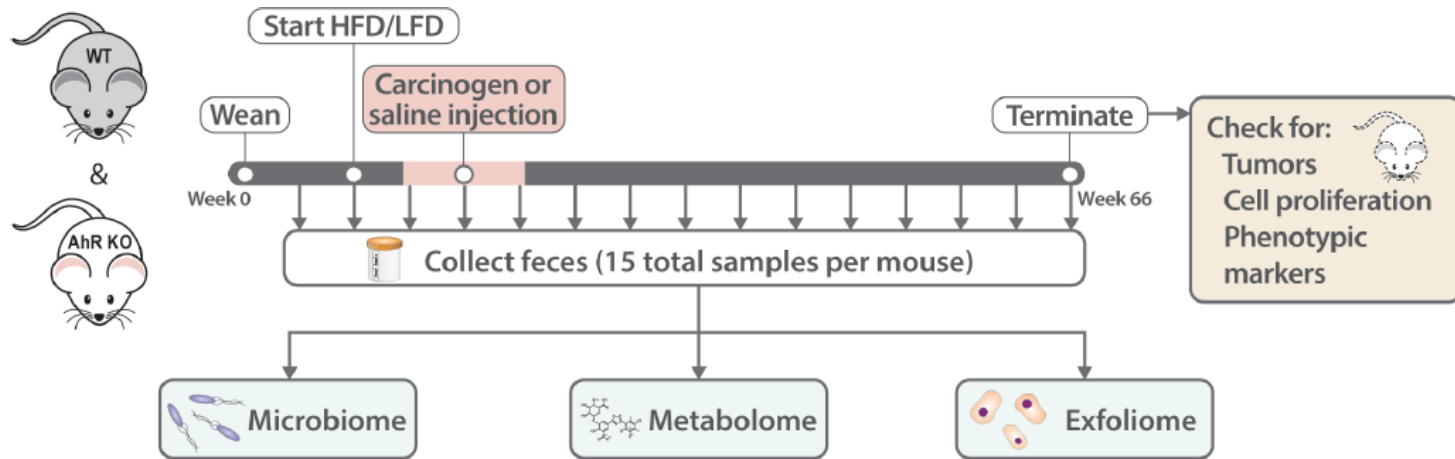
- Zero-inflated counts
- Chromatin accessibility + gene expression

## Problems:

- Integrate these two types of data by using biological knowledge

# Microbiome Multi-Omics

# Microbiome + Metabolome + Exfoliome



## Features:

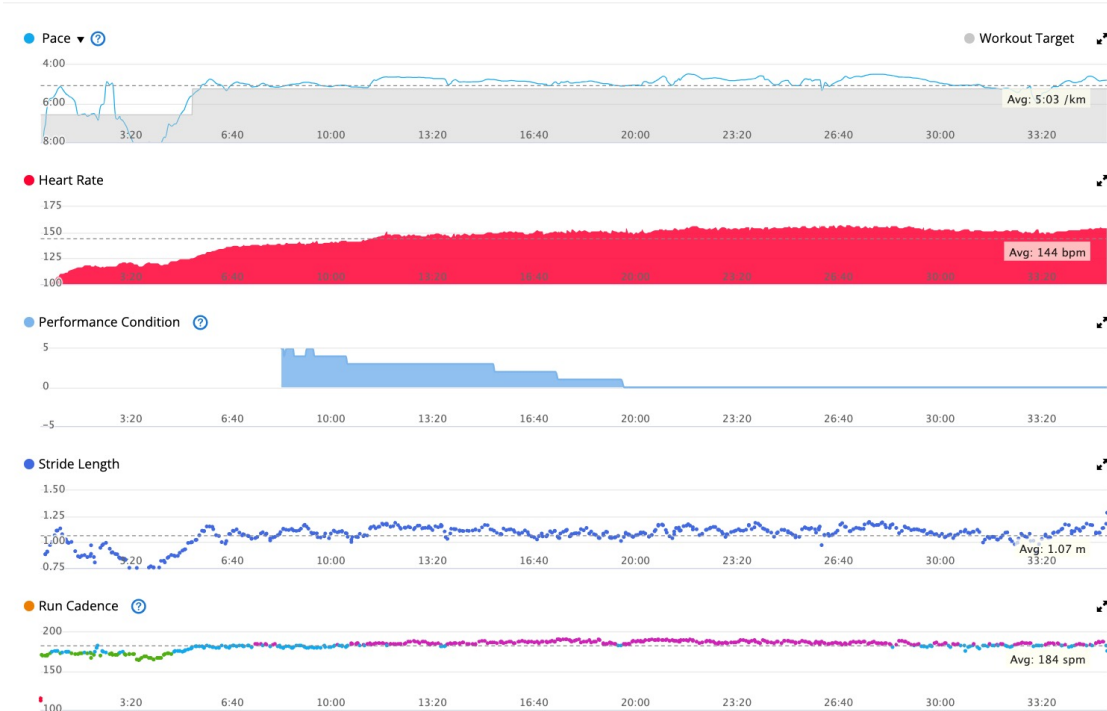
- Microbiome data are zero-inflated and compositional
- Longitudinal data
- Multiple data types

## Problems:

- Find microbial communities
- Estimate microbial networks

# Digital Health

# Wearable Device Data



fenix 6X Pro  
Software: 26.00  
Elev Corrections: Disabled  
Summary Data: Original



HRM-Pro  
Software: 8.8

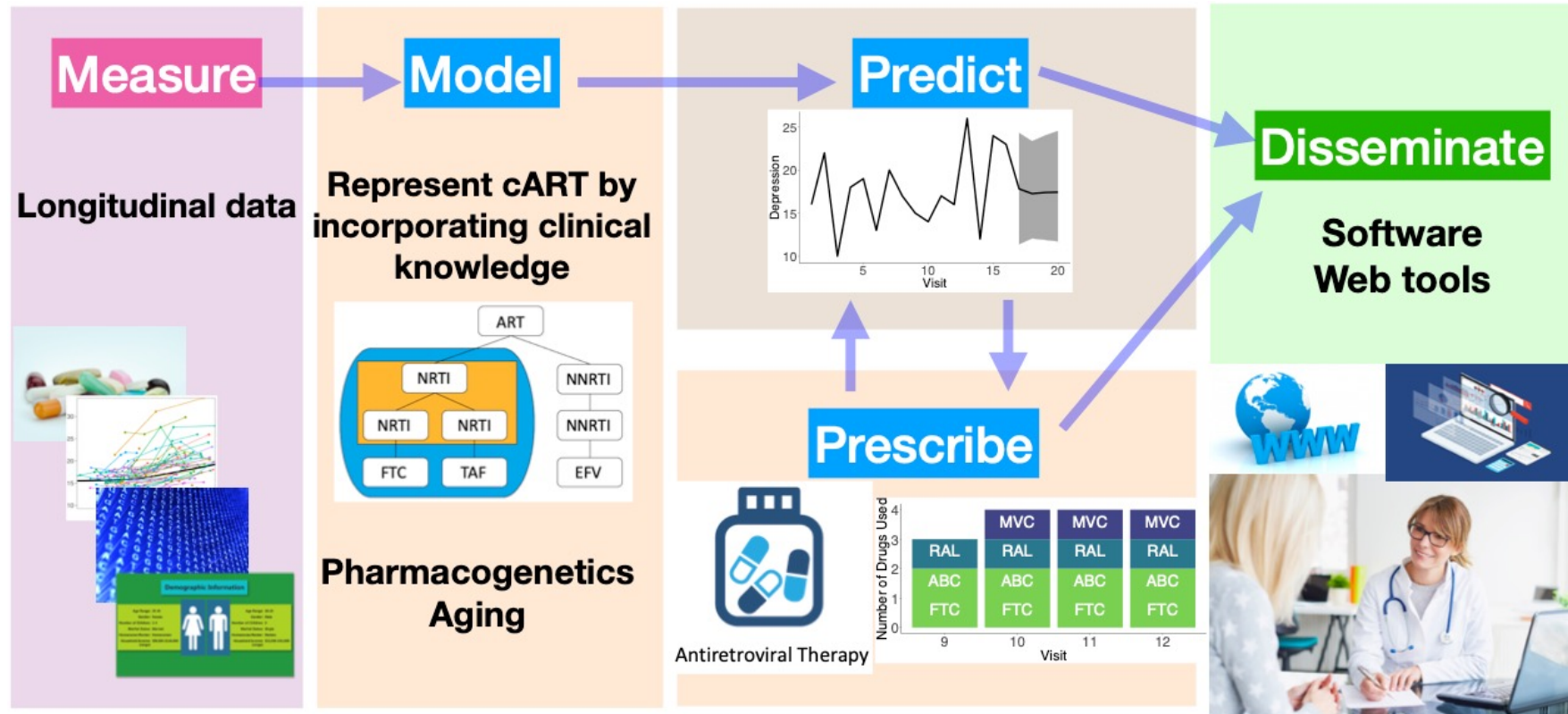
## Features:

- Multivariate functional data are infinite dimensional
- Mixed type (count, categorical, etc)

## Problems:

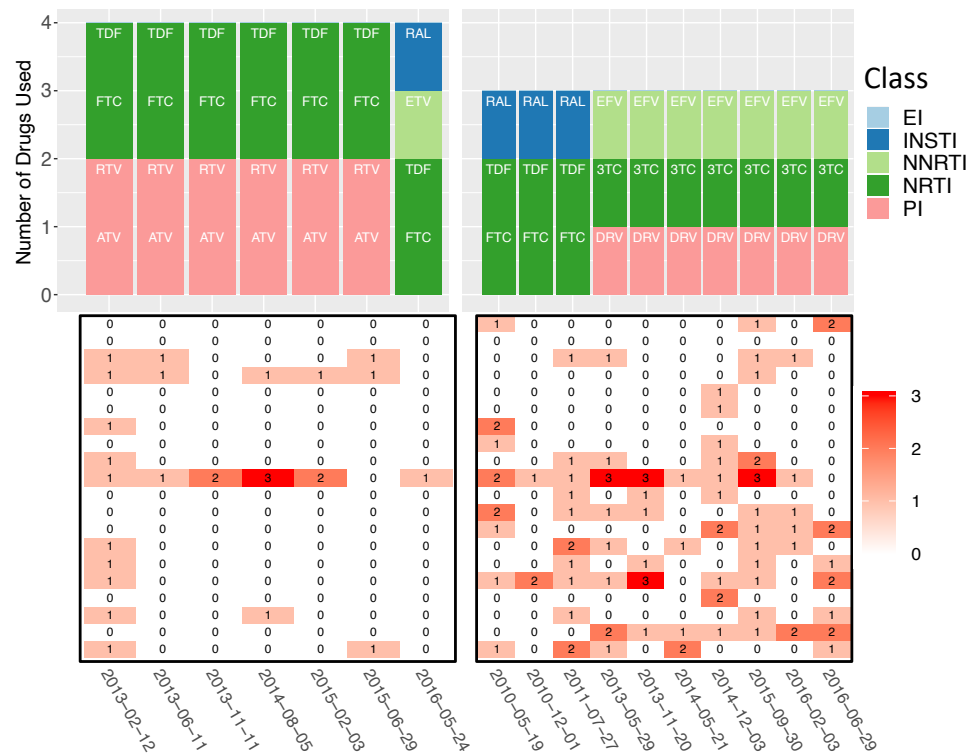
- Identify causal relationships among random functions

# HIV Longitudinal Study





# HIV Longitudinal Study



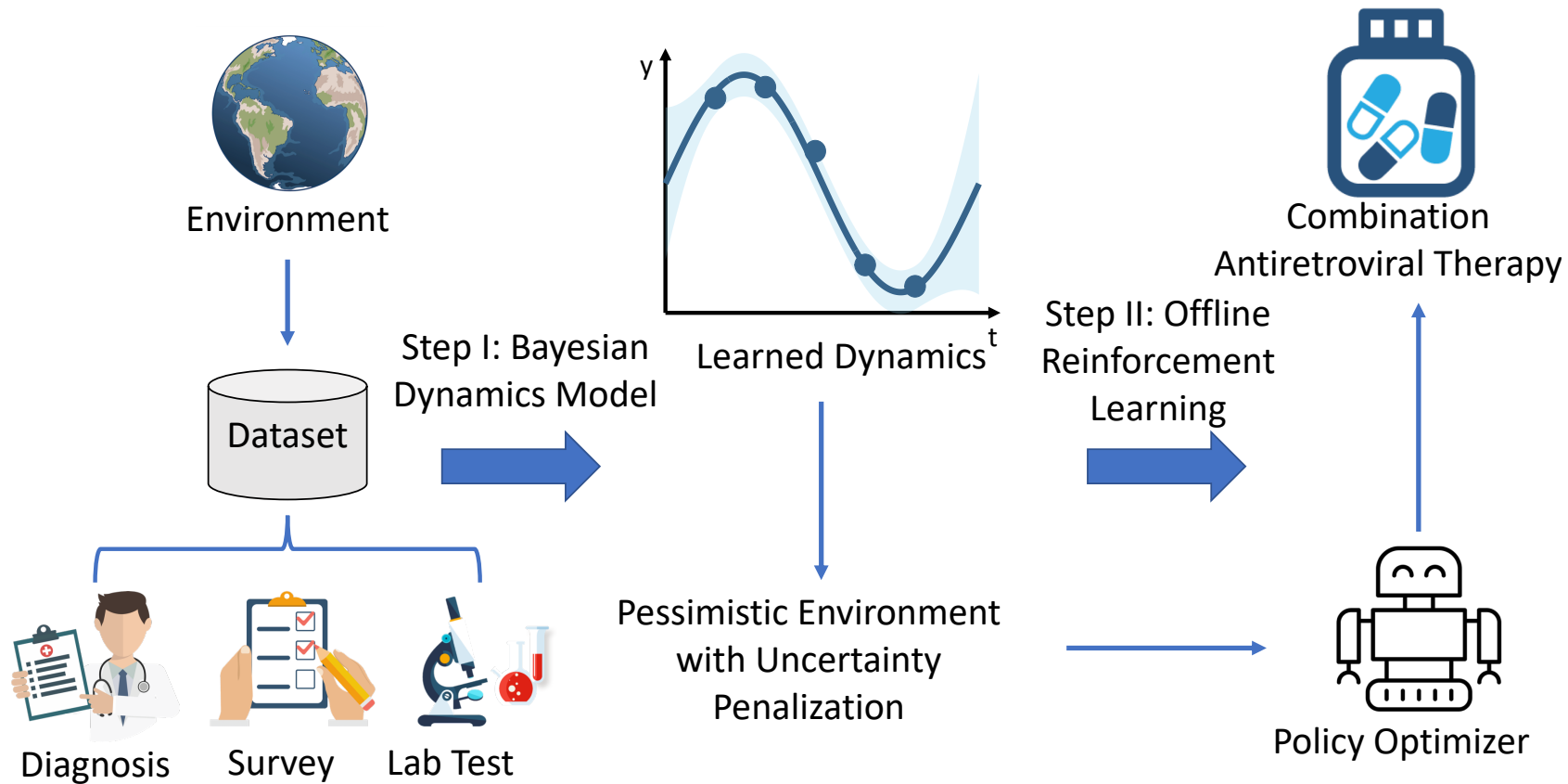
## Center for Epidemiologic Studies Depression Scale (CES-D), NIMH

Below is a list of the ways you might have felt or behaved. Please tell me how often you have felt this way during the past week.

	During the Past			
	Week			
	Rarely or none of the time (less than 1 day )	Some or a little of the time (1-2 days)	Occasionally or a moderate amount of time (3-4 days)	Most or all of the time (5-7 days)
1. I was bothered by things that usually don't bother me.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
2. I did not feel like eating; my appetite was poor.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
3. I felt that I could not shake off the blues even with help from my family or friends.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
4. I felt I was just as good as other people.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
5. I had trouble keeping my mind on what I was doing.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
6. I felt depressed.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
7. I felt that everything I did was an effort.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
8. I felt hopeful about the future.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
9. I thought my life had been a failure.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
10. I felt fearful.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
11. My sleep was restless.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
12. I was happy.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
13. I talked less than usual.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
14. I felt lonely.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
15. People were unfriendly.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
16. I enjoyed life.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
17. I had crying spells.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
18. I felt sad.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
19. I felt that people dislike me.	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
20. I could not get "going."	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

SCORING: zero for answers in the first column, 1 for answers in the second column, 2 for answers in the third column, 3 for answers in the fourth column. The scoring of positive items is reversed. Possible range of scores is zero to 60, with the higher scores indicating the presence of more symptomatology.

# HIV Longitudinal Study



# Non-Gaussian Factor Model for Interpretable Ordinal Time Series Analysis

- $Y_t = (y_{1t}, \dots, y_{pt})'$  observed ordinal responses to the questions at time  $t = 1, \dots, T$
- $y_{jt} \in \{1, \dots, C_j\}$
- $X_t = (x_{1t}, \dots, x_{Lt})'$  latent continuous factors/traits at time  $t$
- Assume  $L \leq p$
- Let  $X_t^{(d)} = \{X_{t-1}, \dots, X_{t-d}\}$

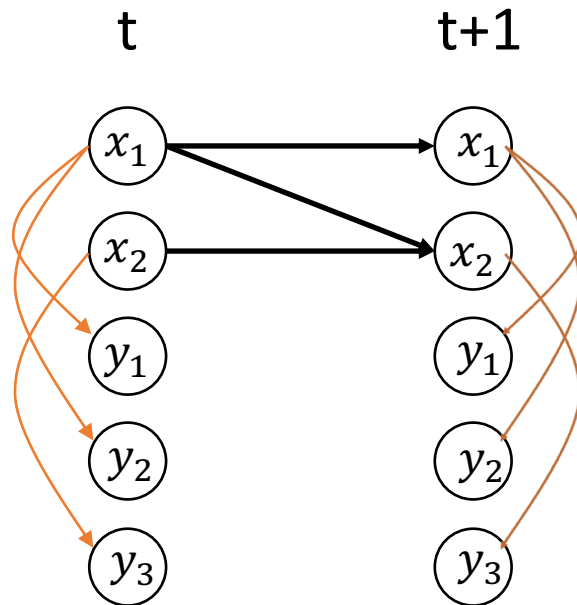
# Non-Gaussian Factor Model for Interpretable Ordinal Time Series Analysis

$$X_t = g\left(X_t^{(d)}, \epsilon_t\right)$$

Causal model

$$\Pr(y_{jt} \leq c | X_T) = \Phi(\eta_c - \beta_j' X_t - \alpha_j)$$

Measurement/factor model



Two sources of non-identifiability:

1. Latent factors
2. Causal graph

Our hypothesis:

If  $\epsilon_t$  is non-Gaussian, both latent factors and the causal graph is identifiable