

ARE YOU EXPLOITING YOUR ASSUMPTIONS?

TOWARDS EXPRESSIVE PRIORS FOR BIOMARKER DISCOVERY AND FUNCTIONAL PREDICTION

Melanie F. Pradier

Harvard University



HDSI

Harvard Data
Science Initiative



CRCS Center for Research on
Computation and Society
at Harvard John A. Paulson School of Engineering and Applied Sciences

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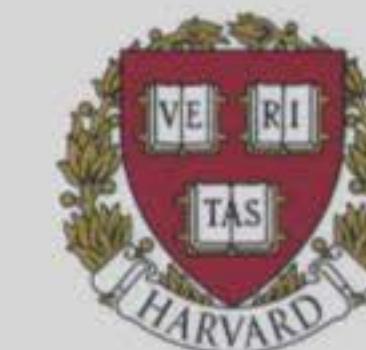
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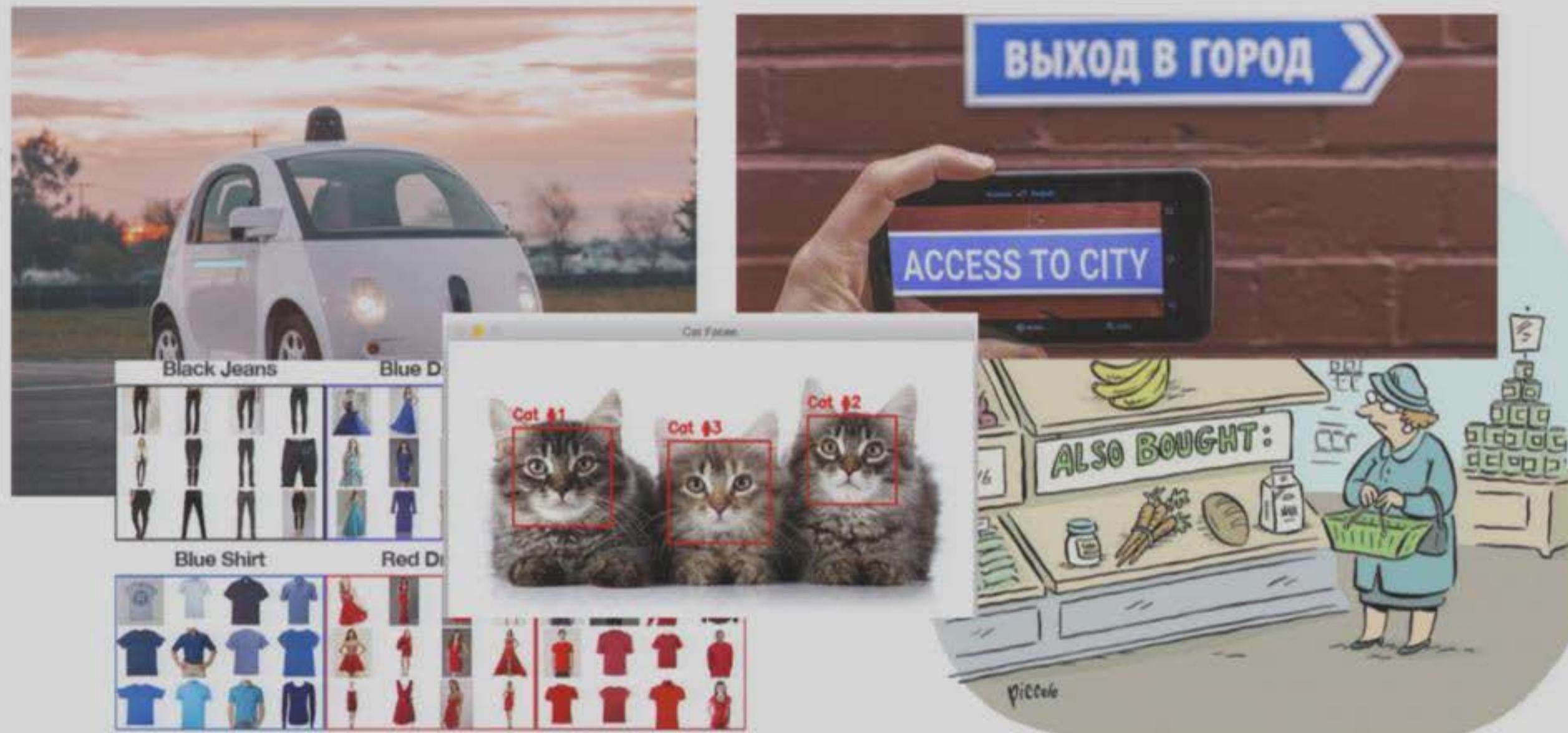
Research in collaboration
with...



UNIVERSITY OF
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ML SYSTEMS ARE TRANSFORMING OUR SOCIETY



A screenshot of a Google search results page. The search query "machine learning" is entered in the search bar. Below the search bar are four filter buttons: "All", "News", "Images", and "Videos". A red oval highlights the text "About 3,000,000,000 results (0.57 seconds)" which is displayed below the filters.

Unveiling Biology with Deep Microscopy

An AI-inspired Revolution in the Life Sciences



Brian Hilbush [Follow](#)

Dec 5, 2019 · 6 min read · *

MAGAZINE · ARTIFICIAL INTELLIGENCE

A.I. breakthroughs in natural-language processing are big for business

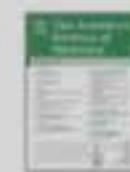


BY JEREMY KAHN

January 20, 2020 8:30 AM EST

The American Journal of Medicine

Volume 132, Issue 7, July 2019, Pages 795-801



Review

Artificial Intelligence Transforms the Future of Health Care

Nazman Noorbakhsh-Sabet MD ^{1,2}, Ramin Zand MD, MPH ^{1,3,4}, Yanfei Zhang PhD ⁵, Vida Abedi PhD ^{1,2} [✉] et al

Article | Open Access | Published: 05 February 2020

Inferring structural variant cancer cell fraction

Nature Communications 11, Article number: 730 (2020) | Cite this article

Research

26 February 2020 | Open Access

[Impact of a deep learning assistant on the histopathologic classification of liver cancer](#)



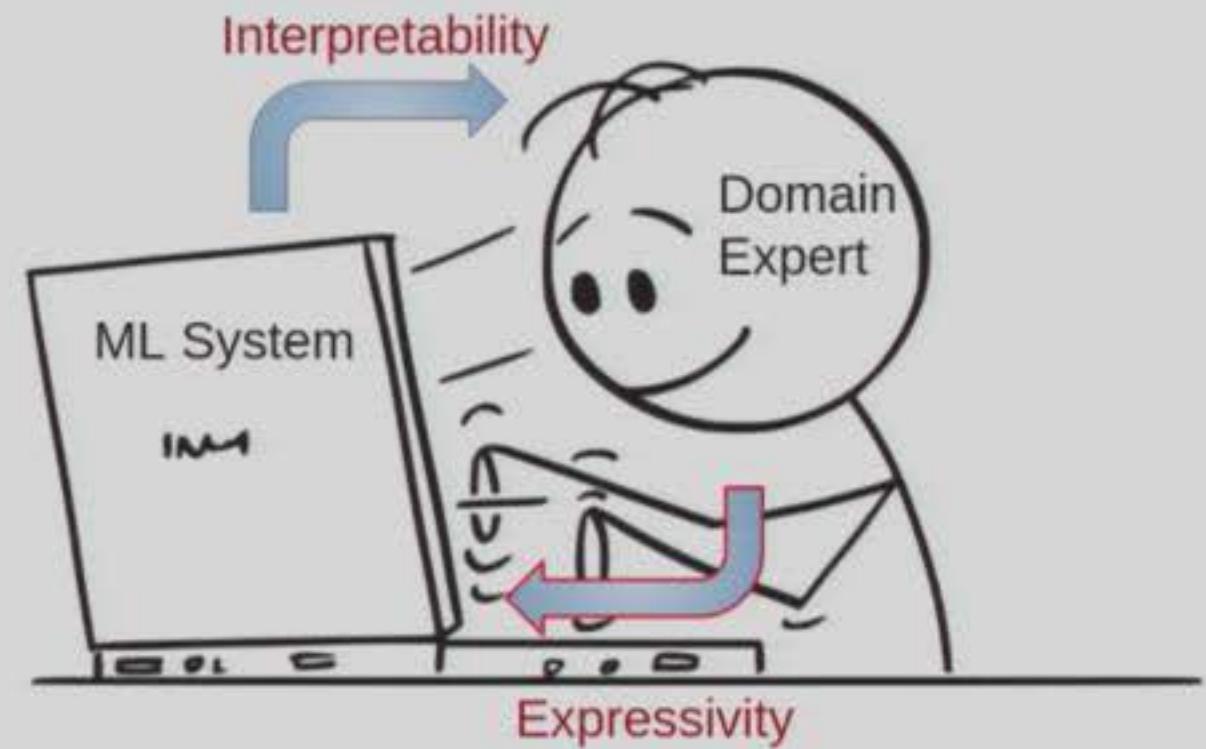
Amirhossein Kiani, Bora Uyumazturk, Jeanne Shen

npj Digital Medicine 3, 23

The image displays four distinct news articles, each with a red oval highlighting a specific term or phrase:

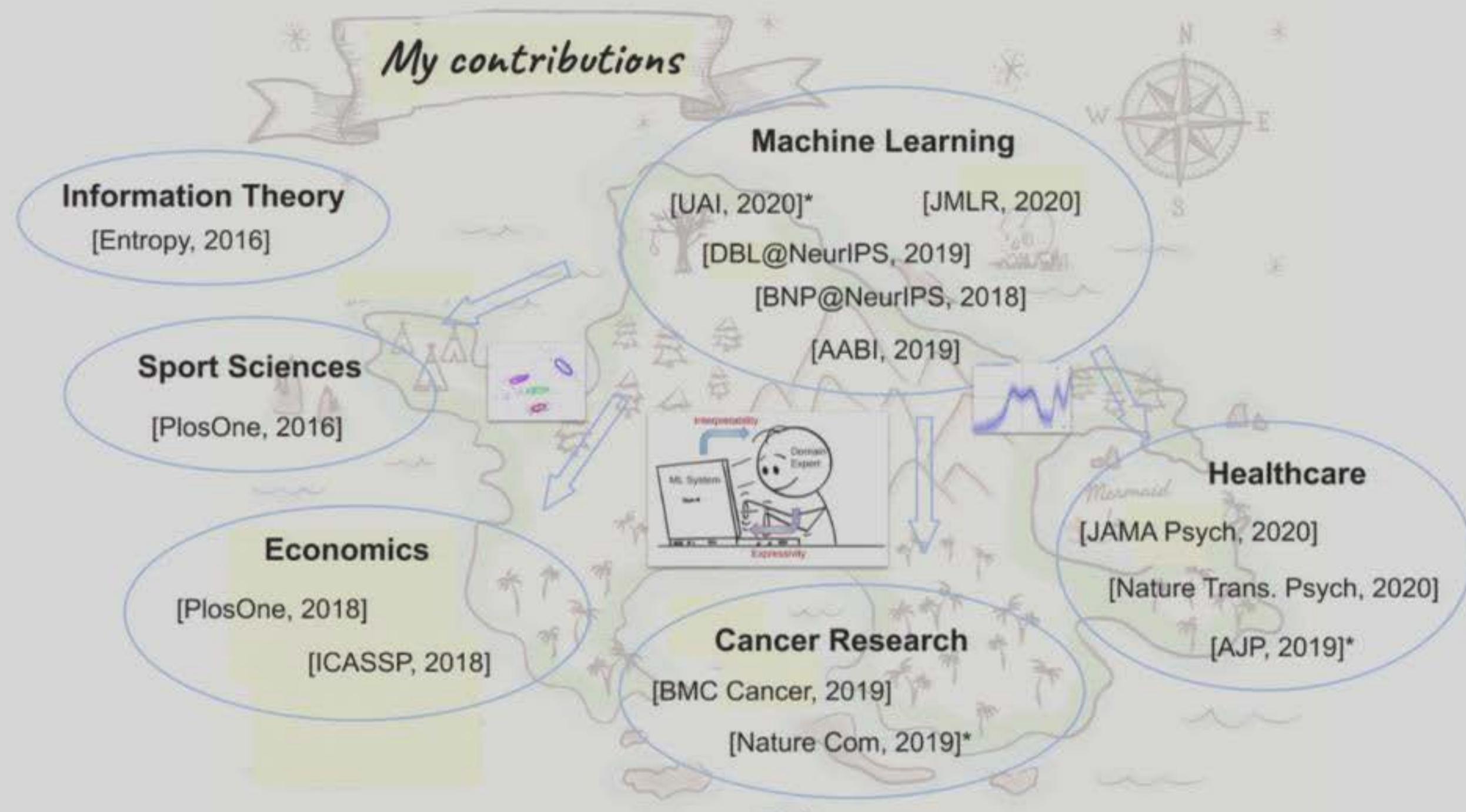
- nature** (top left):
 - NEWS FEATURE** - 09 OCTOBER 2019
 - Why deep-learning AIs are so easy to fool**
 - Artificial-intelligence researchers are trying to fix the flaws of neural networks.
- HDSR** (bottom left):
 - Should We Trust Algorithms?**
 - nature machine intelligence**
 - Perspective | Published: 13 May 2019
 - Stop explaining black box machine learning models for high stakes decisions and use interpretable models instead**
- Artificial Intelligence, Machine Learning, and Bias in Finance: Toward Responsible Innovation** (top right, circled)
 - K Johnson, F Pasquale, J Chapman - Fordham L. Rev., 2019 - HeinOnline
 - Over the last decade, a growing number of digital startups launched bids to lure business
- Improving Fairness in Machine Learning Systems: What Do Industry Practitioners Need?** (middle right, circled)
 - RESEARCH ARTICLE
 - Publication: CHI '19: Proceedings of the 2019 CHI Conference on Human Factors in Computing Systems • May 2019
- Making AI Work with Small Data** (bottom right, circled)
 - TECHNOLOGY AND INNOVATION > DIGITAL TOOLS
 - Alejandro Betancourt
 - FEB 12, 2020
 - The real test of an AI machine is when it can admit to not knowing something
John Naughton

INTERPRETABILITY-EXPRESSIVITY LOOP



INTERPRETABILITY

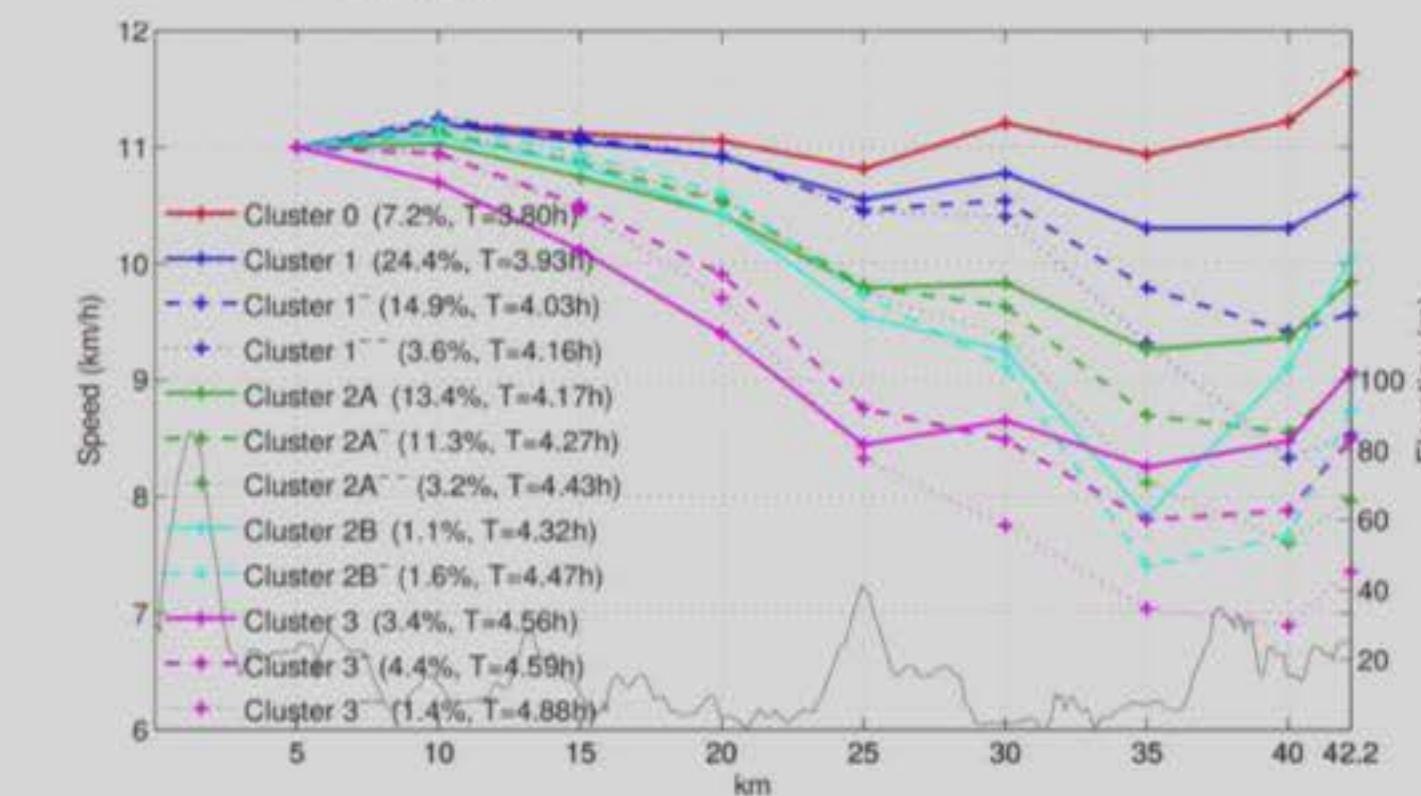
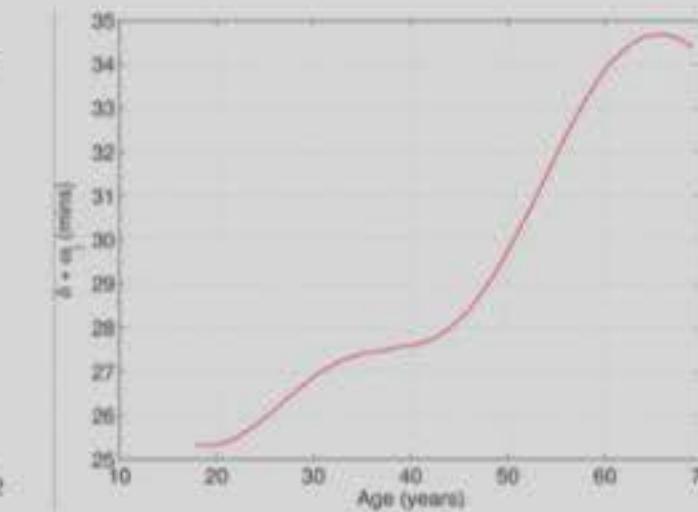
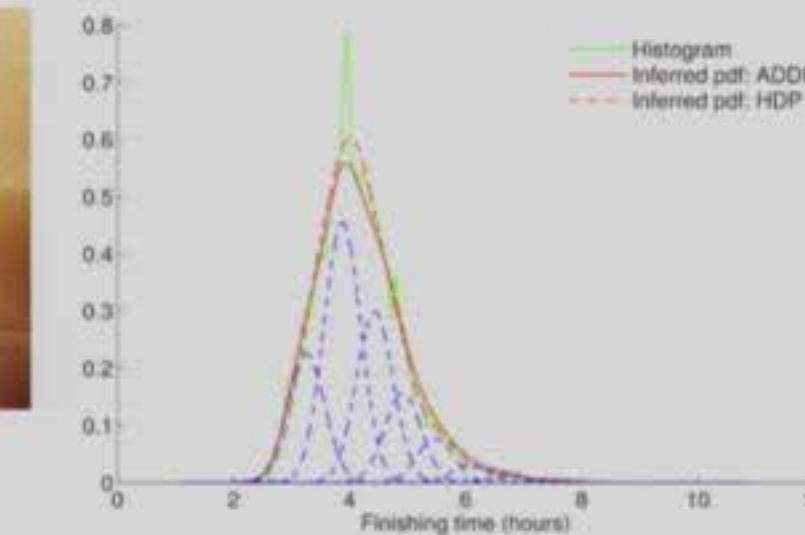
- ▶ “ability to present in understandable terms to a human” (Doshi-Velez et.al, 2017; 2018 EU GDPR)



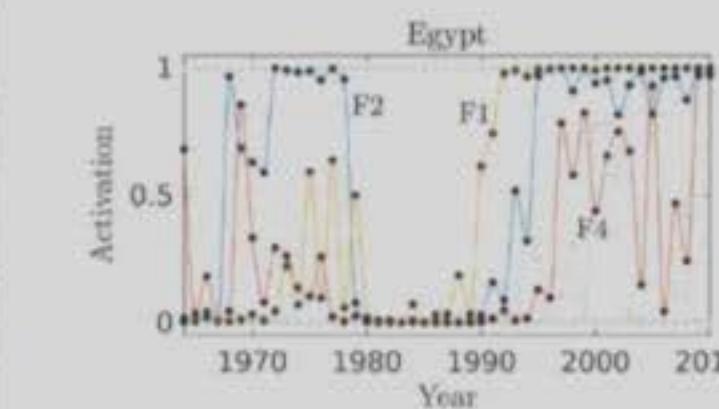
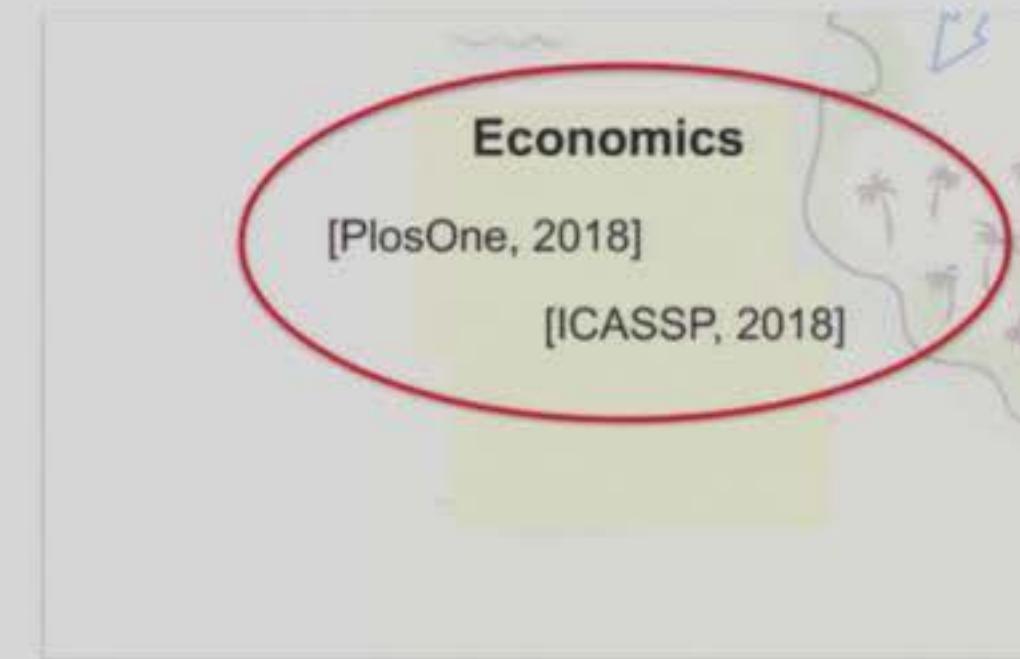
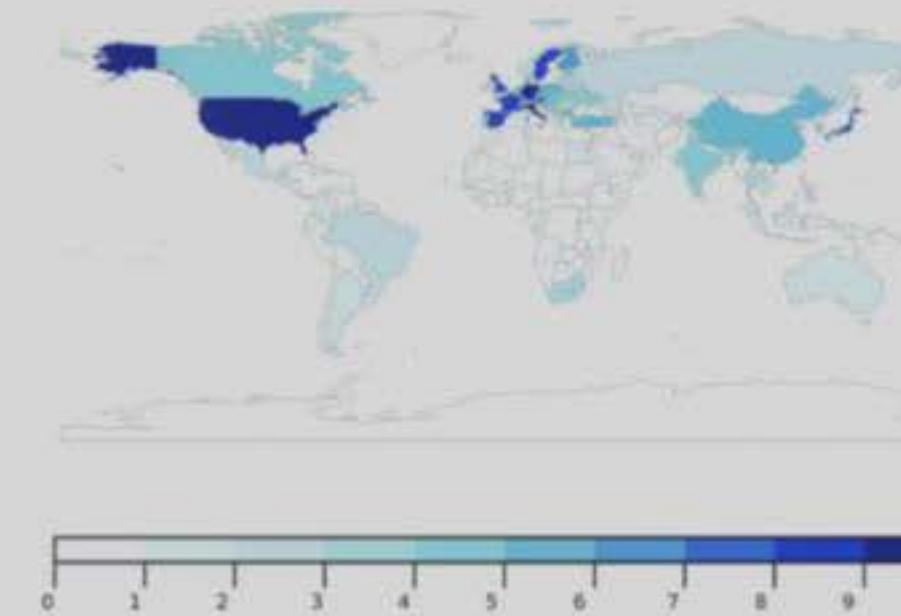


How does aging impact our athletic performance?

M. F. Pradier, F. J. R. Ruiz, and F. Perez-Cruz. **Prior Design for Dependent Dirichlet Processes: An Application to Marathon Modeling.** *PlosONE*. 2016.



Which factors make countries wealthier than others?



M. F. Pradier*, Z. Utkovski*, V. Stojkoski, L. Kocarev and F. Perez-Cruz. **Economic Complexity Unfolded: An Interpretable Model for the Productive Structure of Economies**. *PlosONE*. 2018.

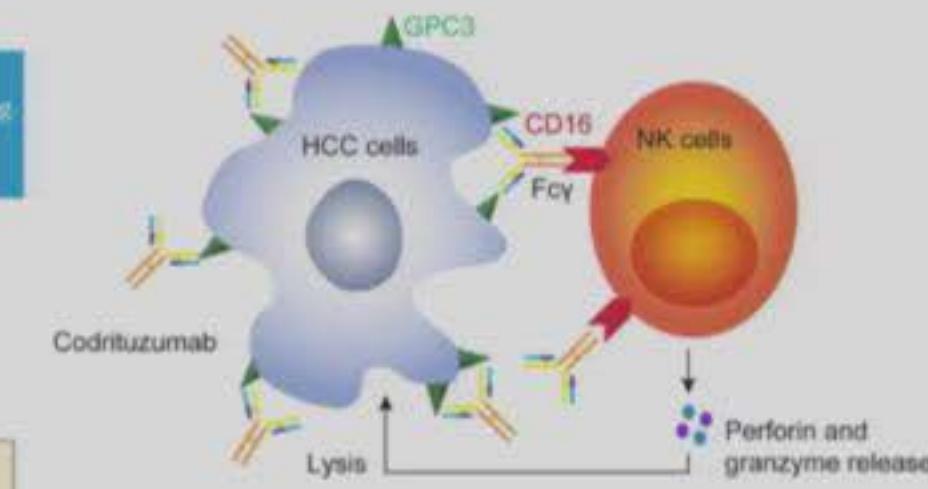
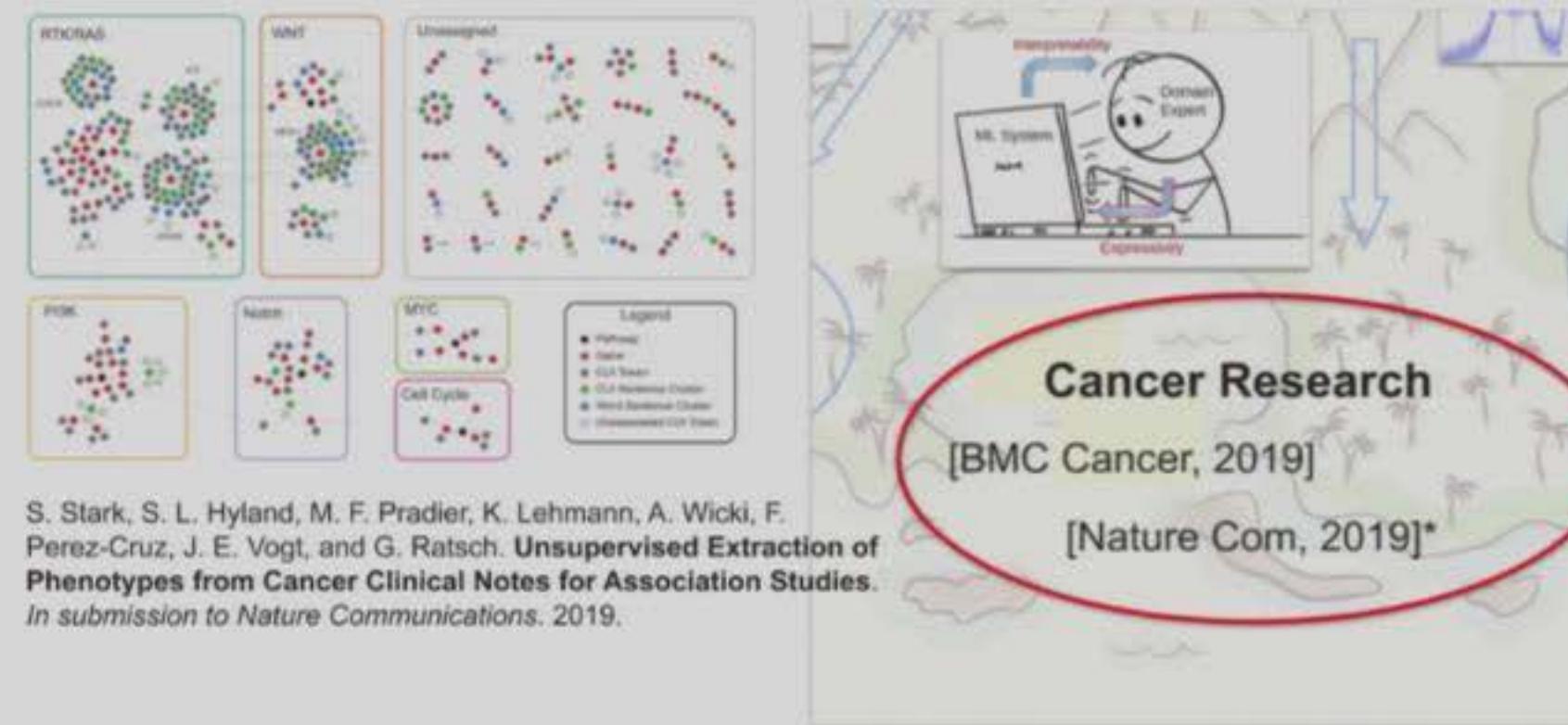
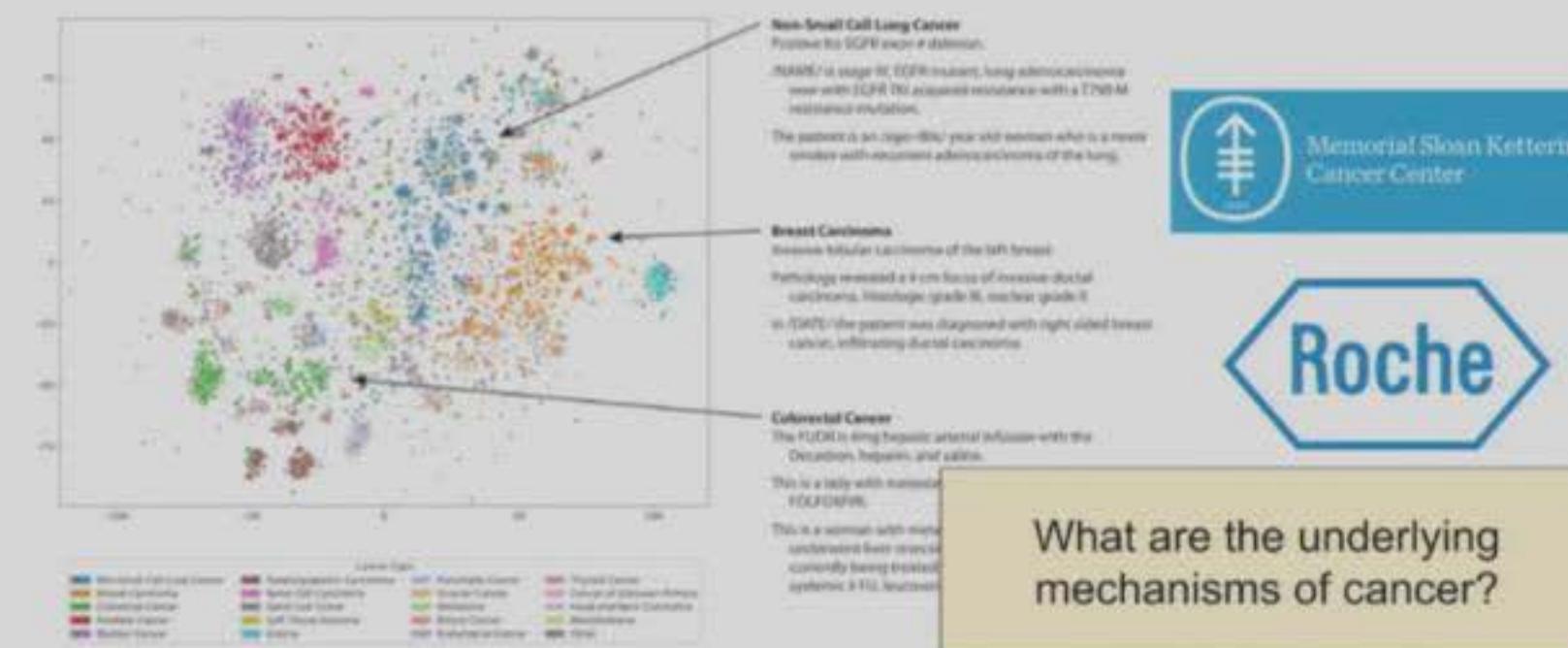
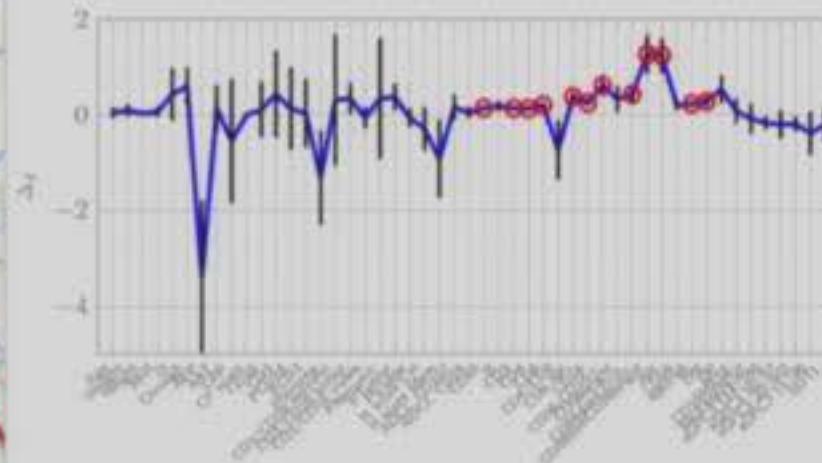
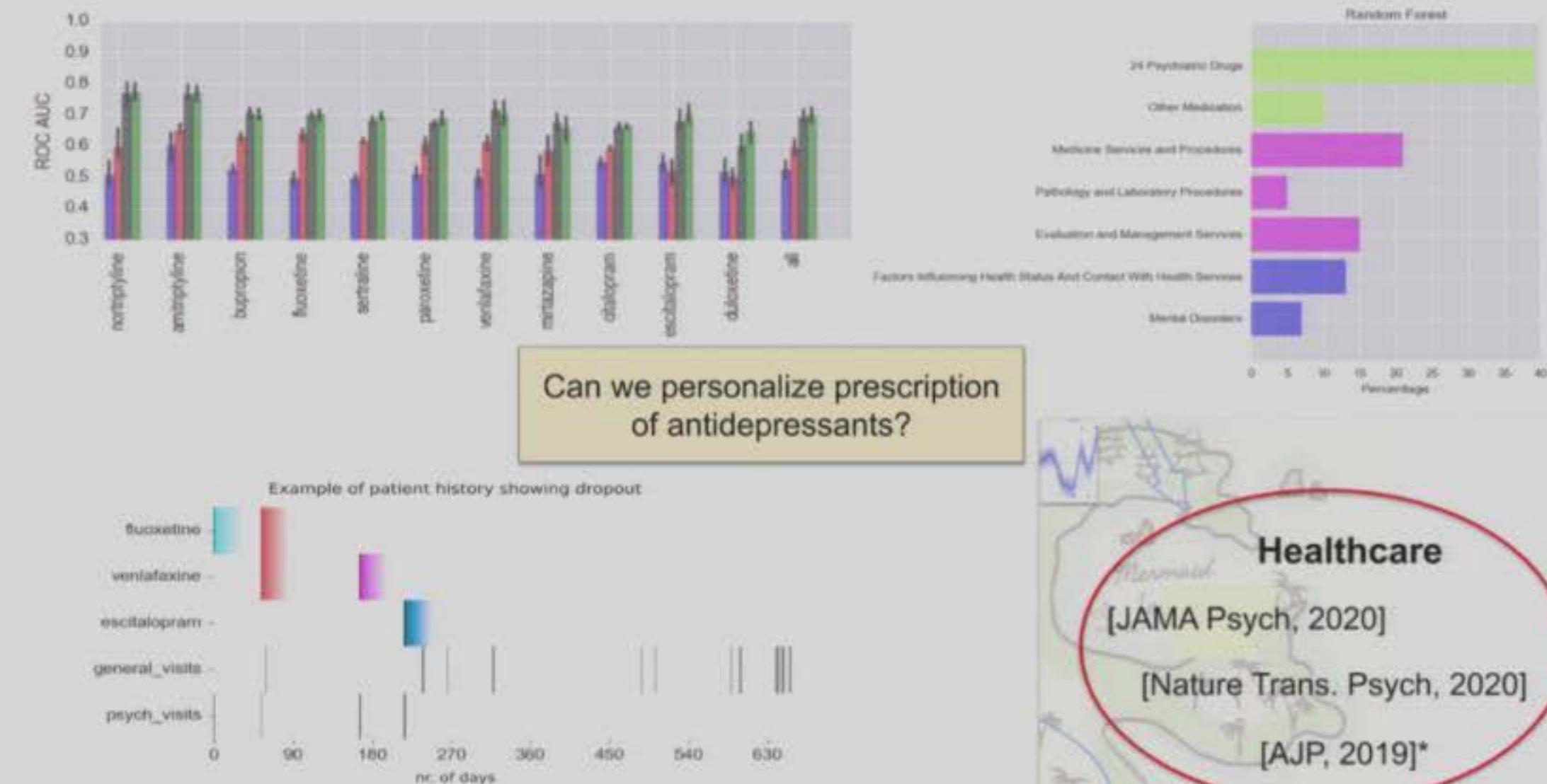


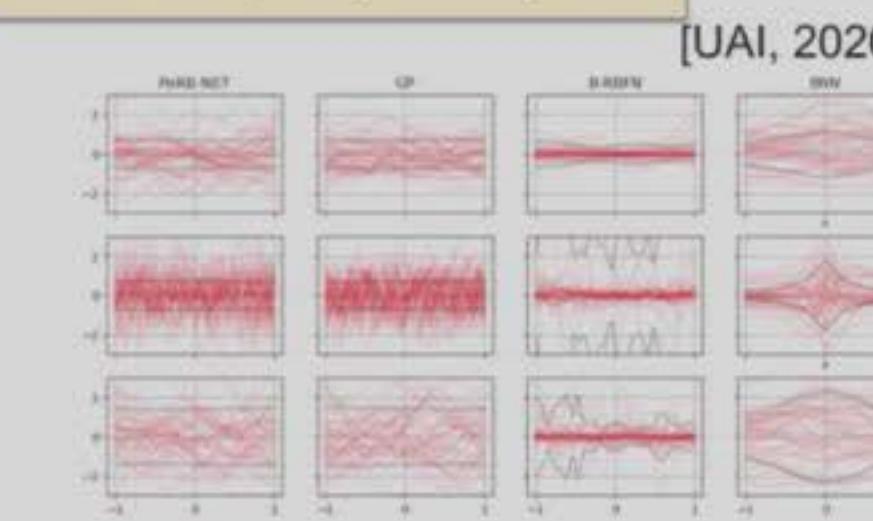
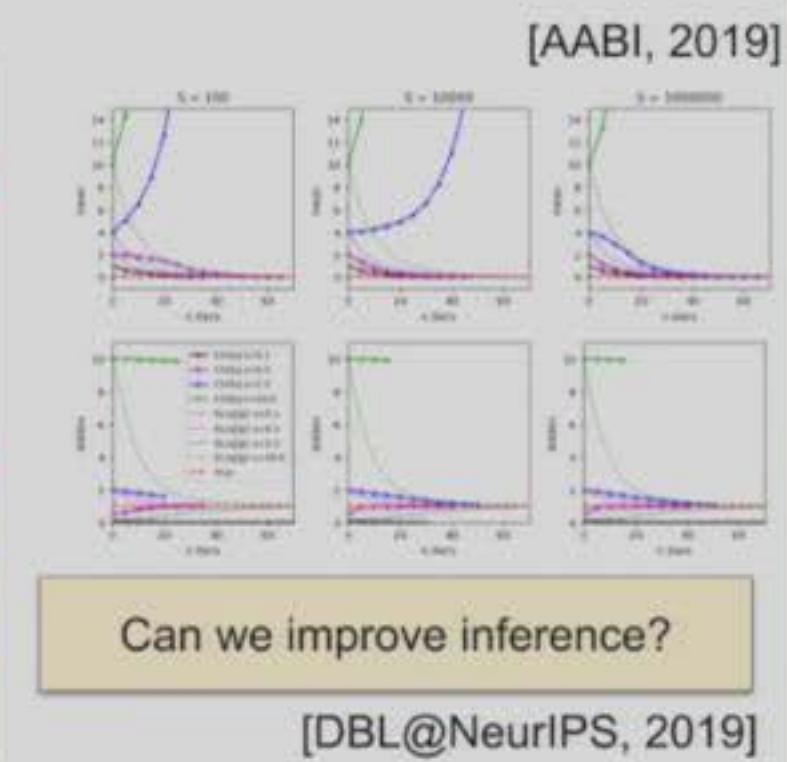
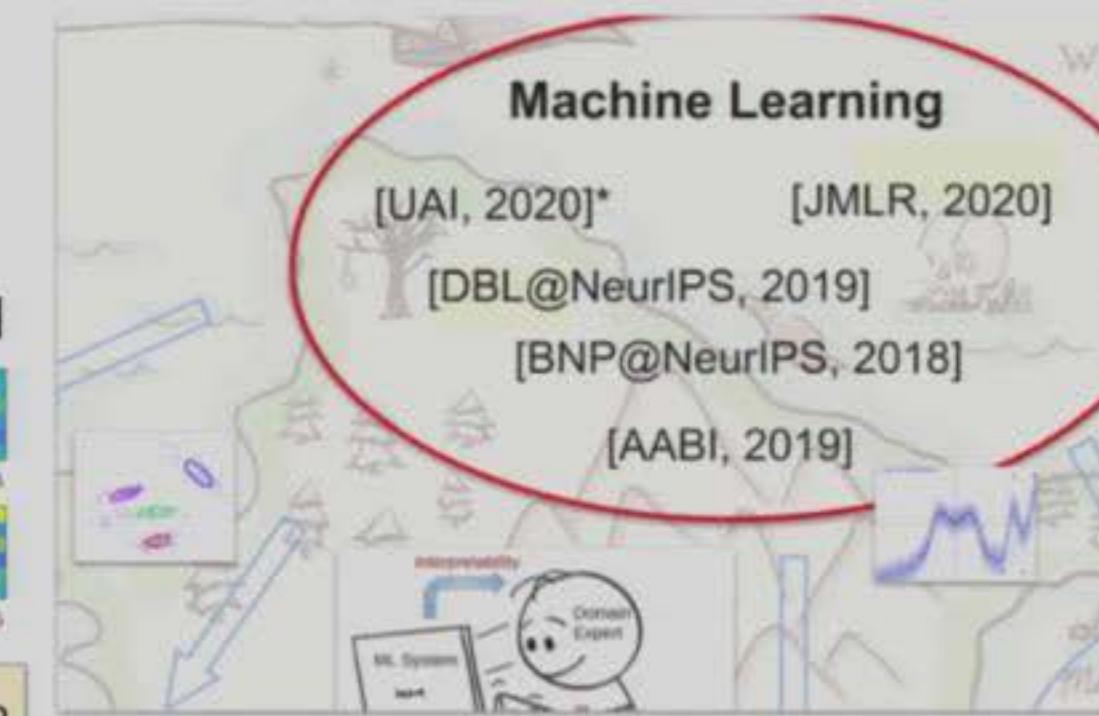
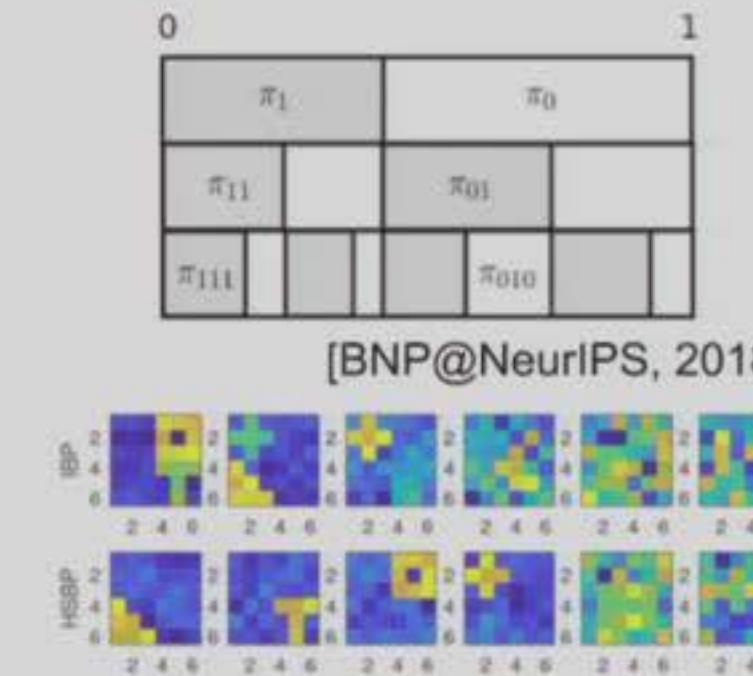
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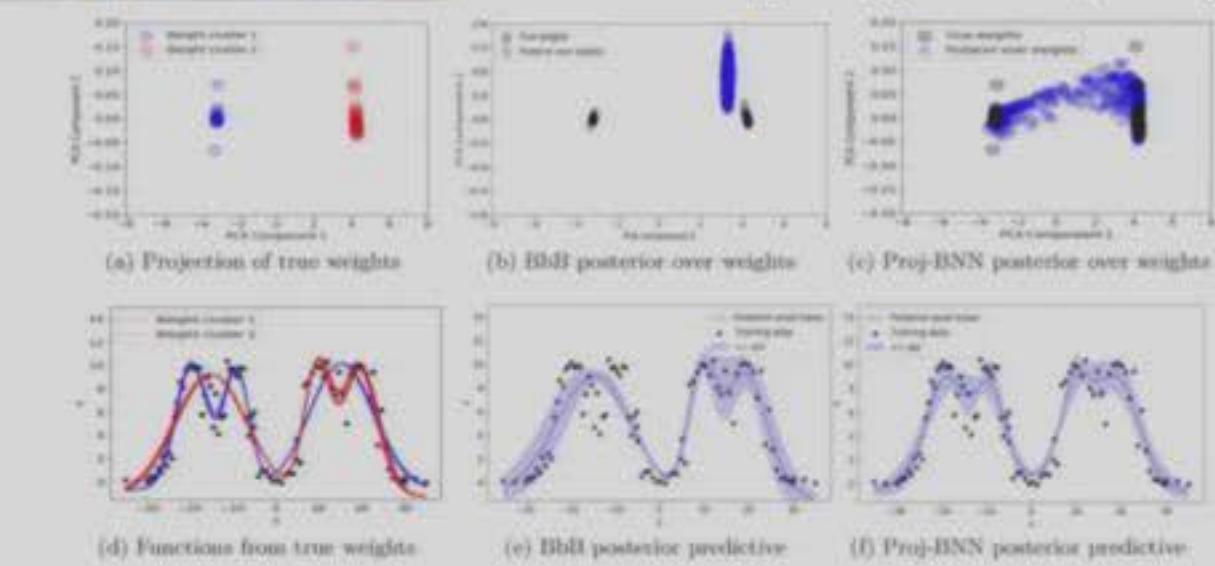
M. F. Pradier, B. Reis, L. Jukofsky, F. Milletti, T. Ohtomo, F. Perez-Cruz, and O. Puig. Case-control Indian Buffet Process identifies biomarkers of response to Codrituzumab. *BMC Cancer*. 2019.

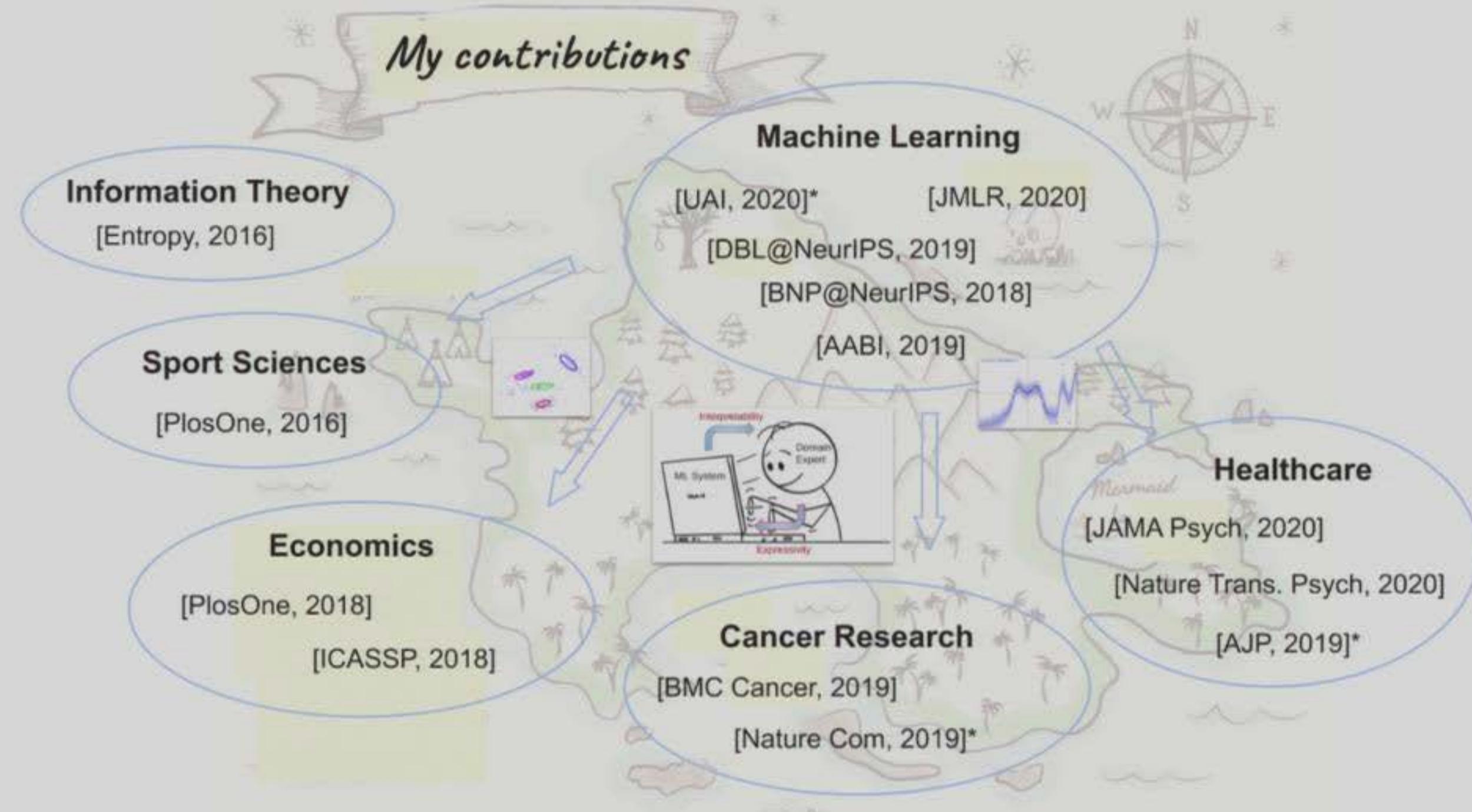


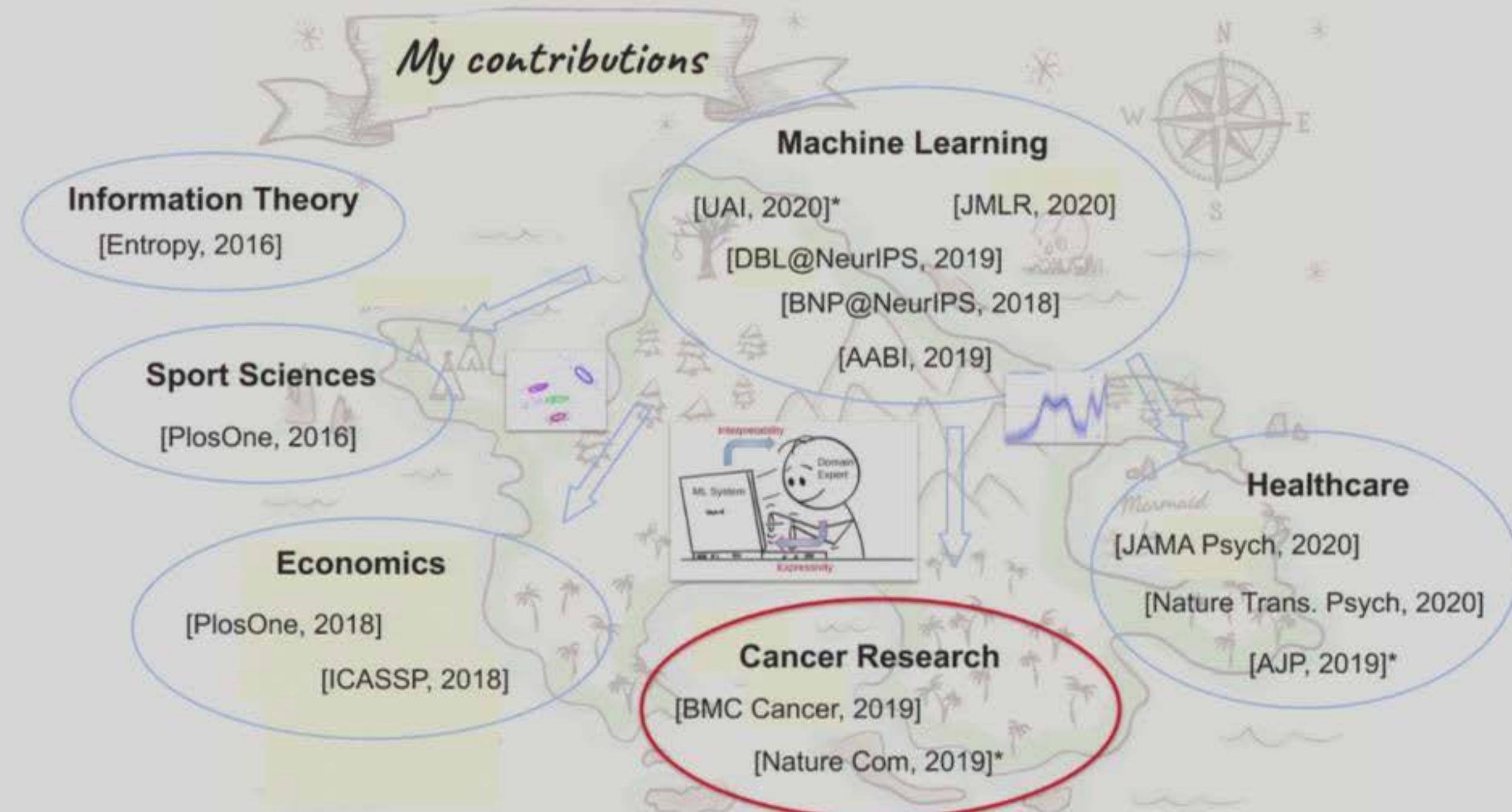
- M. F. Pradier, T. H. McCoy, M. Hughes, R. H. Perlis and F. Doshi-Velez. **Predicting Treatment Discontinuation after Antidepressant Initiation.** *Nature-Trans. Psych.* 2019.
- M. C. Hughes, M. F. Pradier, A. S. Ross, T. H. McCoy, R. H. Perlis and F. Doshi-Velez. **Generating interpretable predictions about antidepressant treatment stability using supervised topic models.** Accepted *JAMA Psychiatry*. 2019.
- M. F. Pradier, M. Hughes, T. H. McCoy, S. Barroilhet, F. Doshi-Velez and R. H. Perlis. **Predicting Transition from Major Depression to Bipolar Disorder after Antidepressant Initiation.** Submitted to *American Journal of Psychiatry*. 2019.



[UAI, 2020]*







OUR FOCUS: BIOMARKER DISCOVERY

DEF: "ANY VARIABLE THAT CAN BE USED AS AN INDICATOR OF A PARTICULAR DISEASE STATE".

Application: Phase II clinical trial for immunotherapy for liver cancer

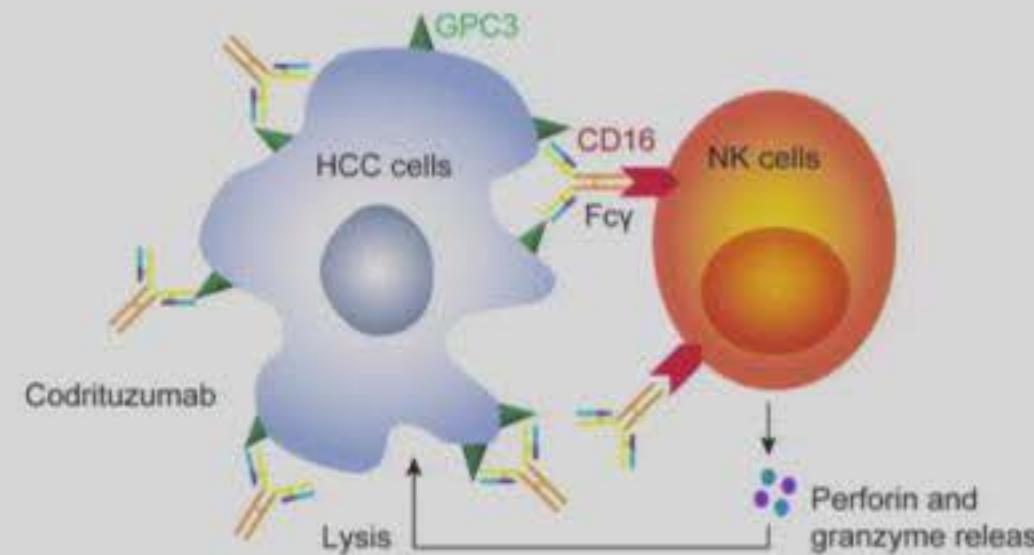


Diagram: Mechanism of codrituzumab-induced antibody-dependent cytotoxicity through the interaction of Fc CD16 in NK cells

[ABOU-ALFA ET.AL, 2016]

- ▶ **Method:** classical statistical tests
- ▶ **Conclusion:** no evidence for treatment effectiveness

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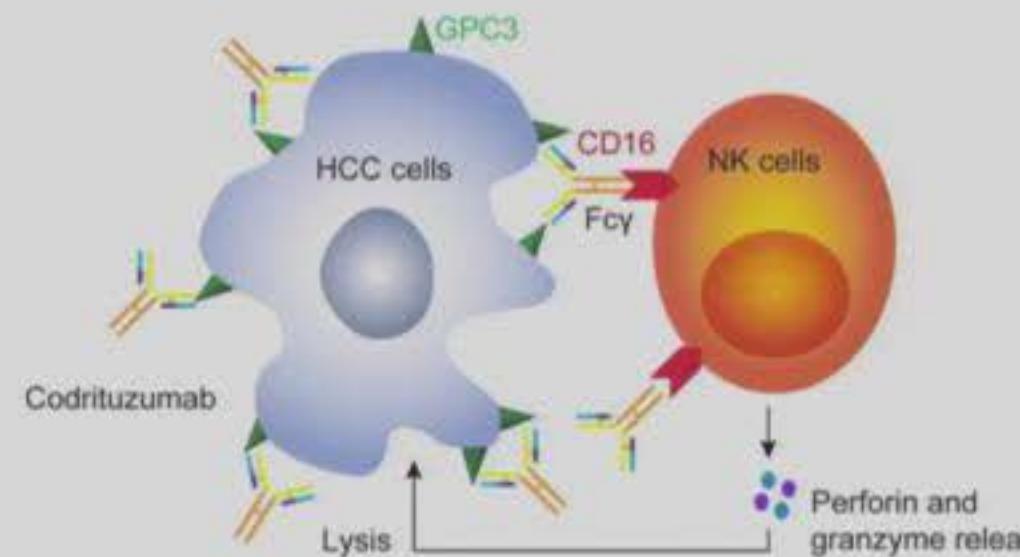
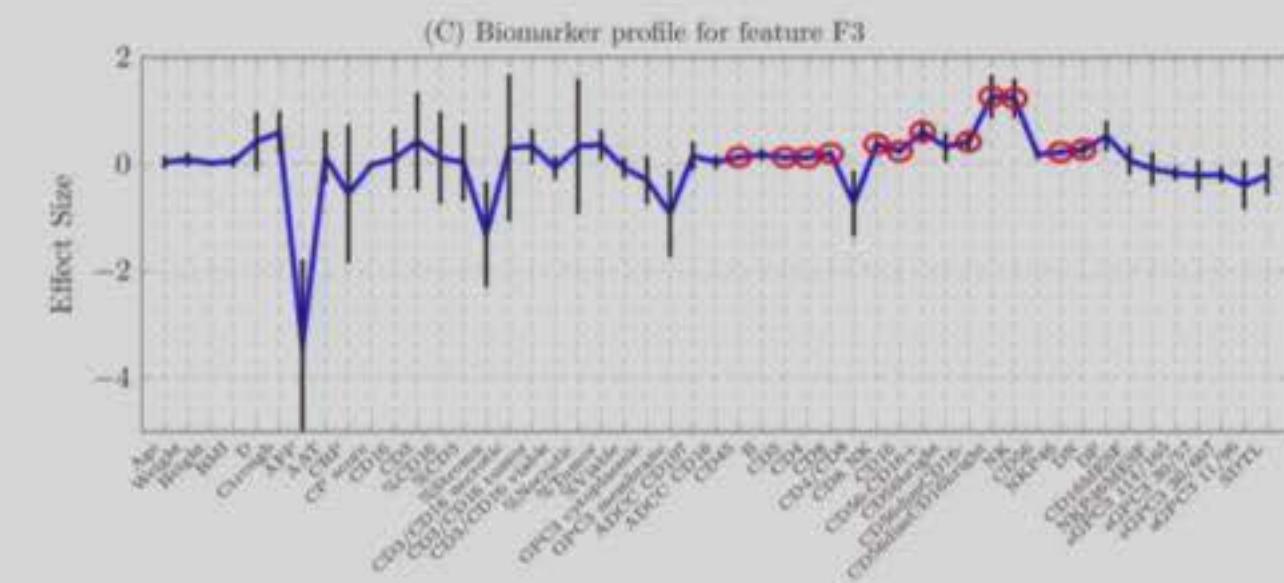


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- ▶ **Conclusion:** no evidence for treatment effectiveness



[BMC CANCER, 2019]

- ▶ **Method:** novel Bayesian approach
- ▶ **Conclusion:** treatment effective for subgroup

WHY IS THIS PROBLEM HARD?

CHALLENGES

WHY IS THIS PROBLEM HARD?

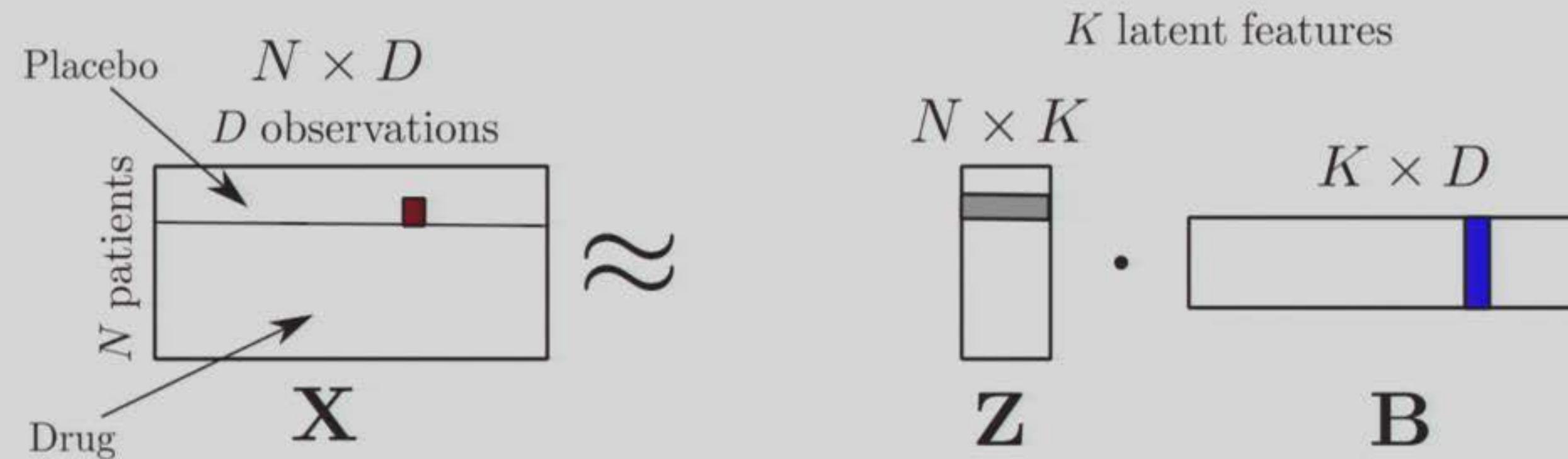
CHALLENGES

1. complex correlations
2. patient heterogeneity
3. few data available
4. very different data types
5. drug effect vs natural response



CHALLENGE 1: HOW TO DEAL WITH COMPLEX CORRELATIONS?

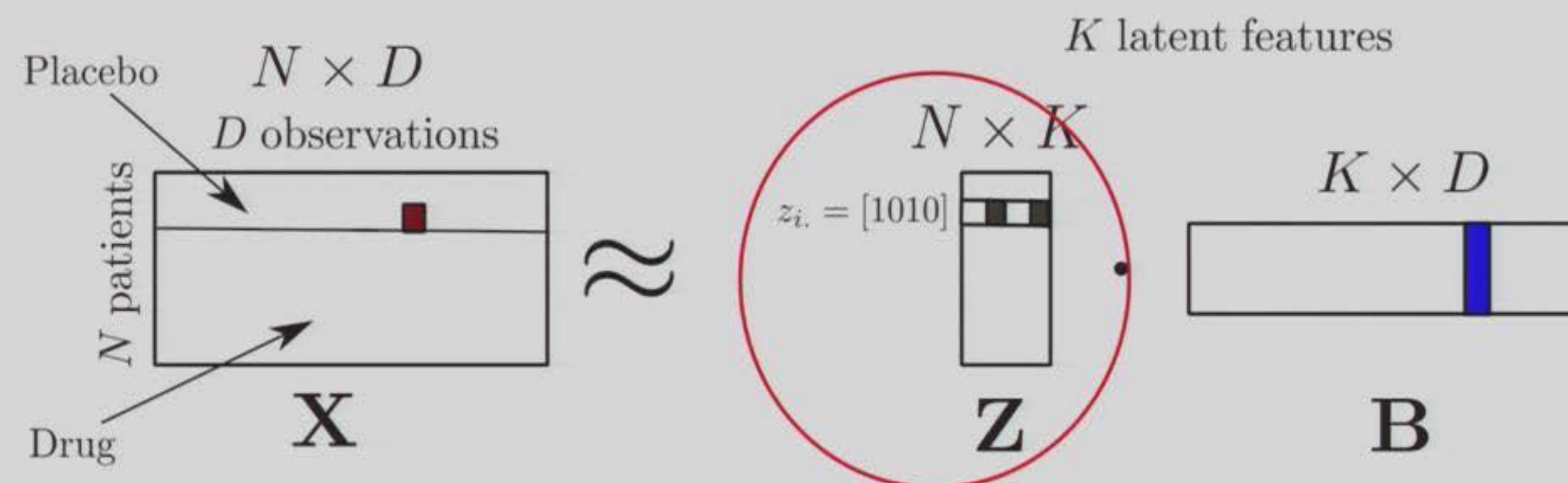
OUR APPROACH: LATENT FEATURE MODEL



$$\blacksquare \quad x_{id} = 173 \text{ ml/dL} = 73 + 0 + 100 \text{ ml/dL}$$

CHALLENGE 2: HOW TO DEAL WITH PATIENT HETEROGENEITY?

OUR APPROACH: BINARY LATENT FEATURE MODEL



- Binary latent feature active: correlation pattern applies to patient

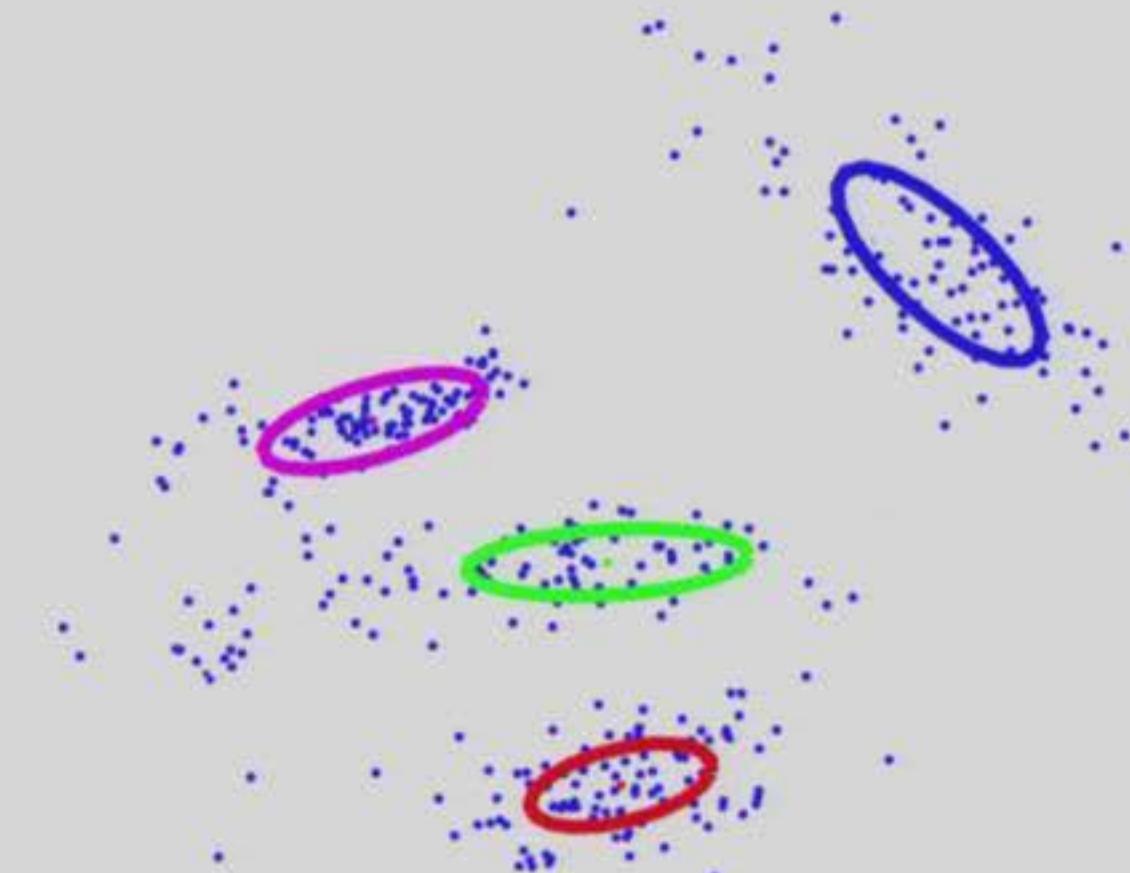
CHALLENGE 3: HOW TO DEAL WITH LITTLE DATA?

OUR APPROACH: **BAYESIAN NONPARAMETRICS**

- ▶ **Bayesian:** to handle uncertainty

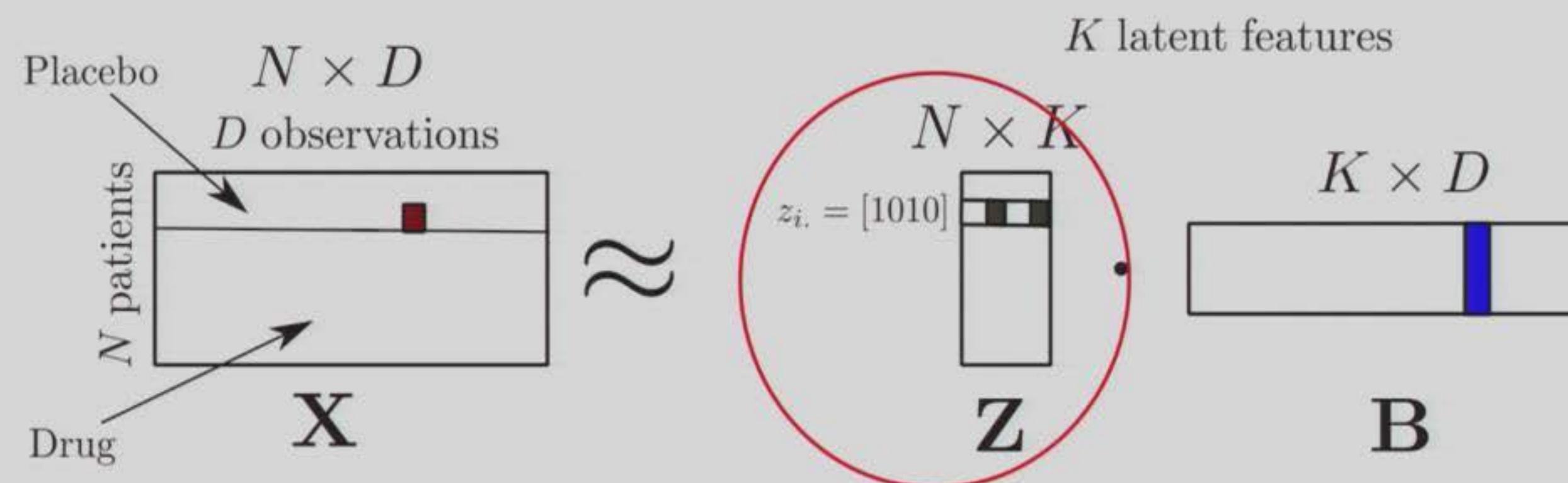
$$p(\text{parameters}|\text{data}) \propto p(\text{data}|\text{parameters})p(\text{parameters})$$

- ▶ **Nonparametric:** to adapt model complexity (hypothesis generation)



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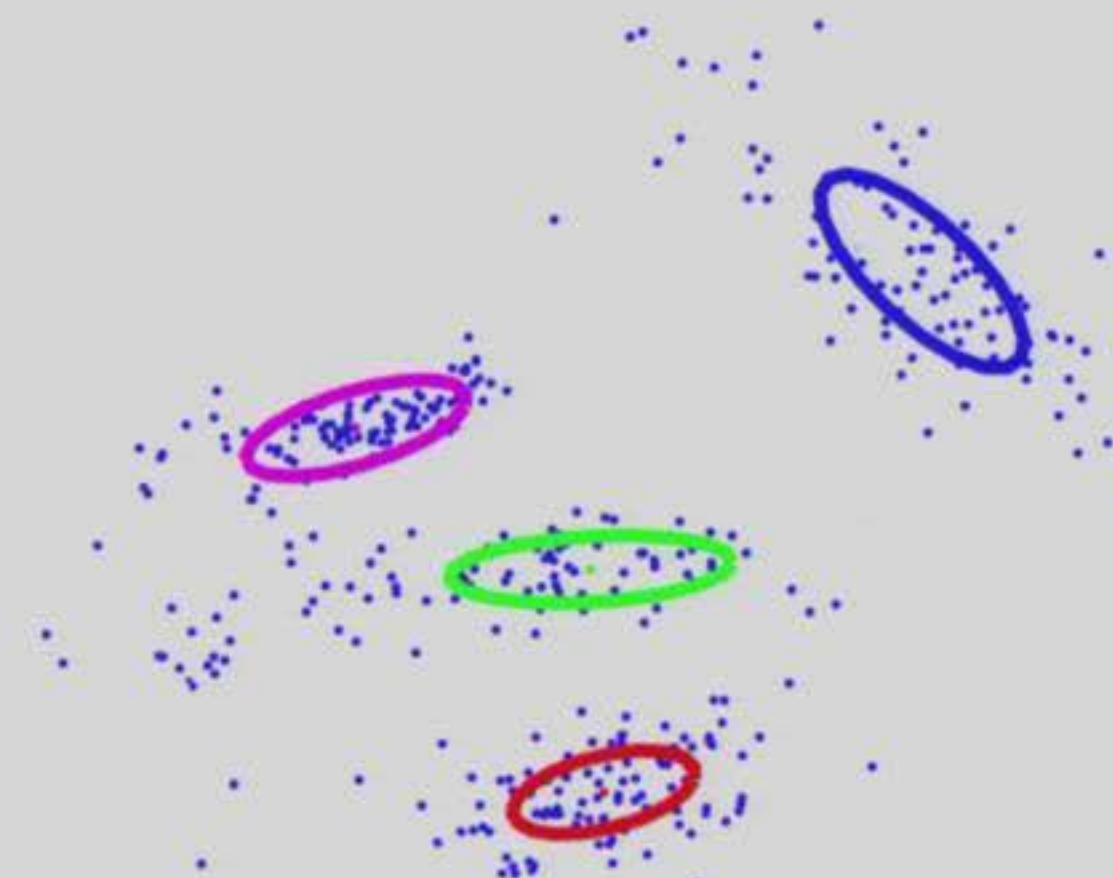
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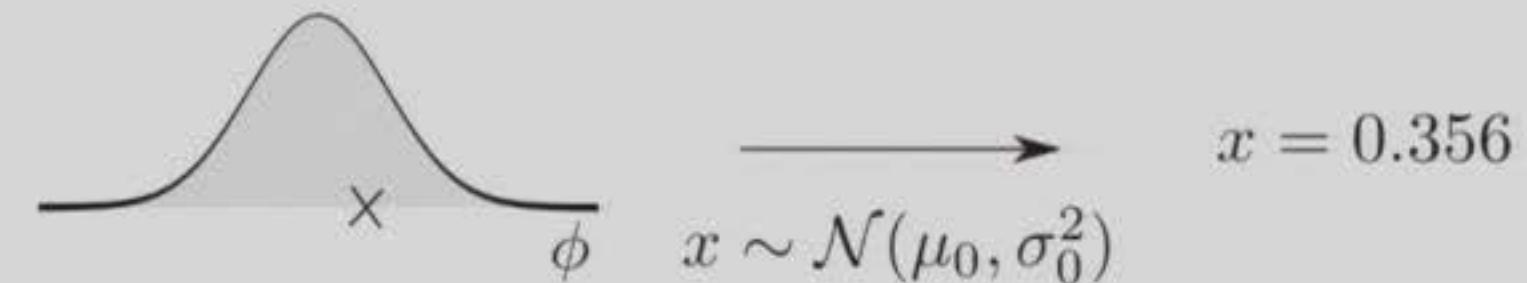
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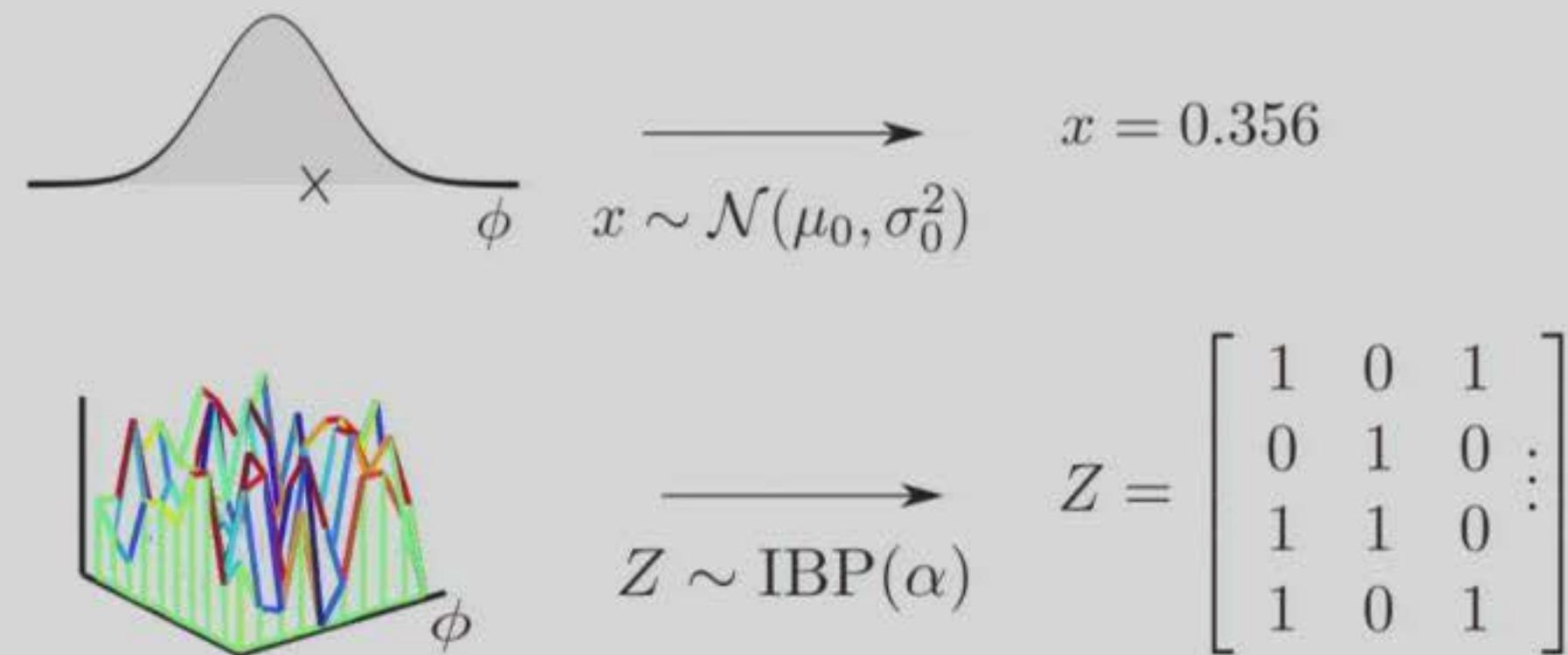
- ▶ Indian Buffet Process [Ghahramani et.al, 2006]



CHALLENGE 3: HOW TO DEAL WITH LITTLE DATA?

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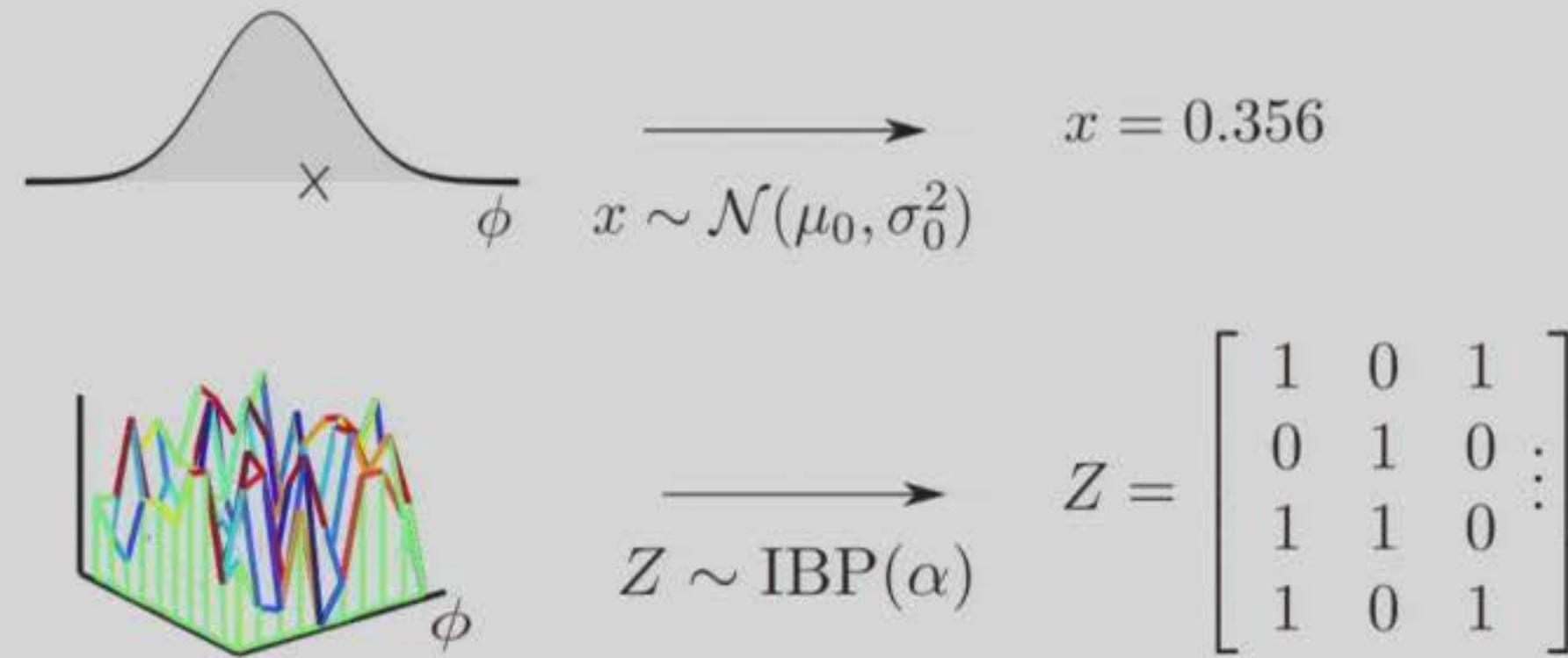
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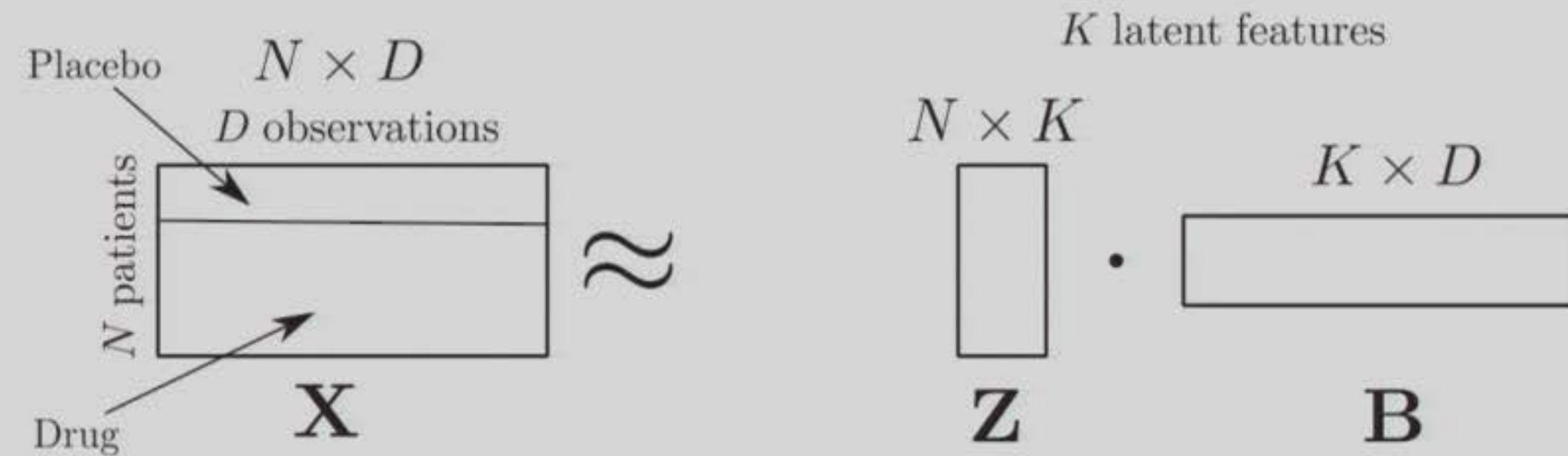
- ▶ Indian Buffet Process [Ghahramani et.al, 2006]



- ▶ Prior over binary matrices with infinite number of columns
- ▶ $Z \sim \text{IBP}(\alpha)$, where α : concentration parameter

CHALLENGE 4: HOW TO DEAL WITH DIFFERENT DATA TYPES?

OUR APPROACH: USE LINK FUNCTIONS



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OUR APPROACH: USE LINK FUNCTIONS

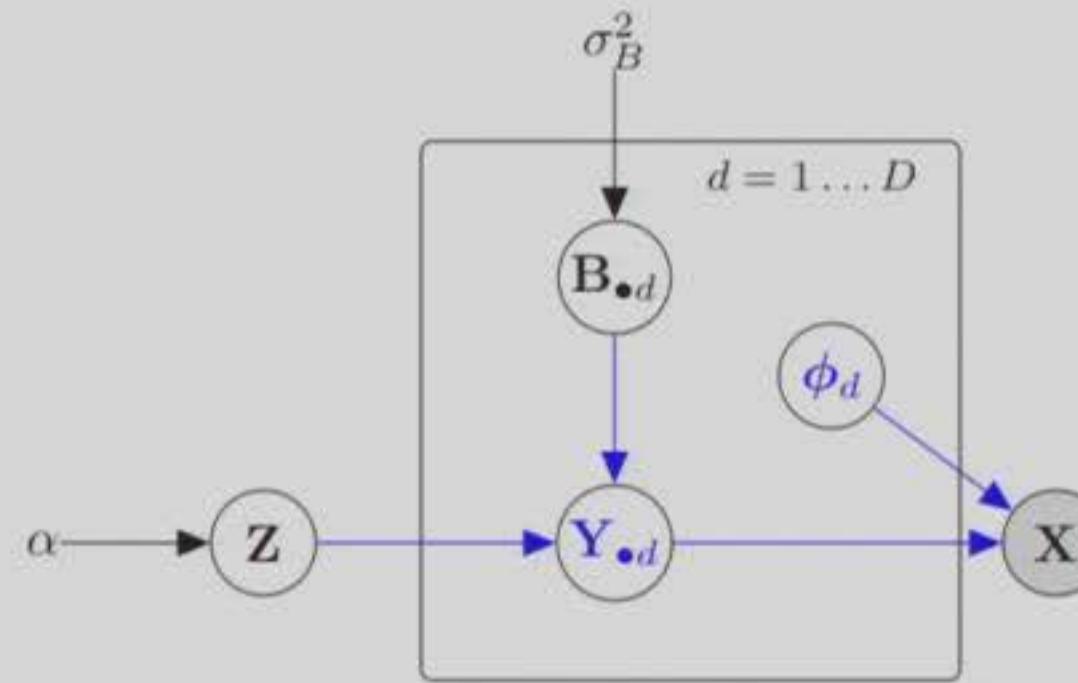
$$\text{Placebo} \quad N \times D \\ N \text{ patients} \quad D \text{ observations} \\ \text{Drug}$$
$$\approx T \left(\begin{matrix} N \times K \\ Z \end{matrix} \cdot \begin{matrix} K \times D \\ B \end{matrix} \right)$$

The diagram illustrates a matrix factorization approach. On the left, a matrix \mathbf{X} is shown with dimensions $N \times D$, representing N patients and D observations. Arrows point from "Placebo" and "Drug" to the top and bottom rows of \mathbf{X} respectively. To the right, the matrix \mathbf{X} is approximated by a product of two matrices, T and $(\mathbf{Z} \cdot \mathbf{B})$. Matrix \mathbf{Z} has dimensions $N \times K$ and matrix \mathbf{B} has dimensions $K \times D$. The label "K latent features" is positioned above the product term.

- ▶ Link functions T_d for each observed feature d
- ▶ Each T_d depends on type of data

SOLVING ALL 4 CHALLENGES TOGETHER

GENERAL LATENT FEATURE MODEL (GLFM) [JMLR, 2020]



$$\begin{aligned}x_{nd} &= T_d(y_{nd}; \phi_d) \\y_{nd} | \mathbf{Z}, \mathbf{B} &\sim \mathcal{N}(\mathbf{Z}_{n\bullet} \mathbf{B}_{\bullet d}, \sigma_y^2) \\B_{kd} &\sim \mathcal{N}(0, \sigma_B^2) \\Z &\sim \text{IBP}(\alpha)\end{aligned}$$

Infinite latent feature model for heterogeneous datasets

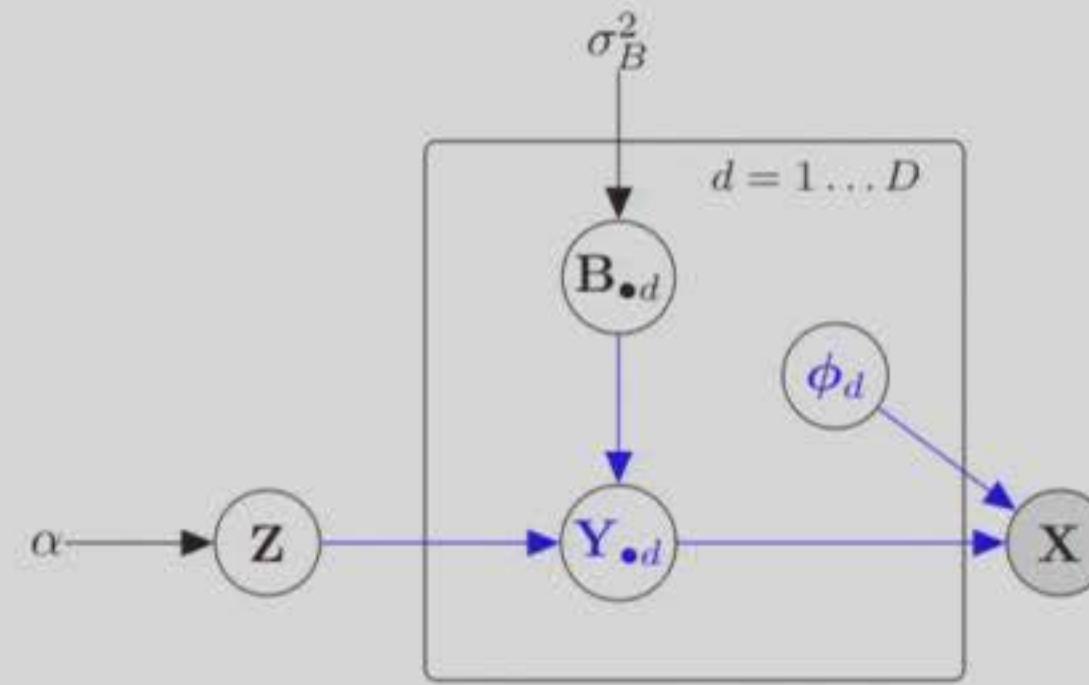
GLFM PACKAGE

- ▶ Open-source python/matlab/R code
- ▶ Discovers complex correlations
- ▶ Is able to deal with data heterogeneity, noisy observations, missing data, different data types
- ▶ Adjusts model complexity

<https://github.com/ivaleraM/GLFM>

SOLVING ALL 4 CHALLENGES TOGETHER

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REVISITING LIST OF CHALLENGES

CHALLENGES

1. complex correlations
2. patient heterogeneity
3. few data available
4. very different data types
5. drug effect vs natural response



CHALLENGE 5: DRUG EFFECT VS NATURAL RESPONSE?

OUR APPROACH: STRUCTURED PRIOR

$$\text{Placebo} \quad N \times D \\ N \text{ patients} \quad D \text{ observations}$$

$\approx T \left(\begin{matrix} N \times K & \\ \textbf{Z} & \end{matrix} \cdot \begin{matrix} K \times D \\ \textbf{B} \end{matrix} \right)$

K latent features

X

Drug

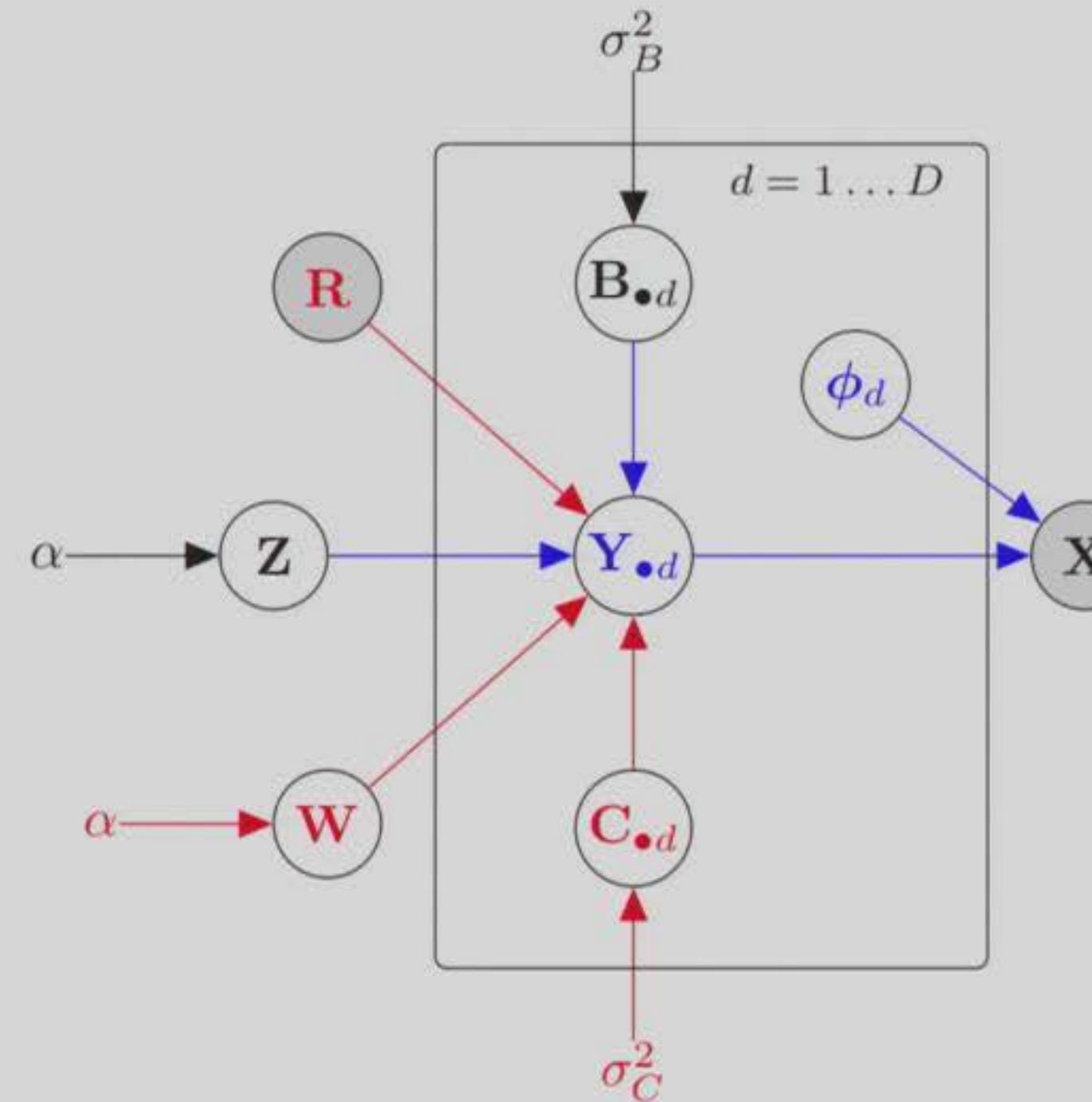
The diagram shows a matrix X representing patient observations. The columns are labeled $N \times D$ and D observations. The rows are labeled N patients and Placebo/Drug. The matrix X is approximated by the product of three matrices: T , Z , and B . Matrix T has dimensions $N \times K$. Matrix Z has dimensions $K \times D$. Matrix B has dimensions $K \times D$. The matrix Z is shown with a vertical bar at the right end labeled '0' (representing zero entries) and a horizontal bar at the bottom labeled 'B' (representing treatment-specific latent features). The matrix B is shown with a horizontal bar at the top labeled 'K latent features'.

TREATMENT-SPECIFIC LATENT FEATURES

- ▶ can only activate for patients in treatment arm
- ▶ number learned automatically

HOW TO DISTINGUISH DRUG EFFECT VS NATURAL RESPONSE?

CASE-CONTROL INDIAN BUFFET PROCESS (C-IBP) [BMC CANCER, 2019]



R_n : drug indicator por patient n

$$x_{nd} = T_d(y_{nd}; \phi_d)$$

$$y_{nd}|Z, W, B, C, R \sim$$

$$\mathcal{N}(Z_n \bullet B_{\bullet d} + \mathbb{1}[R_n = 1] W_n \bullet C_{\bullet d}, \sigma_y^2)$$

$$B_{kd} \sim \mathcal{N}(0, \sigma_B^2)$$

$$Z \sim \text{IBP}(\alpha)$$

$$C_{kd} \sim \mathcal{N}(0, \sigma_C^2)$$

$$W \sim \text{IBP}(\alpha)$$

- ▶ **Inference:** MCMC approach with accelerated Gibbs sampling
- ▶ **Biomarker discovery:** statistical multiple hypothesis testing

RESULTS: SUBPOPULATIONS

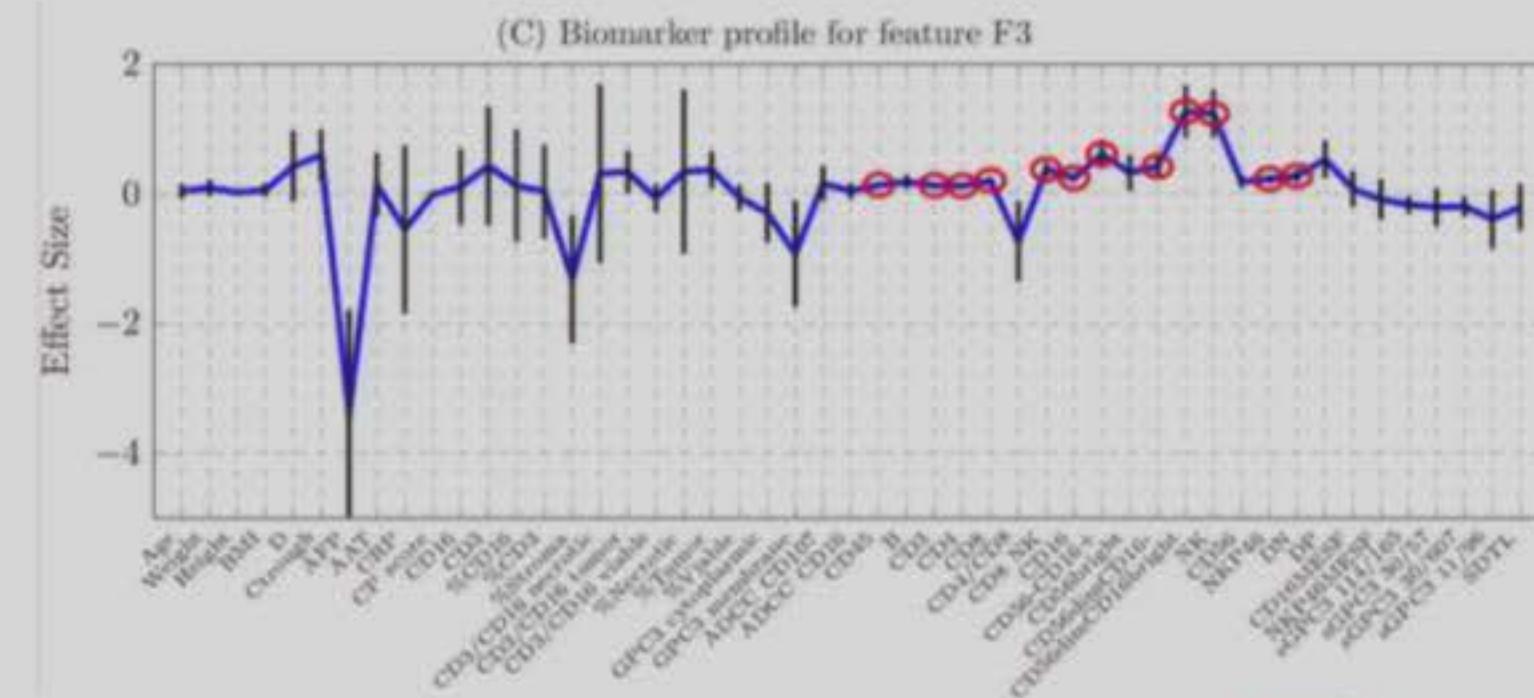
GPC3 Antibody Treatment against Liver Cancer (J. Hepatology. 2016 Apr, Abou-Alfa et.al.)

- ▶ 180 patients: 60 took a placebo, 120 took the drug
- ▶ PFS: Progression Free Survival

Sub-population	Drug Identifier	F1 F2 F3			Size (number of patients)	Mean PFS (months)	Median PFS (months)
		F1	F2	F3			
1.	0	0	0	0	33.37	3.06	1.65
2.	0	0	1	0	4.07	2.29	2.24
3.	0	1	0	0	17.84	2.72	1.81
4.	0	1	1	0	4.72	7.05	7.18
5.	1	0	0	0	51.52	3.22	2.55
6.	1	0	0	1	16.77	4.17	3.65
7.	1	0	1	0	8.38	1.74	1.33
8.	1	0	1	1	2.07	2.69	2.65
9.	1	1	0	0	29.88	3.36	2.03
10.	1	1	0	1	4.90	4.44	4.34
11.	1	1	1	0	4.53	6.31	5.31
12.	1	1	1	1	1.94	10.04	10.01

RESULTS: BIOMARKER DISCOVERY

TREATMENT-SPECIFIC FEATURE F3



M. F. Pradier, B. Reis, L. Jukafsky, F. Milletti, T. Ohtomo, F. Perez-Cruz, and O. Puig. Case-control Indian Buffet Process identifies biomarkers of response to Codrituzumab. *BMC Cancer.* 2019.

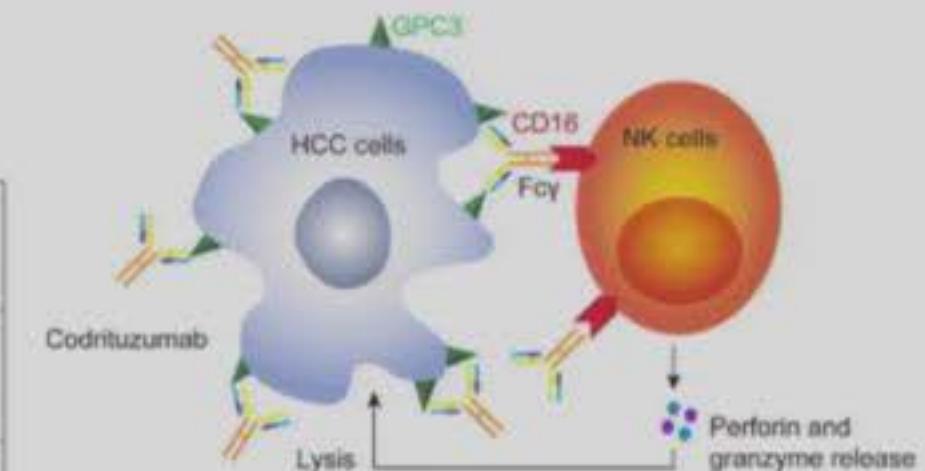


Diagram: Mechanism of codrituzumab-induced antibody-dependent cytotoxicity through the interaction of Fc CD16 in NK cells

What did we find?

- Subgroup for which treatment is especially effective
- Relevant biomarkers (drug acting as expected)

RESULTS: SUBPOPULATIONS

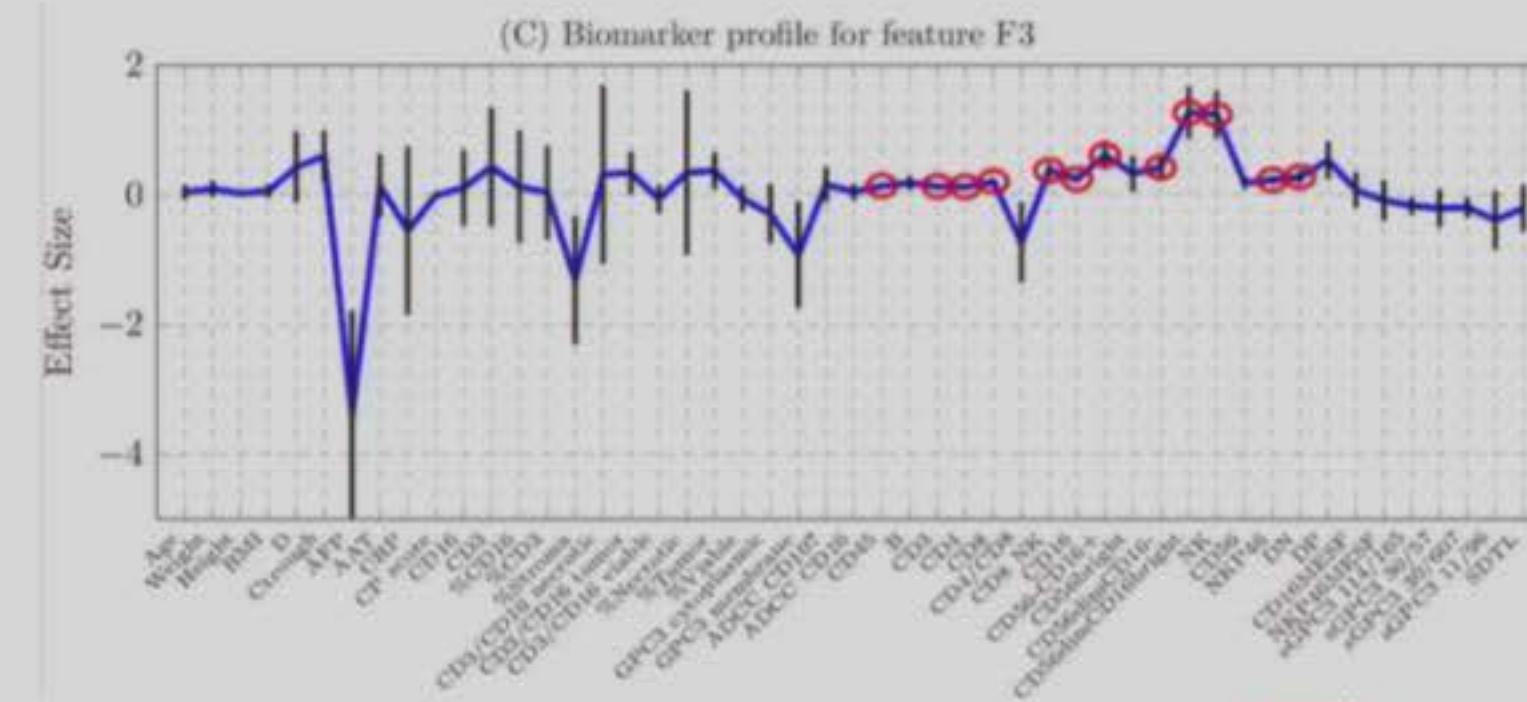
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2.	0	0	1	0	4.07	2.29	2.24
3.	0	1	0	0	17.84	2.72	1.81
4.	0	1	1	0	4.72	7.05	7.18
5.	1	0	0	0	51.52	3.22	2.55
6.	1	0	0	1	16.77	4.17	3.65
7.	1	0	1	0	8.38	1.74	1.33
8.	1	0	1	1	2.07	2.69	2.65
9.	1	1	0	0	29.88	3.36	2.03
10.	1	1	0	1	4.90	4.44	4.34
11.	1	1	1	0	4.53	6.31	5.31
12.	1	1	1	1	1.94	10.04	10.01

RESULTS: BIOMARKER DISCOVERY

TREATMENT-SPECIFIC FEATURE F3



M. F. Pradier, B. Reis, L. Jukofsky, F. Milletti, T. Ohtomo, F. Perez-Cruz, and O. Puig. Case-control Indian Buffet Process identifies biomarkers of response to Codrituzumab. *BMC Cancer.* 2019.

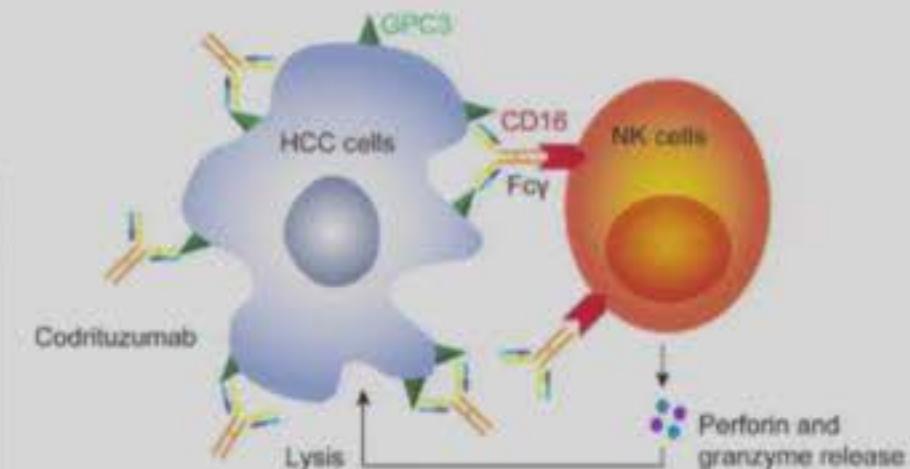
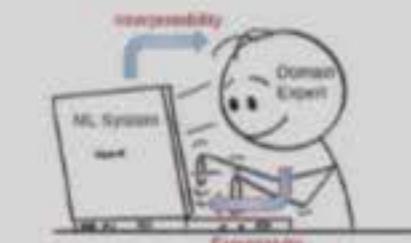


Diagram: Mechanism of codrituzumab-induced antibody-dependent cytotoxicity through the interaction of Fc CD16 in NK cells

What did we find?

- Subgroup for which treatment is especially effective
- Relevant biomarkers (drug acting as expected)

RESULTS: SUBPOPULATIONS

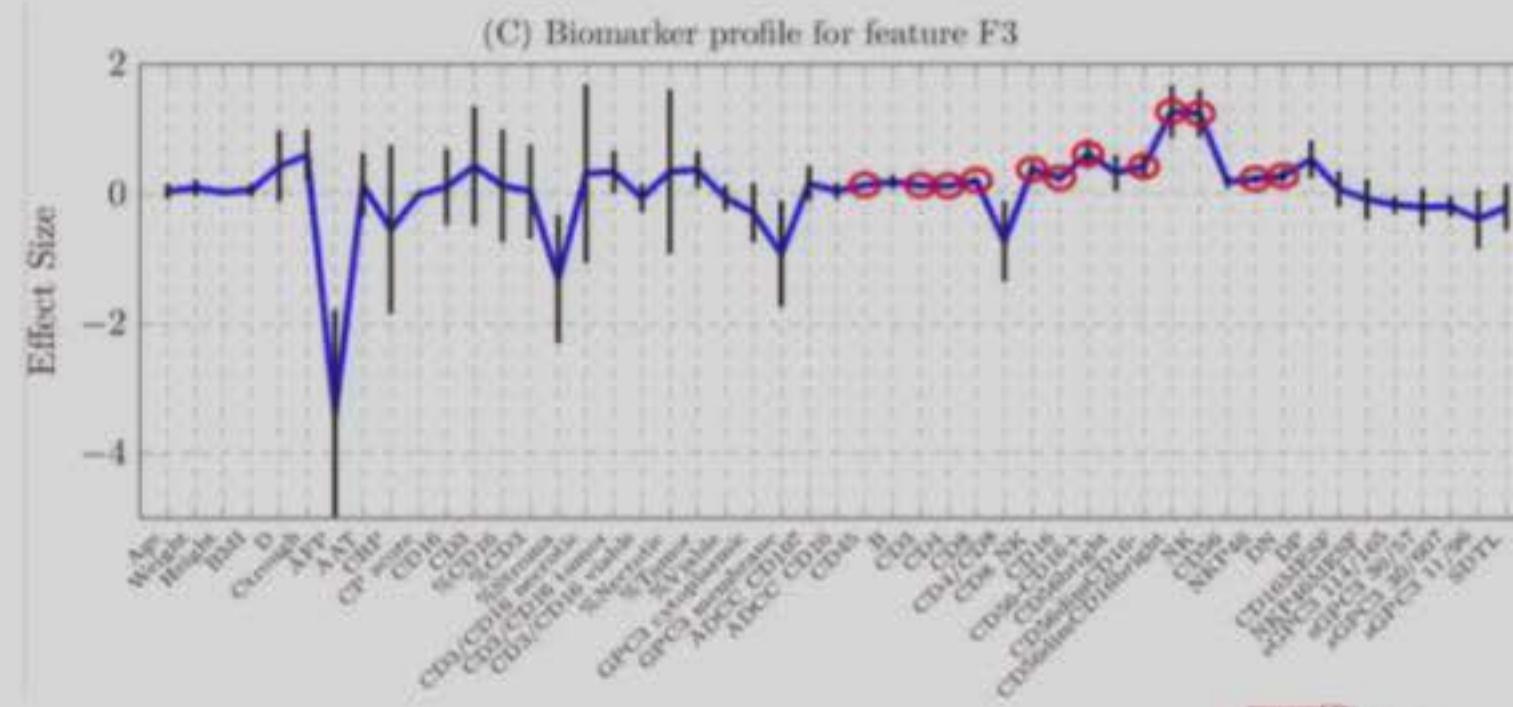
GPC3 Antibody Treatment against Liver Cancer (J. Hepatology. 2016 Apr, Abou-Alfa et.al.)

- ▶ 180 patients: 60 took a placebo, 120 took the drug
- ▶ PFS: Progression Free Survival

Sub-population	Drug Identifier	F1	F2	F3	Size (number of patients)	Mean PFS	Median PFS
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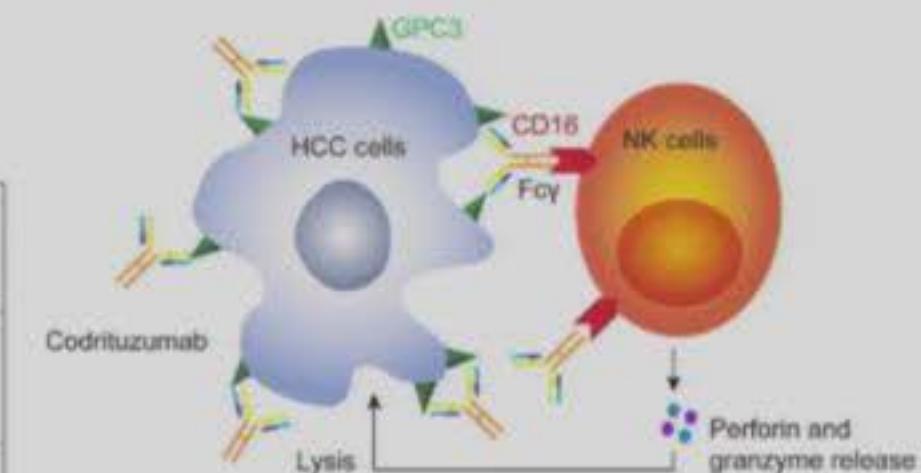


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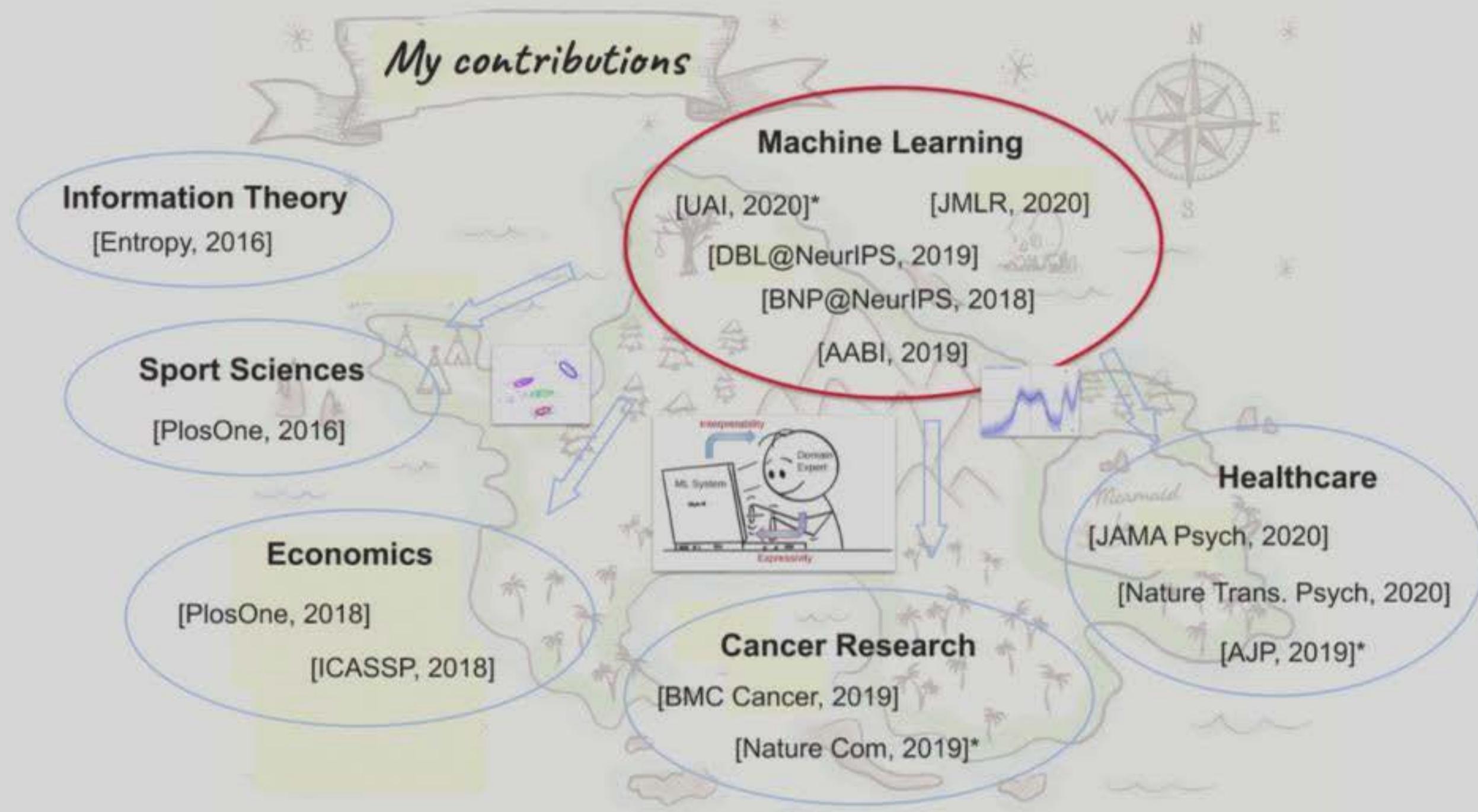
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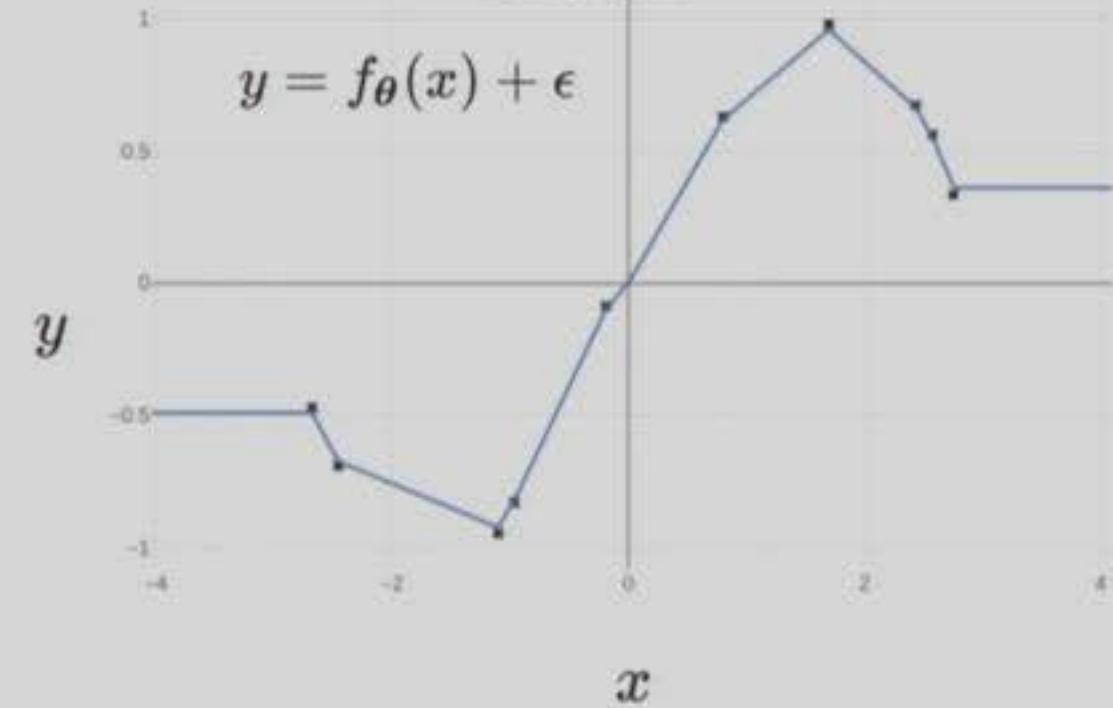
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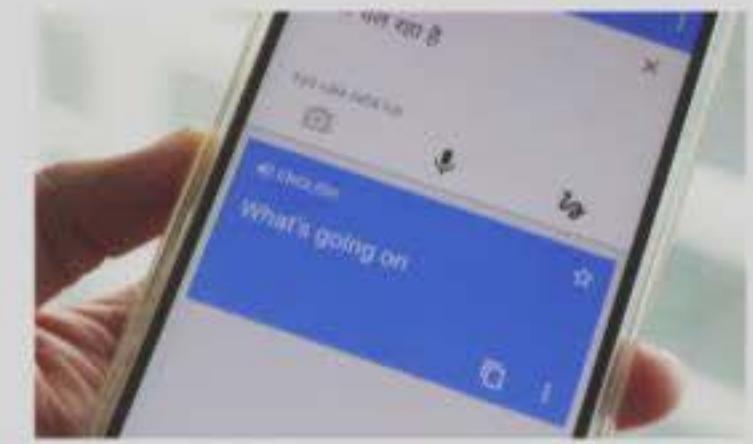
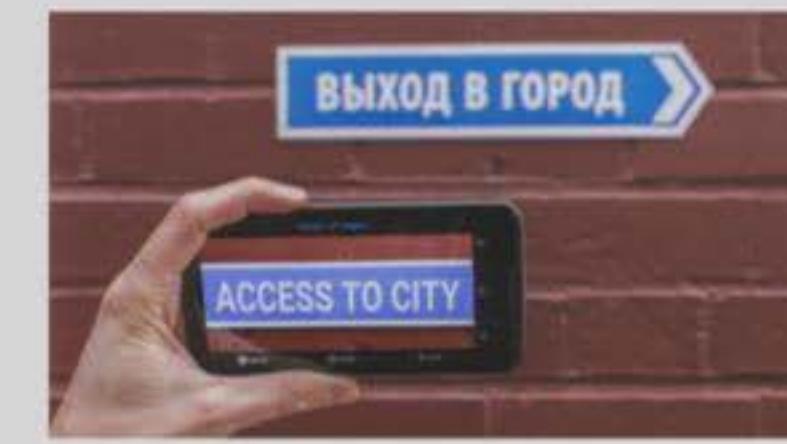
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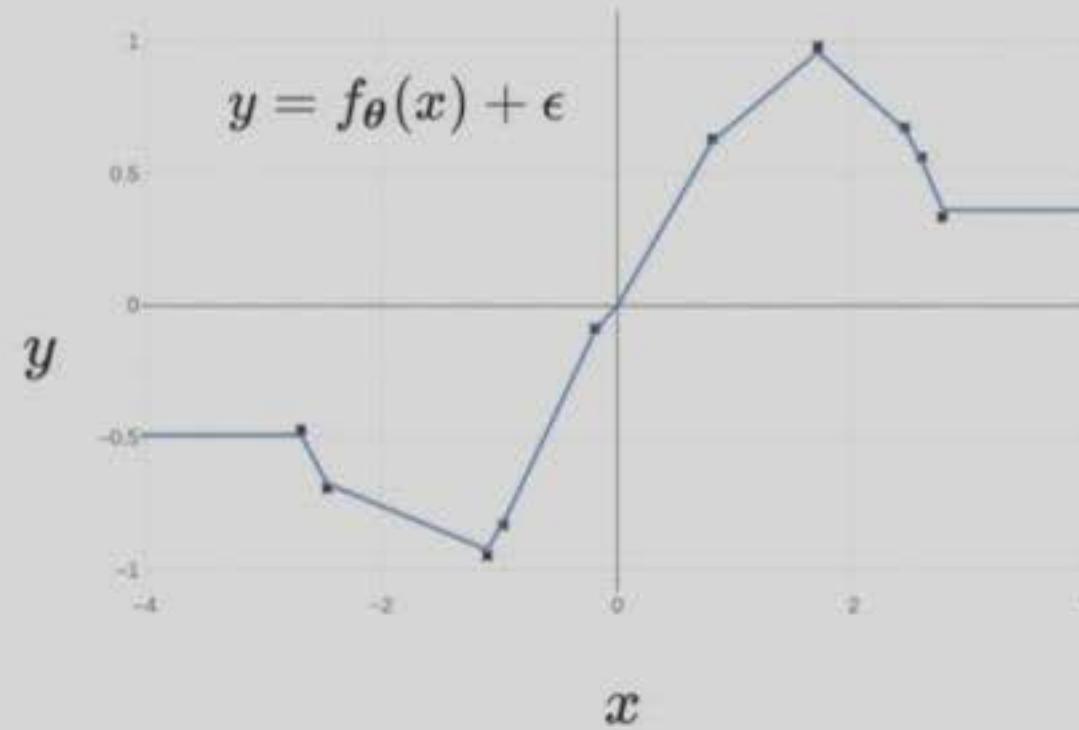
NEURAL NETWORKS (NNs) CAN FIT COMPLICATED TRENDS IN DATA



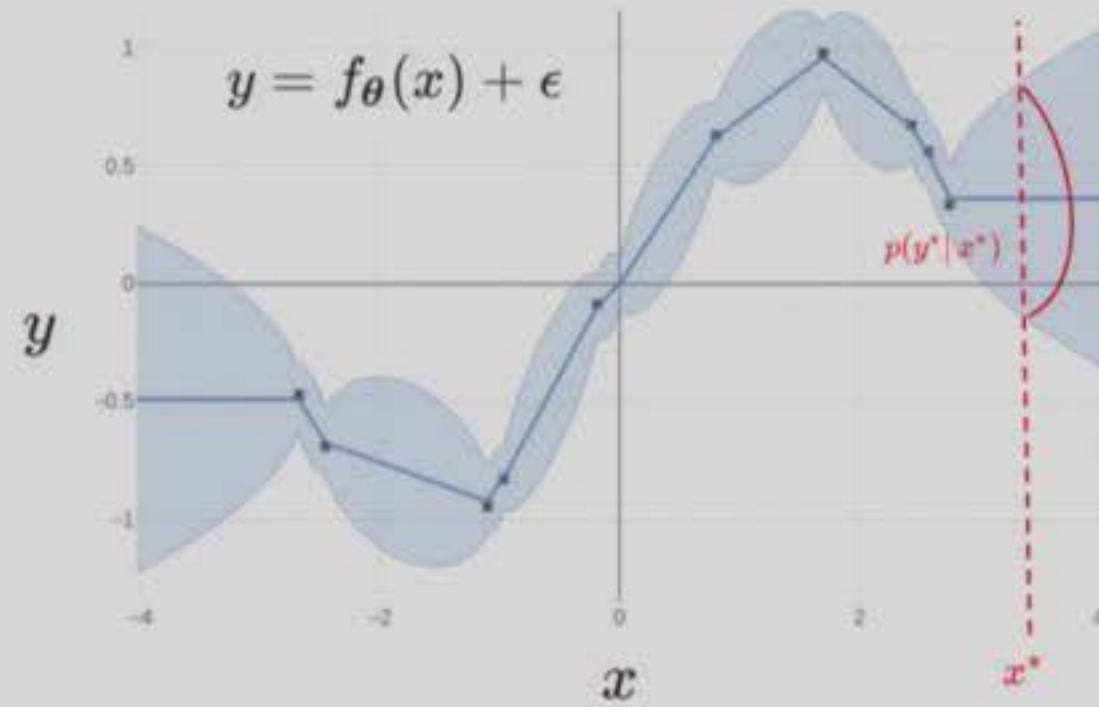
Several success stories...



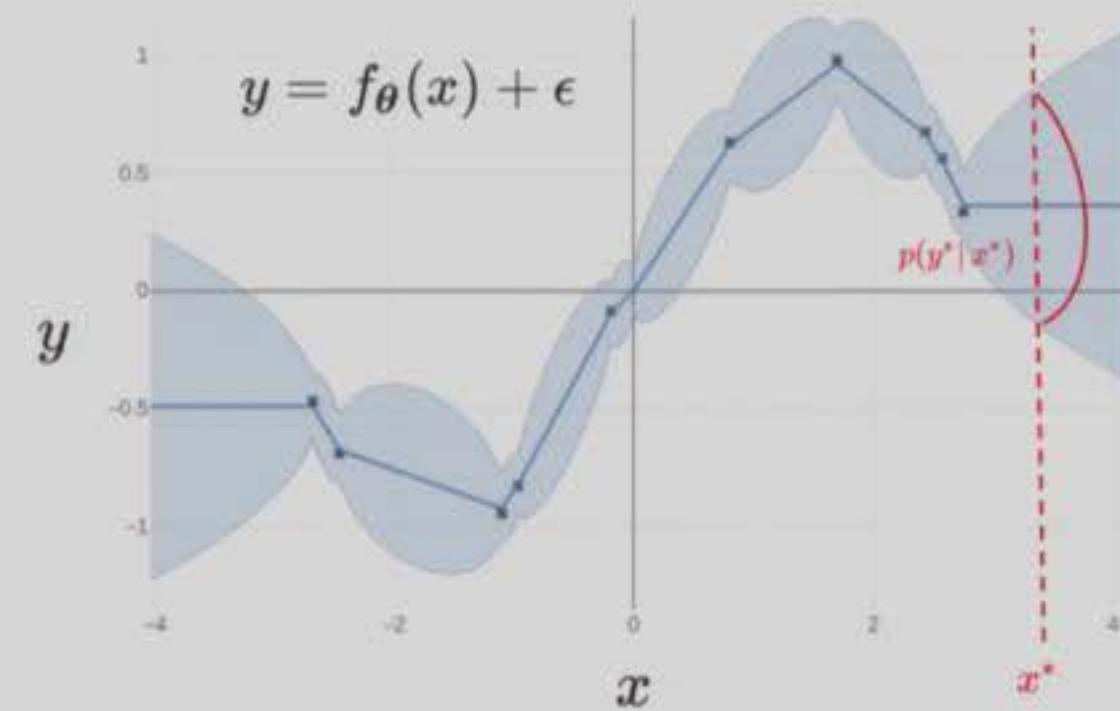
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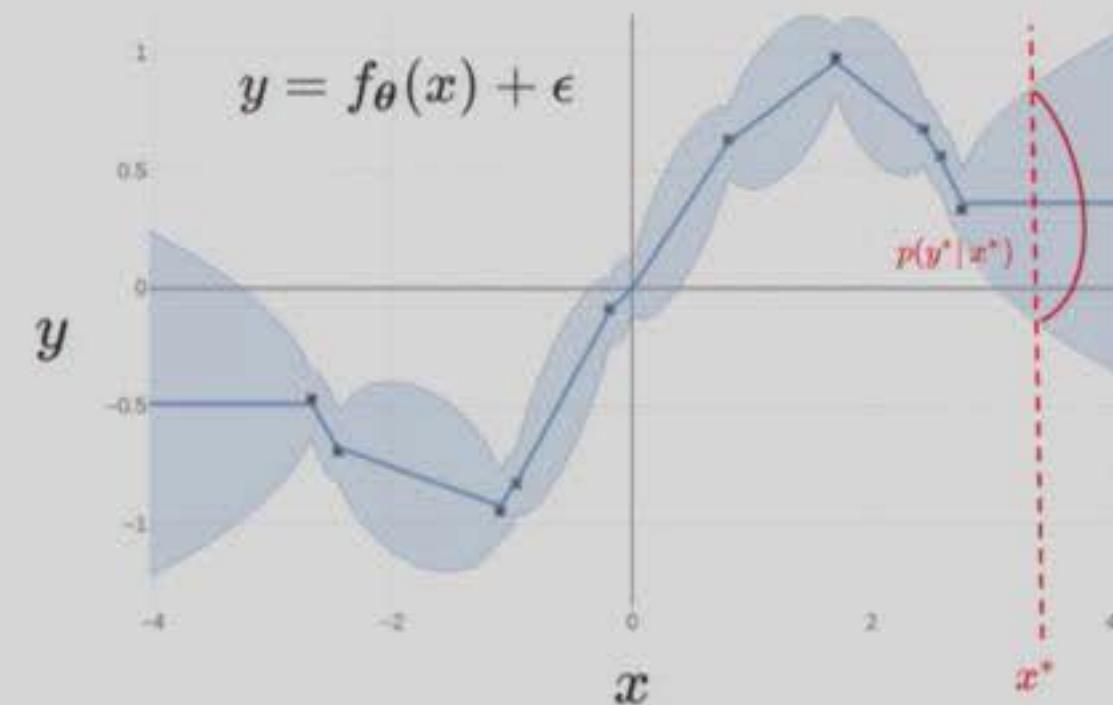
BUT WHAT IF STAKES ARE HIGH?



Uncertainty estimation becomes crucial!



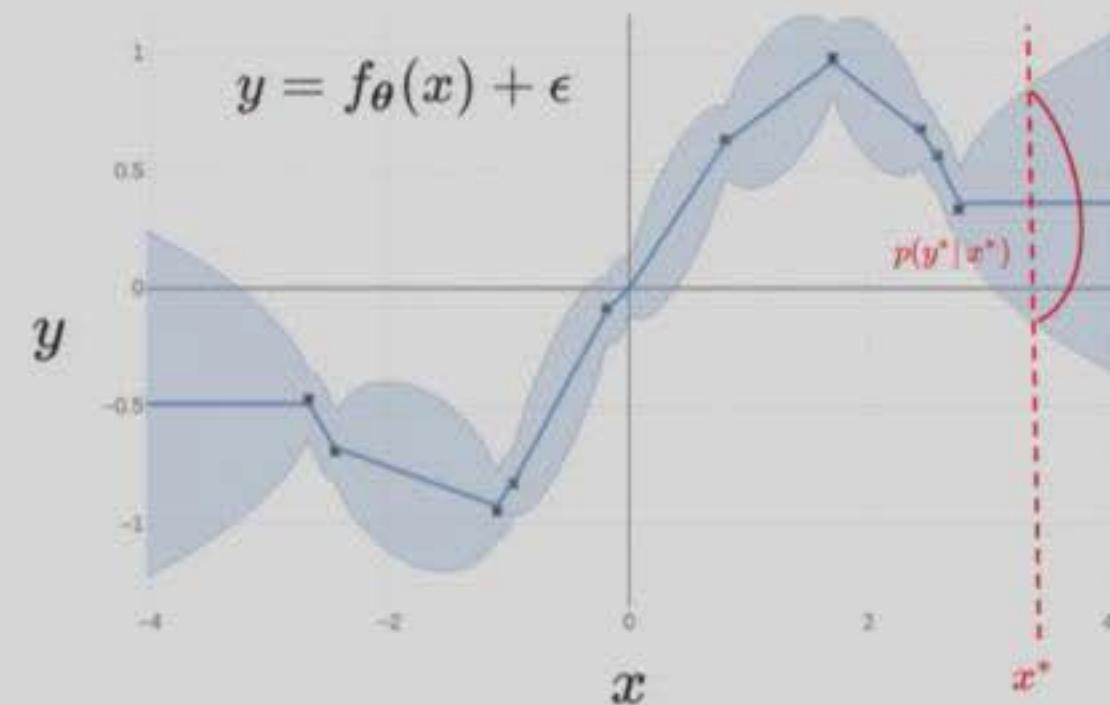
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Some examples of assumptions:

- ▶ Range of heart rate at rest between 60-100 bpm.
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Key point: All of these translate into restrictions on learned functions.

Question: How can we incorporate such desiderata into the model?

AN EASY WAY TO SPECIFY FUNCTIONAL DESIDERATA: GAUSSIAN PROCESSES (GPS)

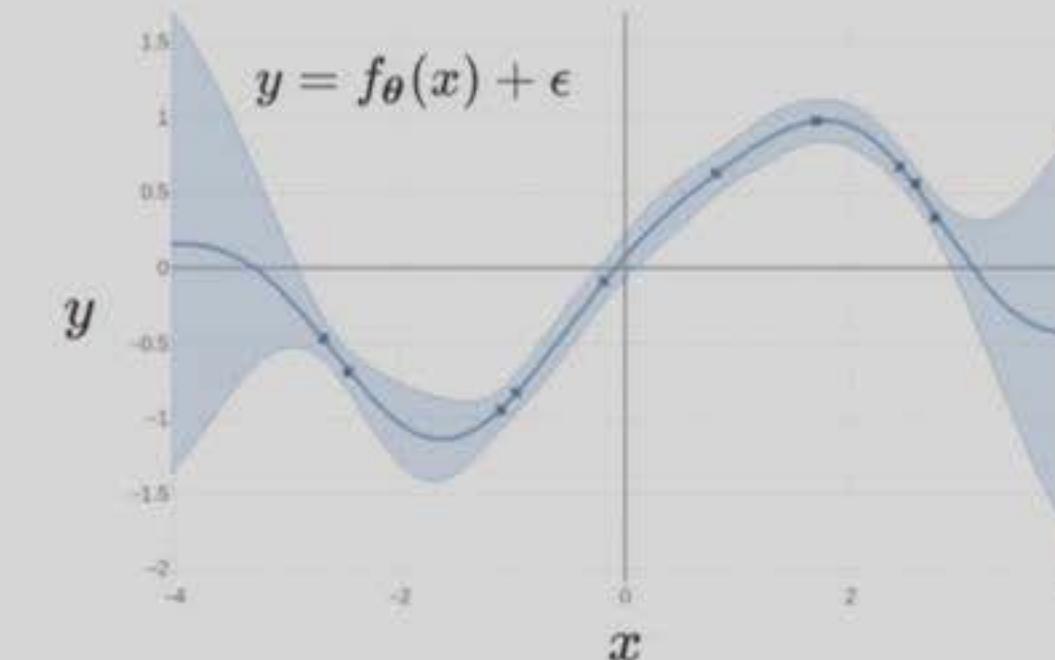
Definition: a Gaussian process is a collection of random variables, any finite number of which have (consistent) Gaussian distributions.

$$f \sim \text{GP}(\mu(\cdot), k(\cdot, \cdot))$$

Example: RBF kernel as covariance function:

$$k(x, x') = \sigma^2 \exp\left(-\frac{(x - x')^2}{2\gamma^2}\right)$$

- ▶ Stationarity
- ▶ Lengthscale
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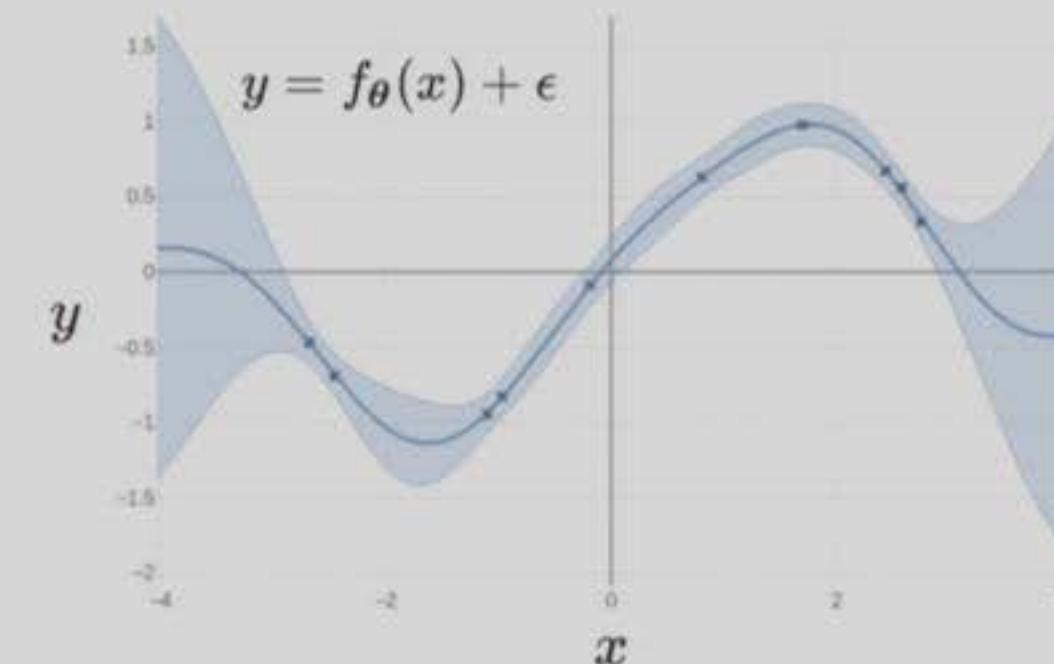
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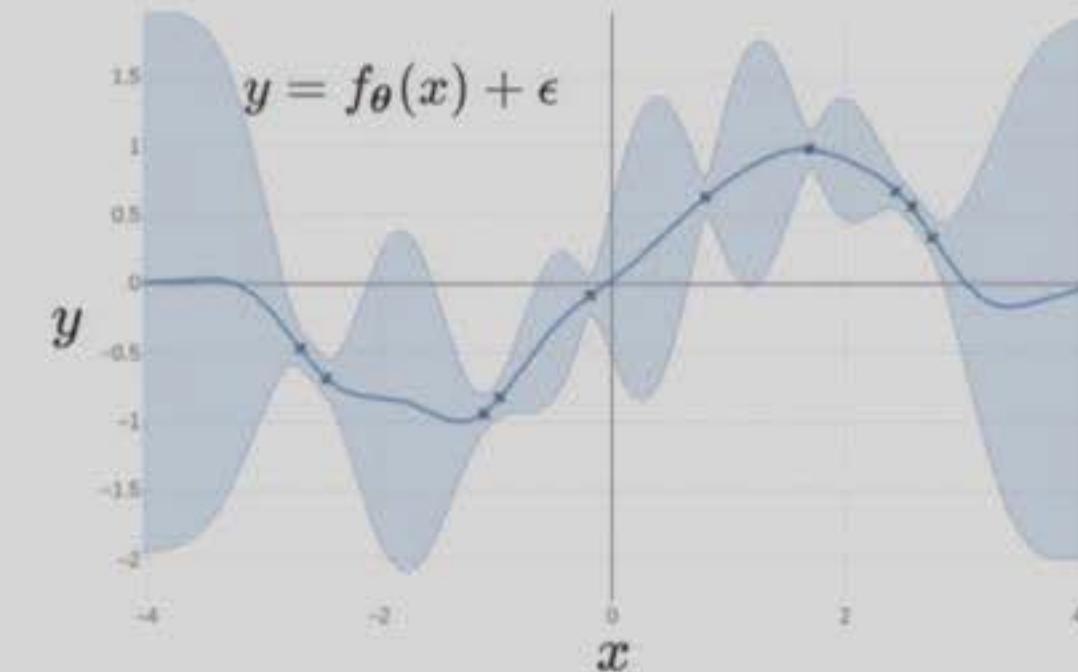
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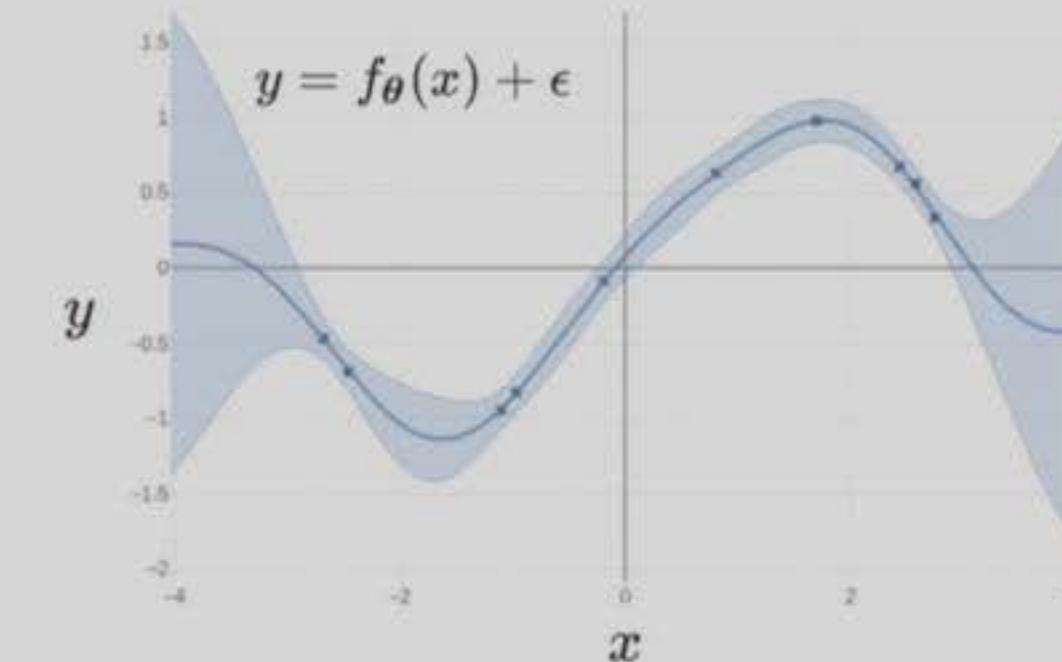
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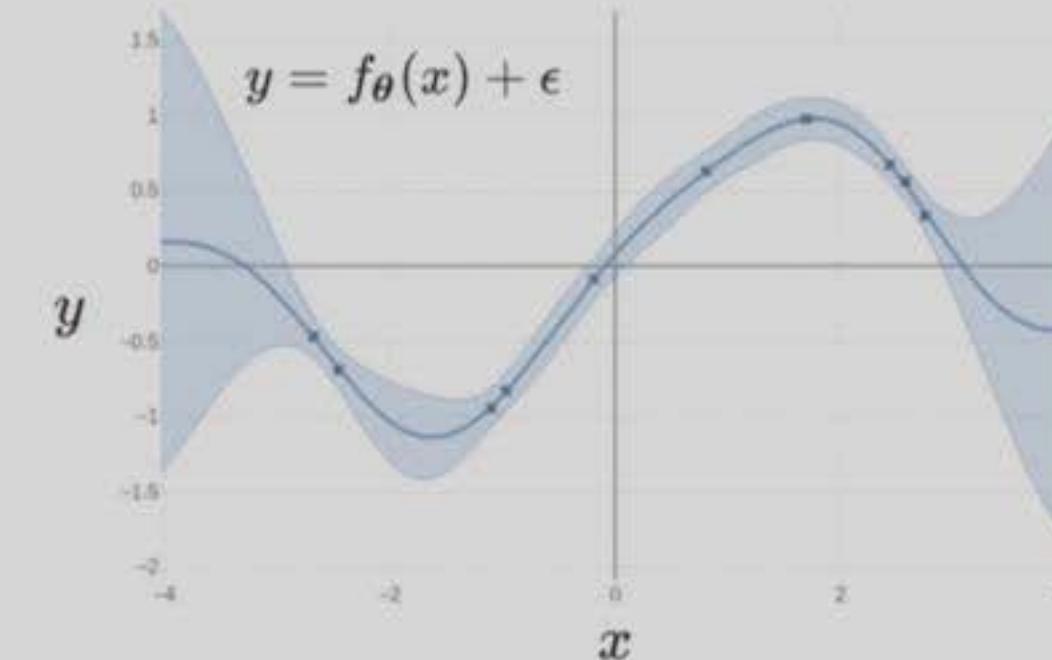
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GPs ARE GREAT, BUT WHAT IF I STILL WANT A NN?

Benefits of NN approaches:

- ▶ widely used (many tools available)
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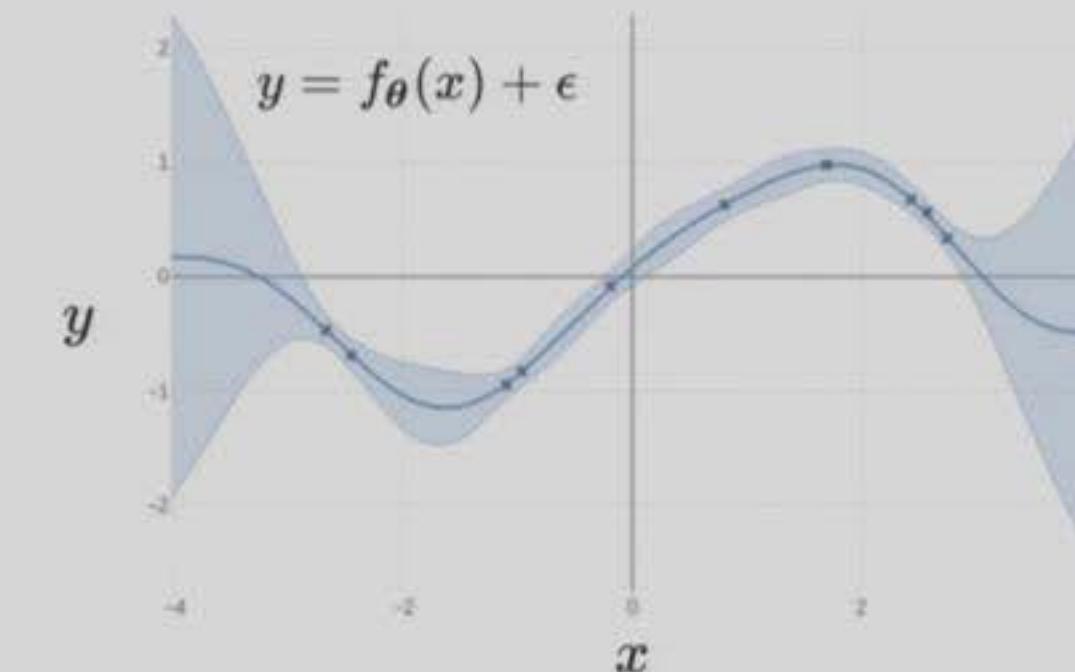
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KEY RESEARCH QUESTIONS:

1. Can we design Bayesian NN priors that encode **stationarity properties** like a GP while retaining the benefits of neural networks?
2. Can we easily specify lengthscale and amplitude variance in a **decoupled** fashion?

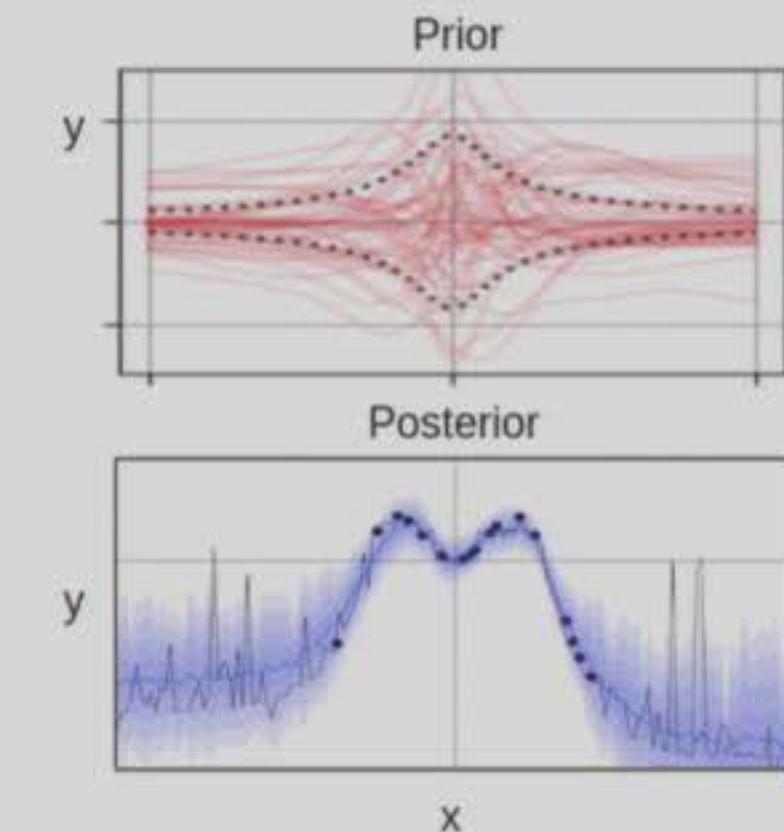
BACKGROUND: BAYESIAN NEURAL NETWORKS

- ▶ Assume prior on network parameters
- ▶ Most common, i.i.d Gaussians

$$\mathbf{y} = f_{\theta}(\mathbf{x}) + \epsilon$$

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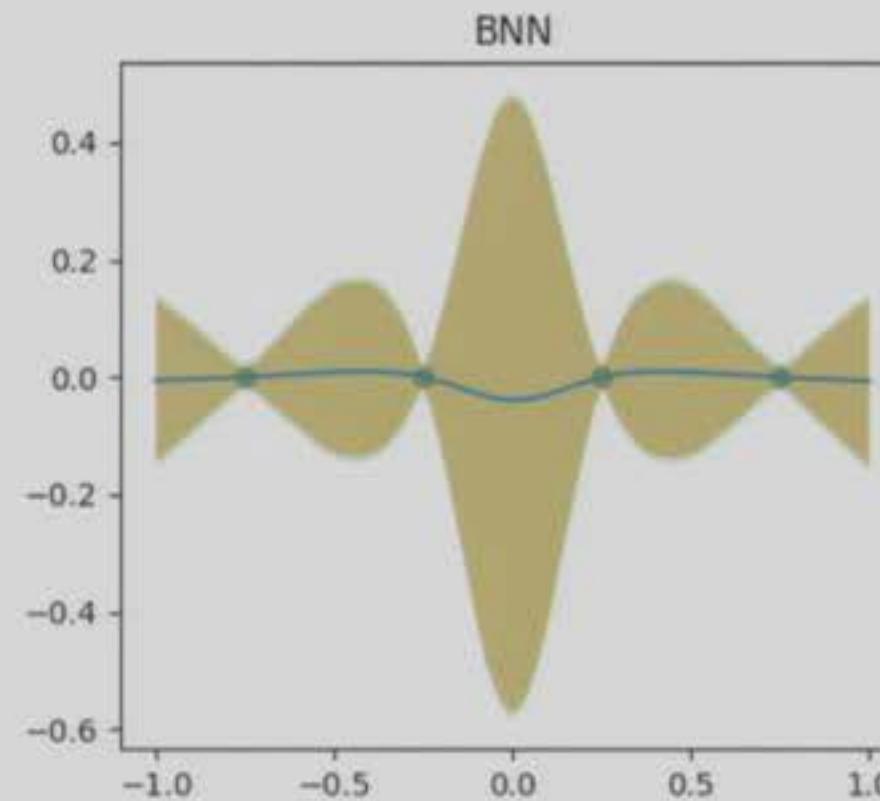
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- ▶ $p(\theta) \implies p(f)$

NOT ONLY HARD TO ENCODE FUNCTIONAL PROPERTIES WITH BNNs; SOME PROPERTIES ARE IMPOSSIBLE TO GET

- ▶ For example, a standard BNN (with RBF activations) is nonstationary in amplitude variance (Williams, 1997)



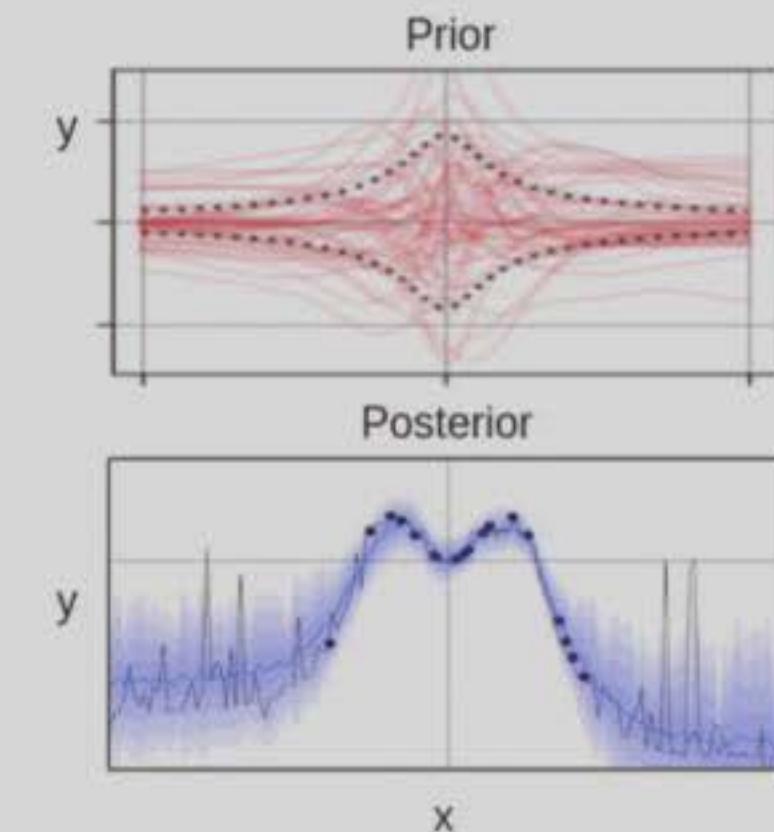
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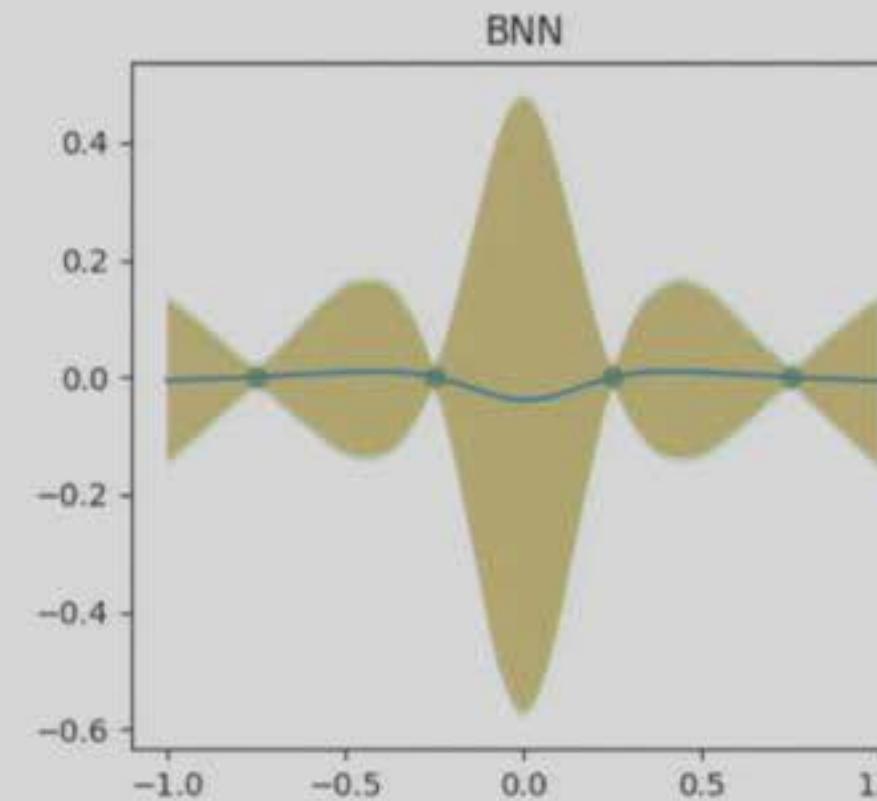


- ▶ $p(\theta) \implies p(f)$

- ▶ But what does a prior over weights mean in function space?
- ▶ Hard to know!

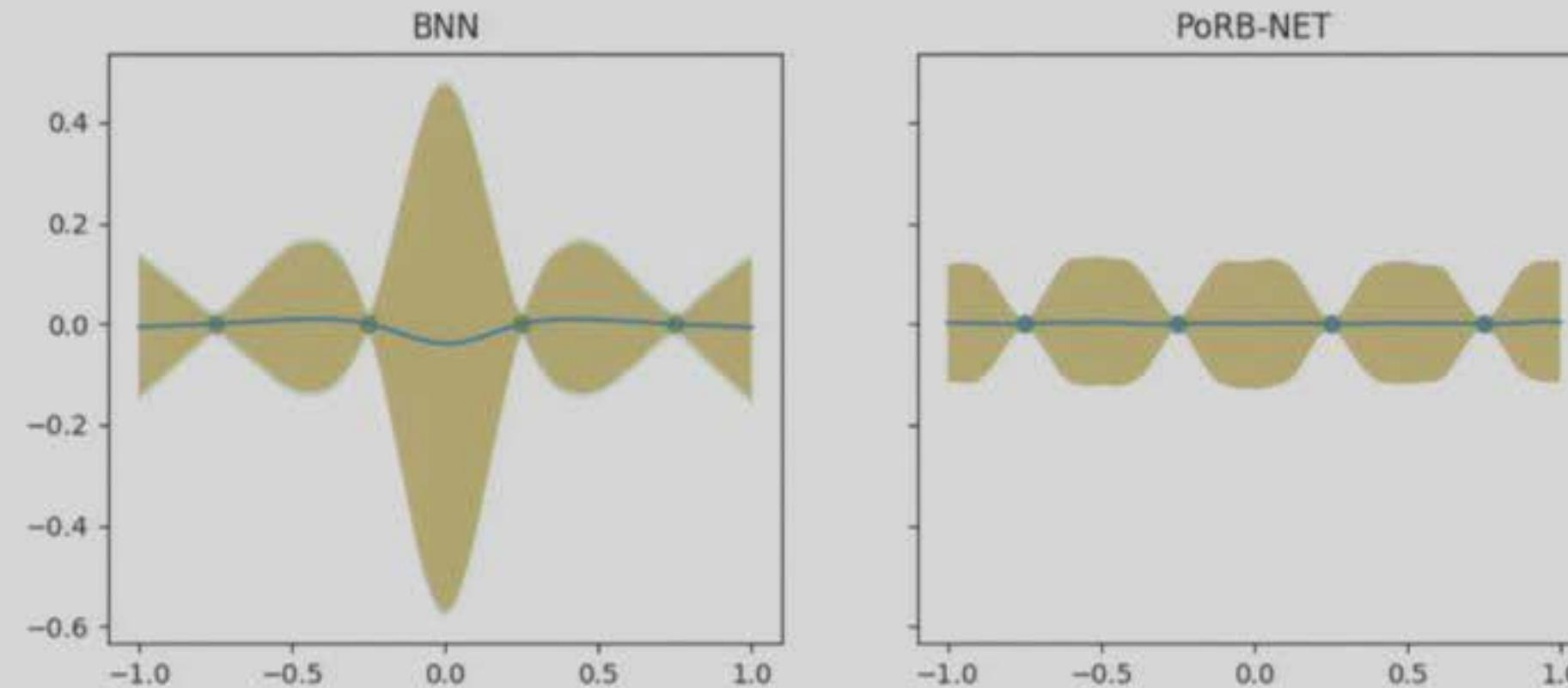
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Question: can we design a Bayesian NN that exhibits stationarity? **Yes!**

RADIAL BASIS FUNCTION NETWORKS (RBFNs)

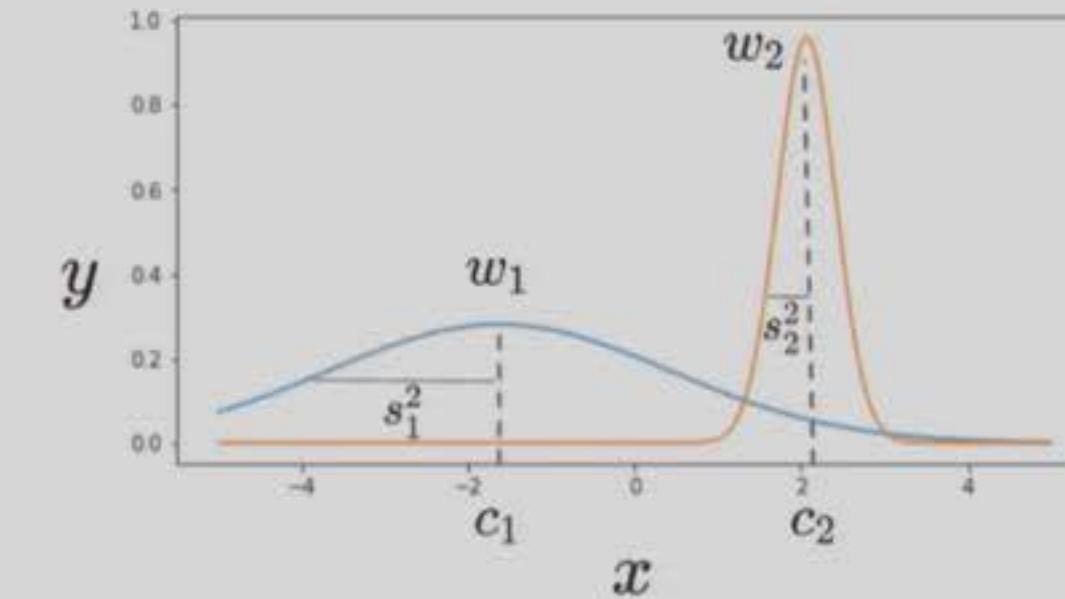
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$$f_{\theta}(x) = b + \sum_{k=1}^K w_k \phi(s_k(x - c_k)),$$

- ▶ s_k : scale
- ▶ c_k : center
- ▶ w_k : output weight
- ▶ b : output bias

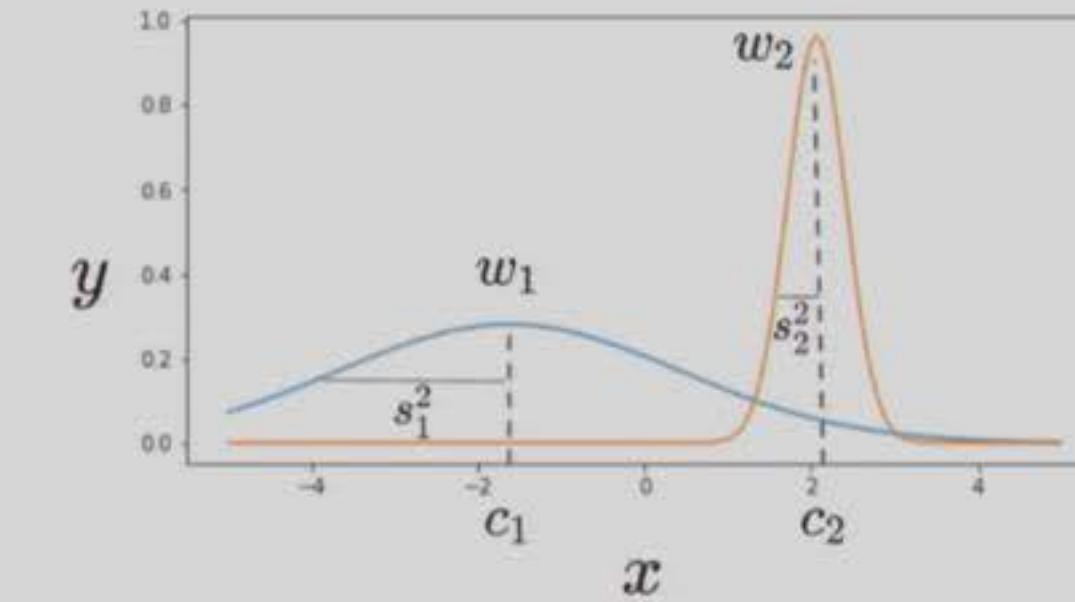


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- ▶ Equivalence to standard BNN exists, but not interpretable!

OUR CONTRIBUTION: POISSON PROCESS RADIAL BASIS FUNCTION NETWORK (PoRB-NET) [UAI, 2020]*

1. We propose an expressive prior for NNs
2. We show desirable properties:
 - 2.1 Stationarity
 - 2.2 Decoupling of lengthscale and amplitude variance
 - 2.3 Consistency
3. We demonstrate successful behavior empirically

(*) submitted

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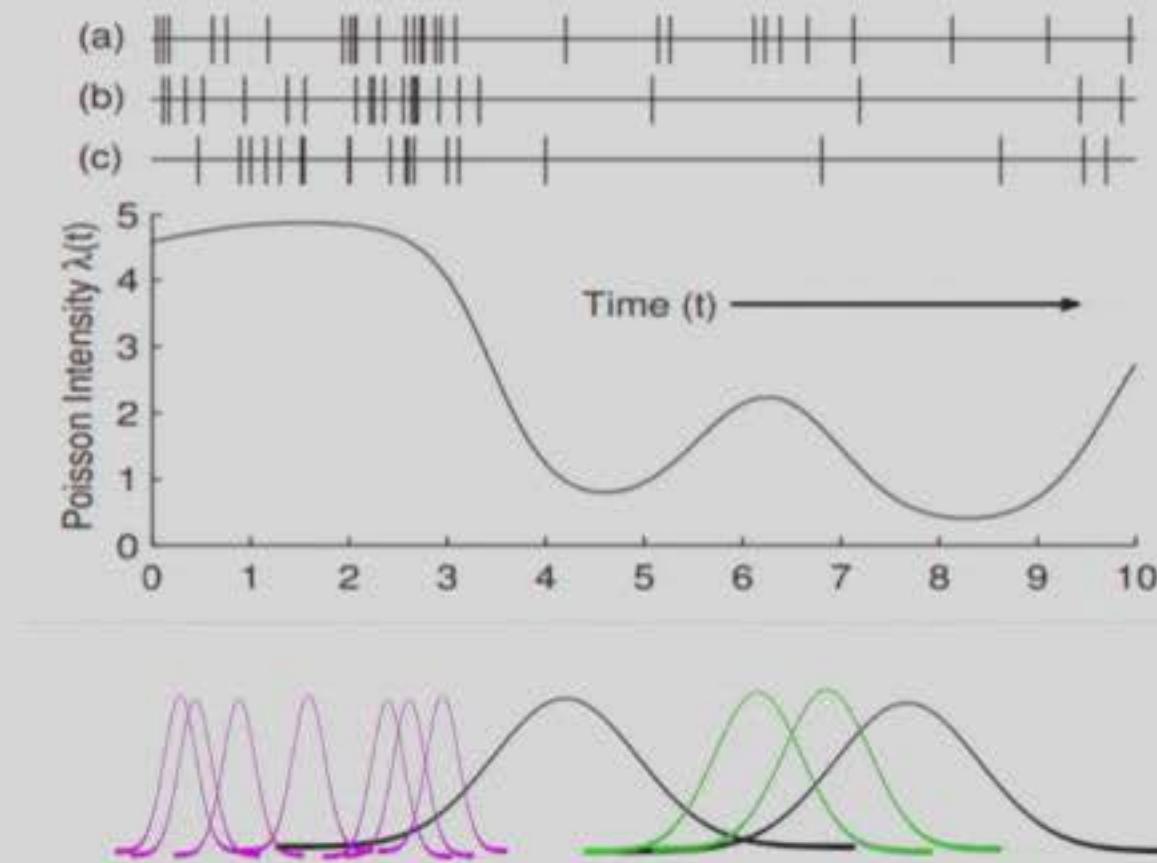
- ▶ Key point: Expressive prior over centers \mathbf{c} and scales $\{s_k\}$.

- ▶ Poisson process priors over \mathbf{c} .

$$\mathbf{c} | \lambda \sim \text{Poisson Process}(\lambda)$$

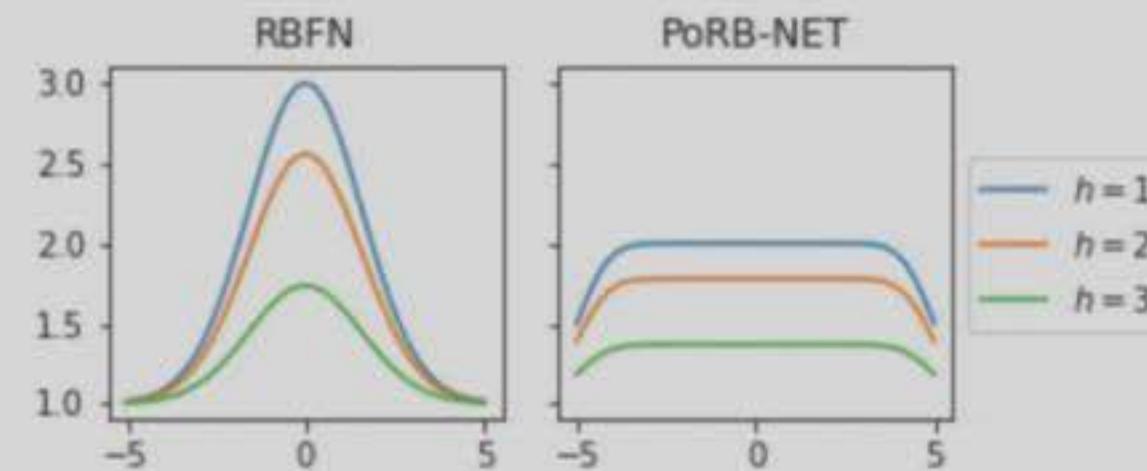
- ▶ Deterministic relationship between \mathbf{c} and $\{s_k\}$.

$$s_k^2 = \lambda^2(c_k)$$



PROPERTIES OF PoRB-NET

1. Stationarity



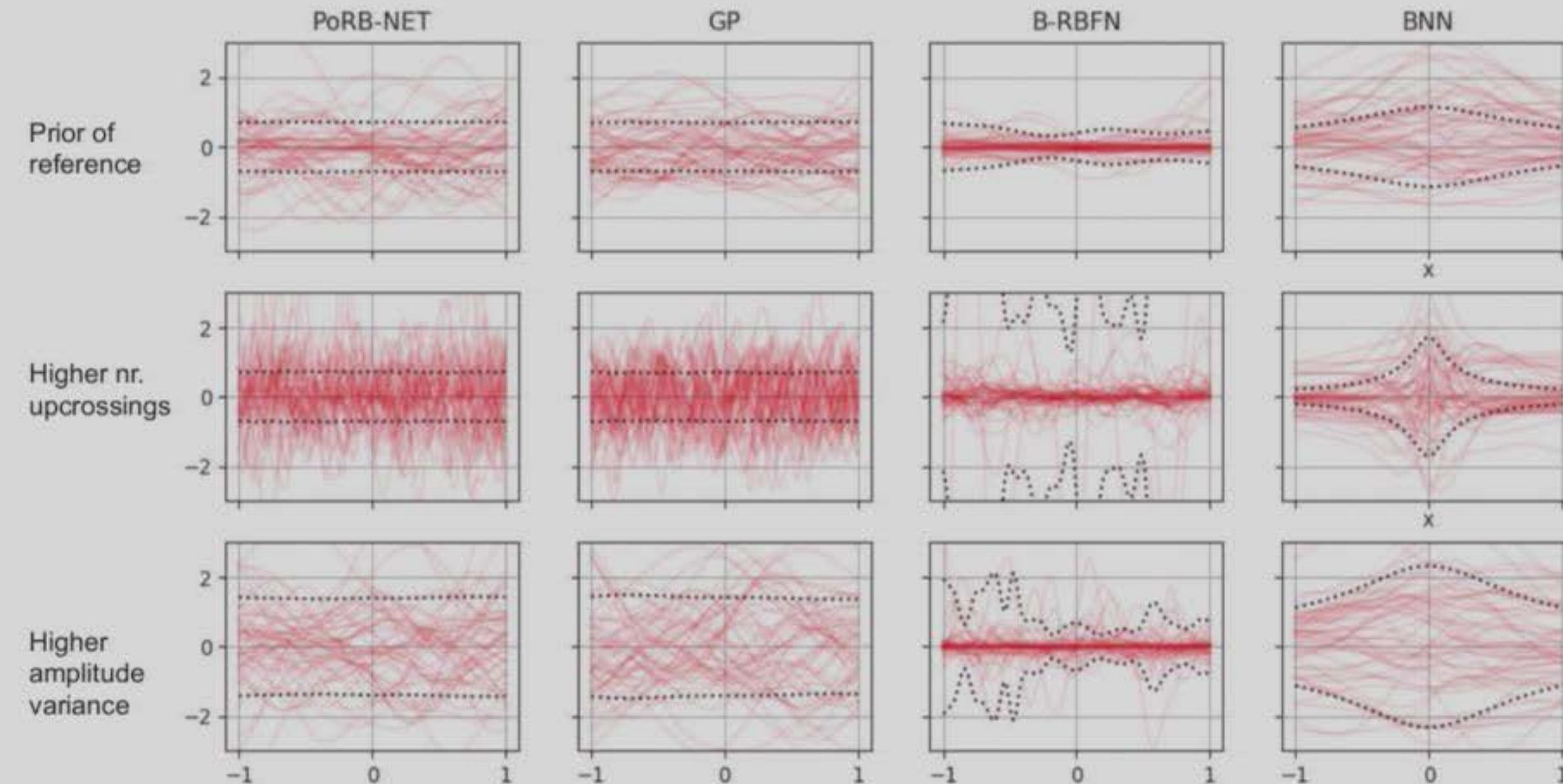
2. Decoupled lengthscale and amplitude variance

$$\text{Cov}(f(x_1), f(x_2)) = \sigma_b^2 + \tilde{\sigma}_w^2 \exp\left\{-\lambda^2 \left(\frac{x_1 - x_2}{2}\right)^2\right\}$$

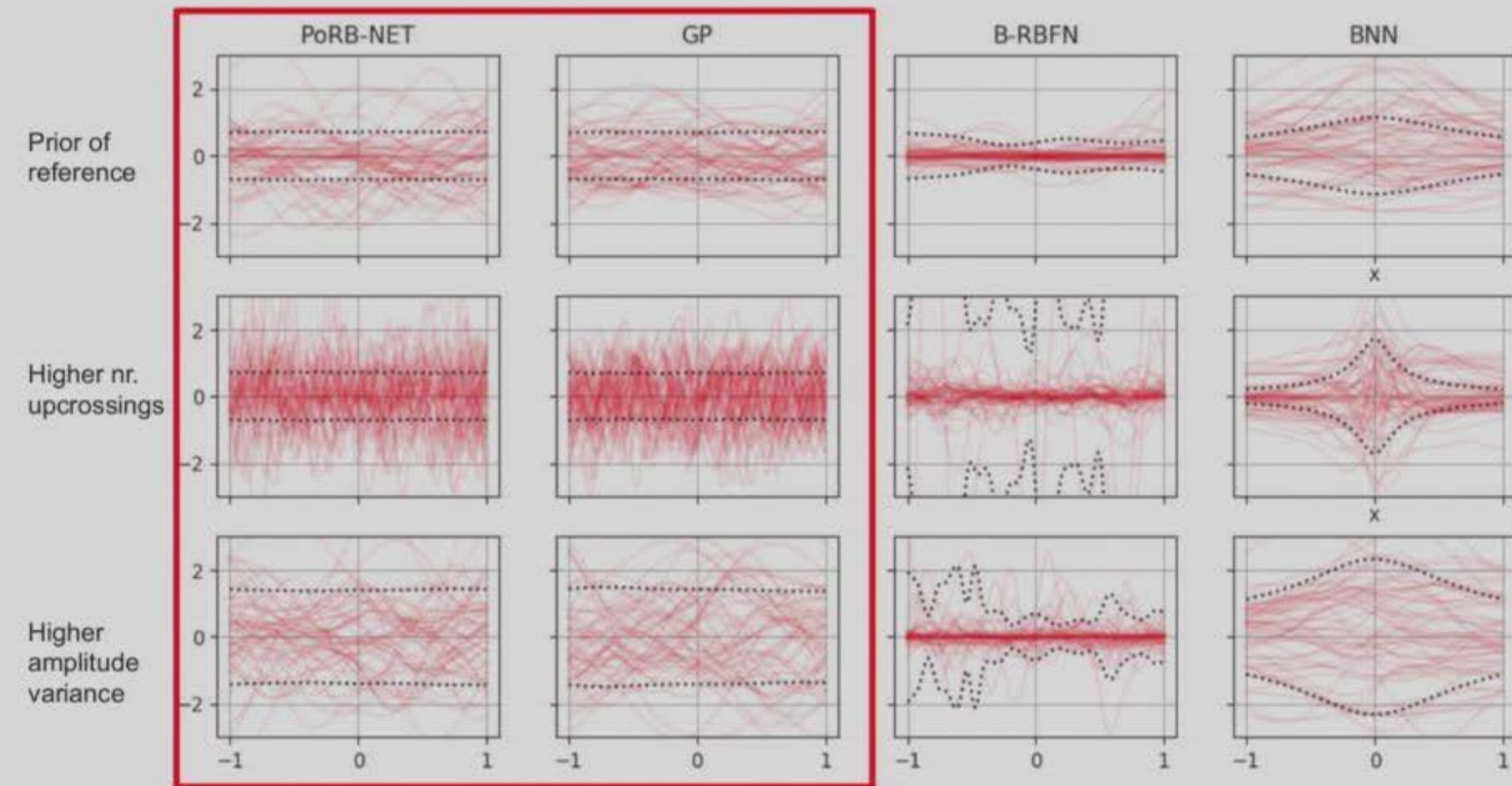
3. Consistency Theorem:

A PoRB-NET with uniform intensity function is Hellinger consistent as the number of observations goes to infinity.

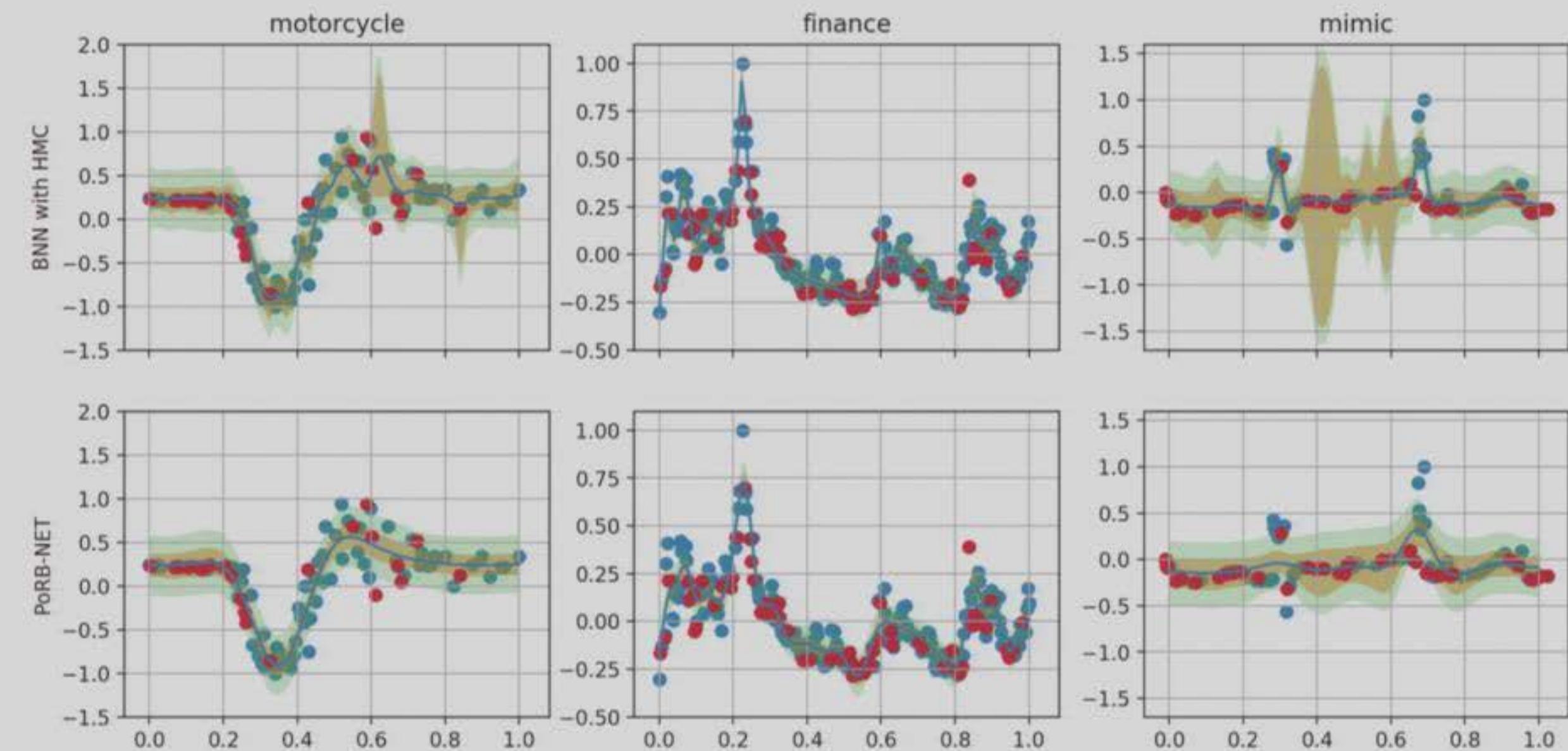
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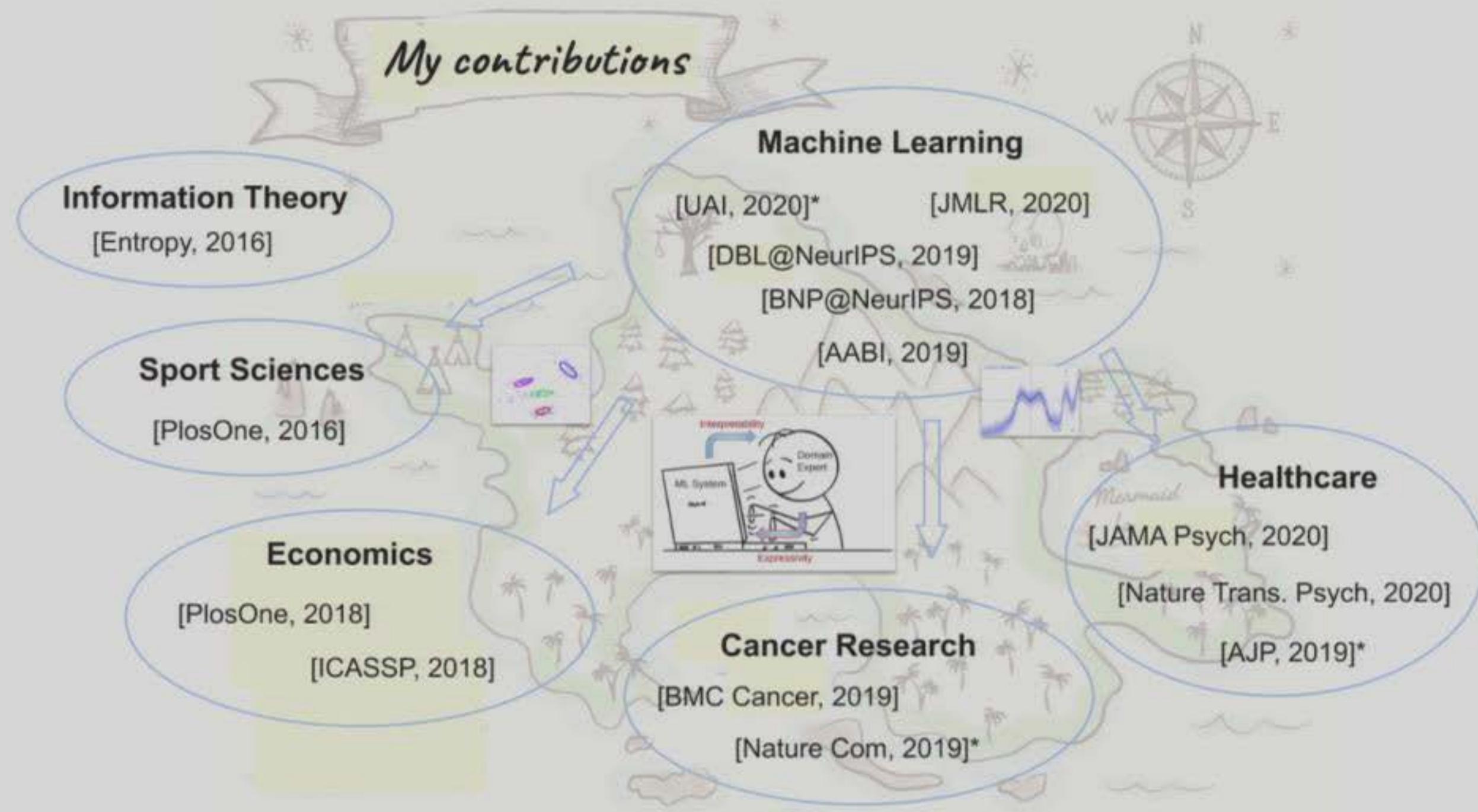


PoRB-NET ALLOWS FOR EASY SPECIFICATION OF LENGTHSCALE AND SIGNAL VARIANCE LIKE A GP



PoRB-NET IS ABLE TO CAPTURE NON-STATIONARY PATTERNS IN REAL SCENARIOS, ADAPTING THE LENGTHSCALE LOCALLY





From the lab to the clinic

- Ongoing user study at MGH, Boston
 - Impact of explanations
 - Usefulness, trust...

Why are these therapies being recommended?

The following patient features had the highest contributions to system.I3's predictions:



Which antidepressant medication would you be most likely to prescribe in this situation?



Patient Details:

Jessica is a 37 year old woman who is married and works part time. She presents with 9 months of depressed mood and lack of appetite. She has a seizure disorder, and current medications include Omeprazole and Celecoxib. Prior treatment with Citalopram had no effect on depressed mood.

System.I3 Recommendation: FLUOXETINE

Top 5 therapies with highest probability for stability:

Therapy	Predicted Stability*	Predicted Dropout Risk**
Fluoxetine	.76	.05
Sertraline	.67	.05
Paxoxetine	.64	.10
Venlafaxine	.60	.14
Vortioxetine	.55	.15

*Stability: continued use of the same medication for at least 3 months

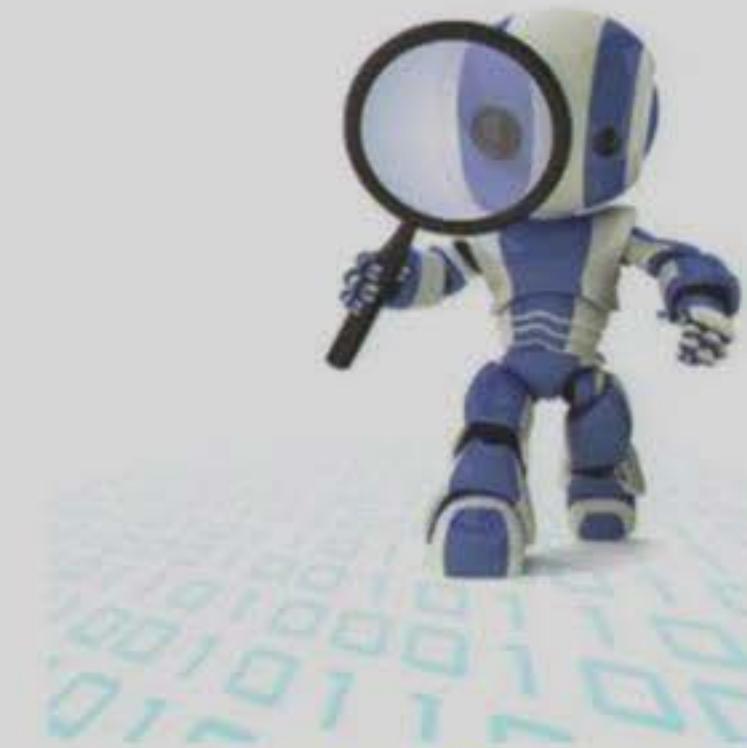
**Dropout: early treatment discontinuation following prescription

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The following rules had the highest contributions to system.I3's predictions:

1. If underweight or lack of appetite, favor weight gain, favor Mirtazapine
2. If underweight or lack of appetite, avoid appetite suppressants, avoid nausea-inducing, avoid SNRIs, avoid Sertraline
3. If lack of response to Paxoxetine, avoid SSRIs

Current and future research agenda



Contact: melanie@seas.harvard.edu

<https://melaniefp.github.io/>

Impactful real-world problems:

- Personalize prescription of antidepressants
- Prognosticate outcomes for in-vitro fertilization
- ...

Useful machine learning methodology:

- How to better quantify model uncertainty?
- How to combine expert knowledge with data-driven evidence?
- How to learn task-meaningful representations?

ACKNOWLEDGEMENTS

Special thanks to:

- Beau Coker
- Finale Doshi-Velez
- All members of DTAK!
- Oscar Puig
- Francesca Milletti
- Fernando Perez-Cruz
- Isabel Valera
- Maria Lomeli
- Zoubin Ghahramani



CRCS Center for Research on
Computation and Society

at Harvard John A. Paulson School of Engineering and Applied Sciences



HDSI

Harvard Data
Science Initiative



THANK YOU FOR LISTENING!

