

# **Randomness and information in biological data**

## **BIO-369**

**Prof. Anne-Florence Bitbol**

**TAs: Cecilia Fruet, Margaret Lane, Benjamin Martin, Shuhao Zhang**



First class

# Outline of the course

## I Randomness in biological processes and biological data

- 1 Randomness and random variables
  - 1.1 Coins and dice: discrete random variables
  - 1.2 Medical testing and conditional probabilities
  - 1.3 Luria-Delbrück experiment: Poisson distribution vs. jackpot distribution
- 2 Importance of thermal fluctuations at the cellular scale
  - 2.1 Thermal fluctuations and associated energy scale
  - 2.2 Strength of various chemical bonds
  - 2.3 Flexibility of biopolymers and biomembranes
- 3 Random walks
  - 3.1 Population genetics
  - 3.2 Protein abundances in single cells
  - 3.3 Importance of random walks in biological systems

# Outline of the course

## II Extracting information from biological data

- 1 Quantifying randomness and information in data: entropy
  - 1.1 Notion of entropy
  - 1.2 Interpretation of entropy
  - 1.3 Entropy in neuroscience data: response of a neuron to a sensory input
- 2 Quantifying statistical dependence
  - 2.1 Covariance and correlation
  - 2.2 Mutual information
  - 2.3 Identifying coevolving sites in interacting proteins using sequence data
- 3 Inferring probability distributions from data
  - 3.1 Model selection and parameter estimation: maximum likelihood
  - 3.2 Introduction to maximum entropy inference
  - 3.3 Predicting protein structure from sequence data
- 4 Finding relevant dimensions in data: dimension reduction
  - 4.1 Principal component analysis
  - 4.2 Beyond principal component analysis
- 5 Introduction to Bayesian inference

# Organization

- **Problem classes:** 3:15pm-5pm on Mondays, room CE1106
- **Lectures:** 10:15am-12noon on Wednesdays, room BS170
- Problem class about *previous* lecture
- **First two weeks:**
  - Monday, Feb. 19, 3:15pm-5pm: lecture 1; Wednesday, Feb. 21, 10:15am-12noon: lecture 2;
  - Monday, Feb. 26, 3:15pm-5pm: problem sets 1&2; Wednesday, Feb. 28, 10:15am-12noon: lecture 3.
- All questions are welcome:
  - During lectures and problem classes
  - Outside of lecture hours, using **Ed Discussion**
- All class material is available on **Moodle**
- Computer-based questions in problems + project: **Jupyter Notebooks with Python3**
- Some quiz questions during lectures, using **TurningPoint**
- **Numerical mini-project** during the semester (40% of the final grade)  
Weeks 10 & 11 (TBC), with two problem sessions on week 10 and one on week 11 devoted to it
- **Written exam** during the exam session (60% of the final grade)  
Mainly classic problems + some coding questions

# A few words about TurningPoint

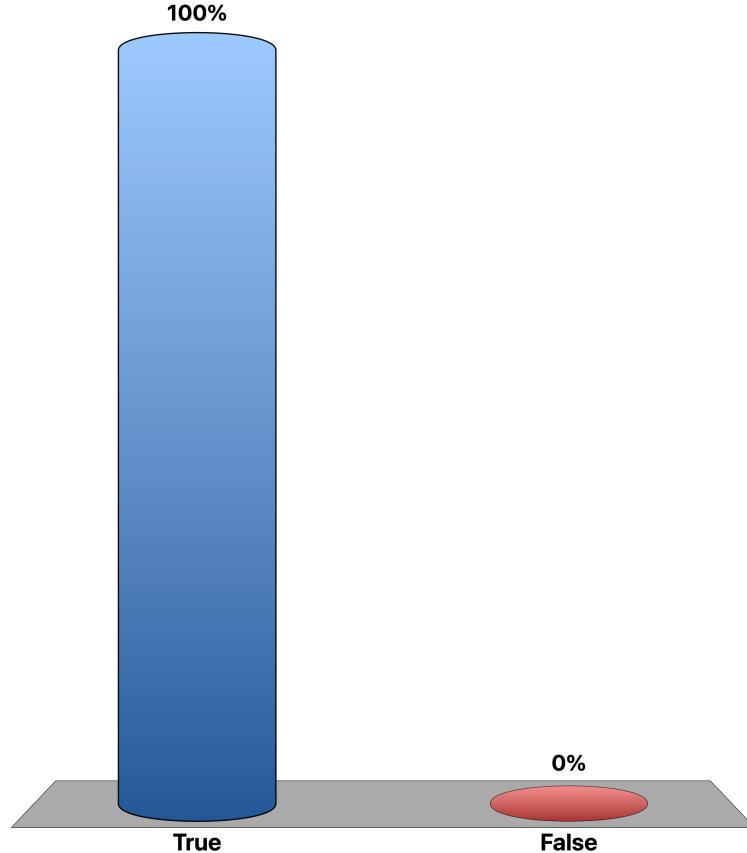
- Need a device connected to the Internet to answer the question
- No need for a TurningPoint account / license – all answers will be **anonymous** (as “guests”)
- **Goals:**
  - Think about a new notion
  - Recall a previously seen notion
  - Encourage active participation
  - Get feedback
- **Not an evaluation**
- Legal information: Data is processed outside Switzerland, which may include the USA or EU countries  
Contract with EPFL = Turning Technologies will not reuse the data collected for any other purpose, provided that you use the “guest mode”

Have you used Python before?

- A. True
- B. False

To answer, please:

- Connect to <http://tppoll.eu>
- Enter the session ID **bio369**
- Select your answer



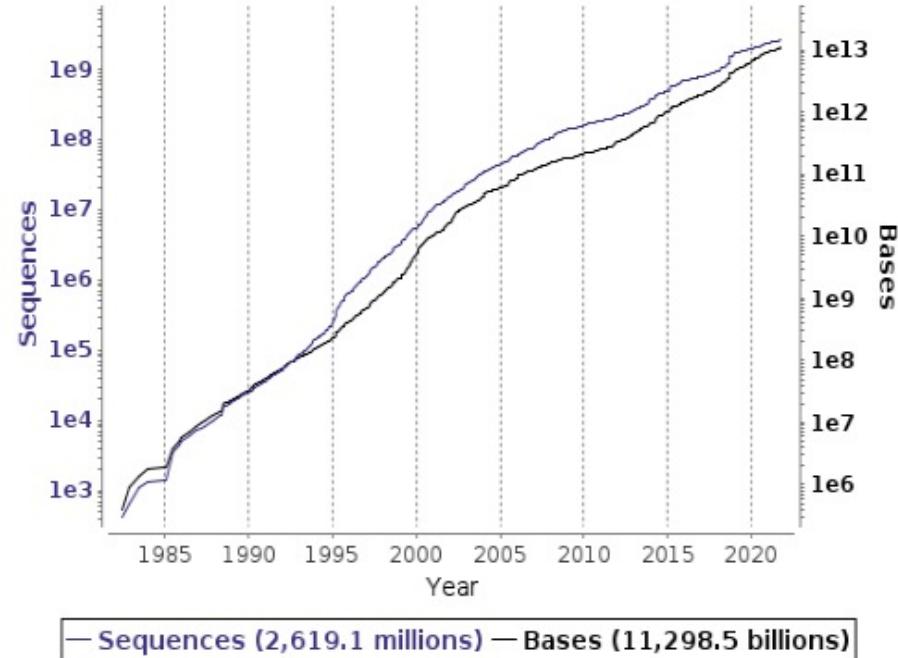
# Motivation

- Biology is becoming more and more a data science**

- Example: sequencing**

- Sequence data**

```
ISHDLKTPITAIILDLMLPGIDG
VSHELKTPPLTSIVILDNLNLPKQDG
VSHELRTPPLTSILVLDLMLPEIGG
ASHELRTPISVIVLLDIMLPGLSG
ISHDLKTPITAIILDLMLPGIDG
ASHELRTPISVIVLLDIMLPGLSG
VSHELRTPPLTSILVLDLMLPEIGG
```



- Accumulating unannotated sequence data (currently  $> 10^9$  sequences)

→ Great opportunity to learn about proteins employing inference, machine learning, statistical physics, information theory

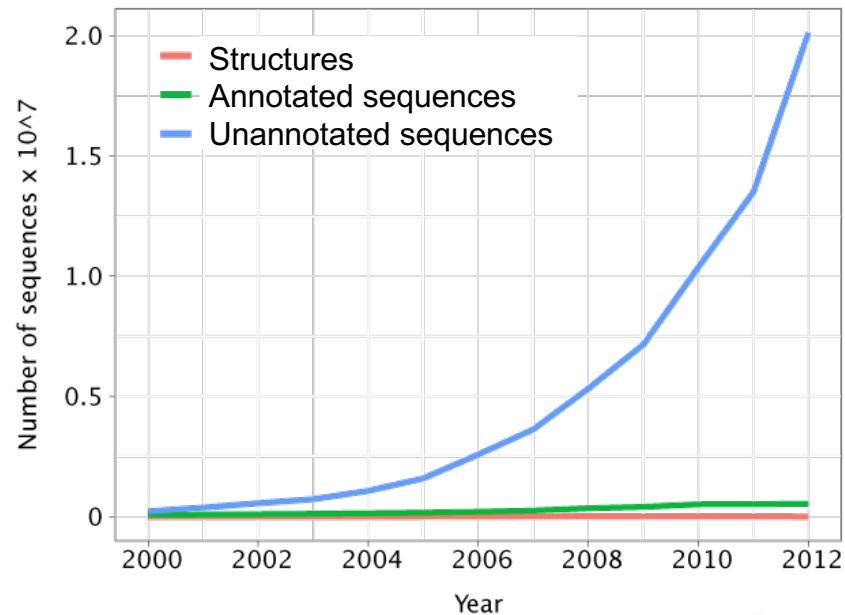
# Motivation

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- Example: sequencing**

- Sequence data**

```
ISHDLKTPITAIILDLMLPGIDG
VSHELRKTPPLTSIVILDNLNLPKQDG
VSHELRTPPLTSILVLDLMLPEIGG
ASHELRTPISVIVLLDIMLPGLSG
ISHDLKTPITAIILDLMLPGIDG
ASHELRTPISVIVLLDIMLPGLSG
VSHELRTPPLTSILVLDLMLPEIGG
```



- Accumulating unannotated sequence data (currently  $> 10^9$  sequences)

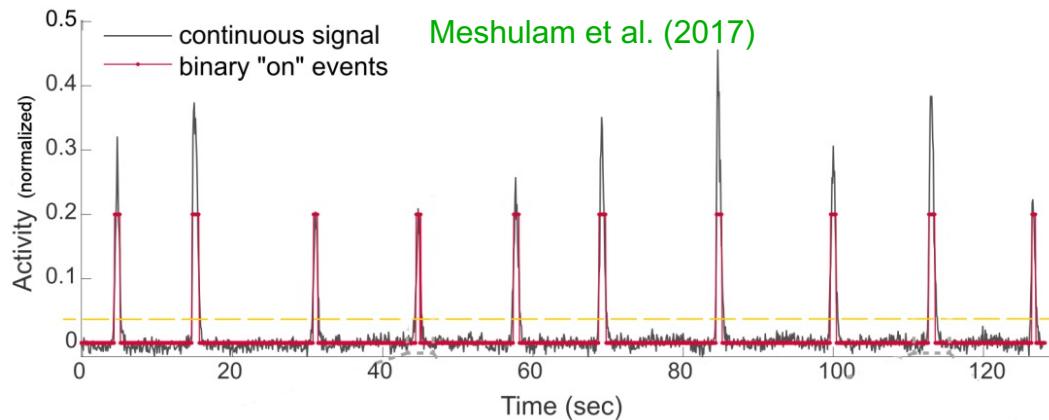
→ **Great opportunity to learn about proteins employing inference, machine learning, statistical physics, information theory**



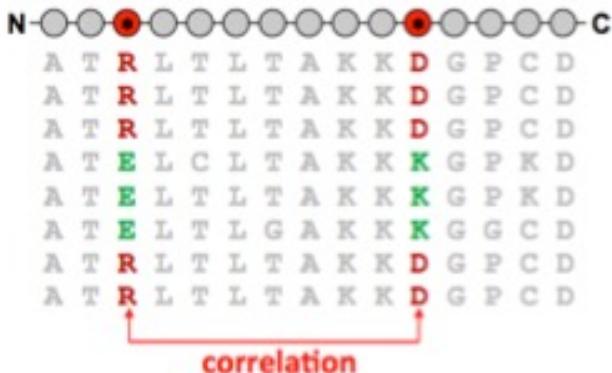
# Motivation

- Biological data can be viewed as sampled from distributions of random variables

- Neuroscience data:



- Protein sequence data:



$$P(\{\sigma_i\}) = \frac{1}{Z} \exp[-E(\{\sigma_i\})].$$

$$E(\{\sigma_i\}) = - \sum_{i=1}^N h_i \sigma_i - \frac{1}{2} \sum_{i,j=1}^N J_{ij} \sigma_i \sigma_j$$

$$P(\alpha_1, \dots, \alpha_L) = \frac{1}{Z} \exp \left\{ - \left[ \sum_{i=1}^L h_i(\alpha_i) + \sum_{i < j} e_{ij}(\alpha_i, \alpha_j) \right] \right\}$$

Weigt, White et al. (2009)

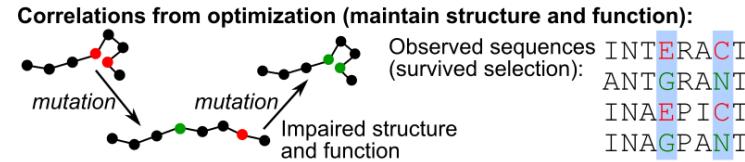
Morcos et al. (2011)

Marks, Colwell et al. (2011)

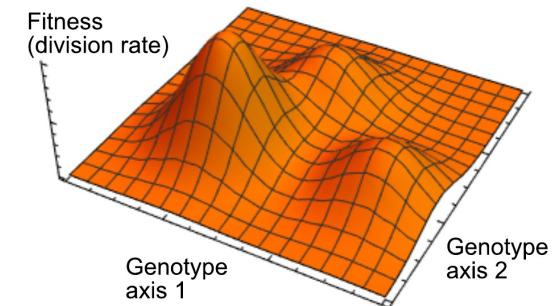
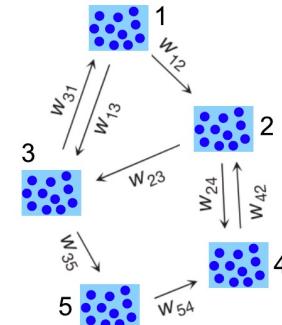
## Optimization & historical contingency



### ■ Aim 1: Understanding how optimization & phylogeny shape protein sequences



### ■ Aim 2: Assessing the predictability of evolution in subdivided populations



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What is the name of the distribution that describes the outcome of a coin flip, including the case where the coin is not fair?

11%

A. Uniform distribution

43%

B. Bernoulli distribution

46%

C. Binomial distribution

0%

D. Poisson distribution

To answer, please:

- Connect to <http://tppoll.eu>
- Enter the session ID **bio369**
- Select your answer

Consider two random variables X and Y.

When can we write, for all x and y,  $P(X=x, Y=y) = P(X=x) P(Y=y)$ ?

0%

A. Always

0%

B. Never

0%

C. When X and Y have the same distribution

78%

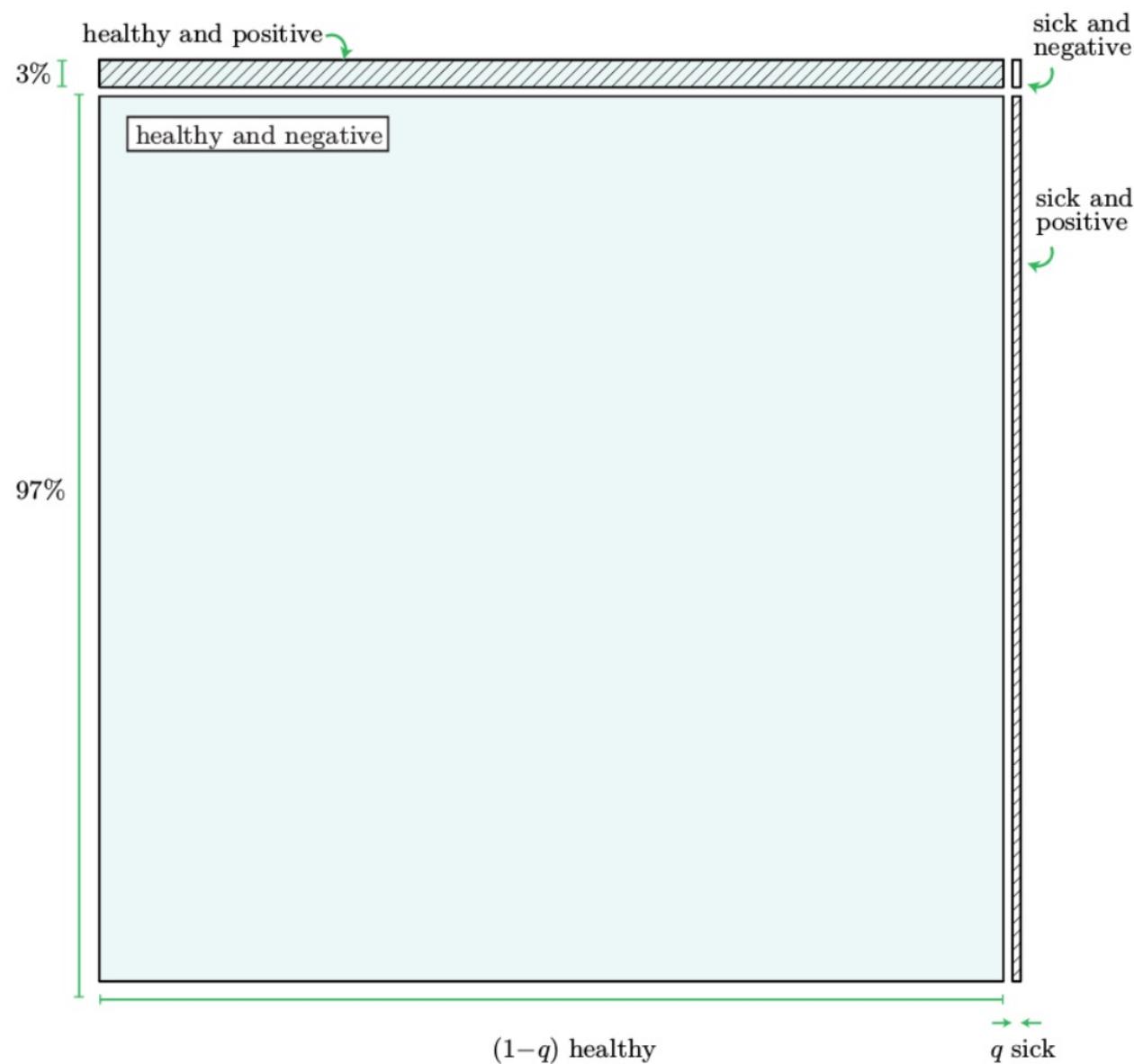
✓ D. When X and Y are independent

22%

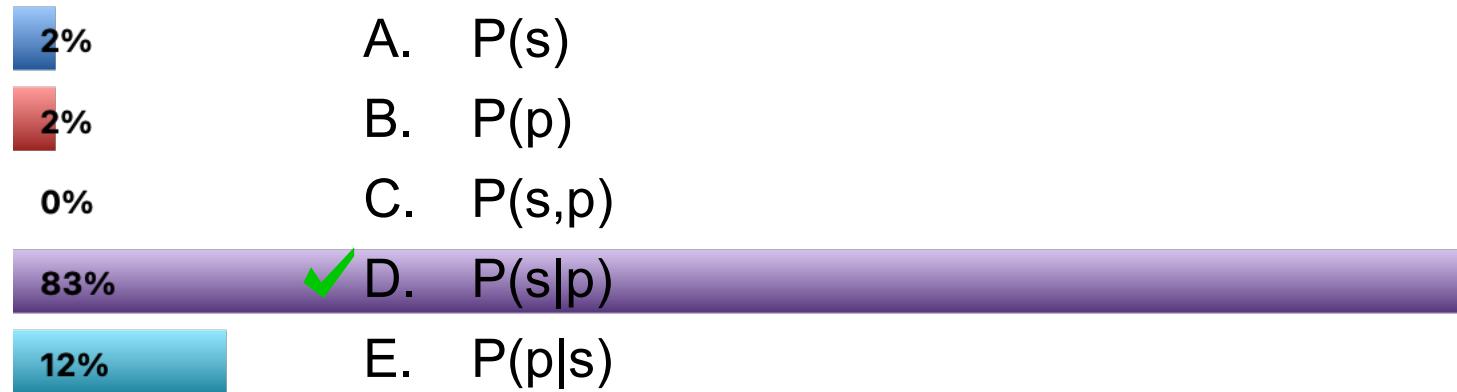
E. When X and Y have the same distribution and are independent

To answer, please:

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- Select your answer



Imagine that you are a doctor. Your patient has taken a test, and tested positive (p). Before taking action, what would you like to know?



To answer, please:

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- Select your answer

All patients are either sick (s) or healthy (h), and all patients are tested (p or n).  
What is  $P(p,s)+P(p,h)$  equal to?

- 0% A.  $P(p)$
- 0% B.  $P(s)$
- 0% C. 1
- 0% D.  $P(p|s)+P(p|h)$
- 0% E. None of the above

To answer, please:

- Connect to <http://tppoll.eu>
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- Select your answer

All healthy (h) patients are tested, and all tests are either positive (p) or negative (n). What is  $P(p|h)+P(n|h)$  equal to?

39%

A.  $P(h)$

0%

B.  $P(n,h)$

55%

✓ C. 1

3%

D.  $P(p,h)+P(n,h)$

3%

E. None of the above

To answer, please:

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- Select your answer