

# HMM for Bioinformatics

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6 marca 2019

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- 4  $P_X(x_i)$  probability.

We can think of entropy as the level of uncertainty associated with a random variable.

# Entropy – Definition

We define the entropy of the random variable  $X \sim P$  as:

$$H(P) = H(X) = - \sum_{i=1}^L P(x_i) \log_2(P(x_i))$$

where  $H(0) = 0$  (convention  $0 \cdot \log_2 0 = 0$ ).

$$H(X, Y) = - \sum_x \sum_y P(x, y) \log P(x, y)$$



$$H(X|Y) = - \sum_x \sum_y P(x, y) \log P(x|y).$$

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- 4  $H(X)$  is concave in  $X$ .

# Chain Rule for entropy

$$H(X, Y) = H(X) + H(Y|X) = H(Y) + H(X|Y).$$

$$M(X, Y) = \sum_x \sum_y P(x, y) \log \frac{P(x, y)}{P(x)P(y)}$$

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## Remark

$$M(X, Y) = H(X) + H(Y) - H(X, Y).$$



## Example – Binary entropy function

$$\mathcal{X} = \{x_1, x_2\}, P(x_1) = p, P(x_2) = 1 - p$$

$$H(p) = -p \log_2 p - (1 - p) \log_2 (1 - p)$$

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- 4 We can think in this case of the entropy as a number of yes/no questions needed to identify an outcome.

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## Kraft inequality

A necessary and sufficient condition on the lengths of codestring  $l_i$  in a uniquely decodable code is:

$$\sum_{i=1}^L 2^{-l_i} \leq 1.$$

The expected length of the optimal uniquely decodable code belongs to the interval  $[H(X), H(X) + 1)$



$$D(P_1|P_2) = \sum_{i=1}^L P_1(x_i) \log \frac{P_1(x_i)}{P_2(x_i)}$$

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## Convention

- 1  $0 \cdot \log \frac{0}{P_2(x_i)} = 0;$
- 2  $P_2(x_i) \log \frac{P_1(x_i)}{0} = \infty$

If  $X$  is a random variable with the probability  $P_1$  on an alphabet of  $L$  symbols, and  $P_2$  is the uniform distribution on this alphabet then

$$D(P_1|P_2) = \log(L) - H(X).$$

# Properties of relative entropy

- 1 T. Inglot, „Teoria informacji a statystyka matematyczne” wykład wygłoszony na XXXVIII Konferencji Statystyka Matematyczna Wisła 2012;
- 2 L. Dębowski, "Information Theory and Statistics", Institute of Computer Science Polish Academy of Science, Warsaw, 2013.