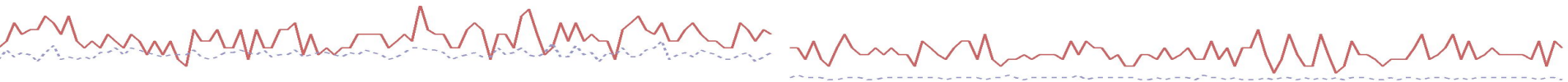


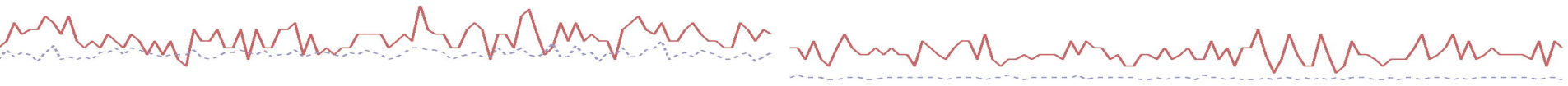
# Machine learning for regression problems

Filippo Biscarini (CNR, Milan, Italy)

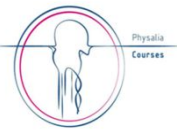
[filippo.biscarini@cnr.it](mailto:filippo.biscarini@cnr.it)



# Introducing the dataset



# DNA methylation and age in European bats



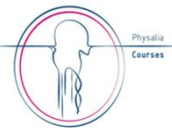
- Wild Bechstein's bats (*Myotis bechsteinii*)
- N = 62
- seven CpG sites ([here](#))
- % methylation
- age: [0-14] years



Source: [https://en.wikipedia.org/wiki/Bechstein%27s\\_bat](https://en.wikipedia.org/wiki/Bechstein%27s_bat)



# DNA methylation and age in European bats



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**RESOURCE ARTICLE**

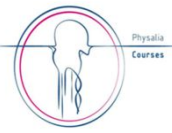
WILEY **MOLECULAR ECOLOGY  
RESOURCES**

## Application of a novel molecular method to age free-living wild Bechstein's bats

Patrick G. R. Wright<sup>1</sup> | Fiona Mathews<sup>2</sup> | Henry Schofield<sup>3</sup> | Colin Morris<sup>3</sup> |  
Joe Burrage<sup>4</sup> | Adam Smith<sup>4</sup> | Emma L. Dempster<sup>4</sup> | Patrick B. Hamilton<sup>1</sup>



# DNA methylation and age in European bats



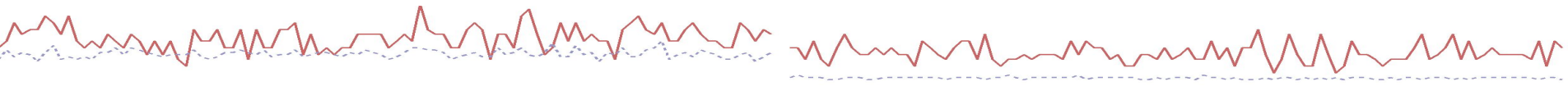
Sample	Age	Age category	CpG 1 TET2	CpG 2 TET2	CpG 3 TET2	CpG 4 TET2	CpG GRIA2 1	CpG GRIA2 2	ASPA 1
BabyBechs_SHW	0	Age 0-3	29	21	26	31	2	2	61
Dd_Juv_Hamgreen	0	Age 0-3	30	21	24	32	1	2	59
A2402	1	Age 0-3	44	38	51	53	1	5	48
A2414-2014	1	Age 0-3	48	36	50	46	1	2	53
...	...	...	...	...	...	...	...	...	...



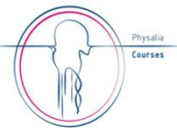
# Regression problems

Demonstration 2.1

→ `linear_regression.Rmd`



# Normalized discounted cumulative gain (NDCG)



NDCG is a **ranking metric** developed in information theory which has been applied to evaluation of genomic selection models

NDCG evaluates the **top** (e.g. 20%) **individuals in the ranking**, which are supposed to be the most relevant when comparing models

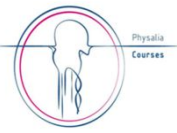
$$DCG@k = \sum_{i=1}^k y[\pi(\hat{y}_i)] \cdot d(i)$$

gain function

discount



# Normalized discounted cumulative gain (NDCG)



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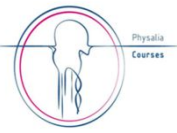
The higher the DCG, the better

However, DCG is difficult to interpret (unbounded)  **NDCG**





# Normalized discounted cumulative gain (NDCG)



NDCG is a **ranking metric** developed in information theory which has been applied to evaluation of genomic selection models

NDCG evaluates the **top** (e.g. 20%) **individuals in the ranking**, which are supposed to be the most relevant when comparing models

$$NDCG(y, \hat{y}) = \frac{\sum_{i=1}^k (y[\pi(\hat{y})]_i \cdot d(i))}{\sum_{i=1}^k (y[\pi(y)]_i \cdot d(i))}$$

NDCG values lie in [0,1]



# Regression problems

Exercise 2.1

→ `linear_regression.Rmd`

