

# Incremental hierarchical clustering for pattern discovery to optimize on-farm milk processing



*I. Alexakis, C. Nickmilder, S. Franceschini, J. Leblois, V. Wolf,  
HoliCow Consortium, H. Soyeurt*

**Interreg**  
North-West Europe



Co-funded by  
the European Union

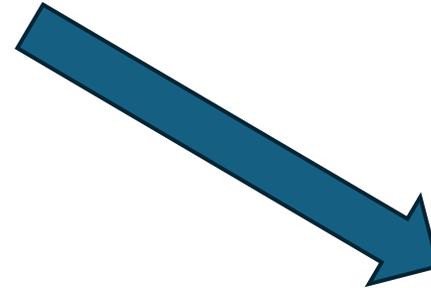
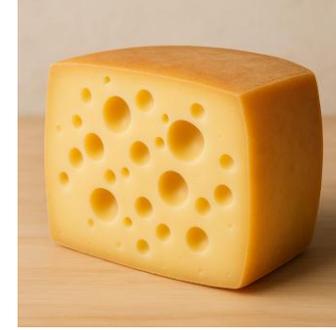
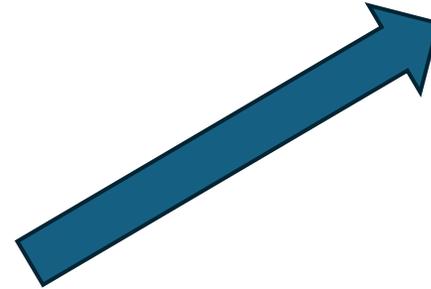
**HoliCow**



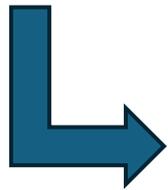
**LIÈGE université**  
**Gembloux**  
**Agro-Bio Tech**



# On-farm milk processing



# How to optimize milk processing



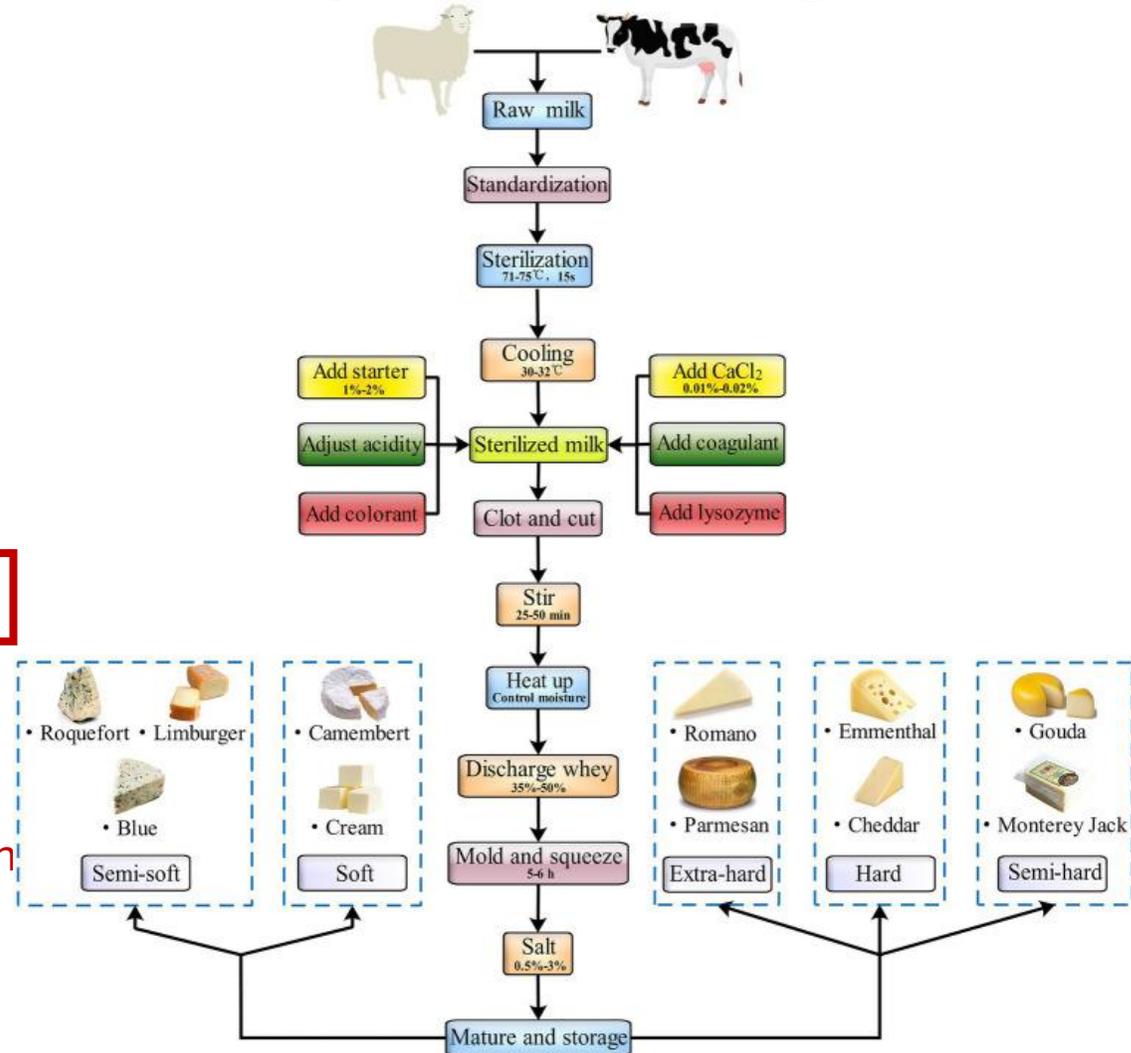
Process optimization



Milk quality optimization



Classifying milk based on th



(Zheng et al., 2021)



# How to do it ?





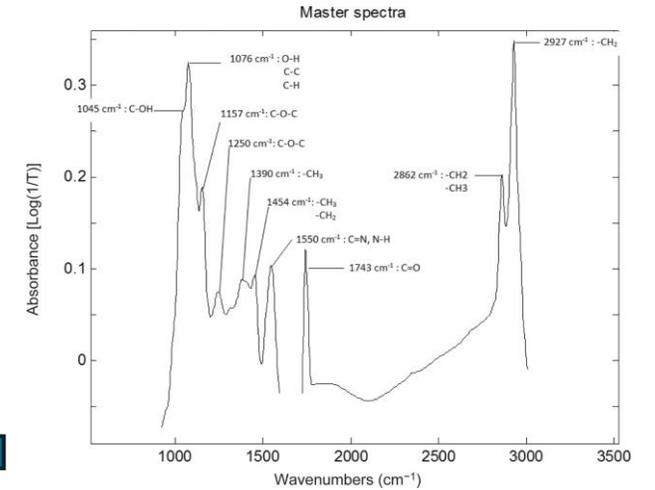
Rapid and cost-effective method,  
generate phenotypes at large scale



Reference laboratory  
measurements



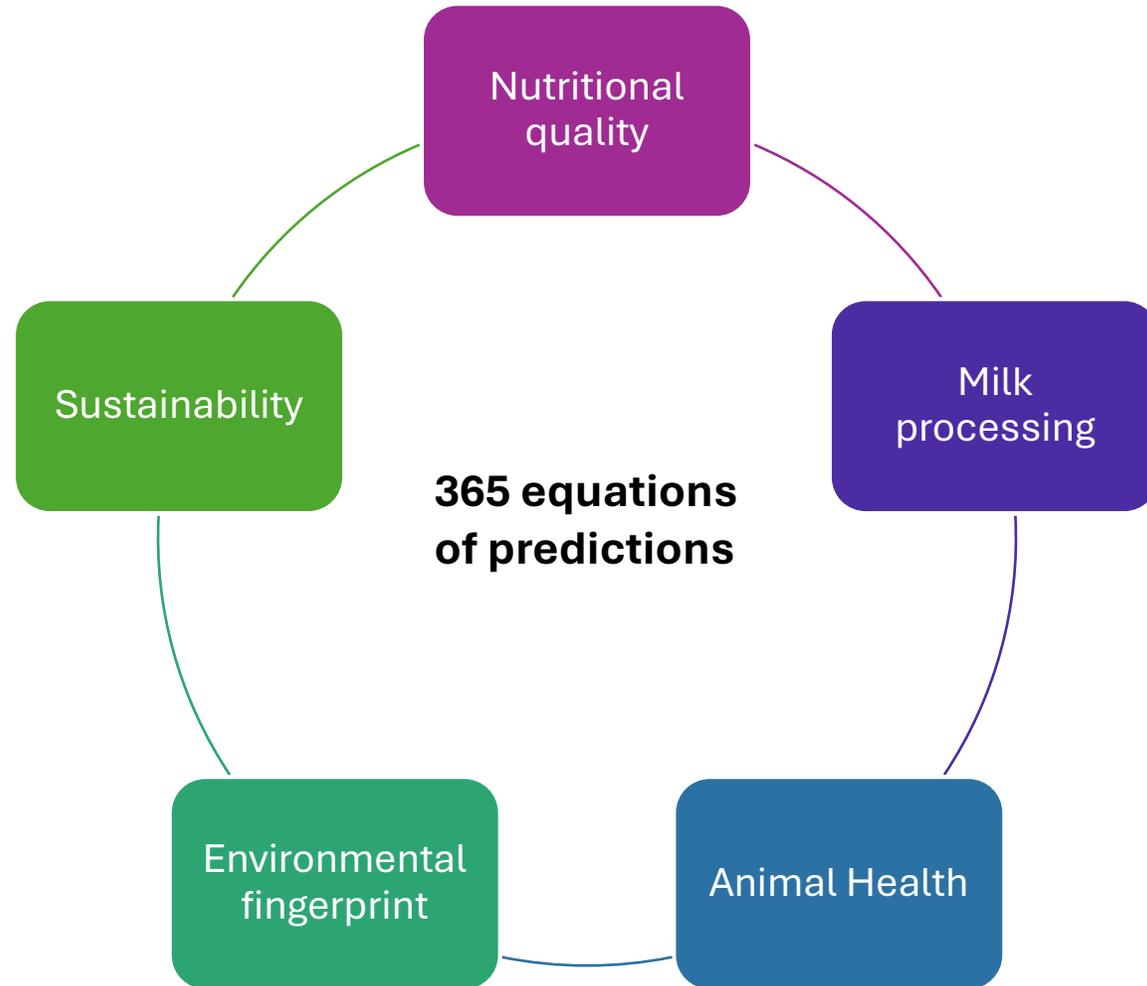
Time-consuming, expensive,  
and often impractical



Milk spectrum

Equations of predictions





Milk processing

26 MIR traits

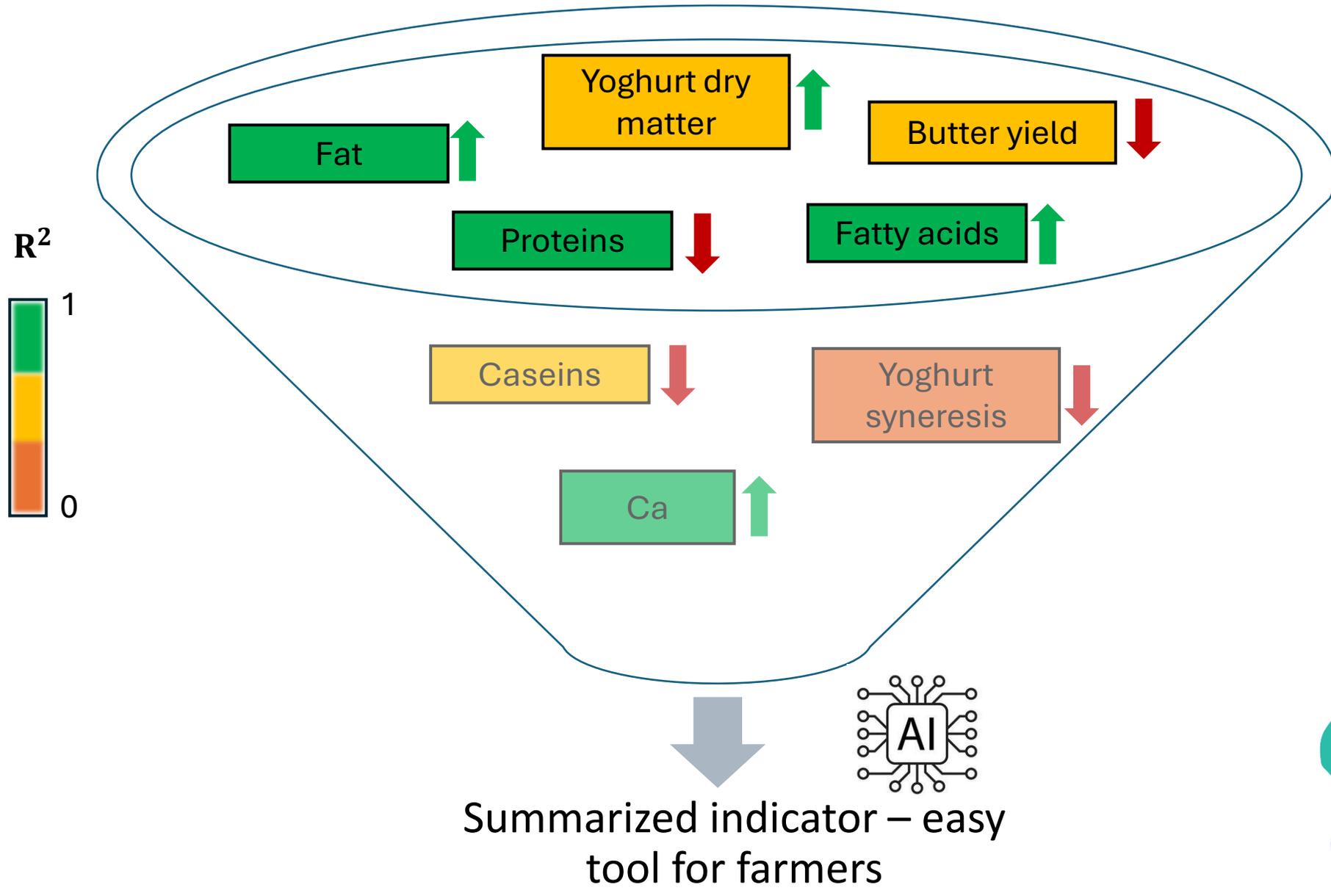
**Chemical composition**

Fat, protein, fatty acids, Ca, ...

**Technological properties**

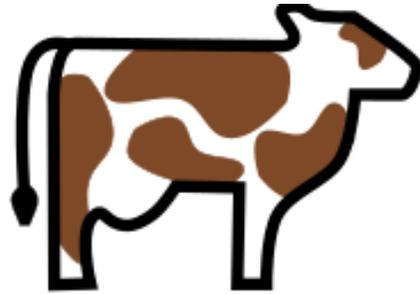
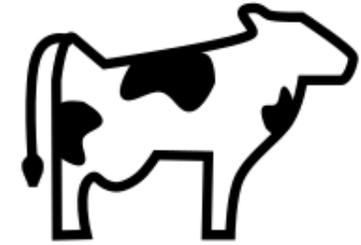
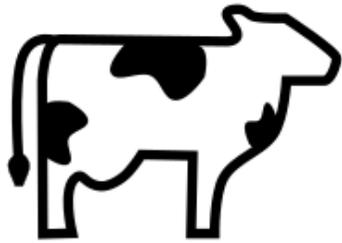
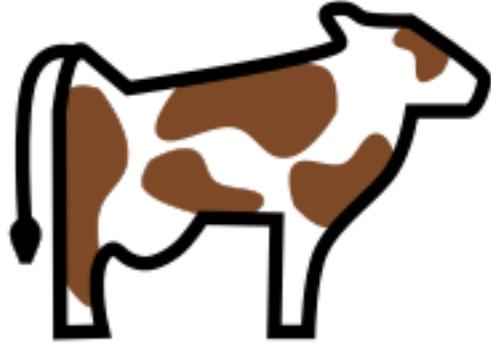
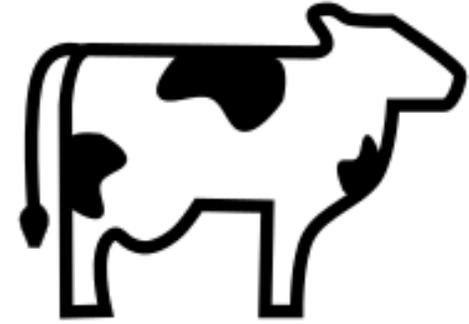
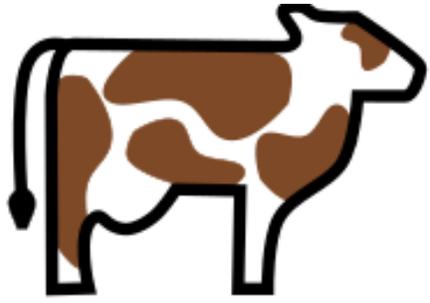
Milk coagulation properties, butter yield, yoghurt syneresis, ...

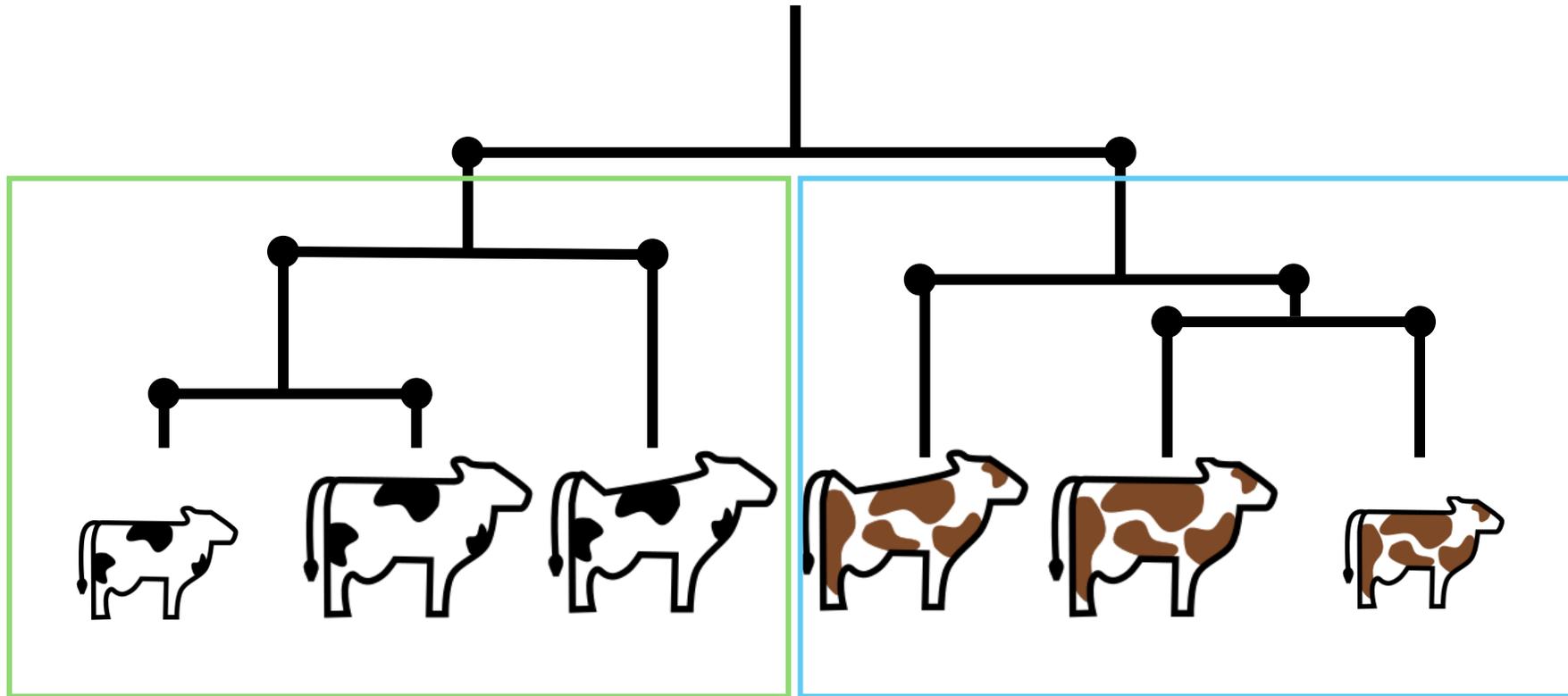
# Too many MIR traits



# Incremental **hierarchical clustering**







**Cluster 1**

Black cows

**Cluster 2**

Brown cows

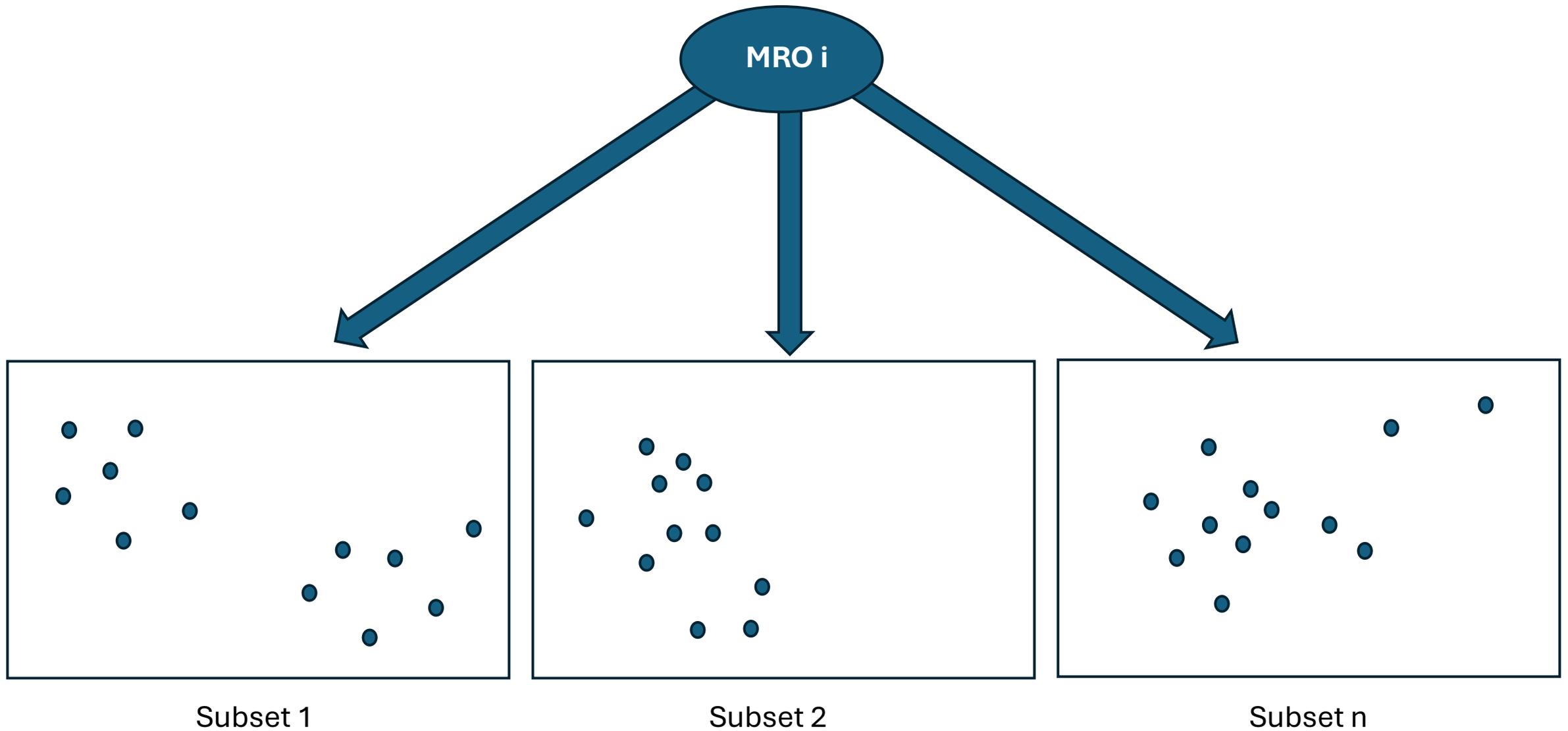


# How to implement it on the entire database ?

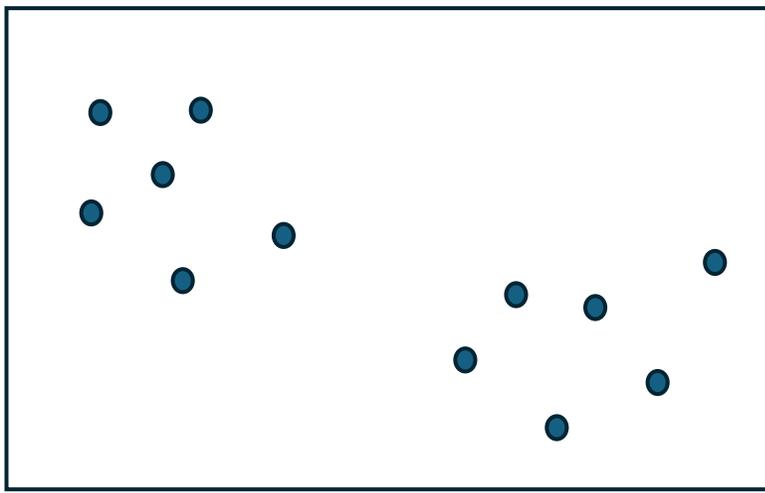


**Incremental hierarchical clustering**

41,000,000 records in 48 milk recording organizations files (MRO)

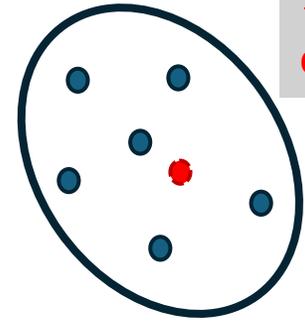


↗ Variability explained over interpretability



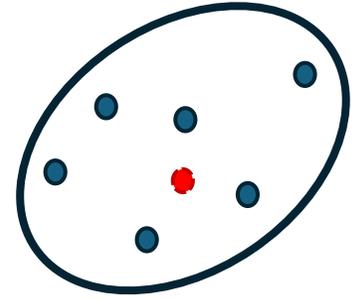
Subset 1

Hierarchical clustering



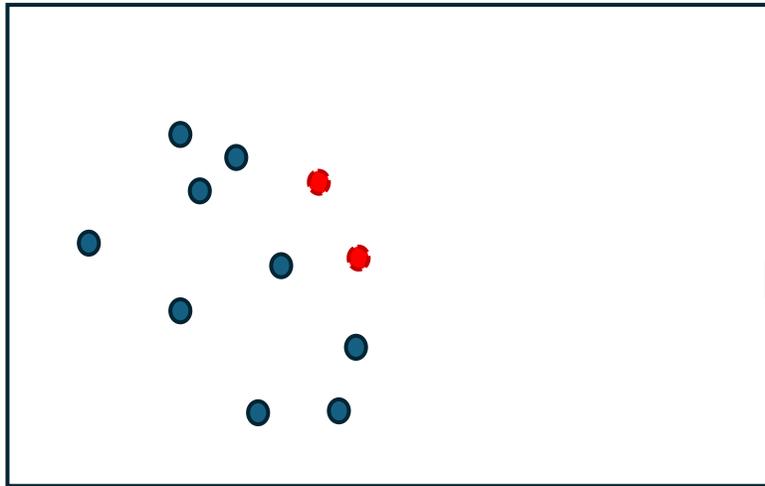
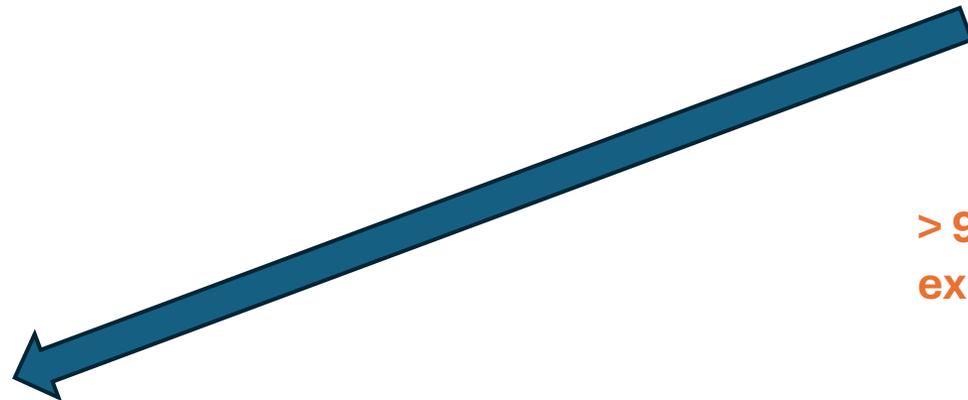
Cluster 1

Centroids



Cluster 2

> 95 % variance explained



Subset 2

Hierarchical clustering



etc.

800 centroids



**800  
centroids**

x 48 MRO = **38,400 centroids**

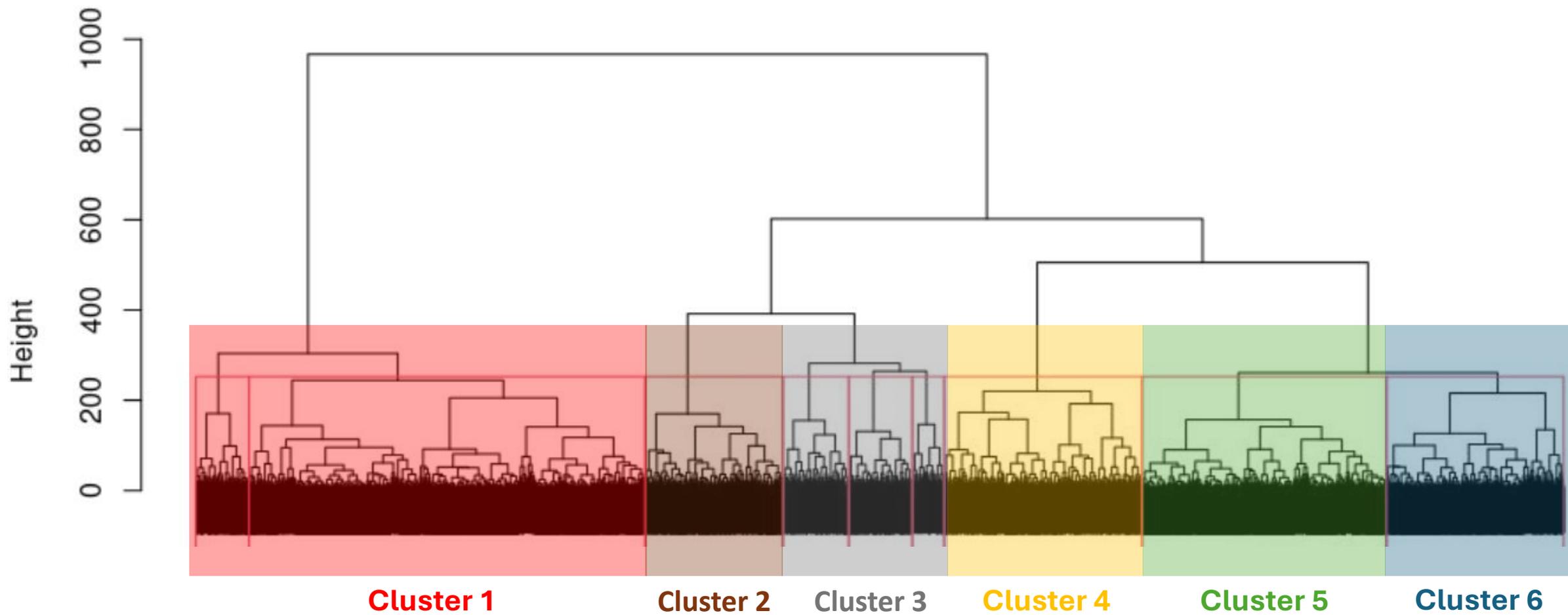
41,000,000 records → 38,400 centroids

**Hierarchical  
clustering**

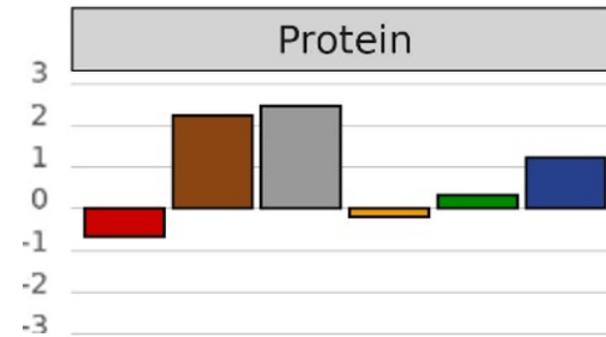
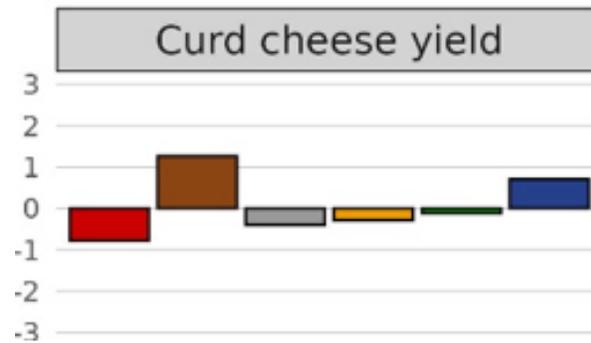
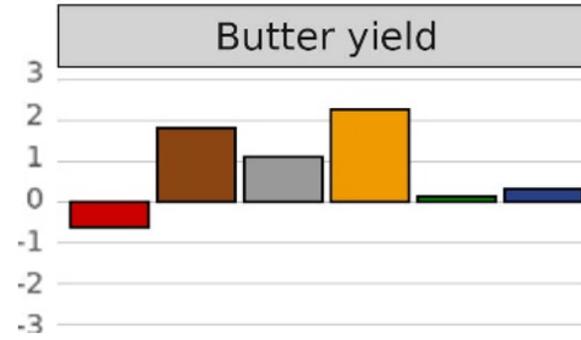
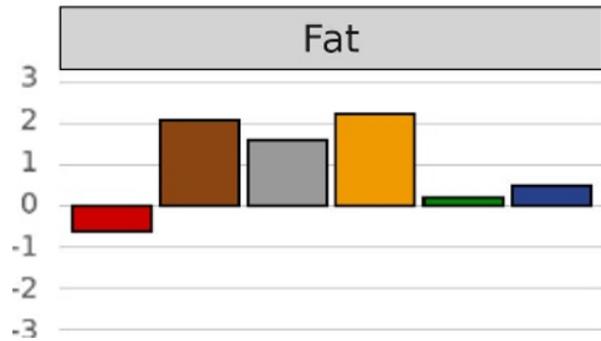
**6 clusters**

↗ Interpretability





# Clusters interpretation



Clusters



On 38,400 centroids !



HOLICOW

Next step ?



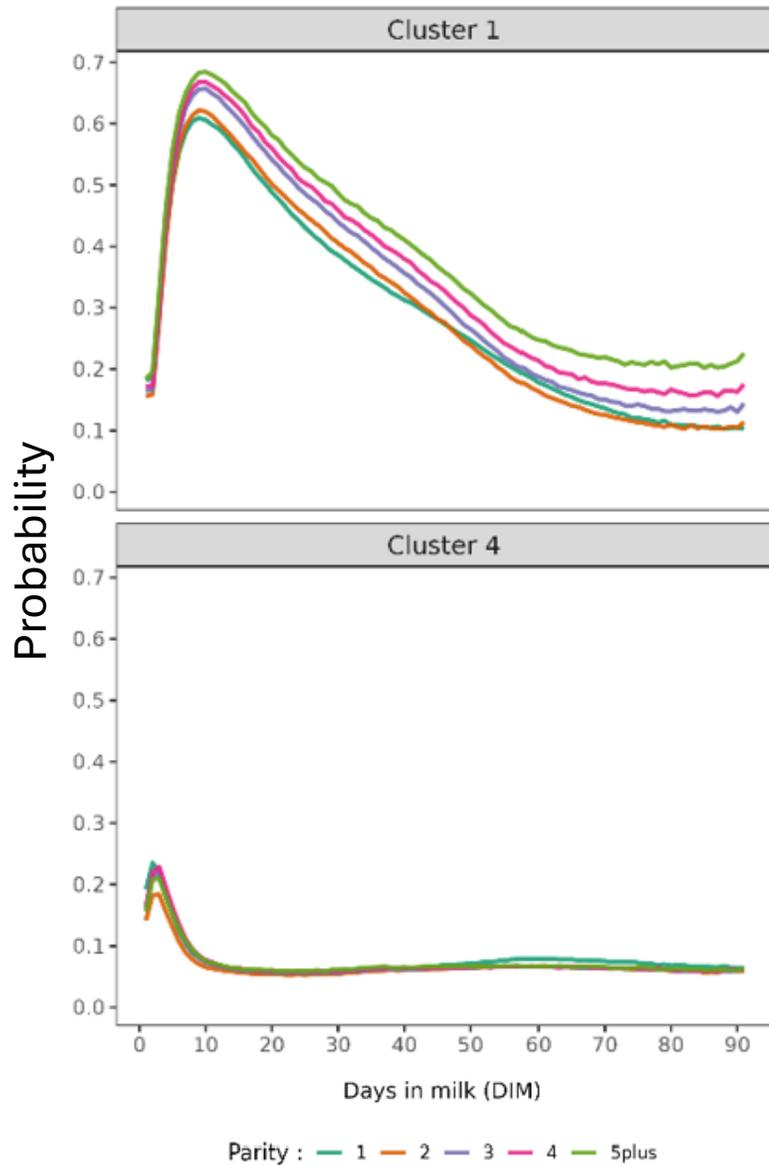
# Extend results to the entire database



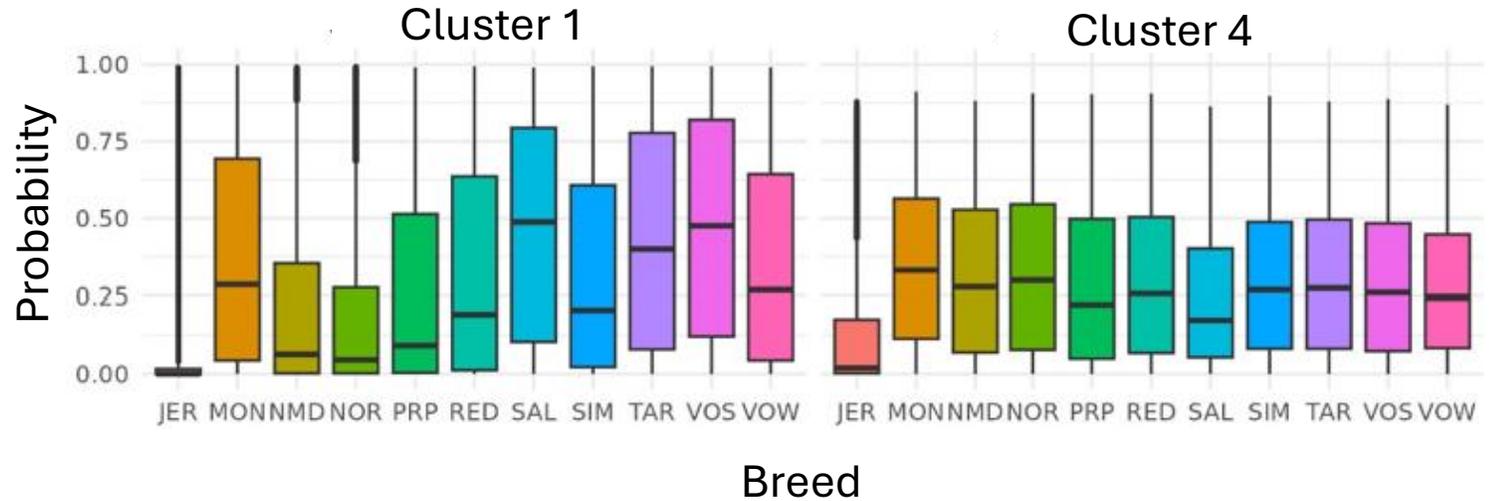
Study temporal, phenotypic,  
and genetic variabilities



## Days in milk (DIM) and Parity effect



## Breed effect



On 41,000,000 records





Contact : [ialexakis@uliege.be](mailto:ialexakis@uliege.be)

Project : <https://holicow.nweurope.eu>



 **LIÈGE université**  
**Gembloux**  
**Agro-Bio Tech**

**Interreg**  Co-funded by  
the European Union  
North-West Europe  

---

**HoliCow**

Avec le soutien de la  
 **Wallonie**