



Optimization of the Measurement Parameters of the Electronic Tongue for the Classification of

Different Fat Content Trappist Cheese

Lueji Regatieri Santos¹, Juan Pablo Aguinaga Bosquez¹, György Bázár², Zoltan Kovacs¹

¹Department of Measurements and Process Control, Institute of Food Science and Technology - Hungarian University of Agriculture and Life Sciences, Budapest, Hungary

²ADEXGO Kft., Balatonfüred, Hungary

INTRODUCTION

Cheese flavor highly depends on both its volatile and nonvolatile compounds such as peptides, amino acids, fats and salts. The characteristics of the endproduct depends on the quality of the milk and the ripening method affecting the microbiological, organoleptic and physical-chemical properties. E-tongue providing simultaneous multi-component quantitative determination along with qualitative discriminatory capacities is considered to be a good alternative to classical laboratory techniques because of its high sensitivity, low cost, simple operation, and inherent portability.

This work aimed to determine the optimal parameters (solutions' concentration and the sequence order) defined by the method, in order to improve the

performance of the method for discrimination of cheese, influenced by selectivity and sensitivity aspects of the sensors array.

MATERIALS AND METHODS

Weight the sample and

prepare dilutions

(Concentrations: 1, 5

and 10%)

Trappista Cheese – 5 Brands



Homogenize with a

mixer

during 1min



Measurement sequences

1st sequence: CH, RI, TE, TO and FI
2nd sequence: RI, CH, TO, FI and TE
3rd sequence: CH, TE, RI, FI and TO



Grid the cheese

Data evaluation using RStudio software (4.1.1 version) in order to accomplish the multivariate data analysis (PCA, LDA) Analysis performed via electronictongue ("α-ASTREE" Liquid and Taste Analyzer - Alpha M.O.S., Toulouse, France)



sieve

Transferred to a volumetric flask



Divided in 3 replicates. One concentration was performed per day: 3 different measurement sequences of 3 replicates from each cheese brand (n = 45)

RESULTS



Summary of LDA confusion tables for classification						
	Cheese		Cheese		Cheese	
	1% concentration		5% concentration		10% concentration	
	Recognition	Prediction	Recognition	Prediction	Recognition	Prediction
st equence	66.29	40.00	84.99	70.48	92.50	82.53
nd equence	98.35	56.78	97.52	87.52	97.52	95.04
rd equence	97.52	80.02	98.76	90.07	94.98	90.02
verage ccuracies	87.39	58.93	93.76	82.69	95.00	89.19

• The results observed from Linear Discriminant Analysis • Even though the best LDA model found was

and Euclidean Distances proves the similarities between RI and CH (lower fat content), in comparison to the other three samples (FI, TE and TO) of high fat content.

10% concentration, it is recommendable to work with the reduced concentration of 5%

considering the integrity of the equipment.

CONCLUSIONS

- This study has shown that discrimination of cheese can be optimized by selecting specific parameters such as:
- Concentration: capability of e-tongue to discriminate between samples improves over higher concentrations (10%).
- Measurement sequence: e-tongue had better performance when samples are analyzed per groups, considering the two with lower fat content first and then the three samples with higher fat content.
- Due to the impact of the matrix, lower concentrations are recommended considering the deterioration of sensors within time.

 According to the best sequence, considering the 10% optimal defined cheese concentration, both second and third sequences perform good, being the 2nd sequence slight better for this concentration (RI, CH, TO, FI and TE).

REFERENCES

Ciosek, P. (2016). Milk and Dairy Products Analysis by Means of an Electronic Tongue. In Electronic Noses and Tongues in Food Science. Elsevier Inc. https://doi.org/10.1016/B978-0-12-800243-8/00021-4

Kovacs, Z., Szöllosi, D., Zaukuu, J.-L. Z., Bodor, Z., Vitális, F., Aouadi, B., Zsom-Muha, V., & Gillay, Z. (2020). Factors Influencing the Long-Term Stability of Electronic Tongue and Application of Improved Drift. Biosensors Article. https://doi.org/doi:10.3390/bios10070074

ACKNOWLEDGEMENTS

The authors acknowledge the Hungarian University of Agriculture and Life Sciences -Doctoral School of Food Science – and Tempus Public Foundation – Stipendium Hungaricum Scholarship – for the support in the presented study.