

Generative Supervised Explainable artificial intelligence for the untargeted analysis of LCMS data

Andrea Jr Carnoli, Melissa Broeren, Ane Arrizabalaga Larrañaga, Bas H.M. van der Velden, Marco H. Blokland

Wageningen Food Safety Research, Wageningen, Netherlands

Introduction

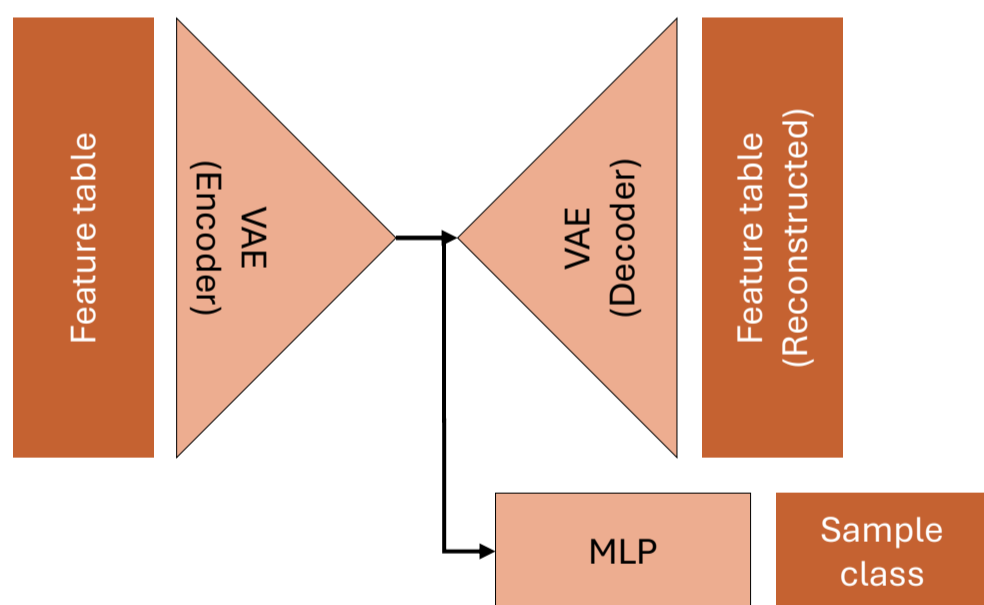
Feature tables obtained from untargeted Liquid Chromatography – High Resolution Mass Spectrometry studies are very common in food safety studies. Artificial Intelligence can be very useful, but its application is constrained by the limited amount of data. In this work we studied how transfer learning can be helpful, even when the tables are derived from completely different experiments

Conclusion

- Transfer Learning is helpful for the development and application of Artificial Intelligence models
- Transfer Learning can be applied even if the experiments are different

Artificial Intelligence

Beta – Branched Variational Autoencoder



Pretraining

Open-Access steroids data

- 1241 bovine urine
- 5% spiked
- Recorded across 3 years



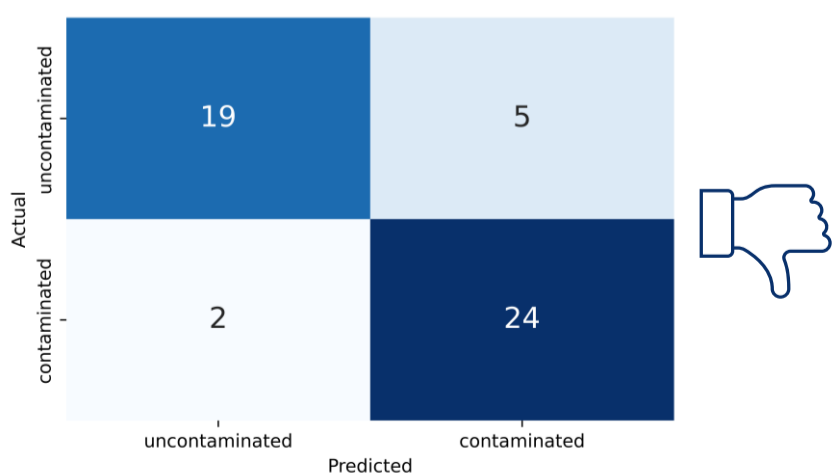
Finetuning

Chicken muscles data

- 156 chicken muscles:
- 50% spiked
- Recorded across 2 days

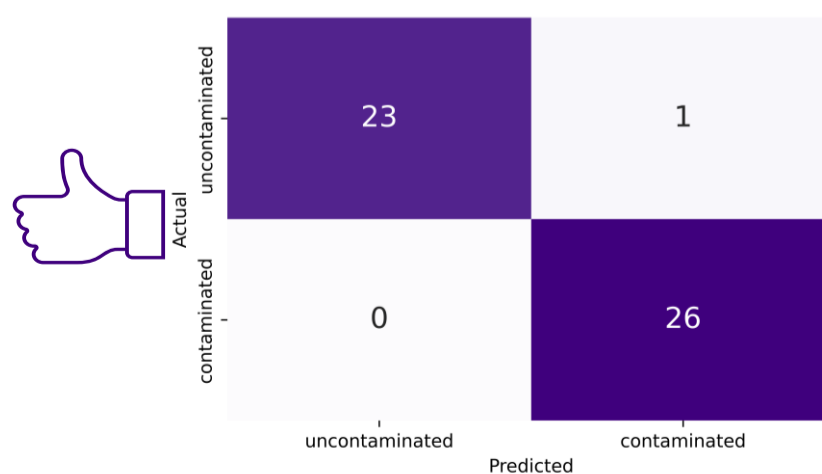
$$Loss_{\beta-BVAE} = \alpha Loss_{rec} + \beta Loss_{KLD} + \gamma Loss_{class}$$

Without transfer learning



- Misleading results (Acc:0.86)
- Shortcut learning (day)

With transfer learning



- Performance improved (Acc:0.98)
- No shortcuts