

OBJECTIVES

- Evaluate the obtainable classification performance from a breath-spiking experiment involving an electronic nose
- Discover how a trusted non-parametric method such as k-Nearest Neighbours (k-NN) performs in a classification task against Long Short Term Memory (LSTM) neural network.

METHODS

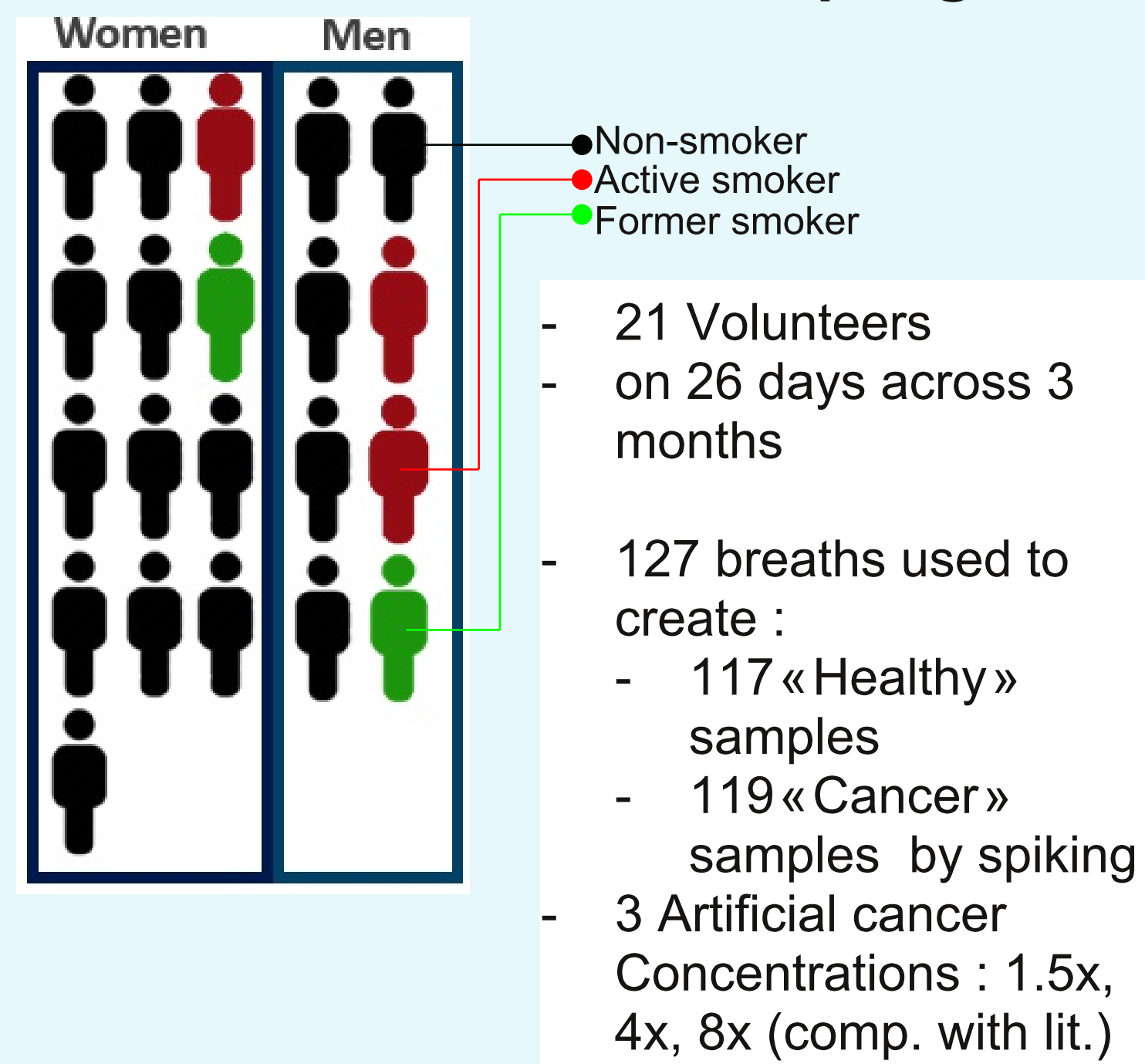
The electronic nose : SAMBre Sensors

- T2603 (Figaro®)
- 3530T, 1430T, 2530T, 8530T (Umwelt Sensor Technik®)
- MP901 (Winsen®)
- BME680 (Bosh®)

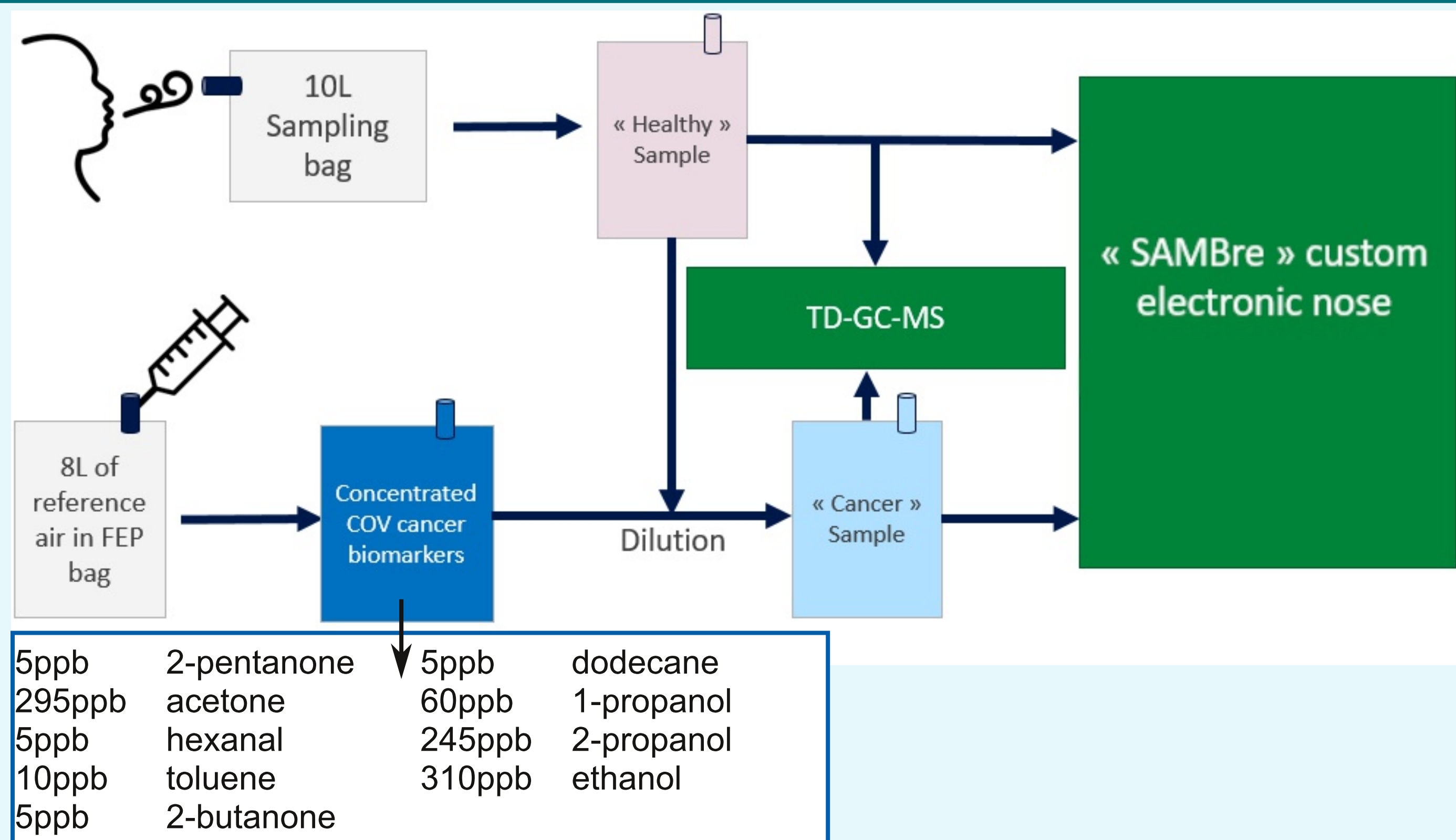
Specifications

- Temperature maintained at 45°C within chamber
- Flow constant at 200mL/min
- Moisture and temperature monitored

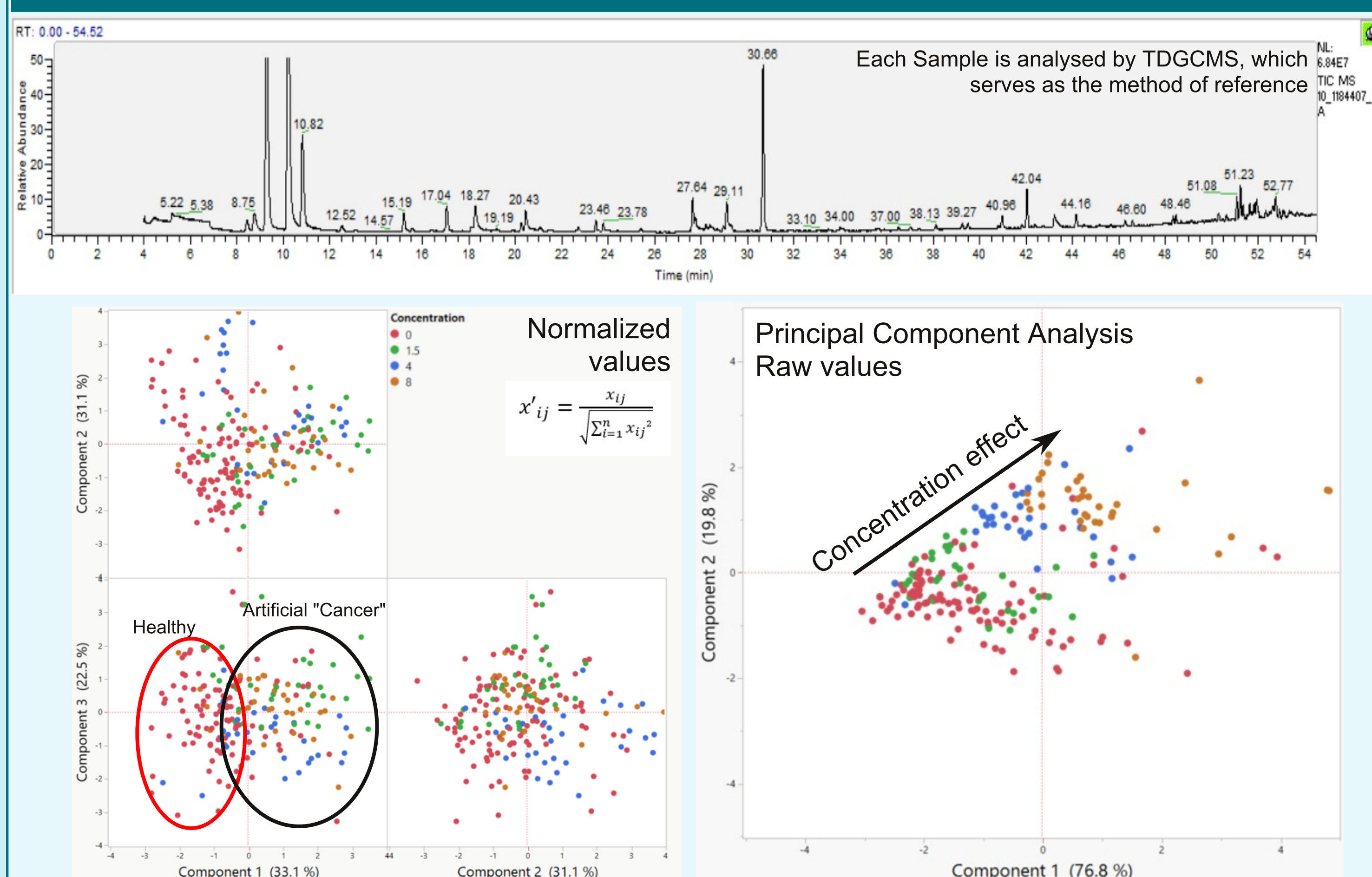
Human breath sampling



TESTING PROCEDURE

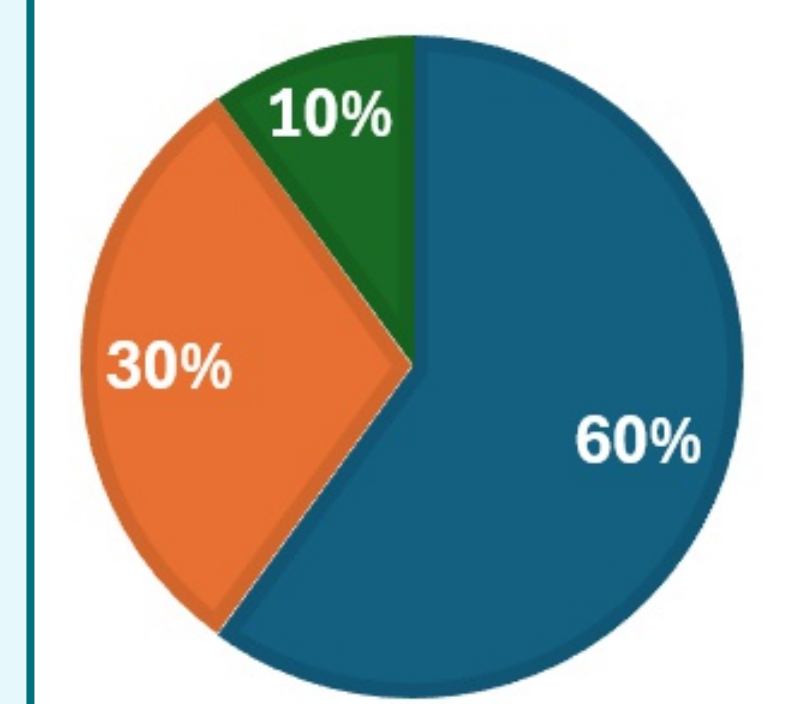


SAMPLE ANALYSIS



MACHINE LEARNING

Training Validation Test



For both models

- Same share of dataset for training, validation and external testing
- External testing created with data from a separate day
- Same initial experiment
- 5-fold cross-validation

LSTM

- Time series of 315 seconds per sample
- 100 memory blocks
- Adam optimizer
- Rescaling to 0-1 range
- Done using Python



k-NN

- 1 data per sample (maximum baseline-corrected conductance increase)
- Best model out of the first 30 values of k is selected for validation and testing
- Done using JMP Software



RESULTS

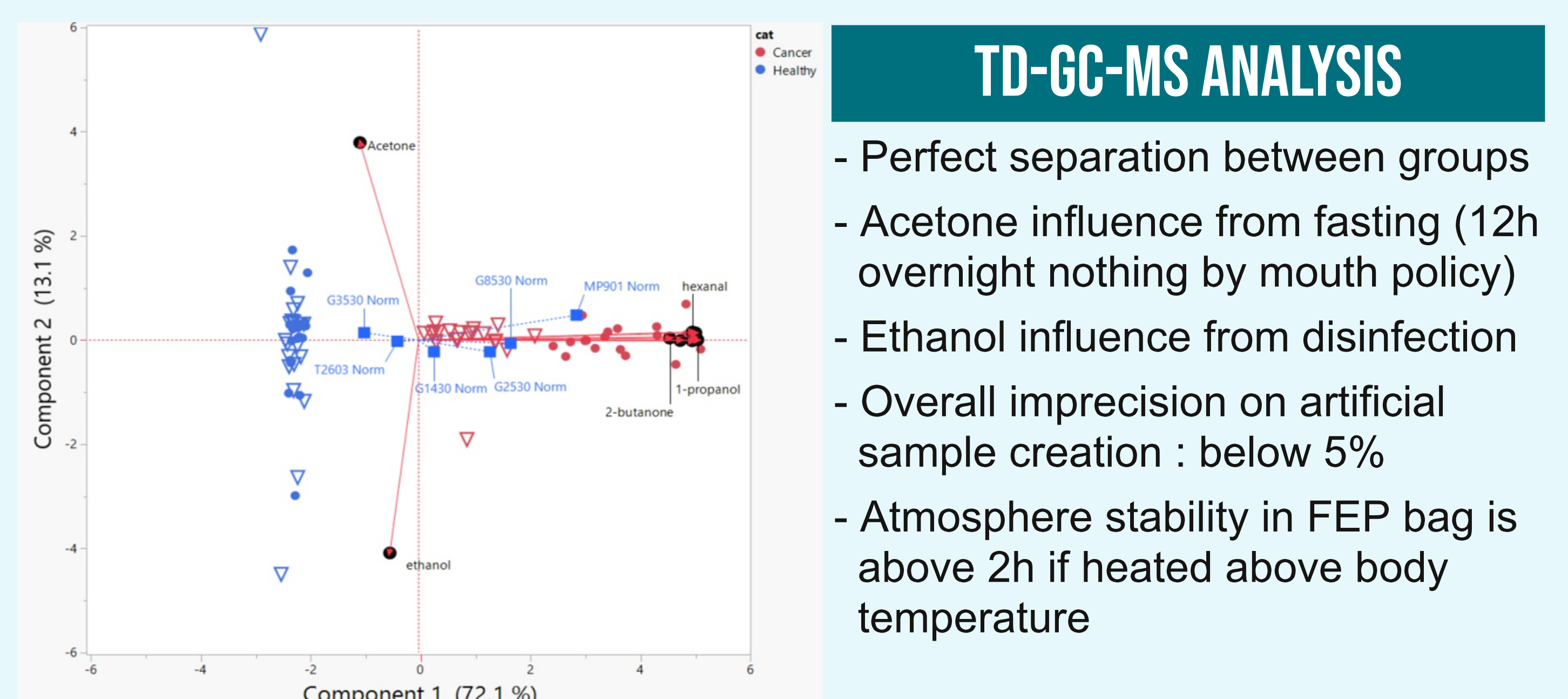
LSTM

- 88±2% accuracy (training)
- 80±5% accuracy (validation)
- 81±4% accuracy (test)
- Using the "time series" dataset on kNN did not produce any performance improvement.
- Underperforming
- No additional information within time series ?

k-NN

- 90±3% accuracy (training)
- 91±2% accuracy (validation)
- 91±2% accuracy (test)
- Best k between 1-16
- Best contributing sensors (to discriminating PC)
 - G2530 (62%)
 - MP901 (23%)
 - G3530 (11%)
 - T2603 (4%)

TD-GC-MS ANALYSIS



CONCLUSIONS

- Data from 246 individual measurements on breath and spiked breath was collected and successfully used to train a k-NN model and LSTM network.
- LSTM did not perform significantly better than k-NN, but this conclusion should be confirmed by training both on future clinical trial data.
- Overall, classification error is low but cancer biomarkers concentrations had to be raised, which underlines the need for a preconcentration step.



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