**Remarks 2018-08-15**

General

* Expectations of the tests should be added (what was expected before the doing the tests? Did they results follow the expectations?)
* Add steps that might be obvious for me, but not for the reader (normalization, classification choice, etc)
* Make plots clearer with for example numbers

Feature selection

* Add 3D plot or pareto front to combine accuracy, features and computation time

TPOT

* Try to add changes with to TPOT on level with: population size, number of pipeline steps, mutation options, timeout – time etc. (Possibly together with Pieter)
* Think of what to do with having an accuracy of lower than 0 (Change the threshold?)
* Ask Koen (BMT) for extra computation power