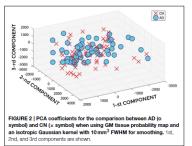
### Main topics

- Machine learning and the «data life cycle»
  - Feature extraction
  - Feature selection
  - Classification
- Validation and classification metrics (or how to stop worrying and setup the model)
- Radiomics and radiogenomics
- Deep-learning techniques
- How to build a model suited for the medical-data domain



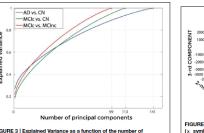


FIGURE 3 | Explained Variance as a function of the number of considered Principal Components, when using GM tissue probability map and no smoothing, for the following comparisons: AD vs. CN, MCIc vs. MCInc.

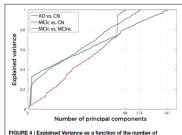


FIGURE 4 | Explained Variance as a function of the number of considered principal components sorted in accordance to their FDR, when using GM tissue probability map and no smoothing, for the following comparisons: AD vs. CN, MCIc vs. CN, MCIc vs. MCInc.

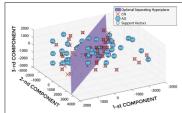
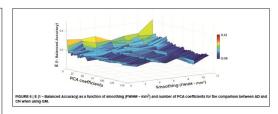
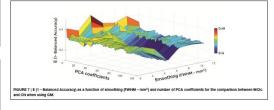
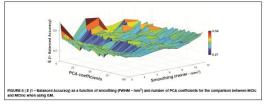


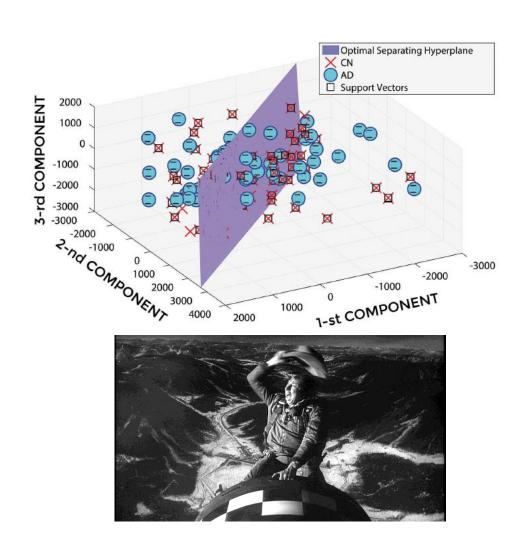
FIGURE 5 I Hyper-plane plane separating AD (o symbol) from C (x symbol) PCA coefficients (3 PCA coefficients), and defined Support Vectors (C symbol), when using GM tissue probability map and an isotropic Gaussian kernel with 10 mm<sup>3</sup> FWHM for smoothing, 1st, 2nd, and GM components are shown.



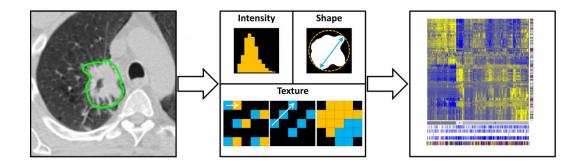




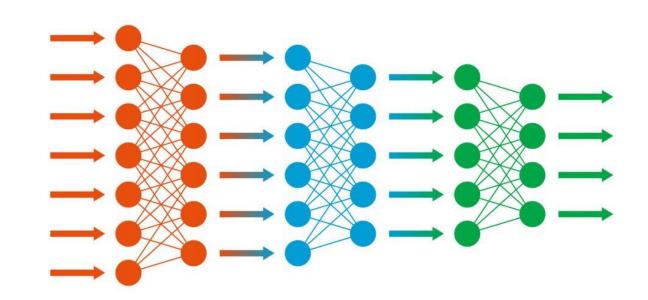
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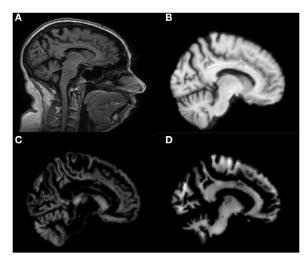
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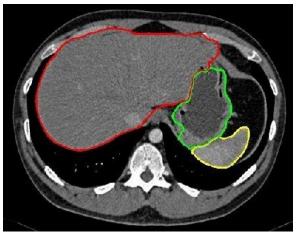


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### Requirements

- Basic skills of Matlab and Python
- Updates @ https://christiansalvatore.github.io/machinelearning-iusspavia





