


# BrainHack 2019

## #Dementia\_Forecast\_Group

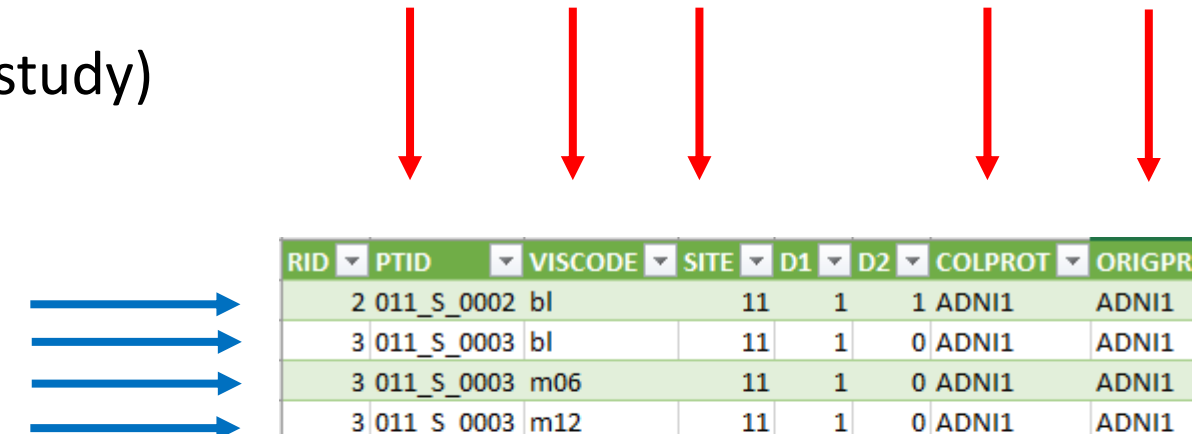
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23/03/2019



# Understanding the data!!

- 1737 patients
- 12741 **instances** (longitudinal study)
- 1906 **features**
- Goal:
  - Diagnosis
  - ADAS-13
  - Ventricles volume/intracranial volume



RID	PTID	VISCODE	SITE	D1	D2	COLPROT	ORIGPR
2	011_S_0002	bl	11	1	1	ADNI1	ADNI1
3	011_S_0003	bl	11	1	0	ADNI1	ADNI1
3	011_S_0003	m06	11	1	0	ADNI1	ADNI1
3	011_S_0003	m12	11	1	0	ADNI1	ADNI1
3	011_S_0003	m24	11	1	0	ADNI1	ADNI1
4	022_S_0004	bl	22	1	0	ADNI1	ADNI1
4	022_S_0004	m06	22	1	0	ADNI1	ADNI1
4	022_S_0004	m12	22	1	0	ADNI1	ADNI1
4	022_S_0004	m18	22	1	0	ADNI1	ADNI1
4	022_S_0004	m36	22	1	0	ADNI1	ADNI1
5	011_S_0005	bl	11	1	0	ADNI1	ADNI1

Group	Var	description
Demo	AGE	Age at BASELINE, not current age
	PTGENDER	Sex
Genotype	APOE4	APOE4 minor allele load
sMRI	STNNCV_*	Regional Volume
	STNNNSA_*	Regional surface area
	STNNMTA_*	Regional thickness average
	STNNNSD_*	Regional thickness standard deviation
PET	*UCBERKELEYAV1451*	Regional PET measures of abnormal Tau
	*UCBERKELEYAV45*	Regional PET measures of abnormal Abeta
	*BAIPETNMRC*	Regional PET measures of glucose metabolism
	FDG / PIB / AV45	Average relative PET measures
DTI	FA_*	Regional fractional anisotropy
	MD_*	Regional mean diffusivity
	AD_*	Regional axial diffusivity
	RD_*	Regional radial diffusivity
CSF	ABETA_*	CSF Amyloid Beta
	TAU_*	CSF Tau
	PTAU_*	CSF phospho-Tau

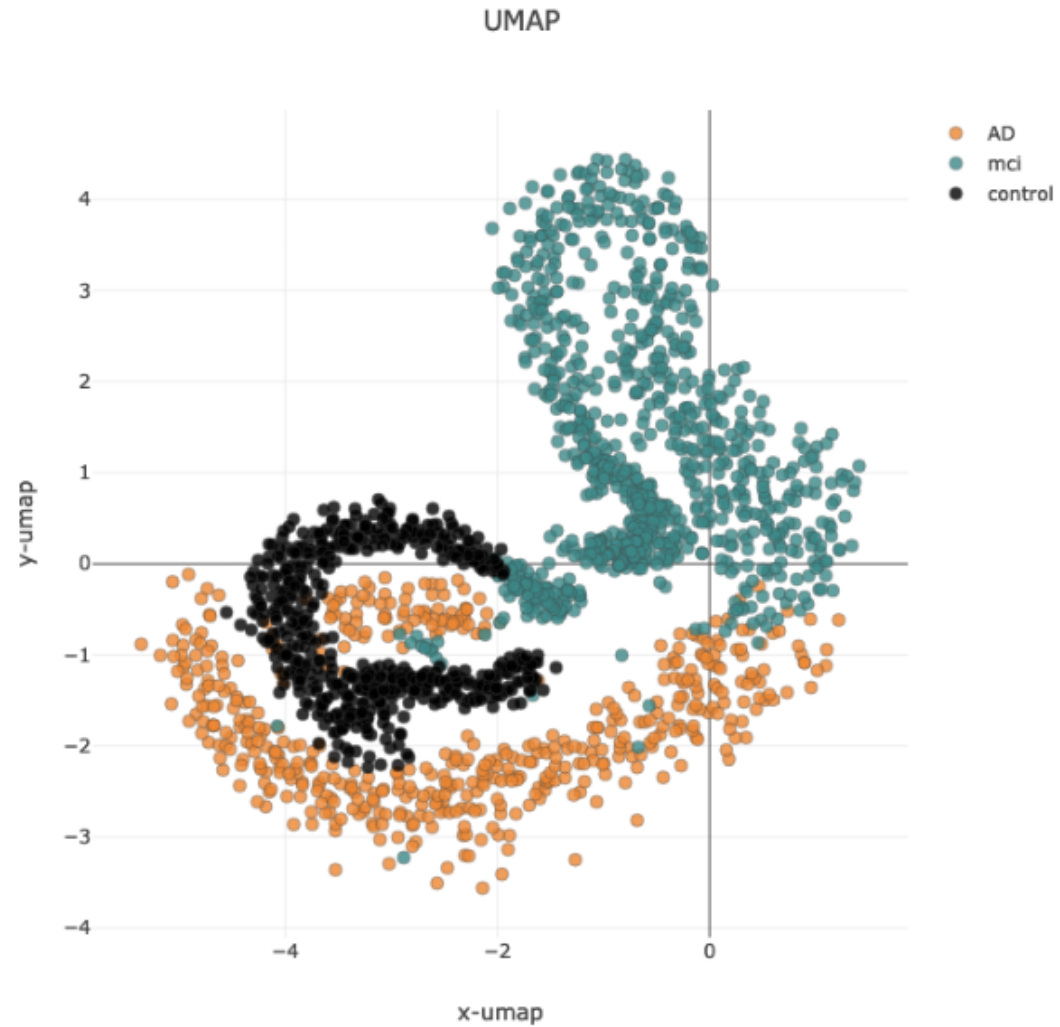
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# Data handling

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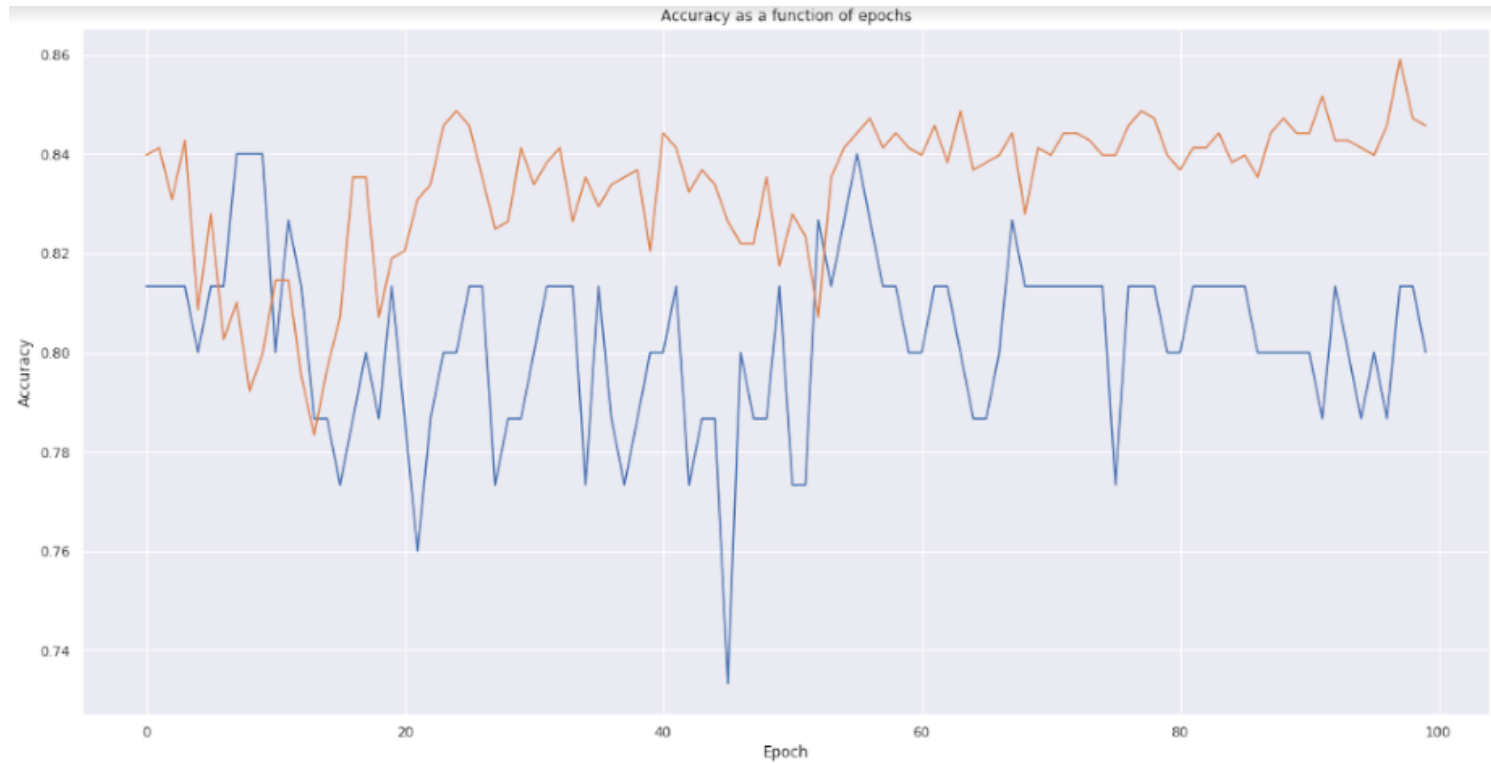
- **Training data:** up to April, 2010
- **Test data:** from April 2010, onwards
- **Missing values:** e.g. for structural MRI:
  - Total = 442 features
  - 19 features not available (unavailable for all patients)
  - 225 incomplete features (available only for some patients)

# Dimensionality reduction



# Training a Neural Net

- Trying to predict the Umap embeddings using a neural network on the test set



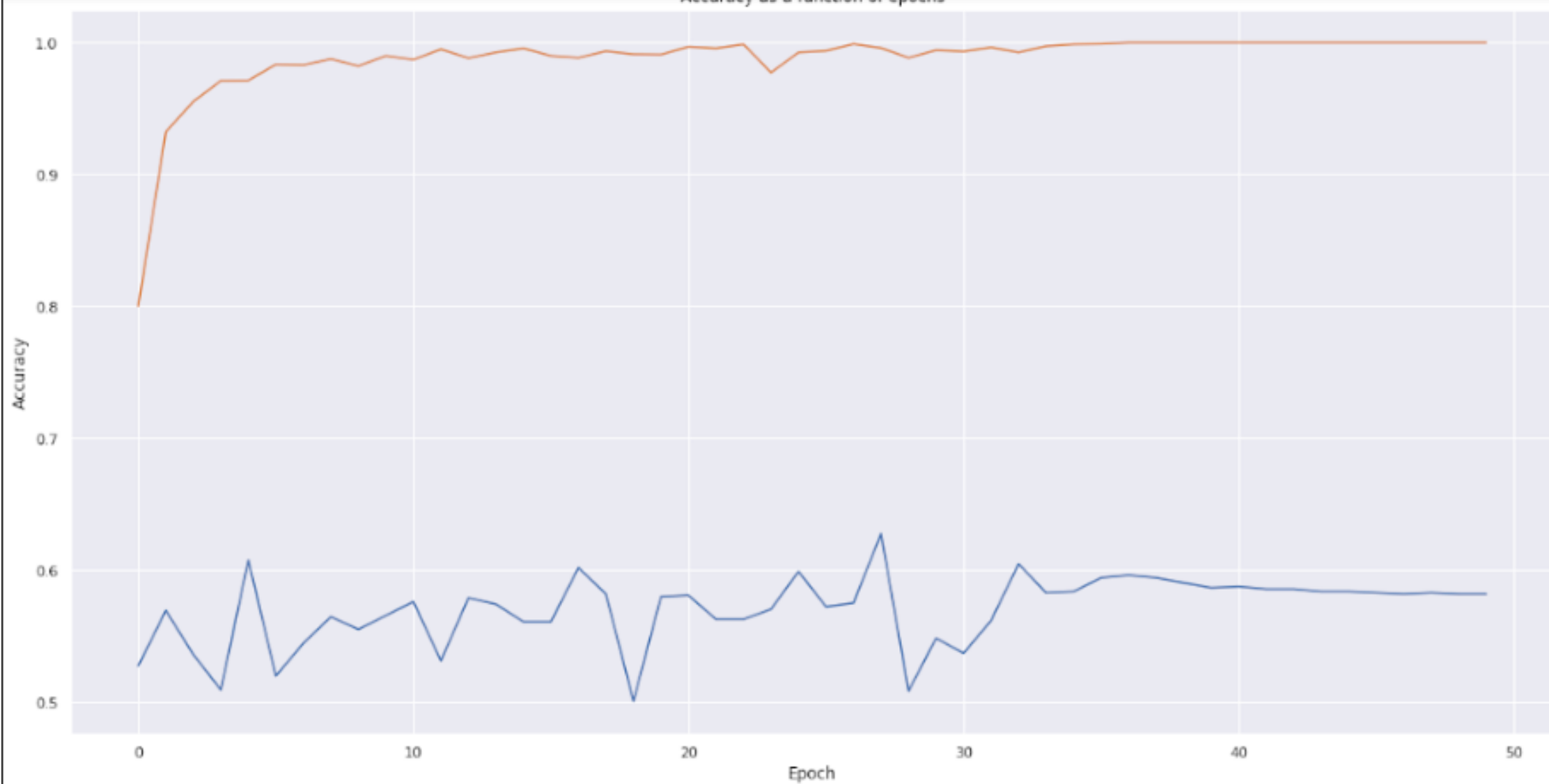
# Trying out classifying the diagnosis

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- Using **only** CSF features with FFNN: overall diagnosis accuracy 57%
- Using the template SVM file provided by challenge organizers:
  - Be careful of dropping the feature to classify!
  - Code not optimized
  - Recommend writing one's own code

Accuracy as a function of epochs

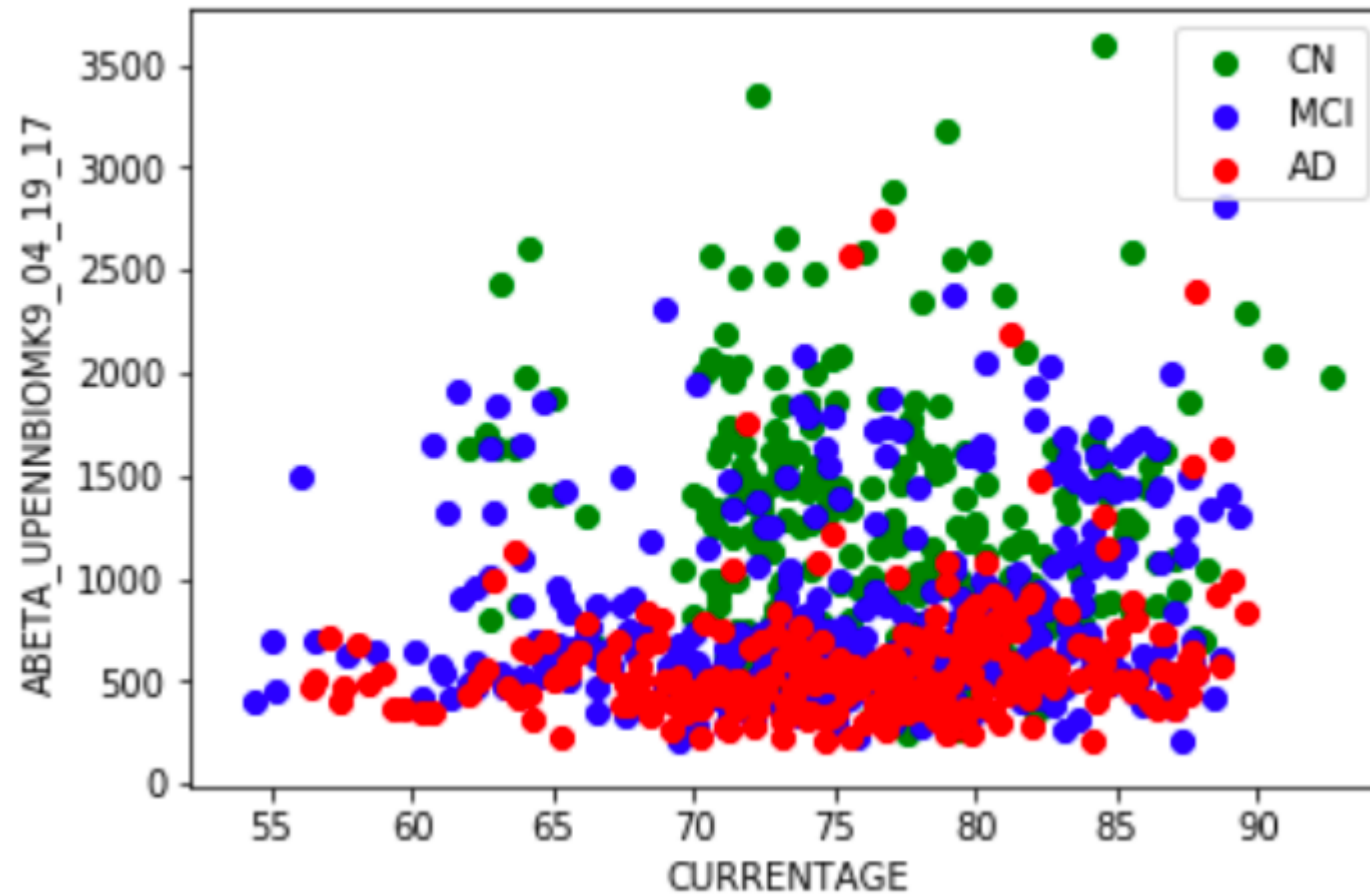
Accuracy as a function of epochs





# Example of variable distribution

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# Predictions with and without age extrapolation

- Classification: random forest
- Regression: SVR

```
AGE
ADAS13
Ventricles
ICV_b1
ST44CV_UCSFFSL_02_01_16_UCSFFSL51ALL_08_01_16
ST105CV_UCSFFSL_02_01_16_UCSFFSL51ALL_08_01_16
ST121TS_UCSFFSL_02_01_16_UCSFFSL51ALL_08_01_16
ABETA_UPENNBIOBK9_04_19_17
TAU_UPENNBIOBK9_04_19_17
PTAU_UPENNBIOBK9_04_19_17
CURRENTAGE
Ventricles_ICV
```

with

```
(9240,)
Evaluate predictions
Diagnosis:
mAUC = 0.679
BAC = 0.621
ADAS:
MAE = 10.859
WES = 9.190
CPA = 0.449
VENTS:
MAE = 8.109e-03
WES = 8.023e-03
CPA = 0.500
```

without

```
mAUC = 0.665
BAC = 0.624
ADAS:
MAE = 8.171
WES = 8.217
CPA = 0.431
VENTS:
MAE = 7.058e-03
WES = 6.898e-03
CPA = 0.500
```

# Trying out XGBOOST

Explore hyperparameter  
optimization with boosting

