



# Python computational pipeline for predictive machine learning modelling of livestock data

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# Summary

- **What you get from this workshop**

- Some (hopefully functional) Python code ... for regression problems (due to time constraints)
  - The code relies on the Python scikit-learn library  
<https://scikit-learn.org/>
- Code available at:
  - [https://github.com/National-Animal-Nutrition-Program/2024ASAS\\_Tulpan](https://github.com/National-Animal-Nutrition-Program/2024ASAS_Tulpan)
- Some information and explanations of what the code does and why

- **Assumptions**

- You know a bit about machine learning
  - If not, read this: *Greener et al. (2021): A guide to machine learning for biologists*  
(<https://www.nature.com/articles/s41580-021-00407-0>)
- You can operate a computer

# Warnings / Disclaimers



- Python code is not optimized or comprehensive
  - It is built to (hopefully) facilitate understanding
  - Sacrificed performance and best programming practices
- Input datasets are assumed to be ready and clean
  - Your job
- The code should only be used for good causes
- If you make money with this code my share is 10% (cash, check or plastic is fine) 😎

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# Python Use

- Follow the instructions provided in the `"Python_usage_instructions.pdf"` file

# Data formatting

- Expectations:
  - Tabular format
  - Last column contains the predictor variable
  - Data was cleaned prior to using the Python script
  - Data includes only numeric values
- Recommended reading:
  - *Browman and Woo (2018) - Data Organization in Spreadsheets* (<https://www.tandfonline.com/doi/full/10.1080/00031305.2017.1375989>)

# Data sets (for this workshop)

- 2 subsets of the data from:

Marshall et al. (2023): A farmer-friendly tool for estimation of weights of pigs kept by smallholder farmers in Uganda

- **Article:** <https://link.springer.com/article/10.1007/s11250-023-03561-z>
- **Data:** <https://data.mel.cgiar.org/dataset.xhtml?persistentId=hdl:20.500.11766.1/FK2/IWXZQH>

# MarshallEtAl2023\_more\_selected\_measurements.csv

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC10205837/>

## Records

754 pigs from Uganda

## Variables

- 6 input variables:
  - household\_id
  - age\_months
  - heartgirth
  - height
  - length
  - body\_condition\_score
- 1 output variable:
  - exact\_weight

MarshallEtAl2023_more_selected_measurements							
	household_id	age_months	heartgirth	height	length	body_condition_score	exact_weight
1							
2	PBM-KML-113	34	140	901	141	4	205
3	PBM-MSK-138	24	0	0	0	4	200
4	PBM-MSK-107	15	130	80	138	4	193.2
5	PBM-MSK-106	41	140	76	141	4	177.2
6	PBM-WKS-401	27	128	85	140	4	170
7	PBM-KML-106	30	121	72	140	4	160
8	PBM-MSK-137	19	124	76	142	4	148
9	PBM-WKS-401	24	122	81	136	3	137.7
10	PBM-MSK-139	18	134	89	147	3	134
11	PBM-MSK-102	20	117	81	149	4	132.9
12	PBM-MSK-142	13	121	80	140	4	131.5
13	PBM-WKS-416	43	120	72	145	3	131.1
14	PBM-HMA-240	12	113	90	137	3	129.5
15	PBM-MSK-107	12	112	78	136	4	127.3
16	PBM-MSK-102	20	122	77	135	4	126.5

# KabululuEtAl2023\_selection.csv

<https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0295433>

## Records

400 pigs from Tanzania

## Variables

- 6 input variables (all numeric):
  - Heart\_girth
  - Body\_length
  - Age\_months
  - Sex
  - District
  - Village
- 1 output variable:
  - Weight\_kg\_by\_scale

KabululuEtAl2023_selection						
HEART_GIRTH	BODY_LENGTH	AGE_MONTHS	SEX	DISTRICT	VILLAGE	WEIGHT_KG_BY_SCALE
118	137	24	Female	Mbeya	Mshewe	91
69	95	6	Female	Mbeya	Njelenje	33
65	97	8	Female	Mbeya	Njelenje	26
66	71	9	Female	Mbeya	Njelenje	20
72	90	12	Male	Mbeya	Njelenje	37
67	93	17	Female	Mbeya	Njelenje	34
78	89	12	Female	Mbeya	Njelenje	42.5
80	90	8	Male	Mbeya	Mjele	37
61	74	7	Female	Mbeya	Mjele	23
60	80	8	Female	Mbeya	Mjele	25
86	115	14	Female	Mbeya	Mjele	59
80	94	9	Male	Mbeya	Mjele	42
72	91	8	Female	Mbeya	Mjele	29
70	90	12	Female	Mbeya	Mjele	28
55	69	8	Female	Mbeya	Mjele	17
76	86	8	Female	Mbeya	Mjele	31
71	79	8	Female	Mbeya	Mjele	39
63	78	8	Female	Mbeya	Mjele	20
57	62	8	Female	Mbeya	Mjele	18
68	80	8	Female	Mbeya	Mjele	26
52	70	6	Female	Mbeya	Mjele	14
63.5	78	10	Male	Mbeya	Mjele	25



# RemusEtAl2020\_lysine\_requirements.csv

<https://www.sciencedirect.com/science/article/pii/S1751731119002660>

## Records

40 pigs from Canada

## Variables

- 8 input variables:
  - Pig\_identification
  - Methionine\_intake
  - Met+Cys\_intake
  - Cys\_intake
  - ADG (avg. daily gain)
  - ADFI (avg. daily feed intake)
  - BW\_initial
  - BW\_final
- 1 output variable:
  - Lys\_intake

RemusEtAl2020_lysine_requirements									
	PIG_identification	Methionine_intake	Met_Cys_intake	Cys_intake	ADG	ADFI	BWinitial	BWfinal	Lysine_intake
1	1	5.56	8.92	3.36	0.85	1.65	25.10	45.60	18.53
2	2	9.51	13.93	4.42	0.96	2.15	27.50	50.45	24.38
3	4	3.06	5.47	2.41	0.45	1.30	24.85	35.65	14.59
4	7	6.97	11.18	4.21	1.01	2.08	27.70	51.95	23.22
5	8	4.52	8.08	3.56	0.91	1.92	25.55	47.46	21.54
6	9	5.45	8.75	3.30	1.02	2.09	25.00	49.45	18.17
7	14	8.85	12.97	4.12	1.08	2.04	25.10	50.90	22.70
8	15	5.62	10.03	4.41	1.10	2.36	25.20	51.60	26.74
9	16	6.00	8.79	2.79	0.69	1.36	22.20	38.70	15.39
10	17	6.18	9.92	3.74	0.85	1.81	22.95	43.45	20.61
11	18	7.71	11.30	3.58	1.01	1.77	22.30	46.65	19.77
12	19	4.46	7.97	3.51	1.03	1.88	25.85	50.45	21.24
13	21	6.40	10.28	3.87	1.01	1.90	25.95	50.25	21.35
14	23	6.40	9.48	3.07	0.99	2.21	25.80	45.05	16.42
15	25	4.48	7.20	2.71	0.79	1.35	23.75	42.60	14.95
16	26	5.86	8.70	2.84	0.94	2.15	30.15	45.75	15.02
17	27	5.10	9.11	4.01	1.08	2.15	27.35	53.25	24.28
18	29	8.83	12.94	4.11	1.04	2.01	24.80	49.80	22.65
19	31	6.25	10.02	3.78	0.88	1.84	27.40	48.50	20.82
20	33	4.37	7.80	3.43	0.90	1.86	24.60	46.10	20.80
21									

# Regression pipeline

1. Data cleaning
2. Data summarization
3. Data visualization
4. Data splitting
5. Data scaling
6. Model initialization (default params)
7. Preliminary model evaluation
8. Overfitting analysis of default models
9. Hyper-parameter optimization
10. Update model hyper-parameters
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# Regression pipeline

## **1. Data cleaning**

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# Data cleaning



- Remove rows with missing values
- Remove duplicate rows
- Remove duplicate columns
- Remove single value columns
- Find and remove outliers (Z-score method)
- Change categorical columns to numeric
- Save cleaned dataset

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# Overall look at the data

- Check the size of the dataset
  - Number of records (rows)
  - Number of variables/features (columns)
- Look at the first few records
- Look at descriptive statistics
  - Check for obvious outliers or extreme values

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# Explore the data visually first

- If feasible/applicable
- Check the distribution of the variables
  - Histograms
  - Scatter-plots
- Check correlations among variables/features



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# Prepare data for modelling

- Separate data into training (80%) and testing (20%)
  - The percentages depend on data size, available time, goals
- Training set:
  - Model construction
  - Model validation
  - Hyper-parameter optimization
- Testing set:
  - Testing the final models

## Golden Rule of Machine Learning

**NEVER EVER** use the testing set during the construction/validation/optimization stage of a model.

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# Scaling your data

- How
  - Transform data to a standardized range
  - [StandardScaler](#), [MinMaxScaler](#), [RobustScaler](#)
- Why
  - Reduces the impact of extreme values
    - ... for algorithms sensitive to outliers or for those relying on normality assumptions
  - Reduces differences in value scales among variables
    - Speeds up convergence and provides equal opportunities for features to influence the outcome variable
  - Helps making more robust models

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# ML Models

- Select models from different categories
  - Tree-based: Decision Tree, AdaBoost, Random Forest
  - Artificial Neural Networks: Multi Layer Perceptron
  - Lazy estimators: K-Nearest Neighbour
  - Linear: Linear Model, LASSO, Ridge
  - Gradient-based: Gradient Boost
- Select more than 2 models
  - Different strengths and weaknesses
  - Different data representations

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# Model evaluation strategies

- K-fold cross-validation
  - Choose K as a function of data size and computing time
    - High K values: small-medium datasets
    - Low K values: large datasets
- Choose your measures/”metrics”
  - Regression
    - Errors: MAE, MSE, RMSE, MAPE, ...
    - Correlation coefficients: Pearson, Spearman, Kendal, Concordance (CCC)
    - $R^2$
  - Classification
    - Confusion matrix-based: F1-score, precision, recall (TPR, sensitivity), accuracy, ... [NOT USED IN THE CURRENT CODE -- NA]

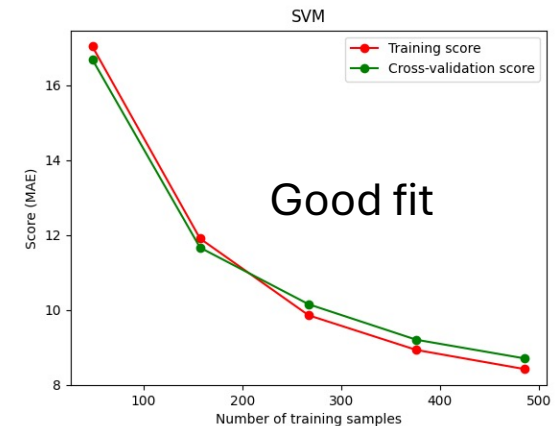
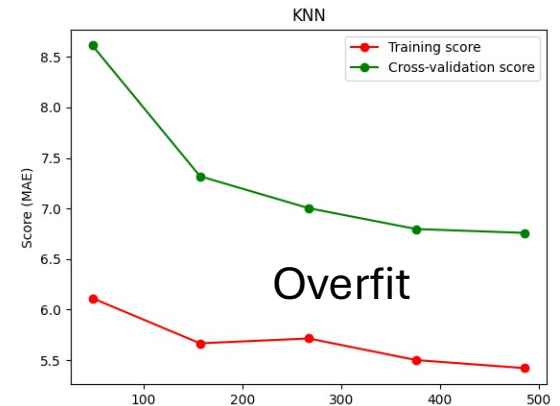
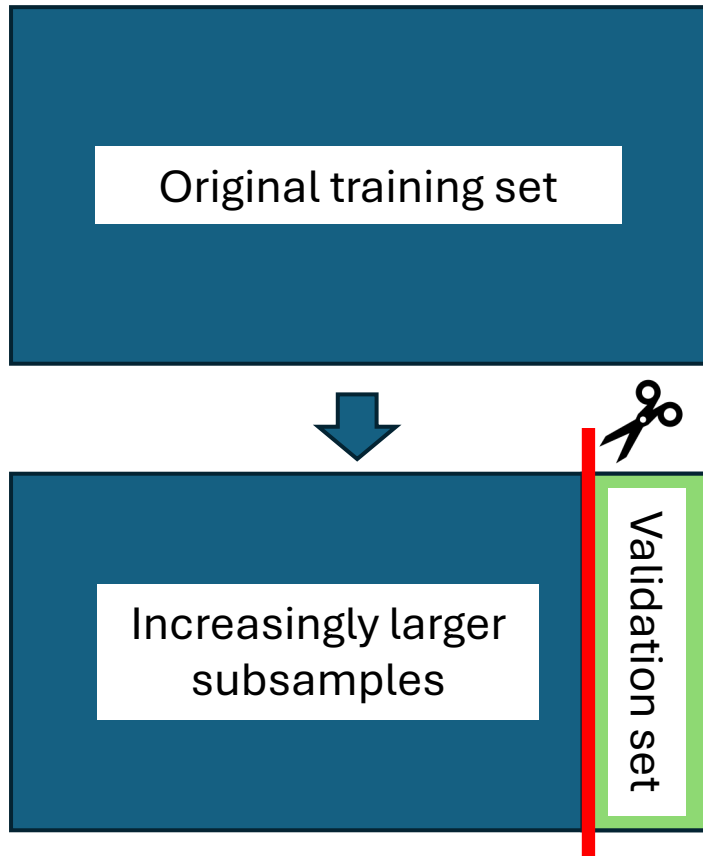


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# Overfitting analysis

- Use learning curves
  - training vs. validation scores for increasing training set sizes



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# Hyper-parameter optimization

- Hyper-parameter = user-tunable parameter

Grid search

Parameter 2	$v_1$	✓	✓		✓
	$v_2$	✓	✓		✓
	...				
	$v_n$	✓	✓		✓
		$v_1$	$v_2$	...	$v_m$
		Parameter 1			

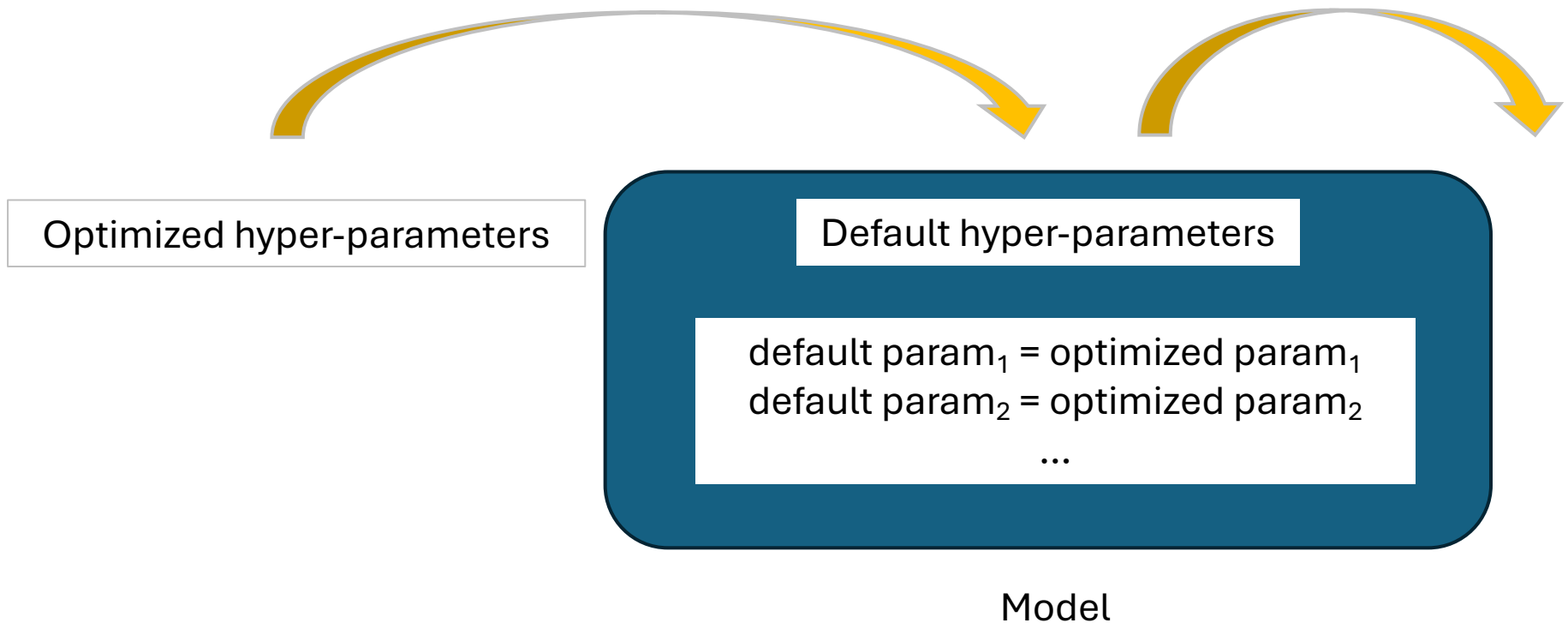
Random search

Parameter 2	$v_1$		✓		✓
	$v_2$				
	...				
	$v_n$	✓			✓
		$v_1$	$v_2$	...	$v_m$
		Parameter 1			

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# Hyper-parameters' update



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# Model evaluation (same as for 7)

- K-fold cross-validation
  - Choose K as a function of data size and computing time
    - High K values: small-medium datasets
    - Low K values: large datasets
- Choose your measures/”metrics”
  - Regression
    - Errors: MAE, MSE, RMSE, MAPE, ...
    - Correlation coefficients: Pearson, Spearman, Kendal, Concordance (CCC)
    - $R^2$
  - Classification
    - Confusion matrix-based: F1-score, precision, recall (TPR, sensitivity), accuracy, ... [NOT USED IN THE CURRENT CODE -- NA]

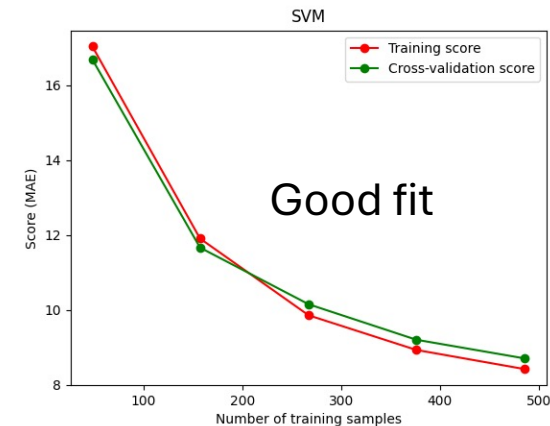
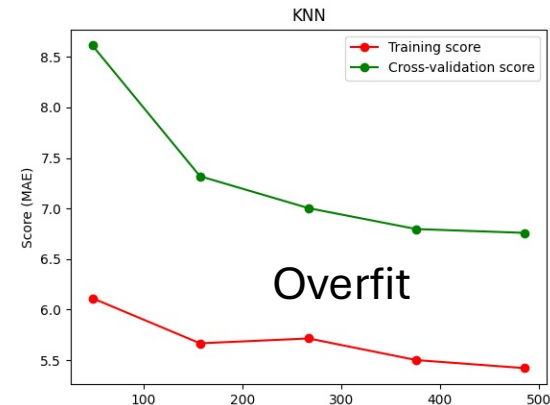
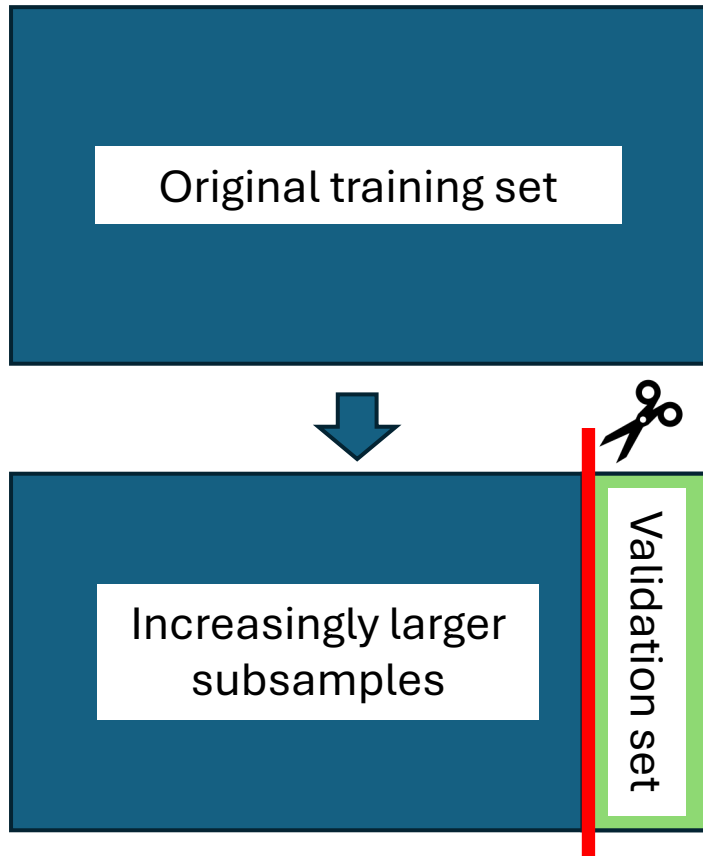


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# Overfitting analysis (same as for 8)

- Use learning curves
  - training vs. validation scores for increasing training set sizes



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# Saving models

- Backup all optimized models
- Can be used later for deployment
- Save time on re-training and re-optimizing hyper-parameters

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# Feature importance

- Use a model-agnostic process
- Permutation Feature Importance (PFI)
  - Shuffle one variable at a time
  - Evaluate each algorithm
  - Idea: if an important variable is shuffled it would hurt the model significantly (poor predictions)
- Other options: [SHAPley values](#)

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# Model evaluation on test sets

- Use various evaluation measures
  - Error-based: MAE, MSE, RMSE, MAPE
  - Correlations: Pearson Product-Moment, Concordance, Spearman
  - (Adjusted) Coefficient of determination

Note: no single evaluation measure captures everything

- Use visual analysis, too
  - Scatter plots (predicted versus true values)
  - QQ plots for prediction errors



# Thank you