# Lecture 9: Cross-validation

Monday, Oct

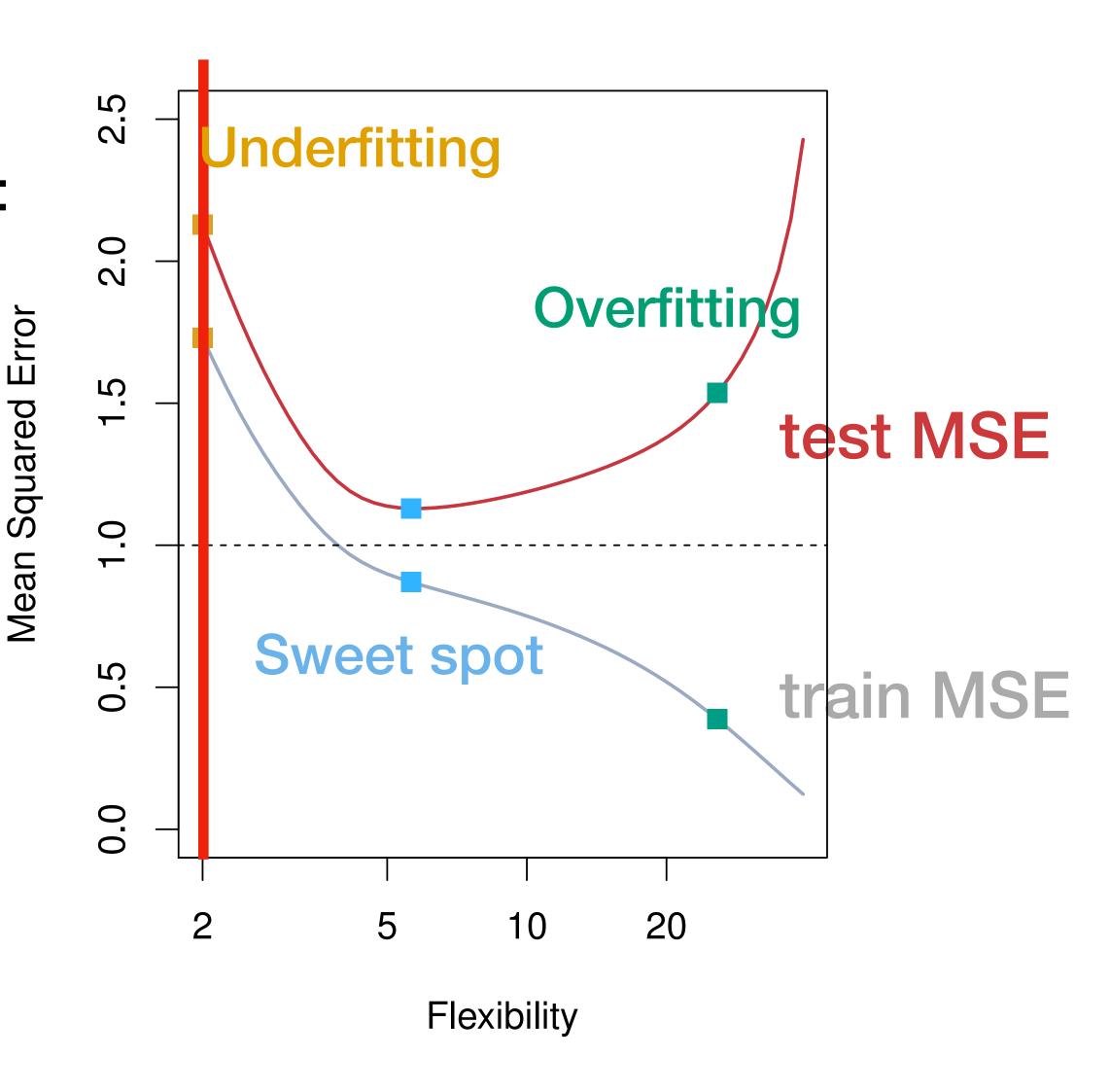
### Model assessment vs. selection

Model assessment: evaluating a model's performance on the test set via metrics such as:

- RSE and (R)MSE (regression)
- Classification error rate (classification)
- $R^2$  and Adjusted  $R^2$

Model selection / comparison: comparing the performance between competing models

 Example: selecting the proper level of flexibility in a model (e.g., number of parameters)



# Resampling

 Resampling: fitting the same model multiple times using different subsets (samples) of the original dataset

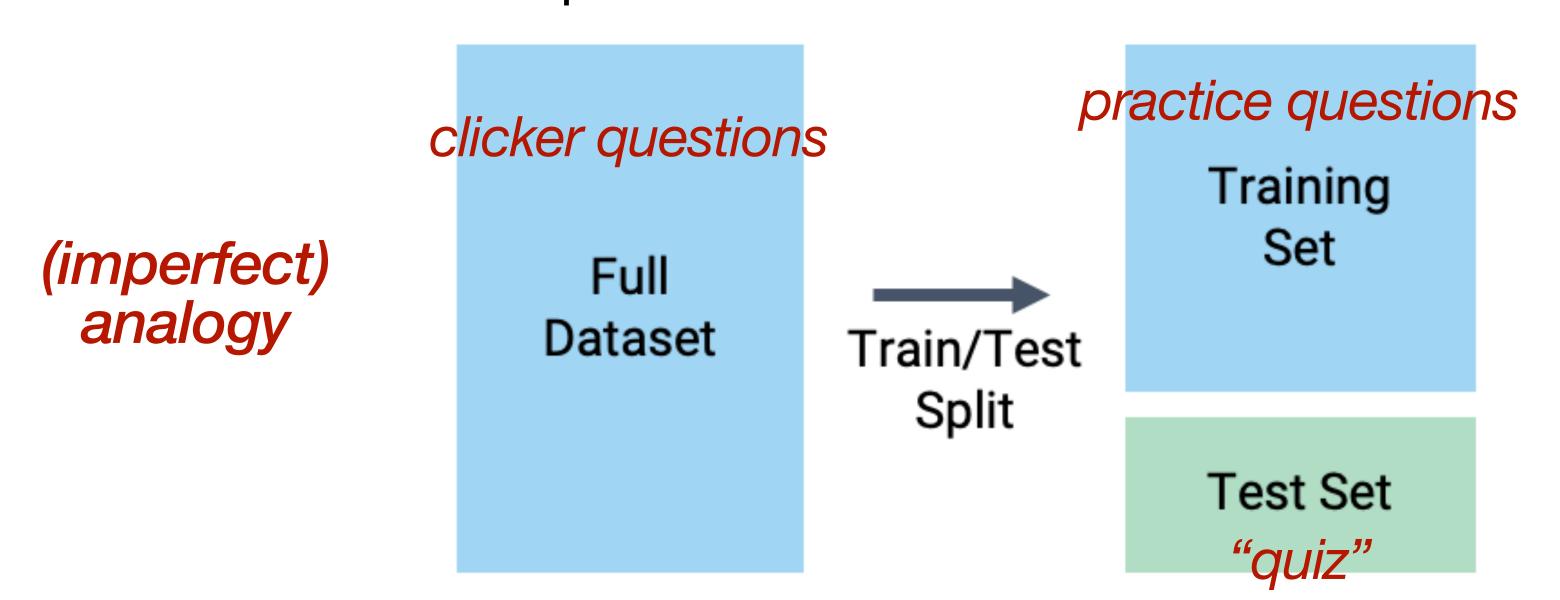
#### Two methods:

- Cross-validation: resampling the data to estimate the model's test error
- Bootstrapping: resampling the data to estimate the uncertainty around population statistics (e.g., standard errors and confidence intervals)
- ⇒ Both are used for model assessment (evaluating model performance) and selection (choosing the best model)

### Train, test, and validation set

### How do we pick the best model?

- So far: split data into train set (to train model) and test set (evaluate model performance)
- However, if we use the test set to <u>fine tune and choose the best model</u>, (e.g., by trying out models w/ different # parameters and comparing them) it is *no longer a true test set*.
- Why? A test set is supposed to be an unbiased evaluation of a final model's fit.
   We don't want test performance to influence model choice!

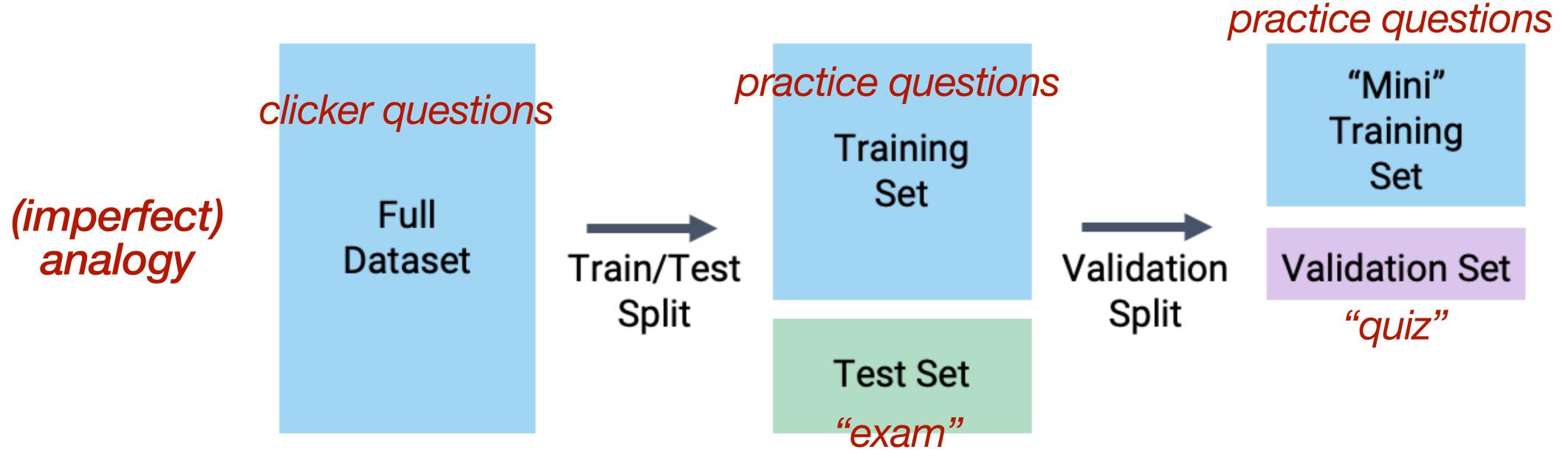


The test set must reflect truly unseen data
("exam") or else the test error becomes biased!

### Train, test, and validation set

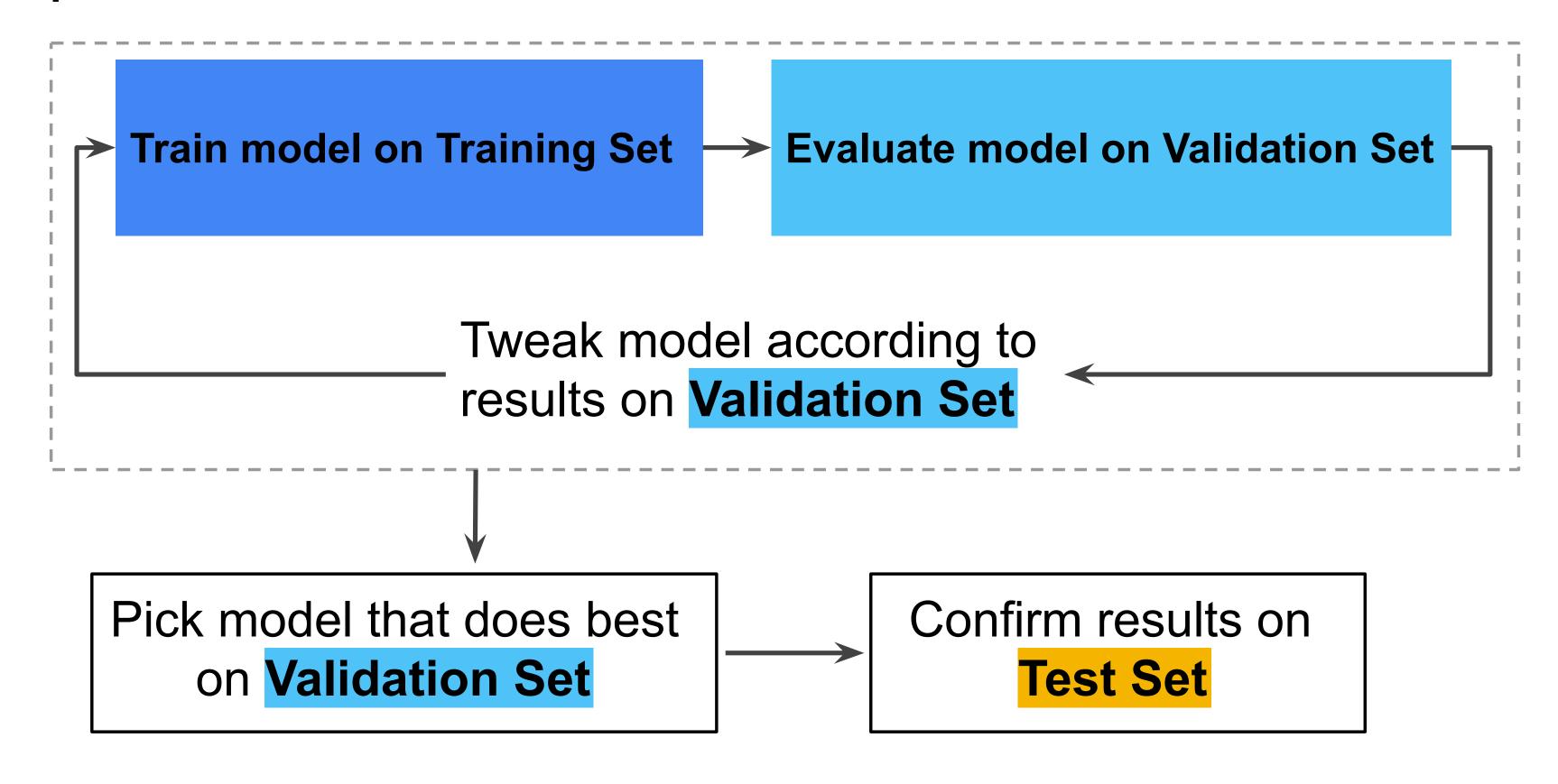
How do we pick the best model?

- Solution: split the dataset into:
  - 1. training set (60-80% of full dataset)
  - 2. validation set (~10-20% of full dataset) (used as an estimate of the test error)
  - 3. test set (~10-20% of dataset)



### Train, test, and validation set

How do we pick the best model?



HW4 Part B: Using validation sets to fine tune your model and test the best one on the held-out test set!

# Validation set approach

### Using the validation set approach:

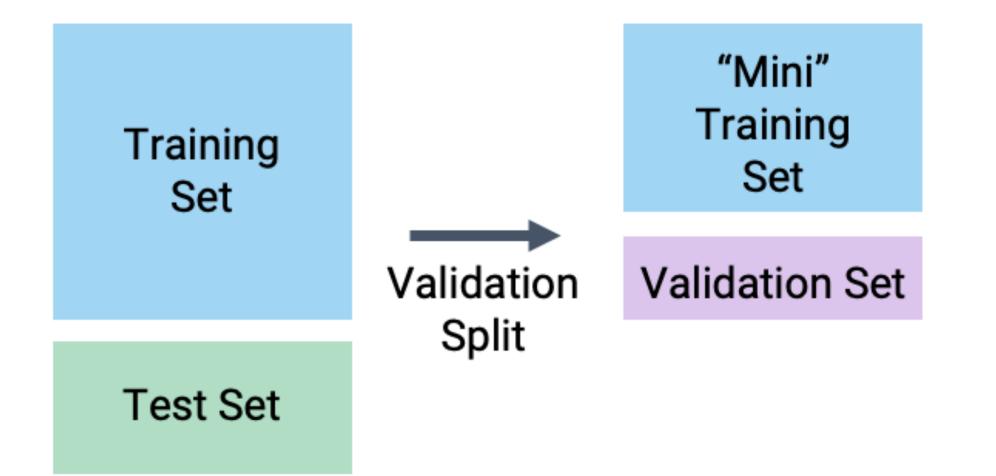
- 1. Split data into train and test sets (50/50%)
- 2. Randomly split train set again into train and validation sets
- 3. Fit model on train and predict responses using the validation set
- 4. Compute the validation set error (e.g., MSE or classification error) provides an estimate of the test error rate
- 5. Repeat #2-4 for many random splits of the training set and/or for many different models of varying flexibility

Variation 1 sales = 
$$\beta_0 + \beta_1 \times TV$$

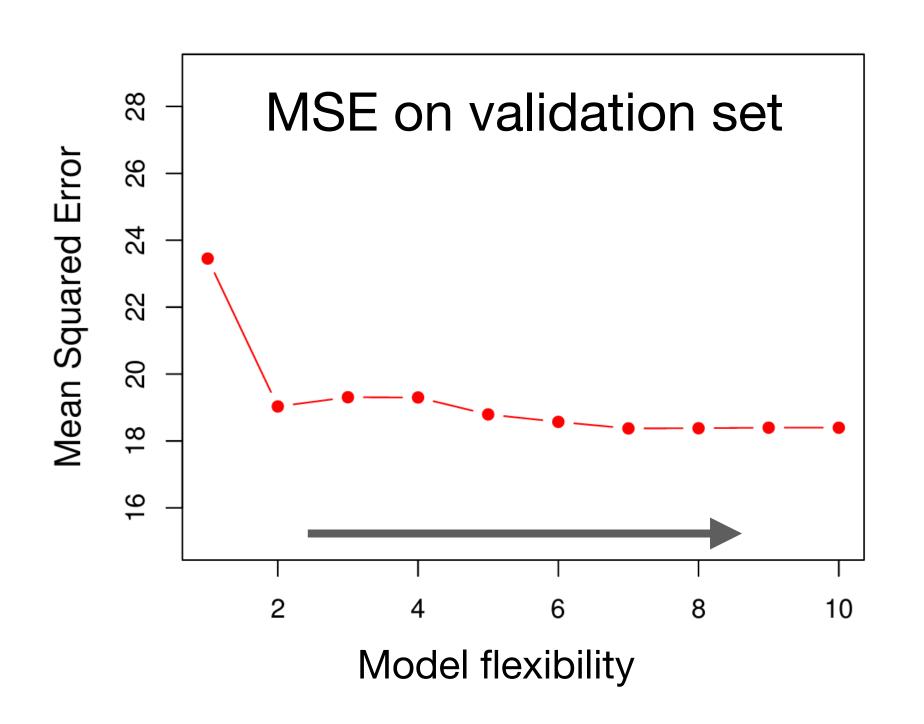
Example: Variation 2 sales = 
$$\beta_0 + \beta_1 \times TV + \beta_2 \times radio$$

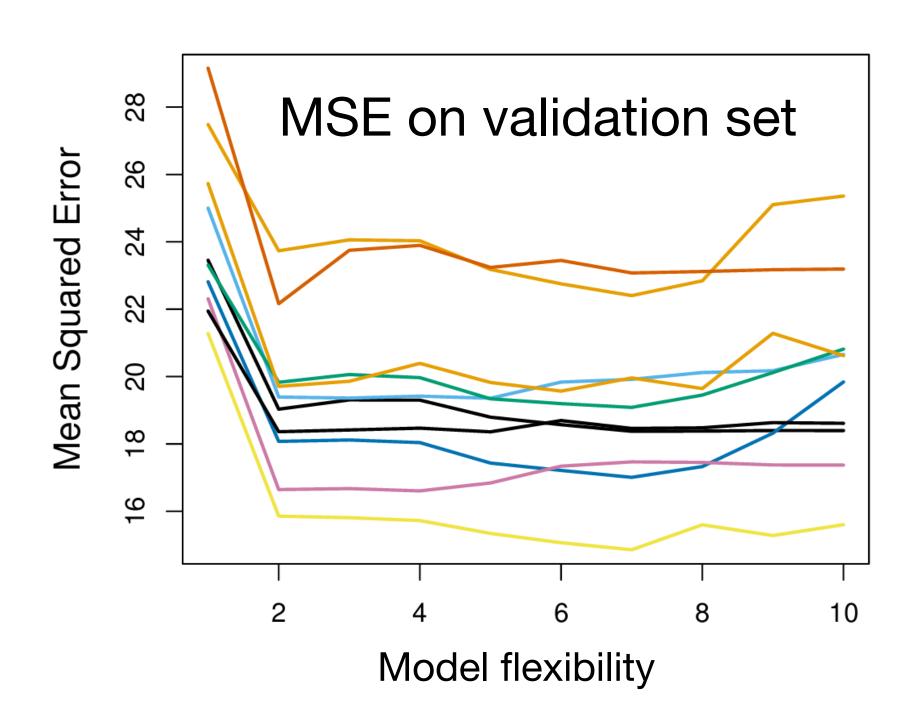
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Variation 3 sales = 
$$\beta_0 + \beta_1 \times TV + \beta_2 \times radio + \beta_3 \times newspaper$$



# Validation set approach





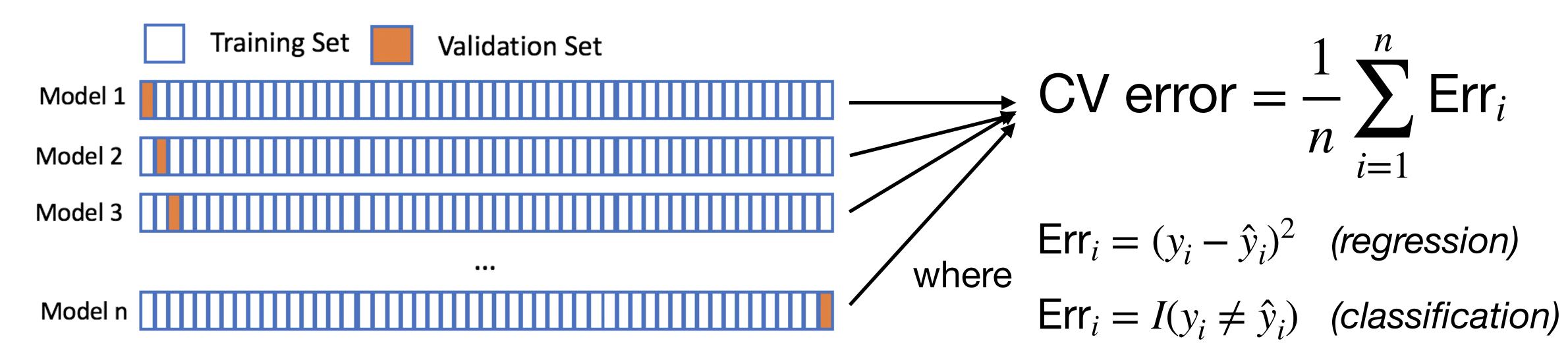
Two drawbacks of the basic validation set approach:

- 1. Validation estimate can be highly variable depending on how data is split
- 2. Only a subset of observations (train set) are used to fit model (even fewer *n* than before)
  - ⇒ Validation set error tends to overestimate test error rate

# Leave-one-out-cross-validation (LOOCV)

### Using the LOOCV approach ("look-v"):

- 1. Split train set into train (n-1 data points) and validation (1 data point) set
- 2. Fit model on <u>n-1 data points</u> and predict the response of the single, held-out <u>validation</u> data point
- 3. Compute the validation error (e.g., MSE or classification error) of the single prediction
- 4. Repeat #2-4, each time holding one data point out as the validation set (leave-one-out)



# Leave-one-out-cross-validation (LOOCV)

### LOOCV advantages:

- Always yields the same CV error because no random splits of data (in contrast to validation set approach)
- Less bias and therefore does not overestimate test error (because we are using almost all of the train data to fit the model, instead of a fraction)

#### LOOCV drawbacks:

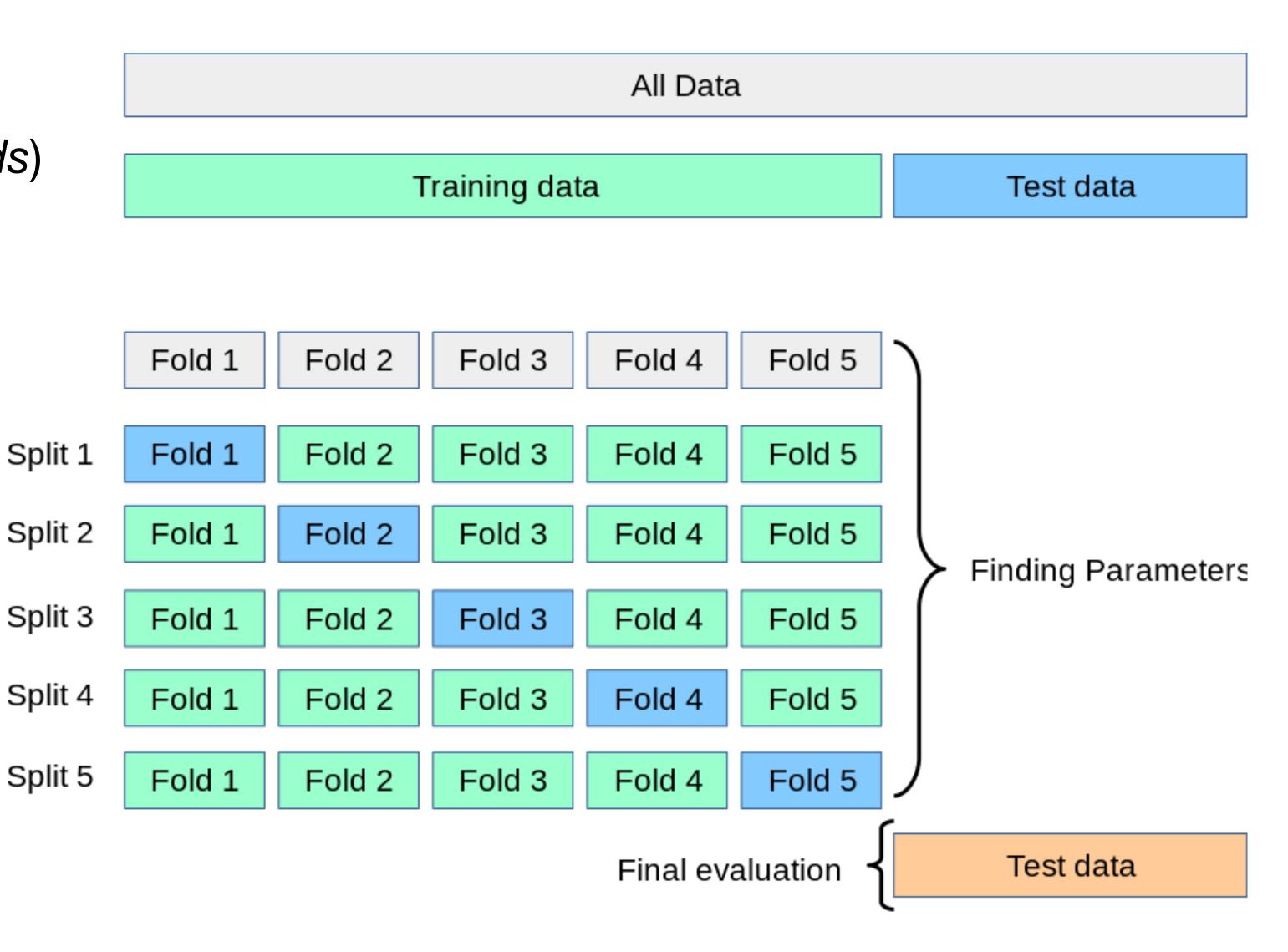
- In most cases, it is computationally expensive (must fit the model *n* times)
  - $\Rightarrow$  Workaround: *k-fold* cross validation (generalization of LOOCV)

### k-fold cross-validation

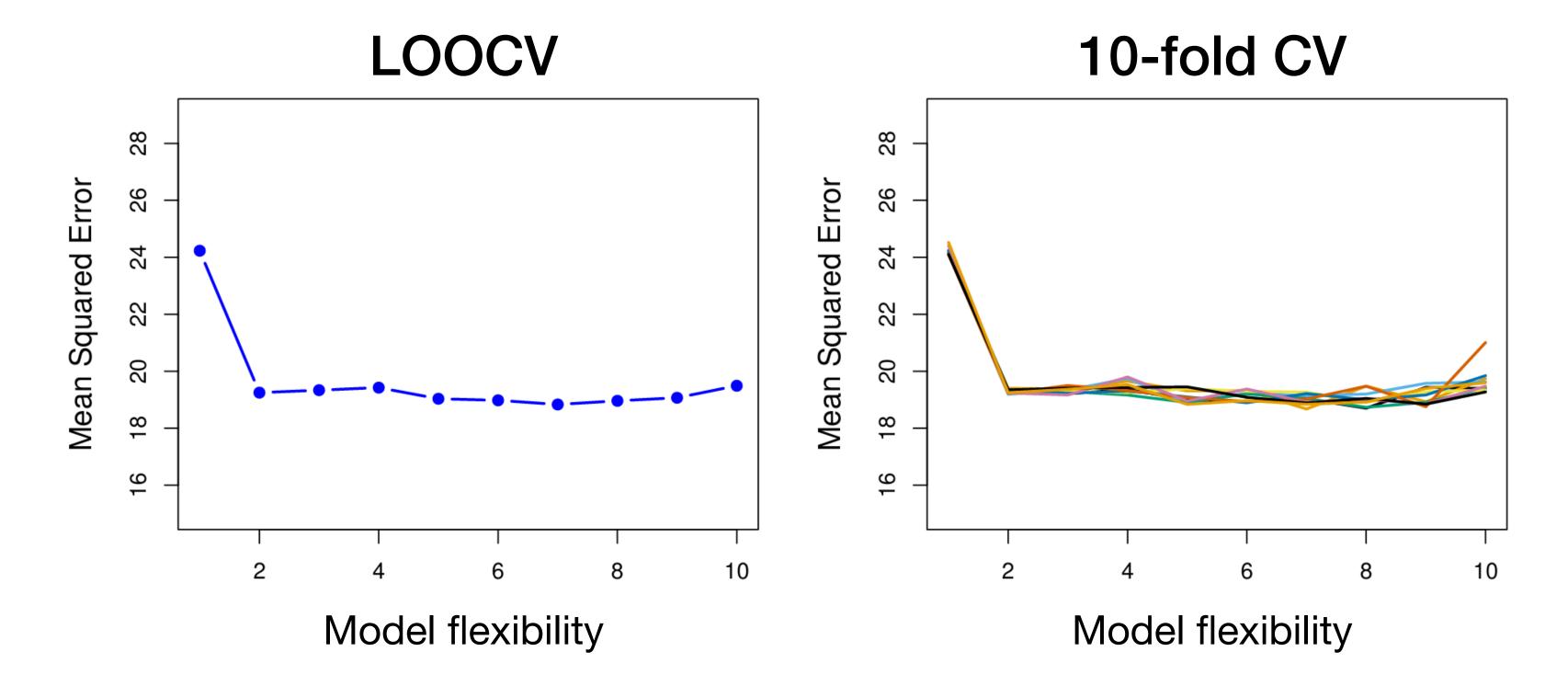
### Using the *k*-fold CV approach:

- 1. Split <u>train set</u> into *k* groups (*folds*) of equal size (*n/k*)
- One fold is treated as the validation set
- 3. Fit model on <u>train</u> and predict responses on <u>validation set</u>
- 4. Compute the validation error
- 5. Repeat #2-4, using the next fold as the <u>validation set</u>

$$CV error = \frac{1}{k} \sum_{i=1}^{k} Err_i$$



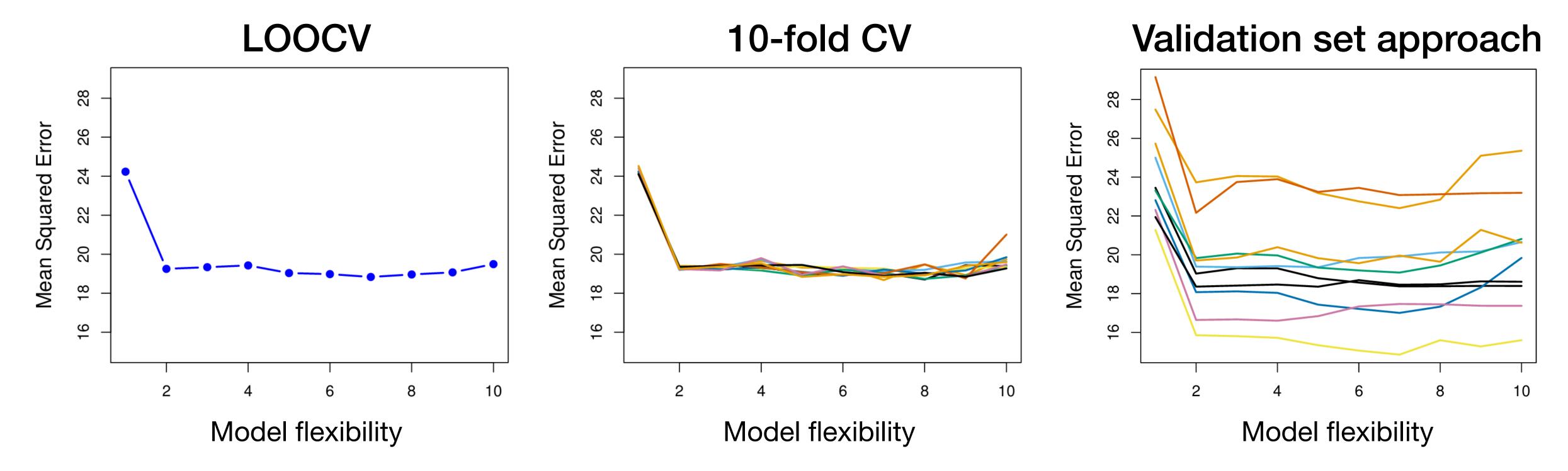
### LOOCV vs. 10-fold CV



Both k-fold CV and LOOCV can be used to assess *any* model, no matter how complex! k-fold advantages:

- Computationally cheaper than LOOCV (only need to fit the model k times)
- Gives more accurate estimates of test error rate...Why? ⇒ bias-variance trade-off!

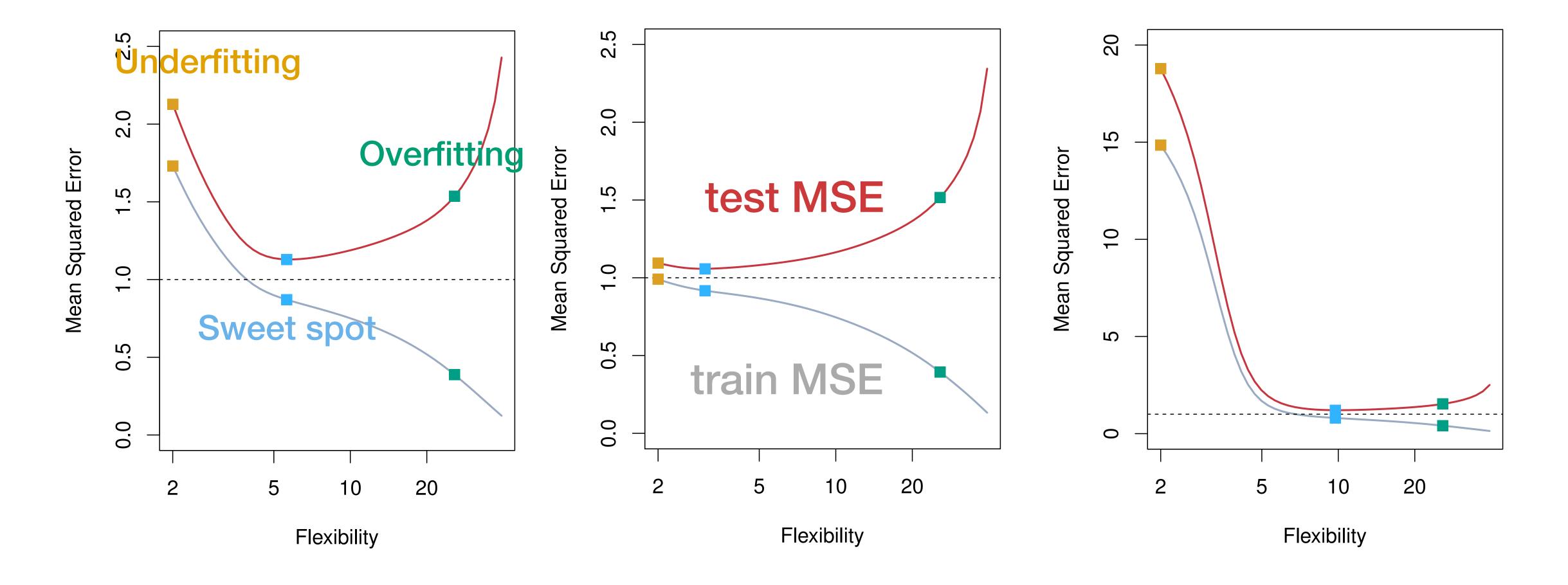
## Comparing cross-validation methods



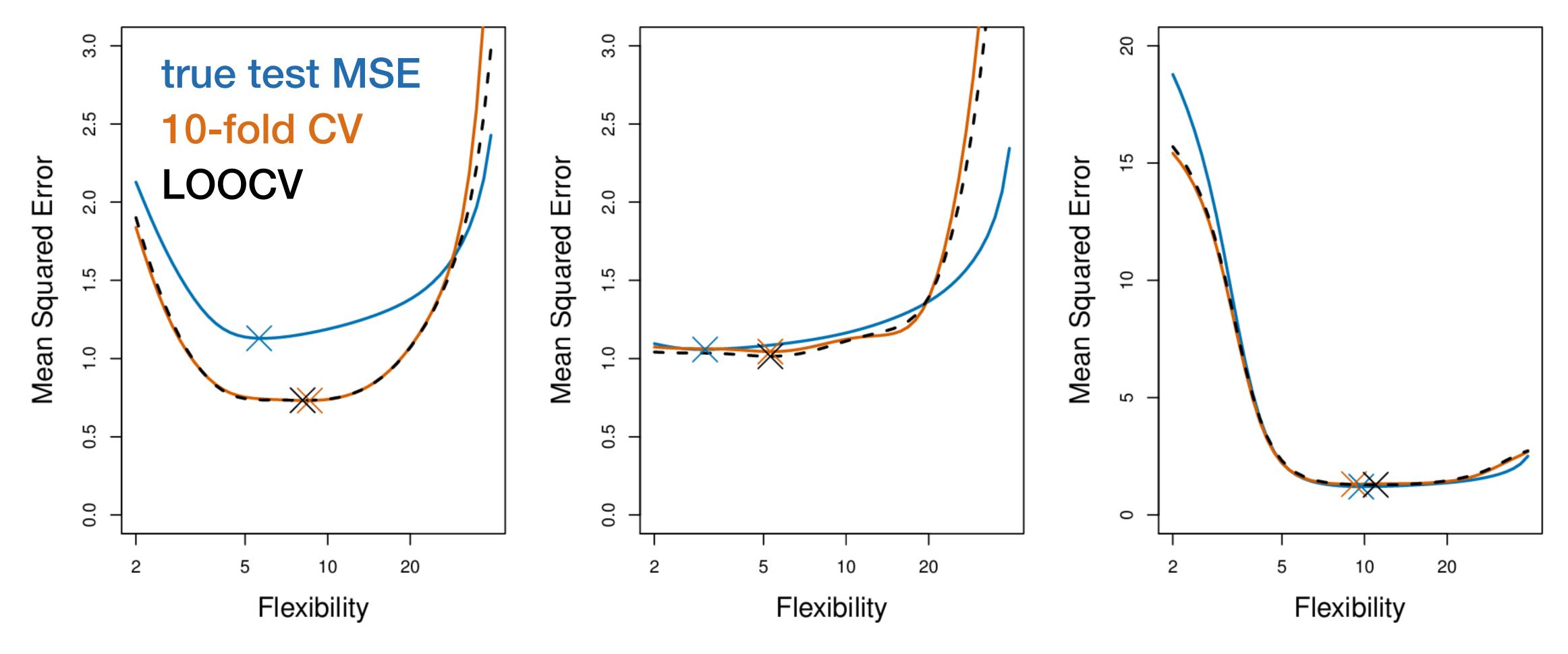
#### Bias and variance in test error estimates:

- In terms of *bias* (accuracy) in test error estimate: LOOCV < *k*-fold CV < validation set
- In terms of <u>variance</u> (noise) in test error estimate: k-fold CV < LOOCV < validation set</li>
- We typically use *k*=5 or *k*=10-fold CV because it hits the sweet spot (not too biased, nor too much variance)

### Comparing cross-validation methods



## Comparing cross-validation methods



- Sometimes we care more about the accuracy of the test error estimate (e.g., MSE)
- Sometimes we only care about finding the optimal model flexibility (X)

# Predictive modeling workflow (so far)

### Using logistic regression as an example:

Fit a multiple logistic regression model to predict a patient's status (1 = alive, 0 = deceased) from the following predictors:

- sex: Factor with levels "Female" and "Male"
- diagnosis: Factor with levels "Meningioma", "LG glioma", "HG glioma", and "Other".
- loc: Location factor with levels "Infratentorial" and "Supratentorial".
- ki: Karnofsky index (0-100, assess a patient's functional ability and prognosis
- gtv: Gross tumor volume, in cubic centimeters.

Let 
$$p = P(\text{status} = 1 \mid \text{sex}, \text{diagnosis}, \text{loc}, \text{ki}, \text{gtv})$$

$$\log\left(\frac{p}{1-p}\right) = \beta_0 + \beta_1 \text{ki} + \beta_2 \text{gtv} + \beta_3 \text{Male[Yes]} + \beta_4 \text{LGglioma[Yes]} + \dots$$
$$\beta_5 \text{Meningioma[Yes]} + \beta_6 \text{OtherDiag[Yes]} + \beta_7 \text{Supratentorial[Yes]}$$

# Predictive modeling workflow (so far)

#### Using multiple logistic regression as an example:

Fit a multiple logistic regression model to predict a patient's status (1 = alive, 0 = deceased) from the following predictors: sex, diagnosis, loc, ki, gtv.

- 1. Split data set into test and train set
- 2. Split train set further into a train and validation set using each of the following approaches:
  - 1. Validation set approach (random 50/50 split)
  - 2. LOOCV
  - 3. 10-fold CV
- Fit the multiple logistic regression model on the training set and predict held-out outcomes on validation set
- 4. Compute the validation error using sklearn.model\_selection.cross\_validate
- 5. **Model selection:** Repeat steps 3-4 for several variations of the logistic regression model (use different subsets of the predictors)
- 6. Plot average classification error vs. number of predictors for each approach
- 7. Report the best model using your chosen cross-validation method

# Upcoming + Reminders

### Assignments:

- Quiz 3 (DUE: TODAY @ 11:59pm)
- Group Project Checkpoint 1 (DUE: TODAY @ 11:59pm)
  - · We will provide written feedback on each video update via Canvas by Friday

Wednesday's topic: Bootstrapping

• Read: ISLP Ch. 5.2