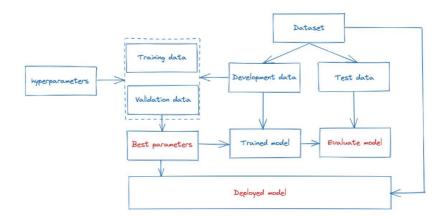
Midterm Preparation



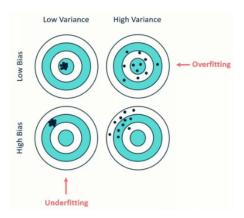
Supervised Learning Framework

▼ Development-Test Split

- Dataset is split into development and test.
- Splitting Strategies:
 - 1. Random Splitting randomly
 - 2. **Stratified Splitting** for highly imbalanced datasets, highly skewed.
 - 3. Structured Splitting prevent data leakage

▼ Hyperparameter Tuning

- Model Complexity vs Model Performance
- Bias-Variance Tradeoff (high error \rightarrow low error; high error \rightarrow good error \rightarrow high error)
 - High Bias: underfitting, models too simple, performs poorly on both training and testing data
 - $\circ \ \ \textbf{High Variance:} \ \text{overfitting, models too strict, performs well on training data but not on test data}$



• Hyper Parameter vs Model Parameters

- $\circ~$ Model Parameters learn from the data (typically with optimization function)
- Hyperparameters you choose!
 - Decision Tree: depth, criterion, min samples, split.
 - Support Vector Machines: kernel width, penalty.
 - Neural Networks: network topology, learning rate.

• Hyperparameter Search

1. **Grid search** – search all in grid layout. – uninformed search strategies.

2. Random search - search random parameters

- a. performs better than gridsearch because there are unimportant parameters that gridsearch might consider and with random you have a chance to explore optimal hyper parameters.
- b. you can combine both → grid search after random search.
- 3. Bayesian optimization informed search strategies.
 - a. sequential, determined at how you perform at each points
- 4. Evolutionary optimization not mentioned

How do we evaluate the hyper-parameters? We could use the test set but we use <u>validation</u> because we tune hyper-parameters on the test set and we only want to use it for the final evaluation. Split development dataset.

Why do you have a validation set? to test the performance of hyper-parameters, as we want to use the test set for the final evaluation.

Model Selection Strategies for Hyperparameter Search

o Three-way holdout

- Development → Train and Validation → Model Selection is the best performance on validation data
- Can get away with this for large data sets.

Leave-one-out CV

Set k = to the number of samples, when you have less data (n data points = n times)

o K-fold CV

- Split development data into k-folds (validation typically 3/5/10)
- Train on k-1 folds, evaluate on last fold (repeat for each k fold) → estimate average model performance → choose best hyperparameter.
- Repeated means you do this multiple times (don't recommend).
- Stratified K-fold CV each fold and test data is stratified.

Why do K-folds? How do you choose one of these?

▼ Data Preprocessing

Numerical Data

- \circ Typically they have different scales \rightarrow
 - StandardScaler \rightarrow (X μ) / σ \rightarrow centered.
 - \underline{MinMax} \rightarrow map min and max to 0, 1.
 - RobustScaler → when data has outlier to not get influenced (standardscaler but with 95th percentile)
 - $\underline{MaxAbsoluteScaler}$ \rightarrow preserve the sign of the data by dividing by the max absolute value.
 - Normalizer → Normalize by the columns, if you have a row, you divide by the norm of the vector → row scaling
- $\circ~$ fit_transform \rightarrow figures out the $\mu\,\sigma$ with development data
- $\circ~$ transform \rightarrow applies it to test data.
- pipeline → give sequence of steps to apply same transformation on test data.
- $\circ~$ scaling with cross validation \rightarrow scale \rightarrow validate.

When does it help? Can set this as a hyperx-parameter with different techniques.

Missing Data

- \circ $\;$ Can be informative \rightarrow make it an indicator.
- Not informative
 - Few data → drop rows
 - Lots of data → drop col
 - Impute using mean or median
 - Label with KNN (nan_euclidean_distance metric), regression models, matrix factorization.

Categorical Data

- \circ **Ordinal Encoding** there is an order
- o One-Hot Encoding no order

- introduces multi-collinearity → drop them (interpretation is very important so might not be good) → instead apply regularization
- some techniques can still take categorical data (tree-based, naive bayes)
- leads to high-dimensional datasets
- Target Encoding map to one column instead of many → map to relationship with average target for that category.
- Combine Numerical and Categorical (make_column_transformer)
 - Why might an NaN appear? Categories that were not fitted before (e.g., OneHotEncoder) are introduced in test/new data.

 Use 'raise error' and 'handle_unknown' to deal with it.

▼ Models - Post Processing

- ▼ Linear Models for Regression
 - <u>Linear Regression:</u> assumptions are *linearity*, *independence*, *homoscedasticity* (noise are randomly distributed around the mean), *normality* (of residuals).
 - transform and observe to see if assumptions are met
 - solve an unconstrained optimization problem → typically there is a closed-form solution.
 - What is the closed form solution? Given X = n (samples)*m (features), w = X transpose X over X transpose y.
 - Typically use gradient descent for unconstrained optimization where there is no closed-form solution.
 - Why do you still need to do an iterative solution even with a closed-form sometimes?
 - If X transpose is large, it can take a long time to invert the matrix. Instead you can do gradient descent. Size of X variables.
 - Inverse may not exist for X (e.g., determinant is 0)
 - Why does it matter that the model is convex? There exists a unique solution and you can use an iterative
 algorithm to find a solution. Performance of the model is stable.
 - find model parameters using loss function.
 - What is the loss functioned used for linear regression? least squares minimization. How does the loss function change with different regularization?
 - What are the problems?
 - Too many predictors (X is not a full rank matrix when $m > n \rightarrow XTX = m*m$ because the rank of the matrix is n)
 - Multi-collinearity one feature is a linear combination of everything else. Check for high condition number for XTX.

• Ridge Regression (L2 norm)

- When m (features) is large compared to n (samples), you must drop features (feature selection and preprocessing.)
- Closed form solution exists!
 - w = (XTX + alpha I) over XTy
 - LSM constrained optimization
 - L2 norm: constrain sum(w^2) ≤ c always lies within a sphere
 - problem is still convex → use lagrangian multipler (unconstrained optimization).
 - \underline{alpha} (hyperparameter) is inversely proportional to \underline{c} .
 - I (identity matrix) is m * m.
- What is the issue? biased (original assumptions of linear regression may not hold).

• Lasso Regression (L1 Norm)

- Applying an sum(abs(w)) < c constraint
- No closed-form solution, must use an iteration to find optimum (Least Angle Regression to solve)
- o Some of the weights are 0 (non-important) Lasso does feature selection.
 - <u>alpha</u> decreases → features increases (becomes active) → coefficient increases
- <u>Elastic-net Regression:</u> Combine Lasso and Ridge with Convex Combination (<u>alpha</u> and <u>L1 ratio</u>)

What is an L2 vs L1 norm, when do you use one over the other? L2 has closed-form solution. L1 does feature selection.

- Contour hits a corner for Ridge (L2), for Lasso (L1) it hits a corner.
- ▼ Linear Models for Classification

▼ Binary Classification

▼ Logistic Regression

- Assume there is a linear relationship y_prediction = sign(wTx + b)
 - find w and b by minimizing a loss function.
 - O-1 loss function
 → not easy to minimize (not differentiable, to make this problem easier define the following...)
 - Hinge Loss max(0, 1-y*f)
 - Log Loss (Logistic Regression get a probability p(y|x) for a label)
 - What is the functional form of a logistic regression? S shape.
 - Maximize the **Log Likelihood** (LL) → convex so you can solve!
 - → variance of Regularized Logistic Regressions
 - · Lasso, Ridge, Elastic-net

▼ SVM

- Maximize the margin classifier (Convex Optimization problem)
 - o s.t.) 2 parallel linear boundaries
 - o s.t.) No points fall between
- Constrained minimization problem: mimize w^2/2 s.t. y(wtx + b) ≥ 1
 - no closed-form solution → can use SMO or other methods.

• Hard-margin vs Soft-margin

- Tradeoff between how much you can push (margin) and allowing no points (error).
- Soft-margin includes a forcing function in the objective function → ζ → Minimize Hinge Loss → introduces upper bound C on alpha in the dual.
- Primal and Dual: the primal is a minimization → dual is maximization.
 - $\circ~$ Primal is solved in \boldsymbol{w} and \boldsymbol{b} (m dimension vector)
 - $\circ~$ Dual is solved on the α (n dimension vector) both has same solution.

How do you choose between primal and dual? make a concious choice based on the size of your dataset (features vs samples).

What is the support vector? When you are making a transition from primal to dual, $\alpha(1 - y(wtx + b)) = 0$, points when α is non-zero and closest to the hyperplane is the support vectors, think holding up the hyperplanes.

- Easily becomes non-linear \rightarrow Kernel Trick.
 - \circ Assume function $\phi(x)$ projects data to high dimensional space
 - Kernel function K(x x) estimates inner product between two points in the projected space (gives you a number)
 - Do SVM but instead $x = K(x \cdot x) = \varphi(x)T\varphi(x)$
 - If K is symmetric and positive definite, you can assume φ exists without finding it (the *trick!*).
 - (RBF) Gaussian kernel shows that it can be eventually projected to infinite dimensional space → at some values of gamma, training error = 0.

▼ Multi-class Classification

• OVO & OVR

- OVR → set of binary classification problems → rank order → choose best.
 - Non-balanced datasets, rarely a region of uncertainty
- OVO → building one against the other → train nC2 classifiers → use classifiers and majority voting to say it belongs to a certain spot.
 - Balanced datasets, region of uncertainty

How do you predict using OVR vs OVO?

Multinomial Logistic Regression

 Another way to do multi-class, find the probability that p(y = class | x) and then find the maximum likelihood function

▼ Decision Trees

- · Greedy algorithm to train
 - choose impure node → estimate reduction in <u>impurity</u> due to every feature → choose feature with maximum reduction → create two child nodes → repeat until stop (can become 100% pure with enough depth).
 - o Splitting for continuous value vs categorical features:
 - Going through all values → measure IG → for numerical check all values; for categorical order by MRR (mean response rate) target encoding, and decide best split.
 - Hyper-parameter to prune or early stop to avoid overfitting.
 - Pruning: reduce error and cost complexity.
 - Early Stop: max depth, nodes, sample_split, IG
- · Non-linear, minimal preprocessing, invariant to scale
- Exhaustive → will always end up at a leaf node

▼ Classification

- Majority Voting by probability.
- Measure Impurity via <u>Gini Index</u> and <u>Entropy</u>.
 - o Probability of split
 - Information Gain takes a weighted average to maximize the reduction in impurity → split.

▼ Regression

- Sample Mean
- Measure Impurity via MSE and MAE
- SSE to maximize reduction in impurity → split.

What is the difference between classification and regression in terms of prediction, measure impurity, and how to decide split?

• Feature Importance: which feature has most impact on IG, its depth and number of samples impacted.

How is the feature importance measured? It is the sum of a probability of a point reaching that node multiplied by its IG at that node.

Can a feature be used more than once to split? Yes, it cycles through all features to determine the optimal split at each node.

▼ Ensemble Methods

- Trees are highly unstable, limited capacity of regression values → build ensembles! → Instead of building 1 tree, build multiple → reduce model variance.
- You have bagging and boosting, difference is the training dataset and how you get the output.

▼ Bagging (Bootstrap Aggregation)

- Sampling w replacement to create training set of same size → build tree with each → average to make final prediction.
- RF
 - Hyper-parameters: # of trees, # of features (sqrt(m) for classification, m for regression)
 - Out of Bag (OOB) error for model selection, average error of a data point, calculated using prediction from trees that do not contain it in their respective bootstrap sample. comes for free!
 - $\circ\hspace{0.1in}$ You already have an amount of trees, and you can add more.

▼ Boosting

- ML algorithms to improve weak learners to stronger ones sequentially.
- Early learners fitting simple models to the data then analyzing data for errors.
- Each tree corrects previous tree.
- Tree finds prediction y → calculates error (y y_prediction) → inputs residual into next tree → repeat until error is
 close to 0 (check with validation set).
- Adaboost (depth 1 (decision <u>stump</u>) → assume average <u>weight</u> → calculate error by summing up the weights → calculate alpha (high means low error) → set weight based on alpha.
- Gradient Boosting Algorithm: ALWAYS building regression trees → Initialize FO Classifier (model predicts the mean of your target) and calculate residual → for iterations, compute pseudo residual →

- Gradient is the residual
- Hyper-parameters Number of trees, learning rate, depth, regularization decision tree parameters, row sampling, column sampling.

What is the gradient in gradient boosted trees? The gradient, derivative of error with respect to F(m), becomes the residual

- GradientBoostingClassifier slow on large data
- HistGradientBoosting faster on large datasets, iterates through all values of histogram.
- <u>XGBoost</u> Fast approximate split finding based on histograms, supports GPU training, sparse data & missing values, adds L1 and L2 penalty on leaf weights, monotonicity
- LightGBM Microsoft version, faster than XGBoost on CPUs, Distributed Training, CLI.
- CatBoost Optimized for categorical features

▼ Model Evaluation

- Evaluation methods for Classification
 - Threshold-based Metrics ← use score to assign label (lower threshold → recall; higher threshold → precision)
 - Classification Accuracy how accurate is your classification misleading for imbalanced datasets Accuracy
 Paradox
 - <u>Precision, Recall & F1-score</u> good for imbalanced datasets <u>positive</u> class is the <u>minority class</u>.
 - Macro (direct average) vs Weighted (average by size) for each class.
 - Weighted Recall = Accuracy

What is the confusion matrix? matrix of actual and predicted positives and negatives.

- Ranking-based Metrics ← use score to rank
 - Average Precision (AP)
 - PR Curve to compare models
 - **AUROC** determines relationship between FPR and TPR (Recall).
 - can say whether the model is good or bad.
- Evaluation metric for Regresstion
 - o R^2, MSE, MAE, MAPE (Percentage Error)
- ▼ Multi-Fidelity Optimization
 - Successive Halving reduce hyper-parameters by halves as you approach the budget
 - **Hyperband** takes care of hyper-parameters by training at different budgets.