Linear & Polynomial Regression

- **LinReg**: model linear data: $\hat{y} = \vec{\theta}^T \vec{x}$ where x_0 =1, linear in params $\vec{\theta}$, OLS normal equation: $\hat{\vec{\theta}} = ({m X}^T {m X})^{-1} {m X}^T \vec{y}$ (slower than GD when m large)
- (+) models linear data well, white-box, feat importance (-) strict assumptions, outliers can skew results
- Assumptions: 1. Linearity feats indep linear wrt targ (pair plot, ypred vs resids) 2. No Multicollinearity hurts coeff interpretability (correlations, VIF)
 - o 3. Normal Residuals (histogram, QQ Plot, stat test) 4. Homoskedastic residual variance equal wrt to ypred (ypred vs residuals, stat test)
 - o 5. No Auto-Correlation resids indep of each other, stationarize data by differencing (ACF plot, Durbin-Watson Test)
- Coeff: mean Δ in depen var for each unit of Δ in indep var, can interpret as feat importance, p-val = prob coeff should be 0
- Cost Func: MSE penalizes outliers more \rightarrow fits them better than MAE, $\text{MSE} = \frac{1}{m} \sum_{i=1}^m (\hat{y}^{(i)} y^{(i)})^2$, $\nabla \text{MSE} = \frac{2}{m} \boldsymbol{X}^T (\boldsymbol{X} \vec{\theta} \vec{y})$
- $m{R^2}$: % of variance explained, compared to model that predicts mean, $m{R^2}=1-rac{ ext{SSE}}{ ext{SST}}$, $m{R_{adj}^2}$ penalizes unnecessary feats to avoid overfitting
- Regularization: limit coeffs to limit overfitting, feats must be scaled, apply by adding term to cost func, α hyperparam controls strength
 - L1 Lasso: diamond loss contours \rightarrow feat selection $\alpha \sum_{i=1}^{n} |\theta_i|$, L2 Ridge, circle loss contours: $\alpha \frac{1}{2} \sum_{i=1}^{n} \theta_i^2$, Elastic Net: weighted mix
- · Polynomial Regression: model nonlinear data, params still linear, add interaction & higher power terms
- Bayesian LinReg: use prior to get posterior distrib for model params: $P(\boldsymbol{\theta}|\boldsymbol{y},\boldsymbol{X}) = \frac{P(\boldsymbol{y}|\boldsymbol{\theta},\boldsymbol{X})\cdot P(\boldsymbol{\theta}|\boldsymbol{X})}{P(\boldsymbol{y}|\boldsymbol{X})}$, preds made from distrib: $\hat{\boldsymbol{y}} \sim \mathcal{N}(\boldsymbol{\theta}^T\boldsymbol{x},\,\sigma^2\boldsymbol{I})$

Logistic & Softmax Regression

- **LogReg**: linear/hyperplane decision boundary, $\hat{p} = \sigma(\vec{\theta}^T \ \vec{x}) = \frac{1}{1 + e^{-\vec{\theta}^T \vec{x}}}$ sigmoid func takes linreg logit ($-\infty$, ∞) and outputs prob [0,1]
- (+) models linearly separable data well, white-box, calcs class proba, feat importance (1 unit Δ in $x_i \to$ depend var Δ by e^{θ_i}) (-) strict assumptions
- Assumptions: 1. No Multicollinearity hurts coeff interpretability (correlations, VIF) 2. Linearity: feats indep linear wrt logit of preds i.e. linreg output
 - \circ 3. No Influential Outliers (Cook's Dist: change in model when outlier removed) 4. Requires larger m when more feats
- Odds $[0,\infty) = \frac{\pi}{1-\pi}$ Log-Odds $(-\infty,\infty) = \log\left(\frac{\pi}{1-\pi}\right) = \operatorname{logit}(\pi) = \eta$ (logit) Sigmoid $[0,1] = \sigma(\eta) = \operatorname{logit}^{-1}(\eta) = \frac{1}{1+e^{-\eta}} = \pi$ (probability)
- Cost Func: Log Loss $J(\vec{\theta}) = -\frac{1}{m} \sum_{i=1}^m \left(y^{(i)} \log(\hat{p}^{(i)}) + (1-y^{(i)}) \log(1-\hat{p}^{(i)}) \right)$ Pred: $\hat{y}^{(i)}$ =1 if $\hat{p}^{(i)} \ge$ 0.5, else $\hat{y}^{(i)}$ =0
- Regularization: L1, L2, & Elastic Net Polynomial LogReg: nonlinear boundary SoftmaxReg: multiclassifier logreg, piecewise linear boundaries
- 1. Calc logit for each class k: $s_k(\vec{x}) = \vec{x}^T \vec{\theta}^{(k)}$, 2. Softmax calcs prob of each logit: $\hat{p}_k = \sigma \big(\vec{s}(\vec{x}) \big)_k = \frac{e^{s_k(\vec{x})}}{\sum_{i=1}^K e^{s_j(\vec{x})}}$, 3. Pred class /w highest prob
- Softmax Cost Func: Cross Entropy, how well class probs match target probs, $J(\mathbf{\Theta}) = -\frac{1}{m} \sum_{i=1}^m \sum_{k=1}^K y_k^{(i)} \log(\hat{p}_k^{(i)})$

Support Vector Machines (SVM)

- SVM: lin/nonlin regressor/binary classifier/outlier detection, finds n dim decision boundary hyperplane maxes margin between 2 classes
- (+) high-dim data, inference memory efficient (only needs SVs) (-) don't scale with data, slow to train, no class probas
- Decision Func: n+1 dim, $h(\vec{x}) = \vec{w}^T \vec{x} + b$, model params are $\vec{w} \& b$ Decision Boundary: n dim, where h=0 Margins: n dim, where h=-1, 1
- Support Vectors: closest samples from each class, determine margin boundaries Classify: plug pts into h, \hat{y} =1 if $h \ge 0$ else \hat{y} =-1
- Soft-Margin: wide margin + limit margin violations C Hyperparam: controls tradeoff, ↓C = wider margin = more violations = more generalizeable
- Cost Func: Hinge Loss = $\max(0, 1 \hat{y}(\vec{w}^T\vec{x} + b))$, is 0 if $\hat{y}(\vec{w}^T\vec{x} + b) \ge 1$ (correct class + outside margin), else linear penalty $1 \hat{y} \cdot h(\vec{x})$
- Kernel Trick: map nonlinearly separable data into higher dim to find hyperplane boundary, kernel func computes dot prod of instances mapped in
- higher dim without transforming them to higher dim, allows to use SVM on transformed feats without transforming them (e.g. polynomial, RBF)

Decision Tree

- DecTree: nonparametric, lin/nonlin regressor/multiclassifier, stochastic if max_feats set, use stop criteria to stop splitting
- (+) white-box, handle outliers/noise/missing, no scaling, inference is log(m), feat importance (-) 90° boundaries, high variance, overfit
- CART: each split is greedy search to find feat k & threshold t_k that results in lowest cost: $J(k,t_k) = \frac{m_{\mathrm{left}}}{m_{\mathrm{node}}} \mathrm{Impurity_{left}} + \frac{m_{\mathrm{right}}}{m_{\mathrm{node}}} \mathrm{Impurity_{right}}$
- $\bullet \ \ \textbf{Impurity} : \textbf{0} \ \text{if node's instances all 1 class, >0} \ \text{if multiple classes} \ \ \textbf{Information Gain} : \\ \textbf{Entropy}_{parent} \big(\frac{m_{left}}{m_{parent}} \\ \textbf{Entropy}_{left} + \frac{m_{right}}{m_{parent}} \\ \textbf{Entropy}_{right} \big)$
- Gini: $G_i=1-\sum_{k=1}^K p_{i,k}^2$ Entropy: $E_i=-\sum_{k=1}^K p_{i,k}\log_2(p_{i,k})$ $p_{i,k}$ = ratio of class k instances among all train instances in ith node
- Classify: leaf node classifies whatever the majority represented label is Class Proba = ratio of train instances of class k in instance's leaf node
- Regression: each split mins MSE, leaf node preds avg targ value of its train instances
- Feat Importance: how much tree nodes that use that feat reduce impurity on avg, each node weighted by num of train samples associated with it

• Regularization: 1. pre-pruning (max depth, max feats, max leafs, min samples per leaf), 2. post-pruning (delete unnecessary nodes)

Random Forest & Gradient Boosting

- RandFor: nonparametric, stochastic, use bagging to train trees, each tree trained on random subset of feats & bootstrapped sample of data
- (+) low variance, handle outliers/noise/missing, no scaling, feat importance (-) slow training (but parallelizeable unlike boosting), black box
- Gradient Boosting: train models sequentially, each model fit on resids of prev model, if tree → shallow <10 splits (shallower than randfor trees)
- Hyperparams: n_trees (too few → underfit, too many → overfit, use early stopping), subsample (% of train data to train each tree → stochastic)
- Regression: pred by summing all tree preds in the ensemble

K-Nearest Neighbors (KNN)

- KNN: supervised, nonparametric, nonlinear regressor/multiclassifier, predict by finding k nearest train samples by dist (instance-based learning)
- (+) handle nonlinear data well, white-box, no training (-) more train data or feats → slower inference, sensitive to noise & outliers, curse of dims
- Assumptions: 1. similar data points are close together in feat space 2. feats are scaled since distance-based
- Inference: compare n feats with all m train samples = $\mathcal{O}(mn)$, then sort calced distances and return k nearest = $\mathcal{O}(m\log m)$
- k Tradeoff: small k (high variance + might overfit + jagged boundary), large k (low variance + high bias + might underfit + smooth boundary)
- Classification: majority label of k nearest neighbors Regression: avg of k nearest neighbors Choose k: either use CV or approx as \sqrt{m}

Naive Bayes

- NBayes: supervised, multiclassifier, probabilistic, estimates posterior $P(y_i|X)$ via naive assumption = feats conditionally independent given y
- (+) works well in high dims (e.g. text), requires less data (doesn't need data demonstrating feat interactions) Assumption: little multicollinearity
- Classification: $P(y_i | \mathbf{X}) = \frac{P(\mathbf{X} \mid y_i) \cdot P(y_i)}{P(\mathbf{X})} = \frac{P(\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \mid y_i) \cdot P(y_i)}{P(\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n)} \rightarrow \text{naive assumption} \rightarrow = P(\vec{x}_1 \mid y_i) \cdot \dots \cdot P(\vec{x}_n \mid y_i) \cdot P(y_i)$
- $P(y_i)$: prior proba of y_i class = $\frac{\text{examples labeled } y_i}{\text{total examples}}$, $P(\vec{x}_j|y_i)$: class conditional = PDF for each \vec{x}_j feat with y_i class (k classes, n feats = kn PDFs)
- PDFs: fit PDF to each feat (e.g. binary → binomial, numeric → gaussian, unknown → kernel density estimation), update prior & PDFs when new data
- . MLE: nbayes estimates priors from data instead of making assumptions about their distributions, it is not bayesian, simply uses MLE to find PDFs

KMeans Clustering

- KMeans: unsupervised, clustering, centroid-based, assigns instance to cluster based on distance from centroid Assumptions: feats are scaled
- (+) guaranteed convergence, relatively fast clustering ($\mathcal{O}(nmk)$ for each centroid update), cluster probas based on distance from centroids
- (-) bad if clusts vary in diameter/density or non-convex, sensitive to cluster init → rerun, k hard to choose, sensitive to outliers & curse of dims
- Algo: 0. rand init k samples as centroids, repeat: 1a. update cluster labels to nearest centroid, 1b. update centroids to center of new clusters
- Convergence: when pts reclassified to same centroid, local or global optima depends on centroid init, rerun and keep model with lowest inertia
- Choose k: 1. Elbow: as k↑ inertia↓, use elbow on k vs inertia graph 2. Silhouettes: compare avg silhouette coeff or cluster knives for each k

Principal Component Analysis (PCA)

- PCA: unsupervised, dim reduction, project data onto max variance orthogonal linear combos of old correlated dims = SVD on covariance matrix
- (+) removes correlated feats (-) new dims are linear combos → less interpretable, not scale/rotation/translation invariant, negative loading scores
- Assumptions: linear data (PCs are linear combos), standardized data, high variance is structure & low variance is noise, no significant outliers
- Principal Axes: orthonormal eigvecs PCs: data projected onto principal axes scale to 1 stdev Loadings: eigvec components = weight of old dims
- Reduce: standardize data → find orthogonal dims of highest variance → keep k PCs based on total variance to preserve (scree plot, keep 95%)