DSA5105 Principles of Machine Learning

AY2024/25 Sem1 By Zhao Peiduo

Supervised Learning

Lecture 1

Empirical Risk Minimization (ERM) The learning process aims to find a function $f \in \mathcal{H}$ that minimizes the empirical risk, $R_{\text{emp}}(f) = \frac{1}{N}\sum_{i=1}^{N} L(y_i, f(x_i))$, where $y_i = f^*(x_i)$.

Common Loss Function Mean squared error (MSE) for regression tasks:$L(y,\hat{y}) = \frac{1}{2}(y-\hat{y})^2$, cross-entropy loss for classification task: $L(y,p) = -\sum_i y_i \log(p_i)$, and huber loss for robust regression: $L_\delta(y,\hat{y}) = \begin{cases} \frac{1}{2}(y-\hat{y})^2 & \text{if } |y-\hat{y}| \le \delta \\ \delta|y-\hat{y}| - \frac{1}{2}\delta^2 & \text{otherwise} \end{cases}$

Softmax Function For a multi-class classification problem with K classes, the softmax function is defined as:$\text{softmax}(z_i) = \frac{\exp(z_i)}{\sum_{j=1}^{K};\exp(z_j)}$

General Ordinary Least Squares Formula Consider $x \in \mathbb{R}^d$ and the new hypothesis space $\mathcal{H}_M = \left\{f : f(x) = \sum_{j=0}^{M-1} w_j \varphi_j(x)\right\}$ Each $\varphi_j : \mathbb{R}^d \to \mathbb{R}$ is called a **basis function** or **feature map**.

We can rewrite the ERM $\min_{w\in\mathbb{R}^M} \frac{1}{2N}\sum_{i=1}^{N}\left(\sum_{j=0}^{M-1} w_j\varphi_j(x_i) - y_i\right)^2$ into $\min_{w\in\mathbb{R}^M} \frac{1}{2N}\|\Phi w - y\|^2$. Solving by setting $\nabla R_{\text{emp}}(\hat{w}) = 0$, we have $\hat{w} = (\Phi^\top\Phi)^{-1}\Phi^\top y$, given invertible $\Phi^T\Phi$.

For cases where $\Phi^T\Phi$ is not invertible, the formula using the Moore-Penrose pseudoinverse is: $\hat{w}(u) = \Phi^\dagger y + \left(I - \Phi^\dagger\Phi\right)u \quad u \in \mathbb{R}^M$

Regularization: To prevent overfitting, regularization techniques add a penalty to the loss function: $\min_{w\in\mathbb{R}^M} \frac{1}{2N}\|\Phi w - y\|^2 + \lambda C(w)\frac{1}{2N}\|\Phi w - y\|^2 + \lambda C(w)$. Minimizing the ERM we get $\hat{w} = \left(\Phi^\top\Phi + \lambda I_M\right)^{-1}\Phi^\top y$ which is always invertible for positive λ.

Common regularization terms:**L_2 (Ridge) regularization:** $\lambda\sum_{j=1}^{p} w_j^2$, **$L_1$ (Lasso) regularization:** $\lambda\sum_{j=1}^{p}|w_j|$

Lecture 2

Reformulation of Ridge Regression We rewrite the regularized least squares solution in another way:
$\hat{w} = \left(\Phi^\top\Phi + \lambda I_M\right)^{-1}\Phi^\top y = \Phi^\top\left(\Phi\Phi^\top + \lambda I_N\right)^{-1} y$

Reformulation of Ridge Regression
$\hat{f}(x) = \sum_{i=1}^{N}\alpha_i\varphi(x_i)^\top\varphi(x)$ $\alpha = (G + \lambda I_N)^{-1}y$ where $G_{ij} = \varphi(x_i)^\top\varphi(x_j)$ is the gram matrix

Mercer's Theorem and SPD Kernels Suppose k is a SPD kernel. Then, there exists a feature space $\mathcal{H}$ and a feature map $\varphi : \mathbb{R}^d \to \mathcal{H}$ such that $k(x,x') = \varphi(x)^\top\varphi(x')$

SPD kernels properties: $K(x,x') = K(x',x)$ (**Symmetry**) For any n and $\{x_1,\dots,x_n\}$, the Gram matrix $G_{ij} = k(x_i,x_j)$ is positive semi-definite. (Recall: a matrix G is positive semi-definite if $c^TGc \ge 0$ for any vector c) (**Positive Semi-definiteness**)

Examples of SPD Kernels: **Linear Kernel:** $K(x,x') = x^\top x'$ **Polynomial Kernel:** $K(x,x') = (1 + x^\top x')^d$

Gaussian RBF Kernel: $K(x,x') = \exp\left(-\frac{\|x-x'\|^2}{2\sigma^2}\right)$

Constructing kernels Given valid kernels $k_1(x,x')$ and $k_2(x,x')$, the following new kernels will also be valid: $ck_1(x,x')$ $f(x)k_1(x,x')f(x')$ $q(k_1(x,x'))$ $\exp(k_1(x,x'))$ $k_1(x,x') + k_2(x,x')$ $k_1(x,x')k_2(x,x')$ $k_3(\varphi(x),\varphi(x'))$ $x^\top Ax'$ $k_a(x_a,x'_a) + k_b(x_b,x'_b)$ $k_a(x_a,x'_a)k_b(x_b,x'_b)$

Lecture 3

SVM Max Margin Formulation $\max_{\mathbf{w},b}\frac{1}{\|\mathbf{w}\|}\min_{i=1,\dots,N}|\mathbf{w}^T\mathbf{x}_i + b|$ subject to $y_i(\mathbf{w}^T\mathbf{x}_i + b) > 0 \quad \forall i$

Optimization Problem $\min_{w,b}\frac{1}{2}\|w\|^2$ subject to: $y_i(w^\top x_i + b) \ge 1 \quad \forall i$. Introducing Lagrange multipliers $\alpha_i \ge 0$, the Lagrangian is: $\mathcal{L}(w,b,\alpha) = \frac{1}{2}\|w\|^2 - \sum_{i=1}^{N}\alpha_i[y_i(w^\top x_i + b) - 1]$

KKT conditions for SVM

1. From Stationarity $\hat{w} = \sum_{i=1}^{N}\hat{\mu}_i y_i x_i, \quad 0 = \sum_{i=1}^{N}\hat{\mu}_i y_i$

2. From Dual Feasibility $\hat{\mu}_i \ge 0$ for $i = 1,\dots,N$

3. From Complementary Slackness$\hat{\mu}_i = 0$ or $y_i(\hat{w}^T x_i + \hat{b}) = 1$

4. The multipliers $\hat{\mu}$ can be found by the dual problem $\max_{\mu\in\mathbb{R}^N}\sum_{i=1}^{N}\mu_i - \frac{1}{2}\sum_{i,j}\mu_i\mu_j y_i y_j (x_i^T x_j)$

Dual formulation of SVM
$\hat{\mu} = \arg\max_{\mu\in\mathbb{R}^N}\sum_{i=1}^{N}\mu_i - \frac{1}{2}\sum_{i,j}\mu_i\mu_j y_i y_j (x_i^T x_j)$ Subject to: $\hat{\mu} \ge 0$ and $\sum_{i=1}^{N}\hat{\mu}_i y_i = 0$
Decision function: $\hat{f}(x) = \text{sgn}\left(\sum_{i=1}^{N}\hat{\mu}_i y_i x_i^T x + \hat{b}\right)$ Complementary slackness: $\hat{\mu}_i = 0$ or $1 = y_i(\hat{w}^T x_i + \hat{b})$

Kernel SVMs

$\hat{\mu} = \arg\max_{\mu\in\mathbb{R}^N}\sum_{i=1}^{N}\mu_i - \frac{1}{2}\sum_{i,j=1}^{N}\mu_i\mu_j y_i y_j k(x_i,x_j)$ Subject to: $\hat{\mu} \ge 0$ and $\sum_{i=1}^{N}\hat{\mu}_i y_i = 0$
Decision function: $\hat{f}(x) = \text{sgn}\left(\sum_{i=1}^{N}\hat{\mu}_i y_i k(x_i,x) + \hat{b}\right)$

Only support vectors satisfying $1 = y_i(\hat{w}^T\varphi_i(x) + \hat{b})$ matter for predictions. This is a **sparse kernel method**.

Lecture 4

Classification and Regression Trees Suppose that the input space is $\mathcal{X}$. A **partition** of $\mathcal{X}$ is a collection of subsets $\mathcal{R}_1,\mathcal{R}_2,\dots,\mathcal{R}_J$ such that $\mathcal{R}_i \cap \mathcal{R}_j = \emptyset$ for $i \ne j$ and $\bigcup_{j=1}^{J}\mathcal{R}_j = \mathcal{X}$

The general decision tree hypothesis space is: $\mathcal{H} = \left\{f : f(x) = \sum_{j=1}^{J} a_j\mathbb{I}_{x\in\mathcal{R}_j}, \{\mathcal{R}_j\} \text{ is a partition of } \mathcal{X}, a_j \in \mathcal{Y}\right\}$ where $\mathbb{I}_{x\in\mathcal{R}_j}$ is an indicator variable returning 1 if x is in $\mathcal{R}_j$.

A decision tree model $f(x) = \sum_{j=1}^{J} a_j\mathbb{I}_{x\in\mathcal{R}_j}$ depends on both a_j and $\mathcal{R}_j$. For regression we take the **average** label values $a_j = y_j = \frac{\sum_i y_i\mathbb{I}_{x\in\mathcal{R}_j}}{\sum_i\mathbb{I}_{x\in\mathcal{R}_j}}$; For classification we take the **mode** label values $a_j = \text{mode}\{y_i : x_i \in \mathcal{R}_j\}$

Loss function for Decision Trees Classification Entropy: $-\sum_{k=1}^{K}\sum_{j=1}^{J} p_{jk}\log p_{jk}$

Gini Impurity: $\sum_{k=1}^{K}\sum_{j=1}^{J} p_{jk}(1-p_{jk})$ where p_{jk} is the proportion of samples in $\mathcal{R}_j$ belonging to class k.

Bagging reduces the variance by aggregating predictions from multiple models trained on different random subsamples of the data:**Regression** $\bar{f}(x) = \frac{1}{m}\sum_{j=1}^{m} f_j(x)$ **Classification** $\bar{f}(x) = \text{Mode}\{f_j(x) : j = 1,\dots,m\}$

Boosting works by training weak learners sequentially, where each learner focuses on correcting the mistakes of the previous ones. The combined model is a weighted sum of the weak learners: $f(x) = \sum_{t=1}^{T}\alpha_t f_t(x)$ where α_t are coefficients based on each learner's performance. Boosting helps reduce bias.

Key Ideas of AdaBoost
1. Initialize with uniform weight across all training samples
2. Train a classifier/regressor f_1
3. Identify the samples that f_1 got wrong (classification) or has large errors (regression)
4. Weight these samples more heavily and train f_2 on this reweighted dataset
5. Repeat steps 3-5

AdaBoost Implementation

Data: $\mathcal{D} = \{(x_i,y_i)\}_{i=1}^{N}$, Initialize $w_i^{(1)} = \frac{1}{N}$ for all $i = 1,\dots,N$;
For $j = 1,\dots,m$ do

1. Obtain f_j from: $f_j = \arg\min_{f\in\mathcal{H}}\sum_{i=1}^{N} w_i^{(j)}\mathbb{I}_{y_i\ne f(x_i)}$
2. Compute combination coefficients: $\delta_j = \frac{\sum_{i=1}^{N} w_i^{(j)}\mathbb{I}_{y_i\ne f_j(x_i)}}{\sum_{i=1}^{N} w_i^{(j)}}$ $\alpha_j = \log\left(\frac{1-\delta_j}{\delta_j}\right)$
3. Update weights:$w_i^{(j+1)} = w_i^{(j)}\exp\left(\alpha_j\mathbb{I}_{y_i\ne f_j(x_i)}\right)$

Return: $\bar{f}(x) = \text{Sign}\left(\sum_{j=1}^{m}\alpha_j f_j(x)\right)$

Cross-Validation In k-fold cross-validation, the data is split into k subsets. The model is trained on $k-1$ subsets and validated on the remaining one. This process is repeated k-times.

Lecture 5

Activation Functions
Sigmoid: $\sigma(z) = \frac{1}{1+e^{-z}}$, $[0,1]$ **ReLU** $\sigma(z) = \max(0,z)$, $[0,\text{inf}]$, leaky ReLU: δzif $z < 0$ instead of 0

Gradient Descent $w^{(t+1)} = w^{(t)} - \eta\frac{\partial L}{\partial w}$

Stochastic Gradient Descent (SGD): $\theta_{k+1} = \theta_k - \eta\frac{1}{B}\sum_{i\in I_B}\nabla\Phi_i(\theta_k)$, less likely to be locally optimal.

Deep Neural Networks (DNNs) Deep neural networks are an extension of shallow networks. The idea is to stack n hidden layers together and forward pass the x sequentially.

Back-propagation Algorithm
Initialize $x_0 = x \in \mathbb{R}^d$.
For $t = 0,1,\dots,T$: $x_{t+1} = g_t(x_t,W_t) = \sigma(W_t x_t)$
Set $p_{T+1} = \nabla_x L(x_{T+1},y)$.
For $t = T,T-1,\dots,1$:

- $\nabla_{W_t}\Phi = p_{t+1}^T\nabla_W g_t(x_t,W_t)$
- $p_t = [\nabla_x g_t(x_t,W_t)]^T p_{t+1}$

Return $\{\nabla_{W_t}\Phi : t = 0,\dots,T\}$.

Back-propagation Example Computation
The model is defined as: $y(x) = v\delta(w_1\delta(w_0 x))$, $x,w_0,w_1,v \in \mathbb{R}$, where δ is the identity function, i.e., $\delta(z) = z$. The loss function is: $L = (y(x)-y)^2$ We want to compute the gradients: $\frac{\partial L}{\partial v}$, $\frac{\partial L}{\partial w_1}$, $\frac{\partial L}{\partial w_0}$

Forward pass: $x_1 = w_0 x$ $x_2 = vw_1x_1$ $L = (x_2-y)^2$

Backward pass:

$$p_2 = \frac{\partial L}{\partial x_2} = 2(x_2-y) \quad \frac{\partial L}{\partial v} = \frac{\partial L}{\partial x_2}\cdot\frac{\partial x_2}{\partial v} = 2(x_2-y)w_1x_1 \qquad \frac{\partial L}{\partial w_1} = \frac{\partial L}{\partial x_2}\cdot\frac{\partial x_2}{\partial w_1} = 2(x_2-y)vx_1$$

$$p_1 = \frac{\partial L}{\partial x_1} = \frac{\partial L}{\partial x_2}\cdot\frac{\partial x_2}{\partial x_1} = 2(x_2-y)vw_1 \qquad \frac{\partial L}{\partial w_0} = \frac{\partial L}{\partial x_1}\cdot\frac{\partial x_1}{\partial w_0} = 2(x_2-y)vw_1x$$

Lecture 6

PCA Algorithm Simplified Flow Center the data → compute sample covariance matrix $S = \frac{1}{N}\sum_{i=1}^{N} x_i x_i^T$ → compute top k eigenvalues and retrieve their corresponding eigenvectors. In feature space we replace all x with ϕ, the feature maps, and compute design matrix as $\Phi_{ij} \leftarrow \Phi_{ij} - \frac{1}{N}\sum_{i=1}^{N}\Phi_{ij}$, where $\Phi_{ij} = \phi_j(x_i)$ is the raw design matrix.

PCA Whitening Transform Principal component scores are given by $Z = XU$ where X is the original features and U is the matrix of eigenvectors. The transformation $X' = XU\Lambda^{-\frac{1}{2}}$, where Λ is the matrix of eigenvalues makes $\text{cov}(X') = I$.

Neural Network AutoEncoders Choose encoder T_{enc} and decoder T_{dec} as:
$T_{enc}(x;\theta) = A\sigma(Wx+b)$ $T_{dec}(x;\phi) = B\sigma(Vx+c)$
$\theta = (A,W,b) \in \mathbb{R}^{m\times q}\times\mathbb{R}^{q\times d}\times\mathbb{R}^q$ $\phi = (B,V,c) \in \mathbb{R}^{d\times q'}\times\mathbb{R}^{q'\times m}\times\mathbb{R}^{q'}$
Given a dataset $\mathcal{D} = \{x_i\}_{i=1}^{N}$, we solve the empirical risk minimization to minimize the distance between x_i and x'_i: $\min_{\theta,\phi}\frac{1}{2N}\sum_i\|x_i - T_{\text{dec}}(T_{\text{enc}}(x_i;\theta);\phi)\|^2$, where the input is used as labels.

Unsupervised Learning

Lecture 7

Clustering partition a dataset $\mathcal{D}$ into disjoint groups $\mathcal{D} = \mathcal{D}_1\cup\mathcal{D}_2\cup\cdots\cup\mathcal{D}_K$, such that data grouped together are similar and dissimilar if they are in different groups.

K-means Algorithm To minimize the distortion $\min_{Z,R}\ J(R,Z)=\frac{1}{2N}\sum_{i=1}^{N}\sum_{k=1}^{K} r_{ik}\|x_i-z_k\|^2$:

Data: $\mathcal{D}=\{x_i\}_{i=1}^N, x_i\in\mathbb{R}^d$ for all i

Hyperparameters: K (number of clusters); stopping criterion (convergence of loss function j for instance)

Initialize $Z\in\mathbb{R}^{K\times d}$, that is, starting with some centroid set (which could determine algo performance)

While stopping criterion not reached do

Assign point to the nearest centroid: $r_{ik}=\begin{cases}1 & \text{if } k=\arg\min_j \|x_i-z_j\|^2\\ 0 & \text{otherwise}\end{cases},\ i=1,\dots,N$

Recompute centroid by taking average of members of cluster: $z_k=\frac{\sum_{i=1}^N r_{ik}x_i}{\sum_{i=1}^N r_{ik}},\ k=1,\dots,K$

Return: centroids Z, assignment matrix R

Depending on the initial condition, we can end up with global/local optimum or saddle point.

Maximum Likelihood Estimation Given dataset x_i, model them as i.i.d. samples from $p(x|\theta)$, θ are parameters to be determined: $\theta_{\text{MLE}}=\arg\max_\phi \log\left(\prod_i p(x_i|\phi)\right)=\sum_i\log(p(x_i|\phi))$

Gaussian Mixture Models (GMMs) Soft probabilistic labeling as compared to K-means (hard deterministic labeling): Model the data as samples from a linear convex combination of K Gaussian:

$p(x)=\sum_{k=1}^K \pi_k\, p_g(x;z_k,\Sigma_k)$, where $p_g(x_i|z,\Sigma)=(2\pi)^{-\frac{d}{2}}|\Sigma|^{-\frac{1}{2}}\exp\left\{-\frac{1}{2}(x-z)^\top\Sigma^{-1}(x-z)\right\}$,

$\pi_k\geq 0,\quad \sum_k\pi_k=1$ (Mixture weights)

Sampling from GMM

- Sample r (one-hot) according to π_k - find a member Gaussian distribution
- Identify the "hot" coordinate of r - Go to that distribution
- Sample x from $p_g(x;z_\ell,\Sigma_\ell)$ - sampling by its mean and covariance matrix

MLE for GMM

Data: $\mathcal{D}=\{x_i\}_{i=1}^N, x_i\in\mathbb{R}^d$ for all i **Hyperparameters:** K (num_clusters); stopping criterion

Initialize: $\pi_k=1/K$, $z_k\in\mathbb{R}^d$, $\Sigma_k\in\mathbb{R}^{d\times d}$ for $k=1,\dots,K$

While stopping criterion not reached do

Update γ_{ik}, the responsibility term: $r_{ik}=\frac{\pi_k p_g(x;z_k,\Sigma_k)}{\sum_{\ell=1}^K\pi_\ell p_g(x;z_\ell,\Sigma_\ell)},\ i=1,\dots,N,\ k=1,\dots,K$

Compute $\{N_k=\sum_{i=1}^N\gamma_{ik}\}$ and Update $\{\pi_k,z_k,\Sigma_k\}$:

$z_k=\frac{1}{N_k}\sum_{i=1}^N\gamma_{ik}x_i,\quad \Sigma_k=\frac{1}{N_k}\sum_{i=1}^N\gamma_{ik}(x_i-z_k)(x_i-z_k)^\top,\quad \pi_k=\frac{N_k}{N}$ for $k=1,\dots,K$

Return: cluster centers, covariances, and mixture coefficients $\{z_k,\Sigma_k,\pi_k\}$, cluster responsibilities or soft assignments $\{\gamma_{ik}\}$

Relationship between K-means and GMM Set $\Sigma_k=\epsilon I_d$, then the responsibility term becomes $\gamma_{ik}=\frac{\pi_k\exp\left(-\frac{1}{2\epsilon}\|x_i-z_k\|^2\right)}{\sum_\ell\pi_\ell\exp\left(-\frac{1}{2\epsilon}\|x_i-z_\ell\|^2\right)}$, and when we take $\epsilon\to 0$ we will get the hard label assignment as K-means.

Reinforcement Learning

Lecture 8

Key Elements of RL Action (action space, policy/chain of actions leading to optimal reward), State (match with action to form transition function, terminal / goal state), Reward (reward function)

Markov Property The future does not depend on the past, i.e. for transition probability $\mathbb{P}(S_{t+1}=s'|S_t=s,S_{t-1}=s_{t-1},\dots,S_0=s_0)$, it is simply $\mathbb{P}(S_{t+1}=s'|S_t=s)$.

Transition Matrix In time homogeneous Markov Chain, The transition probability is independent of time: $\mathbb{P}(S_{t+1}=s'|S_t=s)=P_{ss'}=p(s'|s)$ and $\{P_{ss'}\}$ is the transition matrix.

Markov Decision Process (MDP) $p(s',r|s,a)=\mathbb{P}[S_{t+1}=s',R_{t+1}=r|S_t=s,A_t=a]$, which integrate reward R and action a to the transition probability. It is finite if the action space is discrete finite for each state.

Policy $\pi(a|s)=\mathbb{P}[A_t=a|S_t=s]$, and deterministic policy is a constant mapping from the state space to action space $\pi:\mathcal{S}\to\mathcal{A}$

(Discounted) Return $G_t=\sum_{k=0}^\infty\gamma^k R_{t+k+1}$, $\gamma\in(0,1]$, where γ is the discount factor. The goal is typically to maximize the expected return, and in the context of time homogeneous case, $\mathbb{E}_\pi[G_t|S_t=s]=\mathbb{E}_\pi\left[\sum_{k=0}^\infty\gamma^k R_{t+k+1}\middle|S_t=s\right]$

Dynamic Programming Memoization / caching the result of overlapping optimal substructures. Trade space for cost of recomputation for naive implementations and space usage can be optimized through iterative approach.

Bellman's Equation $v_\pi(s)=\sum_a\pi(a|s)\sum_{s',r}p(s',r|s,a)[r+\gamma v_\pi(s')]$

This can be derived using DP to formulate a recursive solver for value function $v_\pi(s)=\mathbb{E}_\pi[G_t|S_t=s]$ and action value function $q_\pi(s,a)=\mathbb{E}_\pi[G_t|S_t=s,A_t=a]$. and substitute the action value function to the value function.

Bellman's Equation for Finite MDP $v_\pi=\gamma P(\pi)v_\pi+b(\pi)$, where:

- $P(\pi)_{s,s'}=\sum_a\pi(a|s)\sum_r p(s',r|s,a)$ is the state transition matrix for the policy π ($N*N$ matrix with N being number of states, so we only sum over r)
- $b(\pi)(s)=\sum_a\pi(a|s)\sum_{s',r}p(s',r|s,a)\cdot r$ is the expected immediate reward for the policy π ($N*1$ matrix)

The unique solution would thus be $v_\pi=(I-\gamma P(\pi))^{-1}b(\pi)$.

Optimal Policy $\pi_*\geq\pi\quad\forall\pi\neq\pi_*$ Always exist but may not be unique.

Policy Improvement For any two policies π,π', if $\sum_a\pi'(a|s)q_\pi(s,a)\geq\sum_a\pi(a|s)q_\pi(s,a)\forall s$ then $v_{\pi'}(s)\geq v_\pi(s)\forall s$, and we assert π' is better than π.

Bellman's Optimality Equations Given an optimal policy $\pi_*(s)\in\arg\max_a q_*(s,a)$, we can recursively find optimal value and action value functions:

$v_*(s)=\max_{a\in\mathcal{A}}\sum_{s',r}p(s',r|s,a)[r+\gamma v_*(s')]$ $\quad q_*(s,a)=\sum_{s',r}p(s',r|s,a)\left[r+\gamma\max_{a'\in\mathcal{A}}q_*(s',a')\right]$

$v_*(s)=\max_a q_*(s,a)$ Optimal Policy: $\pi_*(s)\in\arg\max_a q_*(s,a)$

Lecture 9

Model-Based vs Model-Free Model-based assumes that the transition function is known: Value iteration, policy iteration. Model-free improves through sampling/ Monte Carlo simulations: Temporal Difference (TD), Q-learning.

Value Iteration Algorithm Start with some value function, update utility estimate iteratively and keep track of corresponding current optimal actions.

Model Parameters: MDP transition probability $p(s',r|s,a)$, discount rate $\gamma<1$

Input: Policy π, Stopping criterion

Initialize $v\in\mathbb{R}^n$

While stopping criterion not reached do

Update value function: $v\leftarrow F(v)=\max_\pi\{\gamma P(\pi)v+b(\pi)\}$ (Make use of P, model-based)

Return v

Reinforcement Learning

Contraction Mapping Theorem Fix point iteration will lead to a unique $v_*\in V$ such that $v_*=F(v_*)$, and v_* can be estimated by starting at arbitrary v_0 and iterate for many times (towards infinity).

Value Iteration Convergence Rate Given error ϵ being the difference between v_K and v_∞, and number of iterations k, to achieve $\mathcal{O}(\epsilon)$ we need $k\sim\mathcal{O}(\log(1/\epsilon))$

Policy Iteration Algorithm Start with some policy, plug into value function and derive a better policy. Repeat until policy does not change.

Model Parameters: MDP transition probability $p(s',r|s,a)$, discount rate $\gamma<1$

Initialize: Deterministic policy $\pi\leftarrow\pi_0$

While $\pi\neq\pi'$ do

$\pi'\leftarrow\pi$ (Set the next policy for evaluation)

$v_\pi\leftarrow(I-\gamma P(\pi))^{-1}b(\pi)$ (Evaluate policy π using P, model-based)

$\pi'\leftarrow\arg\max_\pi\{b(\pi)+\gamma P(\pi)v_\pi\}$ (Retrieve subsequent best policy)

Return v

Its convergence can be shown by $v_{\pi_k}\leq v_{\pi_{k+1}}\leq v_*$ from the algorithm (argmax, minimally equivalent values) and $\pi_{k+1}\neq\pi_k$ if and only if $v_{\pi_{k+1}}>v_{\pi_k}$ (otherwise the while loop ends)

Monte Carlo Method IID sampling and compute expectation: $\mathbb{E}_{x\sim\mu}f(x)\approx\frac{1}{N}\sum_{i=1}^N f(X_i)$ where $X_i\sim\mu$ is IID

Model-free Policy Iteration Instead of relaying on the transition matrix P we sample episodes according current π to get samples of state and reward sequence (S and R), and get the value by averaging discounted rewards. Concretely, update estimation for action-value function by: $q(s,a)\leftarrow\frac{1}{N}\sum_{n=1}^N\left[\sum_{k=0}^\infty\gamma^k R_{k+1}^{(n)}\middle|S_0^{(n)}=s,\ A_0^{(n)}=a\right]$ and perform policy improvement based on this $q(s,a)$.

Temporal Difference (TD) Algorithm

Input: State-reward simulator, initial state sampler, Stopping criterion, Policy π

Initialize: Initial value v_0, Episode length T, Start with $v\leftarrow v_0$;

While stopping criterion not reached do

$s\leftarrow$ InitialStateSampler()(The start of the new episode)

For $t=0$ to $T-1$ do

$a\leftarrow a\sim\pi(\cdot|s)$

$s',r\leftarrow$ StateRewardSimulator(s,a)

$v(s)\leftarrow(1-\alpha)v(s)+\alpha\left(r+\gamma v(s')\right)$

Return v

Q-learning Algorithm

Input: State-reward simulator, initial state sampler, Stopping criterion, Policy π

Initialize: Initial value q_0, Episode length T, Start with $q\leftarrow q_0$

While stopping criterion not reached do

$s\leftarrow$ InitialStateSampler()

For $t=0$ to $T-1$ do

$a\leftarrow a\sim\pi(\cdot|s)$

$s',r\leftarrow$ StateRewardSimulator(s,a)

$q(s,a)\leftarrow(1-\alpha)q(s,a)+\alpha\left[r+\gamma\max_{a'}q(s',a')\right]$

Return q

Graph-Based Methods

Lecture 10

Page Rank Let $L_k\subset\{1,2,\dots,n\}$ denote the set of pages with a link to page k; that is, L_k is the set of page k's backlinks. For each k we require $x_k=\sum_{j\in L_k}\frac{x_j}{n_j}$ where n_j is the number of outgoing links from page j.

The resultant matrix A representing pairwise values of x_k is a column-stochastic matrix with an eigenvalue equal to 1, and 1 is also its largest eigenvalue. For instance: $\text{A}=\begin{bmatrix}0&0&1&\frac{1}{2}\\ \frac{1}{3}&0&0&0\\ \frac{1}{3}&\frac{1}{2}&0&0\\ \frac{1}{3}&\frac{1}{2}&0&0\end{bmatrix}$

Power Iteration Let the eigenvalues of A be $\lambda_1>\lambda_2>\cdots>\lambda_n$ and the associated eigenvectors be $v_1,v_2,\dots,v_n$. Then any vector x_0 can be written as $x_0=c_1v_1+c_2v_2+\cdots+c_nv_n$. Then $A^kx_0=c_1\lambda_1^kv_1+c_2\lambda_2^kv_2+\cdots+c_n\lambda_n^kv_n=c_1\lambda_1^k\left(v_1+\frac{c_2}{c_1}\left(\frac{\lambda_2}{\lambda_1}\right)^k v_2+\cdots+\frac{c_n}{c_1}\left(\frac{\lambda_n}{\lambda_1}\right)^k v_n\right)\to c_1\lambda_1^kv_1$ as $k\to\infty$.

Lecture 11

Graph Laplacian $L_G:=D-W=\sum_{i<j}w_{ij}(e_i-e_j)(e_i-e_j)^\top$, the diagnol entries are the degree of each node and other entries are $-w_{ij}$

Cheeger's cut $h(S)=\frac{\text{cut}(S)}{\min\{\text{vol}(S),\text{vol}(S^c)\}}$ **Normalized cut** (Ncut) $Ncut(S)=\frac{\text{cut}(S)}{\text{vol}(S)}+\frac{\text{cut}(S^c)}{\text{vol}(S^c)}$

$h(S)\leq Ncut(S)\leq 2h(S)$

From Ncut to relaxed balanced cut $Ncut(S)=y^\top L_G y$ $\quad y_i=\begin{cases}\left(\frac{\text{vol}(S^c)}{\text{vol}(S)\,\text{vol}(G)}\right)^{\frac{1}{2}} & \text{if } i\in S,\\ -\left(\frac{\text{vol}(S)}{\text{vol}(S^c)\,\text{vol}(G)}\right)^{\frac{1}{2}} & \text{if } i\in S^c.\end{cases}$

Spectral Clustering for multi-clusters

Given a graph $G=(V,E,W)$, let $v_2,\dots,v_k$ be the eigenvectors corresponding to the second through $(k-1)$th eigenvalues of the normalized Laplacian $\mathcal{L}_G=D^{-\frac{1}{2}}L_GD^{-\frac{1}{2}}$;

Let $\phi_m=D^{-\frac{1}{2}}v_m\in\mathbb{R}^n$;

Consider the map $\phi:V\to\mathbb{R}^{k-1}$ defined as $\phi(v_i)=[\phi_2(i)\quad\cdots\quad\phi_k(i)]^T$, where D is the degree matrix with $D_{ii}=\deg(i)=\sum_{j=1}^n w_{ij}$;

Cluster the n points in $k-1$ dimensions into k clusters using k-means.

For example, given $k=2$, we can compute $\phi_2=D^{-\frac{1}{2}}v_2$, choose a threshold τ (try and error or k-means for $k=2$), and set $S=\{i\in V:\phi_2(i)\leq\tau\}$.

Cheeger's inequality $\frac{1}{2}\lambda_2(\mathcal{L}_G)\leq h_G\leq\sqrt{2\lambda_2(\mathcal{L}_G)}$