CIL Cheat Sheet 2015

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I. LINEAR ALGEBRA PRIMER

A. Equivalent Conditions

For $\mathbf{A} \in \mathbb{R}^{M \times M}$ the following conditions are equivalent:

- A has an inverse A^{-1} ,
- $\operatorname{rank}(\mathbf{A}) = M$,
- range(\mathbf{A}) = \mathbb{R}^M ,
- $null(A) = \{0\},\$
- 0 is not an eigenvalue of A,
- 0 is not a singular value of A

II. NORMS

A. Vector norms

A norm is a function $\|\cdot\|:V\to\mathbb{R}$ quantifying the size of a vector. It must satisfy

- 1) Positive scalability: $||a \cdot \mathbf{x}|| = |a| \cdot ||\mathbf{x}||$ for $a \in \mathbb{R}$
- 2) Triangle inequality: $\|\mathbf{x} + \mathbf{y}\| \le \|\mathbf{x}\| + \|\mathbf{y}\|$ for $\mathbf{x}, \mathbf{y} \in$ V.
- 3) Separability: $\|\mathbf{x}\| = 0$ implies $\mathbf{x} = 0$.
- The most commonly used norms are the *p-norms*:

$$\|\mathbf{x}\|_p := \left(\sum_{i=1}^n |x_i|^p\right)^{1/p}$$

• A special case is the Euclidean norm

$$\|\mathbf{x}\|_2 := \sqrt{\sum_{i=1}^n x_i^2}$$

• The "0-norm" (not really a norm, does not satisfy positive scalability) is $\|\mathbf{x}\|_0 := |\{x_i \mid x_i \neq 0\}|$

B. Matrix norms

We can also define norms on matrices, satisfying the properties described above. $\mathbf{A} \in \mathbb{R}^{M \times N}$:

• Frobenius norm:

$$\|\mathbf{A}\|_F := \sqrt{\sum_{i=1}^M \sum_{j=1}^N a_{ij}^2} = \sqrt{\sum_{i=1}^K \sigma_i^2}, \quad K = \min(M, N) \quad \text{form. (Assumes ordering of singular values } d_k \geq d_{k+1})$$

Only depends on singular values of A

- p-norms for matrices: $\|\mathbf{A}\|_p := \sup\{\|\mathbf{A}\mathbf{x}\|_p : \|\mathbf{x}\|_p = \|\mathbf{A}\mathbf{x}\|_p =$
- Euclidean or spectral norm:

$$\|\mathbf{A}\|_{2} := \sup\{\|\mathbf{A}\mathbf{x}\|_{2} : \|\mathbf{x}\|_{2} = 1\} = \sigma_{1},$$

the largest singular value.

III. STATISTICS

A. Kullback-Leibler Divergence

- Divergence between discrete probability distributions P and $Q: D_{\mathrm{KL}}(P||Q) = \sum_{\omega \in \Omega} P(\omega) \log \left(\frac{P(\omega)}{Q(\omega)} \right)$.
- Properties of the Kullback-Leibler Divergence
 - $D_{\mathrm{KL}}(P||Q) \ge 0$.
 - $D_{KL}(P||Q) = 0$ if and only if P and Q are identical.
 - $D_{\mathrm{KL}}(P||Q) \neq D_{\mathrm{KL}}(Q||P)!$
 - caution: the KL-Divergence is not symmetric, therefore it is not a metric/distance!

IV. DIMENSION REDUCTION

A. Principal Component Analysis (PCA)

Orthogonal linear projection of high dimensional data onto low dimensional subspace. Objectives:

- 1) Minimize error $\|\mathbf{x} \tilde{\mathbf{x}}\|_2$ of point \mathbf{x} and its approxi-
- 2) Preserve information: maximize variance.

Both objectives are shown to be formally equivalent.

- 1) Statistics of Projected Data:
- Mean of the data: sample mean $\bar{\mathbf{x}}$
- Covariance of the data:

$$\Sigma = \frac{1}{N} \sum_{n=1}^{N} (\mathbf{x}_n - \bar{\mathbf{x}}) (\mathbf{x}_n - \bar{\mathbf{x}})^{\top}$$

- 2) Solution: Eigenvalue Decomposition: The eigenvalue decomposition of the covariance matrix $\Sigma = \mathbf{U} \Lambda \mathbf{U}^{\top}$ contains all relevant information.
 - For $K \leq D$ dimensional projection space: Choose K eigenvectors $\{\mathbf{u}_1, \dots, \mathbf{u}_K\}$ with largest associated eigenvalues $\{\lambda_1, \ldots, \lambda_K\}$.

B. Singular Value Decomposition

Theorem (Eckart-Young). Let A be a matrix of rank R, if we wish to approximate A using a matrix of a lower rank K then,

V. CLUSTERING

A. K-Means

- 1) Motivation:
- Given: set of data points $\mathbf{x}_1, \dots, \mathbf{x}_N \in \mathbb{R}^D$
- Goal: find meaningful partition of the data
 - i.e. a labeling of each data point with a unique label

$$\pi: \{1, ..., N\} \to \{1, ..., K\} \text{ or } \pi: \mathbb{R}^D \to \{1, ..., K\}$$

- note: numbering of clusters is arbitrary
- k-th cluster recovered by $\pi^{-1}(k) \subseteq \{1,\ldots,N\}$ or $\subset \mathbb{R}^D$
- 2) Vector Quantization:
- Partition of the space \mathbb{R}^D
- Clusters represented by *centroids* $\mathbf{u}_k \in \mathbb{R}^D$
- · Mapping induced via nearest centroid rule

$$\pi(\mathbf{x}) = \underset{k=1,\dots,K}{\operatorname{argmin}} \|\mathbf{u}_k - \mathbf{x}\|_2$$

- 3) Objective Function for K-Means:
- Useful notation: represent π via indicator matrix **Z**:

$$z_{kn} := [\pi(\mathbf{x}_n) = k]$$

• K-means Objective function

$$J(\mathbf{U}, \mathbf{Z}) = \sum_{n=1}^{N} \sum_{k=1}^{K} z_{kn} \|\mathbf{x}_{n} - \mathbf{u}_{k}\|_{2}^{2} = \|\mathbf{X} - \mathbf{U}\mathbf{Z}\|_{F}^{2},$$

where $\mathbf{X}=[\mathbf{x}_1,\ldots,\mathbf{x}_N]\in\mathbb{R}^{D\times N}$ and $\mathbf{U}=[\mathbf{u}_1,\ldots,\mathbf{u}_K]\in\mathbb{R}^{D\times K}$

- 4) K-means Algorithm: Optimal Assignment:
- Compute optimal assignment Z, given centroids U
 - minimize each column of Z separately

$$\mathbf{z}_{\bullet n}^* = \operatorname*{argmin}_{z_{1n}, \dots, z_{Kn}} \sum_{k=1}^K z_{kn} \|\mathbf{x}_n - \mathbf{u}_k\|_2^2$$

- optimum is attained by mapping to the closest centroid

$$z_{kn}^*(\mathbf{U}) = \left[k = \underset{l}{\operatorname{argmin}} \|\mathbf{x} - \mathbf{u}_l\|_2\right]$$

- 5) K-means Algorithm: Optimal Assignment:
- Compute optimal choice of U, given assignments Z
 - continuous variables: compute gradient and set to zero (necessary optimality condition)
 - look at (partial) gradient for every centroid \mathbf{u}_k

$$\nabla_{\mathbf{u}_k} J(\mathbf{U}, \mathbf{Z}) = \sum_{n=1}^N z_{kn} \nabla_{\mathbf{u}_k} \|\mathbf{x}_n - \mathbf{u}_k\|_2^2 = -2 \sum_{n=1}^N z_{kn} (\mathbf{x}_n - \underline{\mathbf{u}}_k)_{\text{follows from Jensen's inequality (concavity of log-$$

- setting gradient to zero

$$\nabla_{\mathbf{U}} J(\mathbf{U}, \mathbf{Z}) \stackrel{!}{=} 0 \implies \mathbf{u}_{k}^{*}(\mathbf{Z}) = \frac{\sum_{n=1}^{N} z_{kn} \mathbf{x}_{n}}{\sum_{n=1}^{N} z_{kn}}$$

- 6) K-means Algorithm: Analysis:
- Computational cost of each iteration is O(KND)
- K-means convergence is guaranteed
- K-means optimizes a non-convex objective. Hence we are not guaranteed to find the global optimum.
- Finds a local optimum (U, Z) in the following sense
 - for each \mathbf{Z}' with $\frac{1}{2} \|\mathbf{Z} \mathbf{Z}'\|_0 = 1$ (differs in one assignment)
 - $J(\mathbf{U}^*(\mathbf{Z}'), \mathbf{Z}') \ge J(\mathbf{U}, \mathbf{Z})$
 - may gain by changing assignments of ≥ 2 points
- K-means algorithm can be used to compress data
 - with information loss, if K < N
 - store only the centroids and the assignments

B. Mixture Models

1) Gaussian Mixture Model (GMM):

$$p_{\theta}(\mathbf{x}) = \sum_{k=1}^{K} \pi_k \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k),$$

where $\pi_k \geq 0$, $\sum_{k=1}^K \pi_k = 1$.

- 2) Complete Data Distribution:
- Explicitly introduce latent variables in the generative
- Assignment variable (for a generic data point) $z_k \in$ $\{0,1\}, \sum_{k=1}^{K} z_k = 1$
- We have that $\Pr(z_k = 1) = \pi_k$ or $p(\mathbf{z}) = \prod_{k=1}^K \pi_k^{z_k}$
- Joint distribution over (\mathbf{x}, \mathbf{z}) (complete data distribution) $p(\mathbf{x}, \mathbf{z}) = \prod_{k=1}^{K} [\pi_k \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)]^{z_k}$
- 3) Posterior Assignments: Posterior probabilities for assignments

$$\Pr(z_k = 1 \mid \mathbf{x}) = \frac{\pi_k \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{l=1}^K \pi_l \mathcal{N}(\mathbf{x} \mid \boldsymbol{\mu}_l, \boldsymbol{\Sigma}_l)}$$

- 4) Lower Bounding the Log-Likelihood:
- Expectation Maximization
 - maximize a lower bound on the log-likelihood
 - systematic way of deriving a family of bounds
 - based on complete data distribution
- Specifically:

$$\ln p_{\theta}(\mathbf{x}) = \ln \sum_{\mathbf{z}} p_{\theta}(\mathbf{x}, \mathbf{z}) = \ln \sum_{k=1}^{K} p(\mathbf{x}, \theta_k) \pi_k$$
$$= \ln \sum_{k=1}^{K} q_k \frac{p(\mathbf{x}; \theta_k) \pi_k}{q_k}$$
$$\geq \sum_{k=1}^{K} q_k \left[\ln p(\mathbf{x}, \theta_k) + \ln \pi_k - \ln q_k \right]$$

- - can be done for the contribution of each data point (additive)
- 5) Mixture Model: Expectation Step:

$$q_k = \frac{\pi_k \ p(\mathbf{x}; \theta_k)}{\sum_{l=1}^K \pi_l \ p(\mathbf{x}, \theta_l)} = \Pr(z_k = 1 \mid \mathbf{x})$$

6) Mixture Model: Maximization Step:

$$\pi_{k}^{*} = \frac{1}{N} \sum_{n=1}^{N} q_{kn}$$
$$\mu_{k}^{*} = \frac{\sum_{n=1}^{N} q_{kn} \mathbf{x}_{n}}{\sum_{n=1}^{N} q_{kn}}$$

$$\boldsymbol{\Sigma}_k^* = \frac{\sum_{n=1}^N q_{kn} (\mathbf{x}_n - \boldsymbol{\mu}_k^*) (\mathbf{x}_n - \boldsymbol{\mu}_k^*)^\top}{\sum_{n=1}^N q_{kn}}$$

7) AIC and BIC:

- Trade-off: achieve balance between data fit measured by likelihood $p(\mathbf{X} \mid \theta)$ — and complexity. Complexity can be measured by the number of free parameters $\kappa(\cdot)$.
- Different Heuristics for choosing K
 - Akaike Information Criterion (AIC)

$$AIC(\theta \mid \mathbf{X}) = -\ln p_{\theta}(\mathbf{X}) + \kappa(\theta)$$

- Bayesian Information Criterion (BIC)

$$BIC(\theta \mid \mathbf{X}) = -\ln p_{\theta}(\mathbf{X}) + \frac{1}{2}\kappa(\theta)\ln N$$

• Generally speaking, the BIC criterion penalizes complexity more than the AIC criterion.

C. Non-Negative Matrix Factorization

- 1) Non-Negative Matrix Factorization:
- Document-term matrix $\mathbf{X} \in \mathbb{R}^{D \times N}_{\geq 0}$ storing the word counts for each document:

$$\mathbf{X} = \mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N$$

 x_{dn} : Frequency of the d-th word in the n-th document.

• Non-negative matrix factorization (NMF) of X:

$$\mathbf{X} \approx \mathbf{U}\mathbf{Z}$$

- with $\mathbf{U} \in \mathbb{R}_{\geq 0}^{D imes K}$ and $\mathbf{Z} \in \mathbb{R}_{\geq 0}^{K imes N}$
 - * N: number of documents
 - * D: vocabulary size
 - * K: number of dimensions (design choice)
 - * data reduction: $(D+N)K \ll DN$

2) pLSI — Generative Model:

- For a given *document* sample len(document) words by a two-stage procedure:
 - sample a topic according to P(topic | document)
 - sample a word according to $P(\text{word} \mid \text{topic})$
- Key assumption: conditional independence of word and document given topic
- Conditional distribution of a word, given a document:

$$P(\text{word} \mid \text{document}) = \sum_{k=1}^{K} P(\text{word} \mid \text{topic}_k) P(\text{topic}_k \mid \text{document})$$

- Side note: how to sample a "new" document? Can use fully generative model of LDA.
- 3) pLSI Matrix Factorization View:
- Normalize the elements of X so that they correspond to relative frequencies:

$$T := \sum_{d=1}^{D} \sum_{n=1}^{N} x_{dn}, \qquad x_{dn} \leftarrow \frac{x_{dn}}{T}$$

- Matrix Factorization
 - pLSI can be understood as a matrix factorization of the form $\mathbf{X} \approx \mathbf{U}\mathbf{Z}$, with $\mathbf{U} \in \mathbb{R}_{>0}^{D \times K}$, and $\mathbf{Z} \in$

- where additionally we have the constraints:
 - * $\sum_{d=1}^{D} u_{dk} = 1(\forall k)$, identify $u_{dk} \equiv P(\text{word}_d \mid$
 - * $\sum_{k,n} z_{kn} = 1$, identify $z_{kn} \equiv P(\operatorname{topic}_k \mid$ $document_n)P(document_n)$
- 4) pLSI Parameter Estimation:
- Goal: maximize the likelihood of the data under the
- Data: the relative frequencies X
- Probabilistic model: $P(\text{word}_d, \text{document}_n)$ $\sum_{k=1}^{K} P(\text{word}_d \mid \text{topic}_k) P(\text{topic}_k \mid \text{document}_n) =$
- Log likelihood: $\log \mathcal{L}(\mathbf{U}, \mathbf{Z}; \mathbf{X}) = \log P(\mathbf{X}; \mathbf{U}, \mathbf{Z}) = \sum_{d=1}^{D} \sum_{n=1}^{N} x_{dn} \log \sum_{k=1}^{K} u_{dk} z_{kn}$
- 5) EM for pLSI Variational Likelihood:
- Follow similar recipe as for Gaussian Mixture Model
- Reindex the observations in a per token manner with $t = 1, \ldots, T$

 - pairs of word/documents indexes (d_t, n_t) note that $\sum_{t=1}^T f(d_t, n_t) = \sum_{d=1}^D \sum_{n=1}^N x_{dn} f(d, n)$ for arbitrary functions f
- Variational Likelihood

$$\log P(\mathbf{X}; \mathbf{U}, \mathbf{Z})$$

$$= \sum_{t=1}^{T} \log (\mathbf{U}\mathbf{Z})_{d_t n_t} = \sum_{t=1}^{T} \log \left[\sum_{k=1}^{K} u_{d_t k} z_{k n_t} \right]$$

$$\geq \sum_{t=1}^{T} \max_{q \in \mathcal{S}_K} \sum_{k=1}^{K} q_k [\log u_{d_t k} + \log z_{k n_t} - \log q_k]$$

- $\mathcal{S}_K := \left\{ x \in \mathbb{R}^K \mid x \geq 0, \sum_{k=1}^K x_k = 1 \right\}$ (probability simplex)
- 6) EM for pLSI Derivation of E-step:
- Compute the argmin in the variational bound

$$q_t^* = \operatorname*{argmax}_{q \in \mathcal{S}_K} \sum_{k=1}^K q_k [\log u_{d_t k} + \log z_{k n_t} + \log q_k]$$

• Form Lagrangian and differentiate

eument)
$$\frac{\partial}{\partial q_k} \{q_k [\log u_{d_t k} + \log z_{k n_t} - \log q_k - \lambda_t^*]\} \stackrel{!}{=} 0$$

$$\implies q_{tk}^* \propto u_{d_t k} z_{k n_t}, \text{ i.e. } q_{tk}^* = \frac{u_{d_t k} z_{k n_t}}{\sum_{l=1}^K u_{d_t l} z_{l n_t}}$$

- q_{tk}^* = posterior probability that t-th token (i.e. word with index d_t in document with index n_t) has been generated from topic k
- 7) EM for pLSI Derivation of M-step:
- Differentiate lower bound with plugged in optimal choices for q_t^* (t = 1, ..., T)
- M-step solution for U and Z

$$u_{dk}^* = \frac{\sum_{t:d_t=d} q_{tk}^*}{\sum_{t=1}^T q_{tk}^*} \qquad z_{kn}^* = \frac{\sum_{t:n_t=n} q_{tk}^*}{T}$$

VI. SPARSE CODING

A. Optimization

- 1) Coordinate Descent: Idea: Update one coordinate at a time, while keeping others fixed.
 - Algorithm:
 - initialize $\mathbf{x}^{(0)} \in \mathbb{R}^D$
 - for $t = 0, \ldots, \max \text{Iter}$
 - * $d \leftarrow \mathcal{U}\{1, D\}$

*
$$u^* \leftarrow \operatorname{argmin}_{u \in \mathbb{R}} f\left(x_1^{(t)}, \dots, x_{d-1}^{(t)}, u, x_{d+1}^{(t)}, \dots, x_D^{(t)}\right)$$

- $* x_d^{(t+1)} \leftarrow u^*, \quad x_{d'}^{(t+1)} \leftarrow x_{d'}^{(t)} \text{ for } d' \neq d$
- 2) Gradient Descent Method:
- Algorithm:
 - initialize $\mathbf{x}^{(0)} \in \mathbb{R}^D$
 - for $t = 0, \ldots, \max \text{Iter}$

*
$$\mathbf{x}^{(t+1)} \leftarrow \mathbf{x}^{(t)} - \gamma \nabla f(\mathbf{x}^{(t)})$$

- simple to implement
- good scalability and robustness
- stepsize γ usually decreasing with $\gamma \approx \frac{1}{4}$
- 3) Stochastic Gradient Descent:
- Optimization Problem Structure: minimize $f(\mathbf{x}) =$ $\frac{1}{N}\sum_{n=1}^{N}f_n(\mathbf{x})$ with $\mathbf{x} \in \mathbb{R}^D$
- Algorithm:
 - initialize $\mathbf{x}^{(0)} \in \mathbb{R}^D$
 - for $t = 0, \ldots, \text{maxIter}$

*
$$n \leftarrow \mathcal{U}\{1, N\}$$

* $\mathbf{x}^{(t+1)} \leftarrow \mathbf{x}^{(t)} - \gamma \nabla f_n(\mathbf{x}^{(t)})$

- 4) Duality for Constrained Optimization:
- Constrained Problem Formulation (Standard Form): minimize $f(\mathbf{x})$ subject to $g_i(\mathbf{x}) \leq 0$, $i = 1, \ldots, m$, $h_i(\mathbf{x}) = 0, i = 1, \dots, p$
- Unconstrained Problem: minimze $\sum_{i=1}^{m} I_{-}(g_i(\mathbf{x})) + \sum_{i=1}^{p} I_0(h_i(\mathbf{x}))$. I_{-} and I_0 are "brickwall" indicator functions.
- 5) Dual Problem:
- Lagrangian: $L(\mathbf{x}, \boldsymbol{\lambda}, \boldsymbol{\nu}) := f(\mathbf{x}) + \sum_{i=1}^{m} \lambda_i g_i(\mathbf{x}) +$ $\sum_{i=1}^{p} \nu_i h_i(\mathbf{x})$
- Lagrange dual function: $d(\lambda, \nu) := \inf_{\mathbf{x}} L(\mathbf{x}, \lambda, \nu)$
- Lagrange dual problem: maximize $d(\lambda, \nu)$ subject to $\lambda \geq 0$.
 - It is always a lower bound on the primal value $f(\mathbf{x})$ of any feasible x and thus a lower bound on the unknown solution value $f(\mathbf{x}^*)$ of the primal problem.
 - Strong Duality: If the primal optimization problem is convex and under some additional conditions, the solution value of the dual problem is *equal* to the solution value $f(\mathbf{x}^*)$ of the primal problem.
- 6) Convexity:
- Convex Set: A set Q is convex if for any $x, y \in Q$ and any $\theta \in [0, 1]$, we have $\theta \mathbf{x} + (1 - \theta) \mathbf{y} \in Q$.

- Convex Function: A function $f: \mathbb{R}^D \to \mathbb{R}$ is convex if dom f is a convex set and if for all $x, y \in \text{dom } f$, and $\theta \in [0,1]$ we have $f(\theta \mathbf{x} + (1-\theta)\mathbf{y}) \leq \theta f(\mathbf{x}) + (1-\theta)\mathbf{y}$ θ) $f(\mathbf{y})$.
- Convex Optimization: Convex Optimization Problems are of the form $\min f(\mathbf{x})$ s.t. $\mathbf{x} \in Q$ where both f is a convex function and Q is a convex set (note: \mathbb{R}^D is convex). In Convex Optimization Problems every local minimum is a global minimum.

VII. ROBUST PCA