Statistical Learning for Text Data Analytics Text Categorization 2: Clustering

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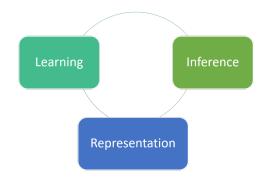
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Spring 2018

*Contents are based on materials created by Noah Smith, Xiaojin (Jerry) Zhu, Eric Xing, Vivek Srikumar, Dan Roth

- Noah Smith. CSE 517: Natural Language Processing https://courses.cs.washington.edu/courses/cse517/16wi/
- Xiaojin (Jerry) Zhu. CS 769: Advanced Natural Language Processing. http://pages.cs.wisc.edu/~jerryzhu/cs769.html
- Eric Xing. 10715 Advanced Introduction to Machine Learning. https://www.cs.cmu.edu/~epxing/Class/10715/lectures/ lecture1.pdf
- Vivek Srikumar. CS 6355 Structured Prediction. https: //svivek.com/teaching/structured-prediction/spring2018/
- Dan Roth. CS546: Machine Learning and Natural Language . http://l2r.cs.uiuc.edu/~danr/Teaching/CS546-16/



- Representation: language models, word embeddings, topic models
- Learning: supervised learning, unsupervised learning, semi-supervised learning, sequence models, deep learning, optimization techniques
- Inference: constraint modeling, joint inference, search algorithms

NLP applications: tasks introduced in Lecture 1

Problem Definition

2 Generative vs. Discriminative Classification

3 General Linear Classification

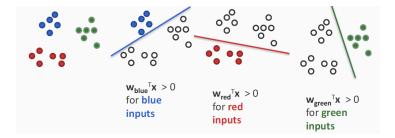
- 4 Unsupervised Learning
- 5 EM Algorithm
- 6 Evaluation of Classification
- 7 Evaluation of Clustering

- Decompose the prediction into multiple binary decisions
 - One-vs-all
 - No theoretical justification
 - Calibration issues: We are comparing scores produced by K classifiers trained independently. No reason for the scores to be in the same numerical range!
 - Might not always work: Yet, works fairly well in many cases, especially if the underlying binary classifiers are tuned, regularized
 - All-vs-all
 - $O(K^2)$ weight vectors to train and store
 - Size of training set for a pair of labels could be very small, leading to overfitting of the binary classifiers
 - Prediction is often ad-hoc and might be unstable. E.g., What if two classes get the same number of votes? For a tournament, what is the sequence in which the labels compete?

4 ∃ > 4

Recall: One-vs-all Classification

- Assumption: Each class individually separable from all the others
- Train K binary classifiers **w**₁, **w**₂, ... **w**_K using any binary classification algorithm we have seen
- Prediction: "Winner Takes All": *label* = arg max_i $\mathbf{w}_i^{\top} \mathbf{x}$



- Rewrite input features and weight vector
 - Define a feature vector for label *i* being associated to input **x**
 - Stack all weight vectors into an *nK*-dimensional vector

$$\phi(\mathbf{x},i) = \begin{bmatrix} \mathbf{0}_n \\ \vdots \\ \mathbf{x} \\ \vdots \\ \mathbf{0}_n \end{bmatrix}_{nK \times 1} \mathbf{w} = \begin{bmatrix} \mathbf{w}_1 \\ \mathbf{w}_2 \\ \vdots \\ \mathbf{w}_K \end{bmatrix}_{nK \times 1}$$

This is called the Kesler construction

For an example with label *i*, we want **w**_i^T**x** > **w**_j^T**x** for all *j*This is equivalent to

$$\mathbf{w}^{\top}\phi(\mathbf{x},i) > \mathbf{w}^{\top}\phi(\mathbf{x},j)$$

or

$$\mathbf{w}^{\top}[\phi(\mathbf{x},i) - \phi(\mathbf{x},j)] > 0$$

- $\bullet\,$ The number of weights is still same as one-vs-all, much less than all-vs-all K(K-1)/2
- Still account for all pairwise label preferences
- Come with theoretical guarantees for generalization
- Important idea that is applicable when we move to arbitrary structures

• "Linear" decision rule

$$\hat{y} = \arg \max_{y \in \mathcal{Y}} \mathbf{w}^{\top} \phi(\mathbf{x}, y)$$

where
$$\phi: \mathcal{V} \times \mathcal{Y} \to \mathbb{R}^d$$

- Parameters: $\mathbf{w} \in \mathbb{R}^d$
- What does this remind you of?

MLE for Multinomial Logistic Regression

• When we discussed log-linear language models, we transformed the score into a probability distribution. Here, that would be

$$P(y|\mathbf{x}) = \frac{\exp(\mathbf{w}^{\top}\phi(\mathbf{x}, y))}{\sum_{y'}\exp(\mathbf{w}^{\top}\phi(\mathbf{x}, y'))}$$

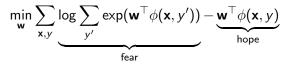
• MLE can be rewritten as a maximization problem:

$$\mathbf{w}^* = \arg \max_{\mathbf{w}} \sum_{\mathbf{x}, y} \underbrace{\mathbf{w}^\top \phi(\mathbf{x}, y)}_{\text{hope}} - \underbrace{\log \sum_{y'} \exp(\mathbf{w}^\top \phi(\mathbf{x}, y'))}_{\text{fear}}$$

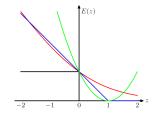
- Recall from language models:
 - Be wise and regularize!
 - Solve with batch or stochastic gradient methods
 - w_i has an interpretation

Log Loss for (\mathbf{x}, y)

 Another view is to minimize the negated log-likelihood, which is known as "log loss":



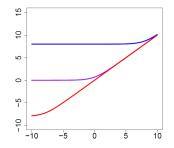
• In the binary case, where the x-axis is the difference in scores between correct and incorrect labels:



All loss functions are considered as upper-bound of "zero-one" loss (error)

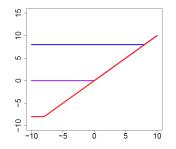
Log Sum Exp

 Below, y-axis plots the log ∑ exp" part of the objective function (with two labels), against x, assuming the other score is one of {8,0, -8}



 $\log(e^{x} + e^{8}), \log(e^{x} + e^{0}), \log(e^{x} + e^{-8})$

• Why not use a hard max instead?



 $\max(x, 8), \max(x, 0), \max(x, -8)$

Hinge Loss for (\mathbf{x}, y)

Average log-loss

$$\min_{\mathbf{w}} \sum_{\mathbf{x}, y} \underbrace{\log \sum_{y'} \exp(\mathbf{w}^{\top} \phi(\mathbf{x}, y'))}_{\text{fear}} - \underbrace{\mathbf{w}^{\top} \phi(\mathbf{x}, y)}_{\text{hope}}$$

$$\min_{\mathbf{w}} \sum_{\mathbf{x}, y} \underbrace{\max_{y'}}_{fear} (\mathbf{w}^{\top} \phi(\mathbf{x}, y'))}_{fear} - \underbrace{\mathbf{w}^{\top} \phi(\mathbf{x}, y)}_{hope}$$

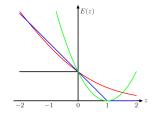
- When two labels are tied, the function is not differentiable
- But it's still sub-differentiable. Solution: (stochastic) sub-gradient descent!

Compare Loss

$$\min_{\mathbf{w}} \sum_{\mathbf{x}, y} \underbrace{\max_{y'}}_{\text{fear}} (\mathbf{w}^{\top} \phi(\mathbf{x}, y')) - \underbrace{\mathbf{w}^{\top} \phi(\mathbf{x}, y)}_{\text{hope}}$$

In binary case:

$$\Rightarrow \min_{\mathbf{w}} \sum_{\mathbf{x}, y} \max\{\mathbf{0}, -y\mathbf{w}^{\top}\mathbf{x}\}\$$



Any thoughts about negative sampling?

$$\min_{\mathbf{w}} \sum_{m=1}^{M} \max_{y'} (\mathbf{w}^{\top} \phi(\mathbf{x}_m, y')) - \mathbf{w}^{\top} \phi(\mathbf{x}_m, y_m)$$

• Stochastic subgradient descent on the above is called the perceptron algorithm

• Pick i_t randomly from $\{1, \ldots, n\}$

•
$$\hat{y}_{i_t} = \arg \max_{y'} \mathbf{w} \phi(\mathbf{x}, y')$$

• $\mathbf{w} \leftarrow \mathbf{w} - \eta \left(\mathbf{w}^{\top} \phi(\mathbf{x}_{i_t}, \hat{y}_{i_t}) - \mathbf{w}^{\top} \phi(\mathbf{x}_{i_t}, y_{i_t}) \right)$

- Suppose that not all mistakes are equally bad
- E.g., false positives vs. false negatives in spam detection
- Let cost(y', y) quantify the "badness" of substituting y' for correct label y
- Intuition: estimate the scoring function so that score(y) − score(y') ∝ cost(y', y)

$$\left(\max_{y'}(\mathbf{w}^{ op}\phi(\mathbf{x},y'))+\operatorname{cost}(y,y')
ight)-\mathbf{w}^{ op}\phi(\mathbf{x},y)$$

• Text classification: many problems, all solved with supervised learners

- Lexicon features can provide problem-specific guidance
- Naive Bayes, log-linear, and linear SVM are all linear methods that tend to work reasonably well, with good features and smoothing/regularization
- Rumor: random forests are widely used in industry when performance matters more than interpretability
- Lots of papers about neural networks, though with hyper-parameter tuning applied fairly to linear models, the advantage is not clear (Yogatama et al. (2015))
- Lots of work on feature design

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Clustering

- Clustering is an unsupervised learning method
- Given items $\mathbf{x}_1, \ldots, \mathbf{x}_M \in \mathbb{R}^d$, the goal is to group them into reasonable clusters
- We also need a pairwise distance/similarity function between items, and sometimes the desired number of clusters
- When documents are represented by feature vectors, a commonly used similarity measure is the cosine similarity

$$sim(\mathbf{x}, \mathbf{x}') = cos(\mathbf{x}, \mathbf{x}') = \frac{\mathbf{x}^{\top} \mathbf{x}'}{||\mathbf{x}|| \cdot ||\mathbf{x}'||}$$

• This similarity has the nice property that document length is implicitly normalized (so that a long document can be similar to a short document)

K-Means Clustering

1 Randomly choose K centers μ_1, \ldots, μ_K

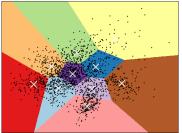
2 Repeat

3 Assign $\mathbf{x}_1, \ldots, \mathbf{x}_M$ to their nearest centers to obtain \hat{y}_m , respectively

4 Update
$$\mu_k = \frac{1}{\sum_m I(\hat{y}_m = k)} \sum_m \mathbf{x}_m I(\hat{y}_m = k)$$

5 Until the clusters no longer change

Step 3 is equivalent to creating a Voronoi diagram under the current centers K-means clustering on the digits dataset (PCA-reduced data)



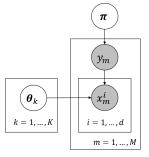
means clustering on the digits dataset (PCA-reduced data Centroids are marked with white cross

http://scikit-learn.org/stable/auto_examples/cluster/plot_kmeans_digits.html < _

- K-means clustering is sensitive to the initial cluster centers
- It is in fact an optimization problem with a lot of local optima
 - To be exact, k-means clustering is a special case of Gaussian Mixture Model (GMM) when the covariance of the Gaussian components tends to zero
- It is of course sensitive to k too
- Both should be chosen with care

Recall Naive Bayes Classifier: A Generative View

Naive Bayes from Class Conditional Unigram Model



Both y_m and $\mathbf{x}_m = (x_m^1, \dots, x_m^d)^T$ are observed variables; π and $\boldsymbol{\theta}_k$ are parameters • For m = 1, ..., M

- Choose $y_m \sim Multinomial(y_m|1, \pi)$
- Choose $N_m = \sum_j^d x_m^j \sim Poisson(\xi)$

• For
$$n = 1, ..., N_m$$

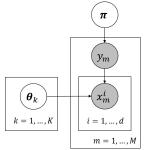
• Choose $v \sim Multinomial(v|1, \theta_{*|y_m}) = \prod_{j=1}^{d} (\theta_{*|y_m}^j)^{v=j}$

Alternative views

- Choose $\mathbf{x}_m \sim Multinomial(\mathbf{X}|N_m, \boldsymbol{\theta}_{*|y_m}) = \begin{pmatrix} N_m \\ \mathbf{x}_m \end{pmatrix} \prod_{j=1}^d (\theta_{*|y_m}^j)^{\mathbf{x}_m^j}$
- Choose $x_m^d \sim Binomial(X|N_m, \theta_{*|y_m}^j) = \begin{pmatrix} N_m \\ x_m^j \end{pmatrix} (\theta_{*|y_m}^j)^{x_m^j} (1 \theta_{*|y_m}^j)^{N_m x_m^j}$

Parameter Estimation (based on Multinomial)

Maximum likelihood of the training set:



$$\begin{aligned} \mathcal{J} &= \log \prod_{m=1}^{M} P_{\boldsymbol{\pi}, \{\boldsymbol{\theta}_k\}}(\mathbf{x}_m, y_m) \\ &= \sum_{m=1}^{M} \log P_{\boldsymbol{\pi}, \{\boldsymbol{\theta}_k\}}(\mathbf{x}_m, y_m) \\ &= \sum_{m=1}^{M} \log P(y_m | \boldsymbol{\pi}) P(\mathbf{x}_m | y_m, \boldsymbol{\theta}_{*|y_m}) \end{aligned}$$

We can formulate a constrained optimization problem

$$\begin{array}{ll} \max & \mathcal{J} \\ s.t. & \sum_{k=1}^{K} \pi_k = 1 \\ & \sum_{j=1}^{d} \theta_k^j = 1 (k = 1, \dots, K) \end{array}$$

Both y_m and $\mathbf{x}_m = x_m^1, \dots, \mathbf{x}_m^d$ are observed variables; π and $\boldsymbol{\theta}_k$ are parameters

It's easy to solve with Lagrange multiplier and arrive at: $|\{y_m=k\}|$

$$\pi_{k} = \frac{\left[\left\{ y_{m} = k \right\} \right]}{M}$$
$$\theta_{k}^{j} = \frac{\sum_{m, y_{m} = k} x_{m}^{j}}{\sum_{m, y_{m} = k} \sum_{j=1}^{d} x_{m}^{j}}$$

What if the documents are not labeled?

In naive Bayes, both y_m and $\mathbf{x}_m = (x_m^1, \dots, x_m^d)^T$ are observed variables; π and θ_k are parameters

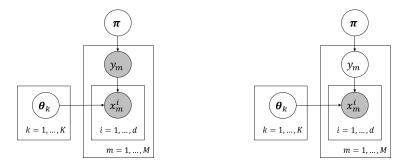


Figure: Native Bayes

Figure: Mixture Model

However, in clustering problems, y_m is not observed (labeled before feeding into machine learning algorithm)

Expectation Maximization (EM) Algorithm

- EM might look like a heuristic method. However, it is not.
- EM is guaranteed to find a local optimum of data log likelihood
- Recall if we have complete data set {x_m, y_m}^M_{m=1} and denote parameter set as Θ = {π, {θ_k}}, the likelihood estimation of native Bayes is

$$\mathcal{J}_{NB}(\Theta) = \log \prod_{m=1}^{M} P_{\boldsymbol{\pi}, \{\boldsymbol{\theta}_k\}}(\mathbf{x}_m, y_m) = \log P(\{\mathbf{x}_m, y_m\}_{m=1}^{M} | \Theta)$$

- However, now $\{y_m\}_{m=1}^M$ are not observed (labeled), so we treat them as hidden variables
- We instead maximize the marginal log likelihood:

$$\mathcal{J}(\Theta) = \log P(\{\mathbf{x}_m\}_{m=1}^M | \Theta)$$

Maximizing the Marginal Log Likelihood

We optimize following objective function:

$$\mathcal{J}(\Theta) = \log P(\{\mathbf{x}_m\}_{m=1}^M | \Theta)$$

= $\sum_{m=1}^M \log P(\mathbf{x}_m | \Theta)$
= $\sum_{m=1}^M \log \sum_{y=1}^K P(\mathbf{x}_m, y | \Theta)$
= $\sum_{m=1}^M \log \sum_{y=1}^K P(y | \Theta) P(\mathbf{x}_m | y, \Theta)$
= $\sum_{m=1}^M \log \sum_{y=1}^K P(y | \pi) P(\mathbf{x}_m | y, \theta_{*|y})$

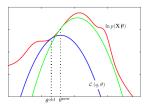
Compared to supervised learning:

$$\begin{aligned} \mathcal{J}_{NB}(\Theta) &= \log \prod_{m=1}^{M} P_{\boldsymbol{\pi}, \{\boldsymbol{\theta}_k\}}(\mathbf{x}_m, y_m) \\ &= \sum_{m=1}^{M} \log P_{\boldsymbol{\pi}, \{\boldsymbol{\theta}_k\}}(\mathbf{x}_m, y_m) \\ &= \sum_{m=1}^{M} \log P(y_m | \boldsymbol{\pi}) P(\mathbf{x}_m | y_m, \boldsymbol{\theta}_{*|y_m}) \end{aligned}$$

- It's more complicated with a summation inside the log!
- If we try to maximize the marginal log likelihood by setting the gradient to zero, we will find that there is no longer a nice closed form solution, unlike the joint log likelihood with complete data

EM Algorithm: General Idea

- EM is an iterative procedure to maximize the marginal log likelihood $\mathcal{J}(\Theta)$
- It constructs a concave, easy-to-optimize lower bound $\mathcal{J}(\Theta) \geq Q(\Theta, \Theta^t)$, where Θ is the variable and Θ^t is the previous, fixed, parameter
- The lower bound has an interesting property $Q(\Theta^t, \Theta^t) = \mathcal{J}(\Theta^t)$
- Therefore the new parameter Θ^{t+1} that maximizes $Q(\Theta^t, \Theta)$ is guaranteed to have $Q \ge \mathcal{J}(\Theta^t)$. Since Q lower bounds \mathcal{J} , we have $\mathcal{J}(\Theta^{t+1}) \ge \mathcal{J}(\Theta^t)$

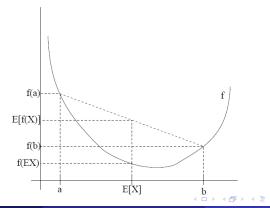


Lower Bound $Q(\Theta, \Theta^t)$

• The lower bound is obtained via Jensens inequality (concavity of log function)

$$\log \sum_i P_i f_i(x) \ge \sum_i P_i \log f_i(x)$$

which holds if the p_i 's form a probability distribution



Lower Bound $Q(\Theta, \Theta^t)$ (Cont'd)

The lower bound is obtained via Jensens inequality (concavity of log function)

$$\log \sum_i P_i f_i(x) \ge \sum_i P_i \log f_i(x)$$

which holds if the p_i 's form a probability distribution

• Then the lower bound can be derived:

$$\begin{aligned} \mathcal{J}(\Theta^{t}) &= \sum_{m=1}^{M} \log \sum_{y=1}^{K} P(\mathbf{x}_{m}, y | \Theta^{t}) \\ &= \sum_{m=1}^{M} \log \sum_{y=1}^{K} q_{\mathbf{x}_{m}, y}(\Theta) \frac{P(\mathbf{x}_{m}, y | \Theta^{t})}{q_{\mathbf{x}_{m}, y}(\Theta)} \\ &\geq \sum_{m=1}^{M} \sum_{y=1}^{K} q_{\mathbf{x}_{m}, y}(\Theta) \log \frac{P(\mathbf{x}_{m}, y | \Theta^{t})}{q_{\mathbf{x}_{m}, y}(\Theta)} \\ &\doteq Q(\Theta, \Theta^{t}) \end{aligned}$$

where $\sum_{y=1}^{K} q_{\mathbf{x}_m, y}(\Theta) = 1$ is some distribution



$$\sum_{m=1}^{M} \log \sum_{y=1}^{K} q_{\mathbf{x}_{m}, y}(\Theta) \frac{P(\mathbf{x}_{m}, y | \Theta^{t})}{q_{\mathbf{x}_{m}, y}(\Theta)} \geq \sum_{m=1}^{M} \sum_{y=1}^{K} q_{\mathbf{x}_{m}, y}(\Theta) \log \frac{P(\mathbf{x}_{m}, y | \Theta^{t})}{q_{\mathbf{x}_{m}, y}(\Theta)}$$

- To make the bound tight for a particular value of Θ, we need for the step involving Jensens inequality in our derivation above to hold with equality
- For this to be true, we know it is sufficient that the expectation be taken over a constant-valued random variable $\frac{P(\mathbf{x}_m, y | \Theta^t)}{a_{\mathbf{x}_m} \ v(\Theta)} = c$
- This is easily done by choosing $q_{\mathbf{x}_m,y}(\Theta) \propto P(\mathbf{x}_m,y|\Theta^t)$
- Since $\sum_{y=1}^{K} q_{\mathbf{x}_m, y}(\Theta) = 1$, we have (considered as E-step)

$$q_{\mathbf{x}_m, y}(\Theta) = \frac{P(\mathbf{x}_m, y | \Theta^t)}{\sum_{y=1}^{K} P(\mathbf{x}_m, y | \Theta^t)} = P(y | \mathbf{x}_m, \Theta^t)$$

• The equation holds in the inequality iff $q_{\mathbf{x}_m, y} = P(y | \mathbf{x}_m, \Theta^t)$

M-step

• In M-step, we maximize the lower bound

$$\begin{aligned} Q(\Theta^t, \Theta) &= \sum_{m=1}^{M} \sum_{y=1}^{K} q_{\mathbf{x}_m, y} \log \frac{P(\mathbf{x}_m, y|\Theta)}{q_{\mathbf{x}_m, y}} \\ &= \sum_{m=1}^{M} \sum_{y=1}^{K} q_{\mathbf{x}_m, y} \log \frac{P(y_m | \boldsymbol{\pi}) P(\mathbf{x}_m | y_m, \boldsymbol{\theta}_{*|y_m})}{q_{\mathbf{x}_m, y}} \end{aligned}$$

• Now we can set the gradient of Q w.r.t. π and θ_k 's to zero and obtain a closed form solution

$$\pi_{k} = \frac{\sum_{m} \mathbf{q}_{\mathbf{x}_{m,y}}}{M} \theta_{k}^{j} = \frac{\sum_{m} \mathbf{q}_{\mathbf{x}_{m,y}} \mathbf{x}_{m}^{j}}{\sum_{m} \sum_{j=1}^{d} \mathbf{q}_{\mathbf{x}_{m,y}} \mathbf{x}_{m}^{j}}$$

• Compared to naive Bayes:

$$\pi_{k} = \frac{|\{y_{m} = k\}|}{M}$$
$$\theta_{k}^{j} = \frac{\sum_{m, y_{m} = k} x_{m}^{j}}{\sum_{m, y_{m} = k} \sum_{j=1}^{d} x_{m}^{j}}$$

EM Algorithm

- Repeat
 - E-step: compute posterior of hidden variables

$$q_{\mathbf{x}_m,y} = P(y|\mathbf{x}_m,\Theta)$$

• M-step: parameter estimation by maximizing the lower bound

$$\begin{aligned} \pi_{k} &= \frac{\sum_{m} q_{\mathbf{x}_{m}, y}}{M} \\ \theta_{k}^{j} &= \frac{\sum_{m} q_{\mathbf{x}_{m}, y} x_{m}^{j}}{\sum_{m} \sum_{j=1}^{d} q_{\mathbf{x}_{m}, y} x_{m}^{j}} \end{aligned}$$

• Until the convergence of the objective function

- Randomly choose K centers μ_1, \ldots, μ_K
- Repeat
 - Assign x₁,..., x_M to their nearest centers to obtain ŷ_m, respectively
 - Update $\mu_k = \frac{1}{\sum_m I(\hat{y}_m = k)} \sum_m \mathbf{x}_m I(\hat{y}_m = k)$
- Until the clusters no longer change

In practice, K-means is cheaper. We can run multiple times to find good initialization to mixture models.

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Convergence of EM Algorithm

• E-step: With $q_{\mathbf{x}_m, y}(\Theta) = P(y|\mathbf{x}_m, \Theta^t)$, the equation holds, which leads

 $Q(\Theta^t,\Theta^t)=\mathcal{J}(\Theta^t)$

• M-step: Since Θ^{t+1} maximizes $Q(\Theta^t, \Theta)$, we have

$$Q(\Theta^t,\Theta^{t+1})\geq Q(\Theta^t,\Theta^t)=\mathcal{J}(\Theta^t)$$

• On the other hand, Q is lower bound of \mathcal{J} , we have:

$$\mathcal{J}(\Theta^{t+1}) \geq Q(\Theta^t,\Theta^{t+1}) \geq Q(\Theta^t,\Theta^t) = \mathcal{J}(\Theta^t)$$

- This shows EM algorithm always increase the objective function (log likelihood)
- By iterating, we arrive at a local maximum of it

- EM is general and applies to joint probability models whenever some random variables are missing
- EM is advantageous when the marginal is difficult to optimize, but the joint is easy
- To be general, consider a joint distribution P(X, Z|Θ), where X is the collection of observed variables, and Z unobserved variables
- The quantity we want to maximize is the marginal log likelihood

$$\mathcal{J}(\Theta) = \log P(X|\Theta) = \log \sum_{Z} P(X, Z|\Theta)$$

which we assume difficult to optimize

A More General View of EM (Cont'd)

• One can introduce an arbitrary distribution over hidden variables Q(Z)

$$\begin{aligned} \mathcal{J}(\Theta) &= \log P(X|\Theta) = \log \sum_{Z} P(X, Z|\Theta) \\ &= \sum_{Z} Q(Z) \log P(X|\Theta) \\ &= \sum_{Z} Q(Z) \log \frac{P(X|\Theta)Q(Z)P(X,Z|\Theta)}{P(X,Z|\Theta)Q(Z)} \\ &= \sum_{Z} Q(Z) \log \frac{P(X,Z|\Theta)}{Q(Z)} + \sum_{Z} Q(Z) \log \frac{P(X|\Theta)Q(Z)}{P(X,Z|\Theta)} \\ &= \sum_{Z} Q(Z) \log \frac{P(X,Z|\Theta)}{Q(Z)} + \sum_{Z} Q(Z) \log \frac{Q(Z)}{P(Z|X,\Theta)} \\ &= F(Q,\Theta) + KL[Q(Z)||P(Z|X,\Theta)] \end{aligned}$$

• Note $F(Q, \Theta)$ is the right hand side of Jensen's inequality

- If KL > 0, $F(Q, \Theta)$ is a lower bound of $\mathcal{J}(\Theta)$
- First consider the maximization of F on Q with Θ^t fixed
 - F(Q,Θ) is maximized by Q(Z) = P(Z|X,Θ^t) since J(Θ) is fixed and KL attends its minimum zero (E-Step)

• Next consider the maximization of F on Θ with Q fixed as above

• Note in this case $F(Q, \Theta) = Q(\Theta^t, \Theta)$ (M-Step)

- Generalized EM (GEM) finds Θ that improves, but not necessarily maximizes, $F(Q, \Theta) = Q(\Theta, \Theta^t)$ in the M-step. This is useful when the exact M-step is difficult to carry out. Since this is still coordinate ascent, GEM can find a local optimum.
- Stochastic EM: The E-step is computed with Monte Carlo sampling. This introduces randomness into the optimization, but asymptotically it will converge to a local optimum.
- Variational EM: Q(Z) is restricted to some easy-to-compute subset of distributions, for example the fully factorized distributions Q(Z) = ∏_i Q(z_i). In general P(Z|X, Θ), which might be intractable to compute, will not be in this subset. There is no longer guarantee that variational EM will find a local optimum.

Yogatama, D., Kong, L., and Smith, N. A. (2015). Bayesian optimization of text representations. In *EMNLP*, pages 2100–2105.

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