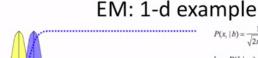
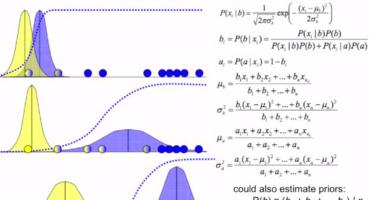
# Expectation-Maximization Algorithm





$$P(b) = (b_1 + b_2 + ... b_n) / n$$
  
 $P(a) = 1 - P(b)$ 

## **Expectation-Maximization Algorithm**

A general technique for finding maximum likelihood estimators in latent variable models is the expectation-maximization (EM) algorithm.

E-Step
 Estimate the missing variables in the dataset.

 Calculate the expectation of complete-data log-likelihood:

$$Q(\theta|\theta^{(t)}) := E[\log P(y_{obs}, y_{mis}|\theta)|y_{obs}, \theta^{(t)}]$$

• M-Step Maximize the parameters of the model in the presence of the data. Find  $\theta^{(t+1)}$  by maximizing  $Q(\theta|\theta^{(t)})$ 

$$\theta^{(t+1)} := \operatorname*{argmax}_{\theta} \mathcal{Q}(\theta|\theta^{(t)})$$

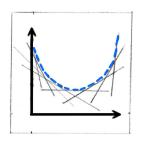
Iterate the above 2 steps until convergence.

# Why EM works?

- Lemma 1: Jensen's inequality
- Proposition 1: Ascent property of EM
- Theorem 1: Convergence property of EM

### Convex function

Upper envelop and supporting lines



$$g(x) \ge a_0x + b_0$$
;  $g(x_0) = a_0x_0 + b_0$ .

Supporting line at  $x_0$  touches g(x) at  $x_0$ , but below g(x) at other places.

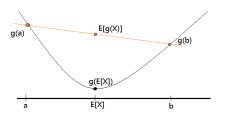
## Jensen's inequality

$$P(X = a) = P(X = b) = 1/2.$$

$$E(X) = (a+b)/2, g(E(X)) = g((a+b/2).$$

$$E(g(X)) = (g(a) + g(b))/2.$$

 $E(g(X)) \ge g(E(X))$ . Note: g(x) is convex



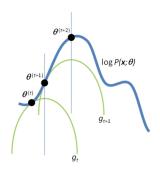
$$x_0 = E(X)$$
.  $g(x_0) = a_0x_0 + b_0$  (supporting line at  $x_0$ )  $g(x) \ge a_0x + b_0$ .  $E(g(X)) \ge E(a_0X + b_0) = a_0E(X) + b_0 = a_0x_0 + b_0 = g(E(X))$ .

## Ascent property of EM

Let  $\ell(\theta|Y_{obs}) := \log P(Y_{obs}|\theta)$ , which is the observed-data log-likelihood. Then the EM iterations satisfy

$$\ell(\theta^{(t+1)}|Y_{obs}) \ge \ell(\theta^{(t)}|Y_{obs})$$

# EM Algorithm: Evidence Lower Bound and Convergence



#### Theorem

Under some conditions, the sequence  $\{\theta^{(t)}\}$  defined by the EM iteratations converges to a stationary point of the observed-data  $\log$ -likelihood  $\log(P(y_{obs}|\theta))$ .

### Revisit the mixture model

- Observed variables  $X = (X_1, X_2, ..., X_n)$ An observation of X is called an incomplete data set
- Unobserved variables Z = (Z<sub>1</sub>, Z<sub>2</sub>,..., Z<sub>k</sub>).
   (An observation (X,Z) is called a complete data set, but we never have a complete dataset)
- Parameters  $\theta = (\pi, \mu, \sigma)$ Cluster probabilities:  $\pi = (\pi_1, \dots, \pi_k)$ Cluster means:  $\mu = (\mu_1, \dots, \mu_k)$ Cluster standard deviation:  $\sigma = (\sigma_1, \dots, \sigma_k)$
- Complete data likelihood  $P(X = i, Z = j | \theta) = \pi_j N(x_i | \mu_j, \sigma_j^2)$

### Revisit the mixture model

- **1** Choose initial  $\theta^{old} = (\pi^0, \mu^0, \sigma^0)$
- ② Expectation step:

$$\log(P(X=i,Z=j|\theta)) = \log(\pi_j) + \log(N(x_i|\mu_j,\sigma_j^2))$$

$$p(z = j | x = i, \theta^{old}) = \gamma_i^j = \frac{\pi_j^{old} N(X_i | \mu_j^{old}, \sigma_{j,old}^2)}{\sum_{c=1}^k \pi_c^{old} N(X_i | \mu_c^{old}, \sigma_{c,old}^2)}$$

$$Q(\theta, \theta^{old}) = \sum_{i=1}^{n} \sum_{j=1}^{k} \gamma_{i}^{j} [\log(\pi_{j}) + \log N(x_{i}|\mu_{j}, \sigma_{j}^{2})]$$

Maximization step:

$$\theta^{\textit{new}} = \operatorname*{argmax}_{\theta} \textit{Q}(\theta, \theta^{\textit{old}})$$

• Let  $\theta^{old} = \theta^{new}$ , go to step 2, until convergence

# Maximization Step

• 
$$\pi_j^{\text{new}} = \frac{\sum_{i=1}^n \gamma_i^j}{n}$$

$$\bullet \ (n_j^{new} = n * \pi_j^{new})$$

$$\bullet \ \mu_j^{\text{new}} = \frac{\Sigma_{i=1}^n \gamma_i^j x_i}{n_i^{\text{new}}}$$

• 
$$\sigma_{j,\text{new}}^2 = \frac{1}{n_j^{\text{new}}} \sum_{i=1}^n \gamma_i^j (x_i - \mu_j^{\text{new}})^2$$

for each  $j = 1, \ldots, k$ 

### More examples

- Bivariate binary data
- Multinomial distribution with cell probabilities
- Coin flipping (A paper from Nature Biotechnology)