

# Speech Features and Speaker Classification

**CSC401/2511 – Natural Language Computing – Winter 2024**

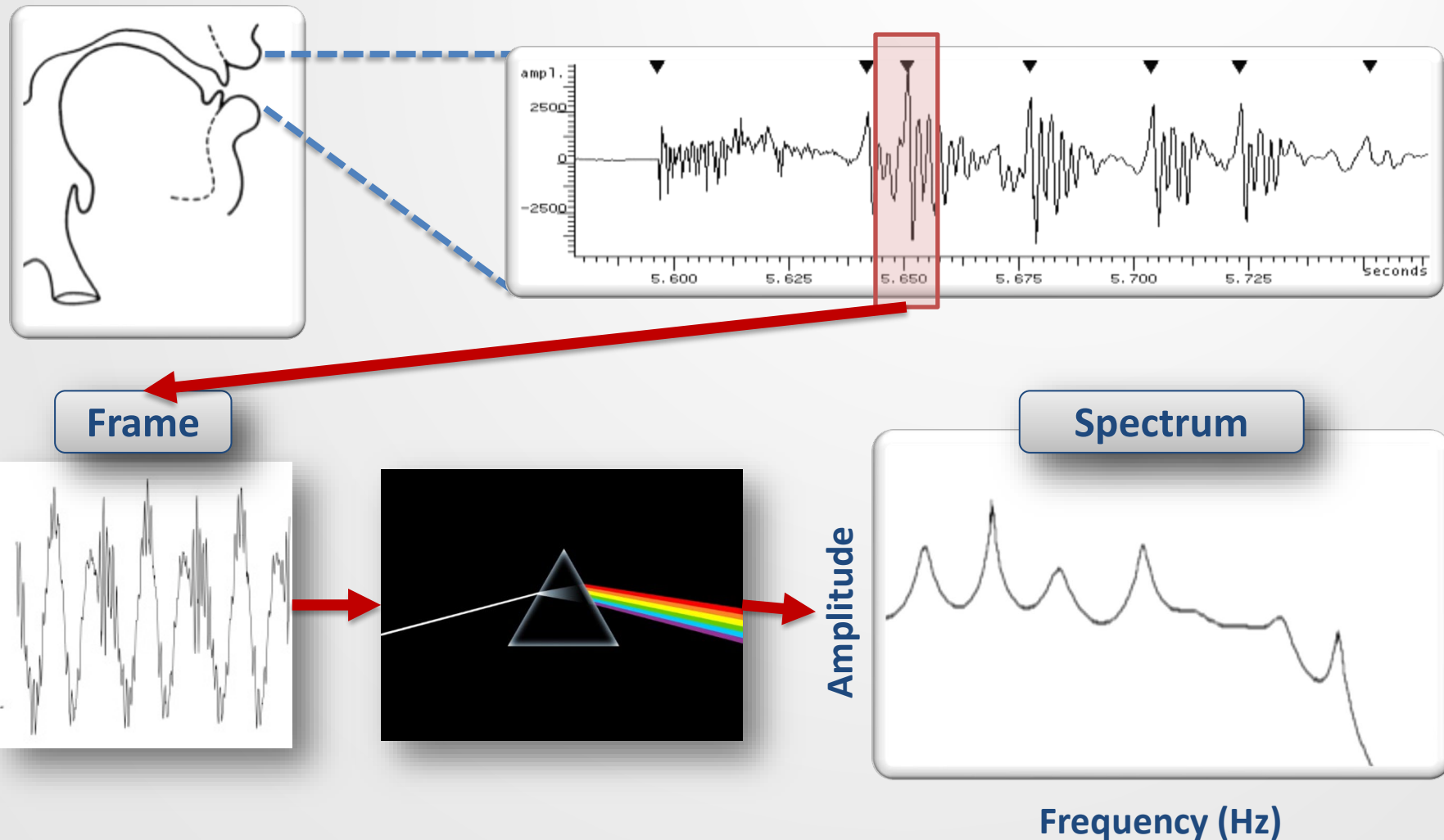
**Lecture 9**

# Contents

- Today we will
  - Define some common feature vectors for speech processing
  - Use them as input to a GMM-based speaker classification system
- All of this is part of A3

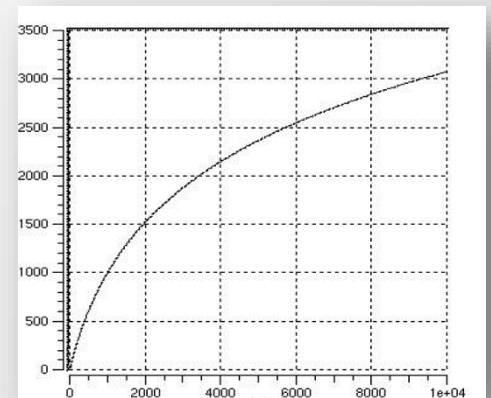
# SPEECH FEATURES

# Recall the spectrogram pipeline



# Problems with spectrograms

- As input to speech systems, spectrograms are...
- **Too big**
  - The discrete signal is usually 16,000 samps/sec
  - 100 frames/sec x 400 samps/frame = 40,000 samps/sec!
- **Too linear**
  - Pitch perception is log-linear (recall Mels)
  - Lots of coefficients wasted on high frequencies
- **Too entangled**
  - Speaker and phoneme info is correlated

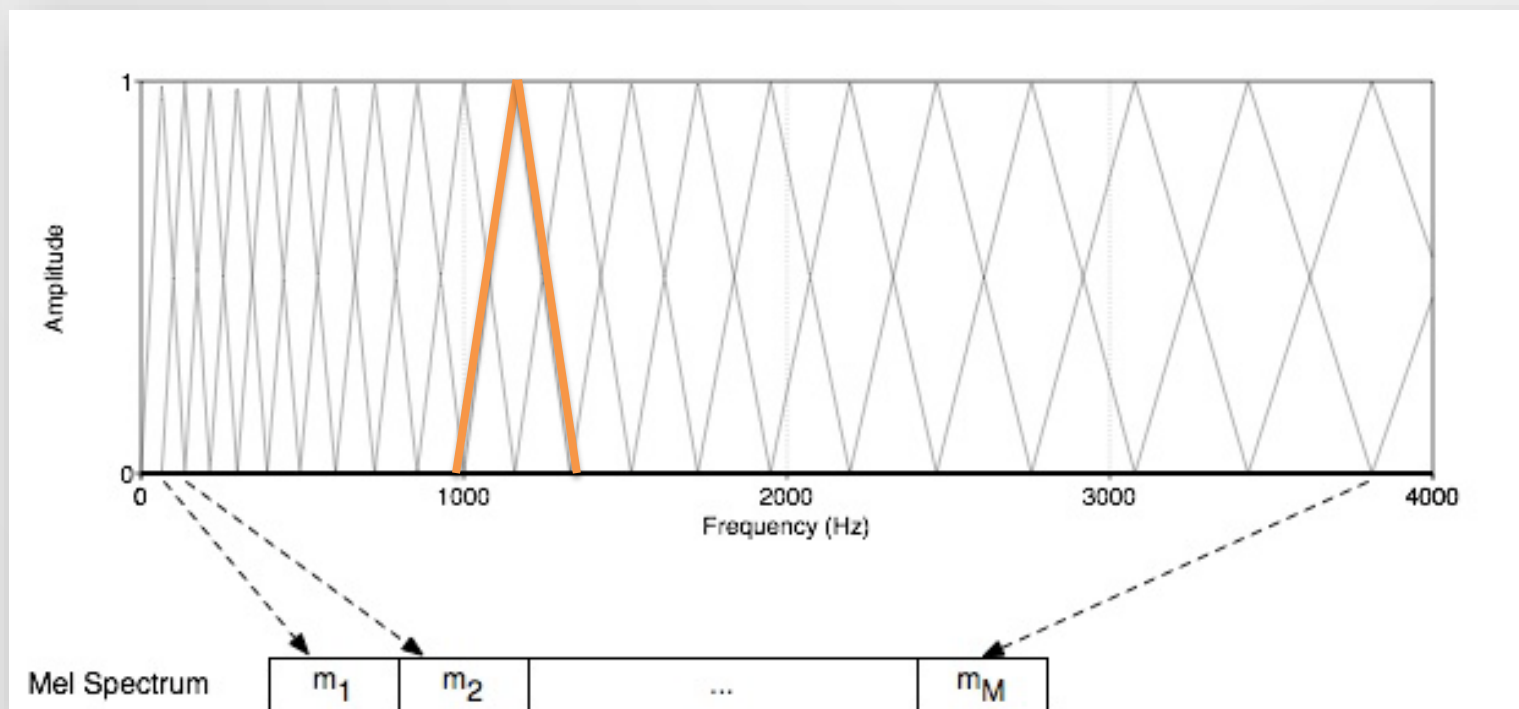


# Filtering

- To reduce the size of the spectra, we **filter** it with **filters** from a **filter bank**
- Each filter is a signal whose spectrum  $F_m \in \mathbb{R}^N$  picks out small a range (or **band**) of frequencies
- The bands of the  $M$  filters are overlapping and span the spectrum
- A **filter coefficient** is computed as the **log** of the dot product of the **magnitude** of the frame  $X_t$  and filter  $F_m$  spectra:
$$c_{t,m} = \log \sum_{n=1}^N |X_t|[n]|F_m|[n]$$
- If there are  $T$  frames, this gives us a real-valued feature matrix of size  $T \times M$ 
  - $M = 40$  is a lot smaller than 400!

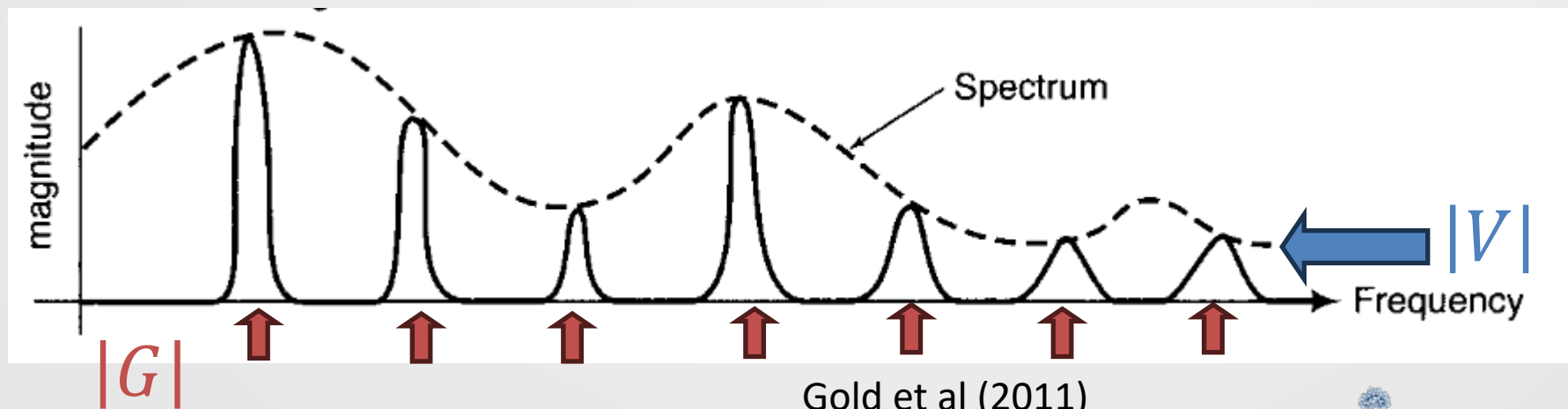
# The mel-scale filter bank

- The mel-scale triangular overlapping filter bank, or **f-bank**, is a popular choice
- The filter's vertices are arranged along the mel-scale
  - Ascending frequency = wider bands



# The source-filter model

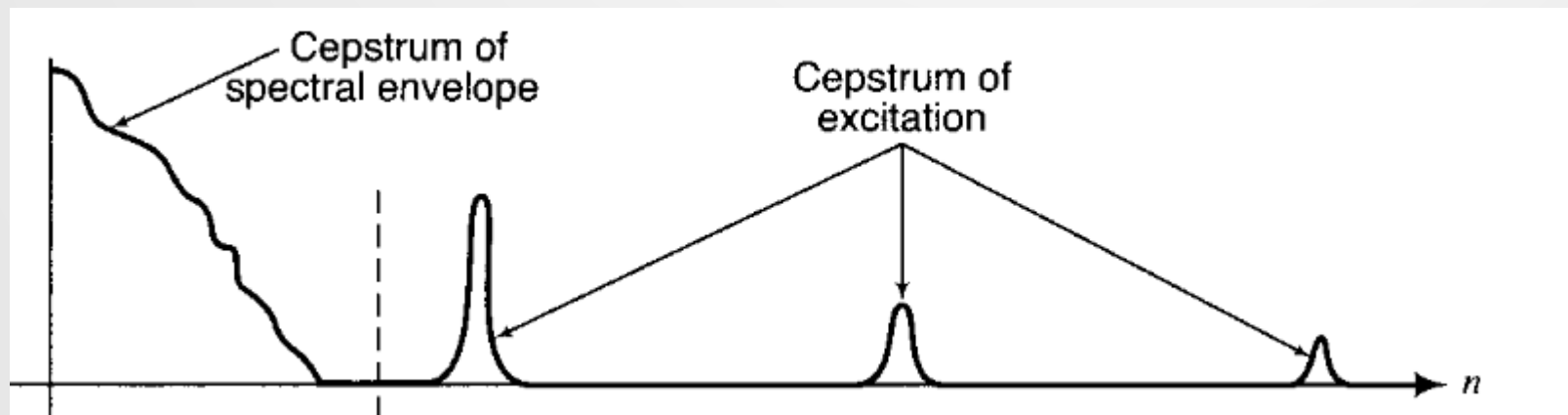
- In vowels, the sound signal emitted from the glottis  $g$  is filtered by the vocal tract  $v$
- The **source-filter model** of speech assumes
$$|X[n]| = |G[n]||V[n]|$$
- $|V|$  is responsible for the smooth shape (envelope)
- $|G|$  is responsible for all the bumps (F0 harmonics)





# The cepstrum

- We can get at  $|V|$  by computing the **cepstrum**  $\hat{x}$
- The cepstrum is  $\log|X|$  transformed by the inverse DFT
- Because  $\log|X| = \log|G| + \log|V|$ , and  $DFT^{-1}$  is linear
$$\hat{x}[n] = \hat{g}[n] + \hat{v}[n]$$
- $DFT^{-1} \approx DFT$ , so  $\hat{x}$  is like the spectrum of  $\log|X|$
- $|V|$  is slower-moving than  $|G|$ , so  $\hat{v}[n]$  is higher for lower  $n$  (lower frequency of frequency)



Gold et al (2011)

# Mel-Frequency Cepstral Coefficients

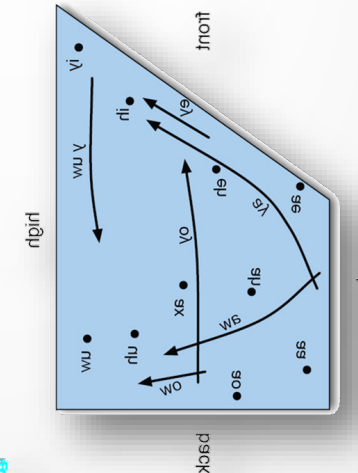
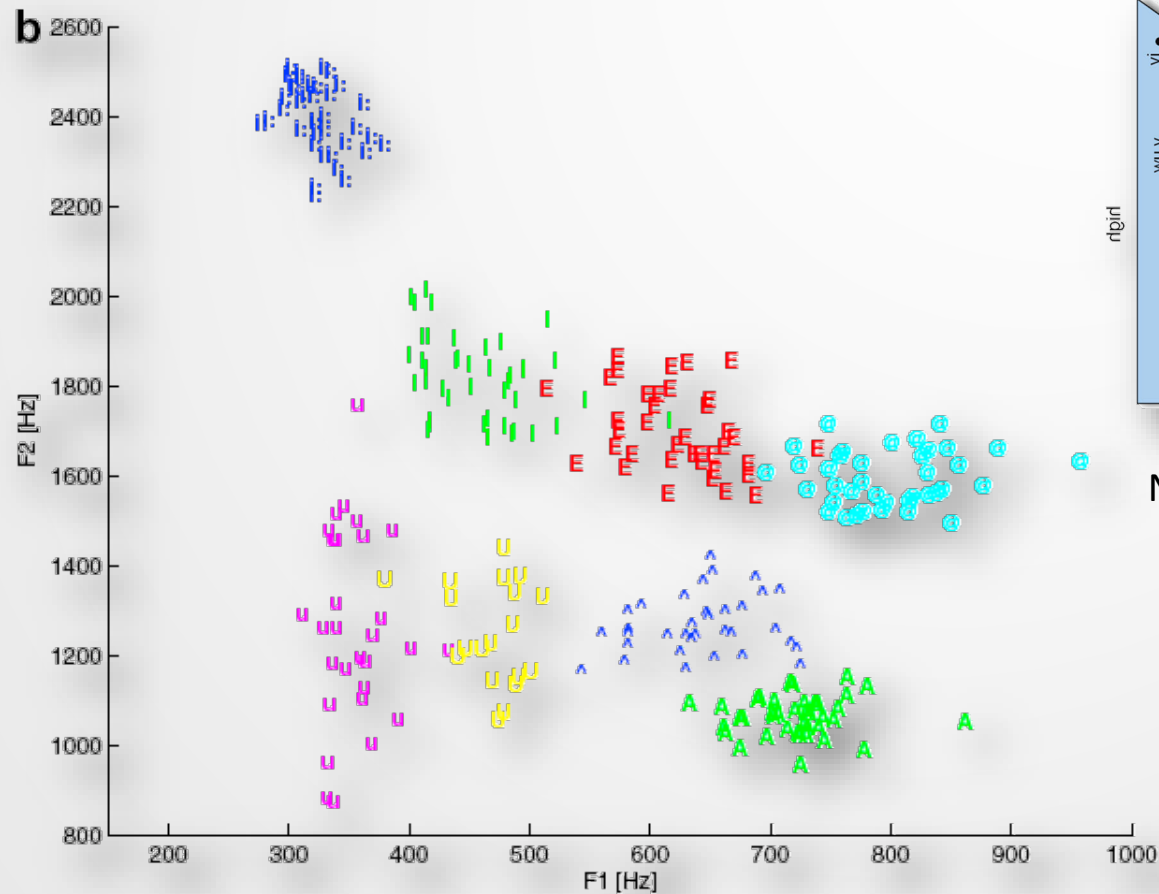
- **MFCCs** are the coefficients of the cepstrum of F-bank coefficients
- Altogether



- MFCCs are useful for models which can't handle speaker correlations themselves, like (diagonal) GMMs
- F-banks are better for those which can, like NNs

# GAUSSIAN MIXTURES

# Classifying speech sounds



Note: The vowel trapezoid's dimensions were physical

- Speech sounds can cluster. This graph shows vowels, each in their own colour, according to the 1<sup>st</sup> two formants.

# Classify speakers by cluster attributes

- Similarly, all of the speech produced by one **speaker** will cluster differently in the **Mel space** than speech from another speaker.
  - We can  $\therefore$  decide if a given observation comes from one speaker or another.

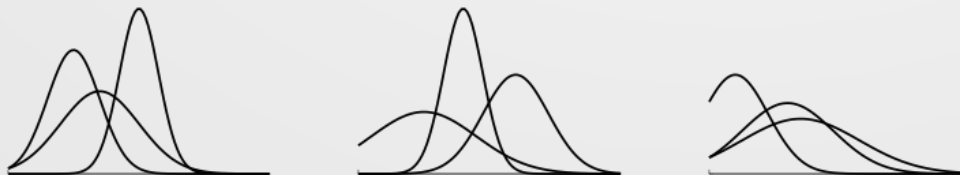
		Time, $t$			
		0	1	...	T
MFCC	1			...	
	2			...	
	3			...	
	...	...	...	...	...
	42			...	

Observation matrix

$$P(\text{orange bar} \mid \text{woman on phone}) > P(\text{orange bar} \mid \text{man in uniform})$$

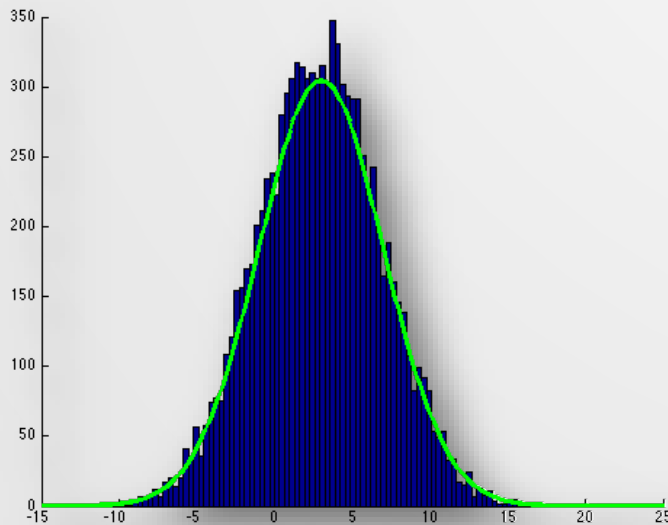
# Speaker classification

- **Speaker classification:**  $n$ . picking the most likely speaker among several speakers given only acoustics.
- Each **speaker** will produce speech according to **different** probability distributions.
  - We train a statistical model, given annotated data (mapping utterances to speakers).
  - We choose the speaker whose model gives the highest probability for an observation.



# Fitting continuous distributions

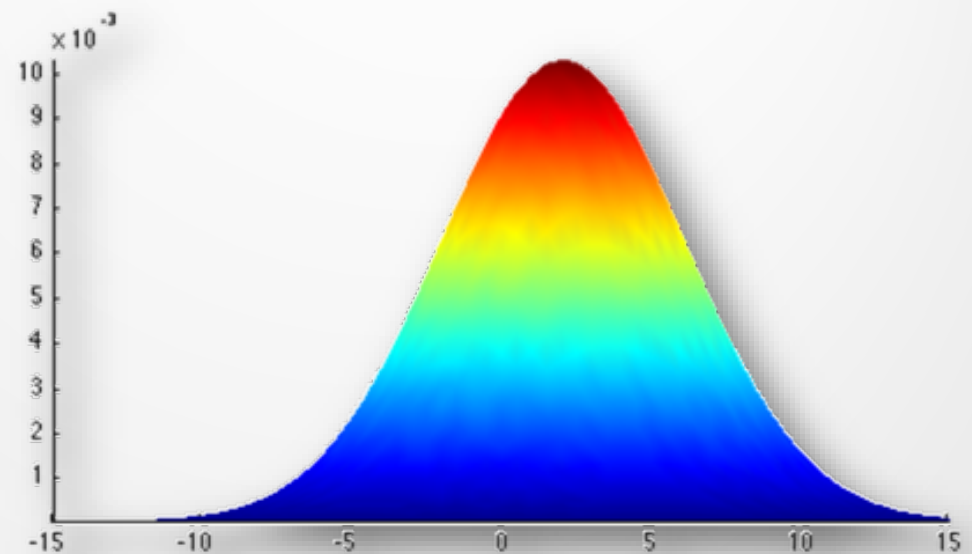
- Since we are operating with **continuous** variables, we need to **fit continuous probability** functions to a **discrete number** of observations.
- If we *assume* the 1-dimensional data in **this histogram** is Normally distributed, we can fit a continuous Gaussian function simply in terms of the mean  $\mu$  and variance  $\sigma^2$ .



# Univariate (1D) Gaussians

- Also known as **Normal** distributions,  $N(\mu, \sigma)$

- $$P(x; \mu, \sigma) = \frac{\exp\left(-\frac{(x-\mu)^2}{2\sigma^2}\right)}{\sqrt{2\pi}\sigma}$$



- The parameters we can modify are  $\theta = \langle \mu, \sigma^2 \rangle$ 
  - $\mu = E(x) = \int x \cdot P(x)dx$  (**mean**)
  - $\sigma^2 = E((x - \mu)^2) = \int (x - \mu)^2 P(x)dx$  (**variance**)

*But we don't have samples for all  $x$ ...*



# Maximum likelihood estimation

- Given data  $X = \{x_1, x_2, \dots, x_n\}$ , MLE produces an estimate of the parameters  $\hat{\theta}$  by maximizing the **likelihood**,  $L(X, \theta)$ :

$$\hat{\theta} = \underset{\theta}{\operatorname{argmax}} L(X, \theta)$$

where  $L(X, \theta) = P(X; \theta) = \prod_{i=1}^n P(x_i; \theta)$ .

- Since  $L(X, \theta)$  provides a **surface** over all  $\theta$ , in order to find the **highest likelihood**, we look at the derivative

$$\frac{\delta}{\delta \theta} L(X, \theta) = 0$$

to see **at which point** the likelihood **stops growing**.


# MLE with univariate Gaussians


- Estimate  $\mu$ :

$$L(X, \mu) = P(X; \mu) = \prod_{i=1}^n P(x_i; \theta) = \prod_{i=1}^n \frac{\exp\left(-\frac{(x_i - \mu)^2}{2\sigma^2}\right)}{\sqrt{2\pi}\sigma}$$

$$\log L(X, \mu) = -\frac{\sum_i (x_i - \mu)^2}{2\sigma^2} - n \log(\sqrt{2\pi}\sigma)$$

$$\frac{\partial}{\partial \mu} \log L(X, \mu) = \frac{\sum_i (x_i - \mu)}{\sigma^2} = 0$$

$$\mu = \frac{\sum_i x_i}{n}$$


- Similarly,  $\sigma^2 = \frac{\sum_i (x_i - \mu)^2}{n}$  

# Multivariate Gaussians

- When data is  **$d$ -dimensional**, the input variable is

$$\vec{x} = \langle x[1], x[2], \dots, x[d] \rangle$$

the **mean** is

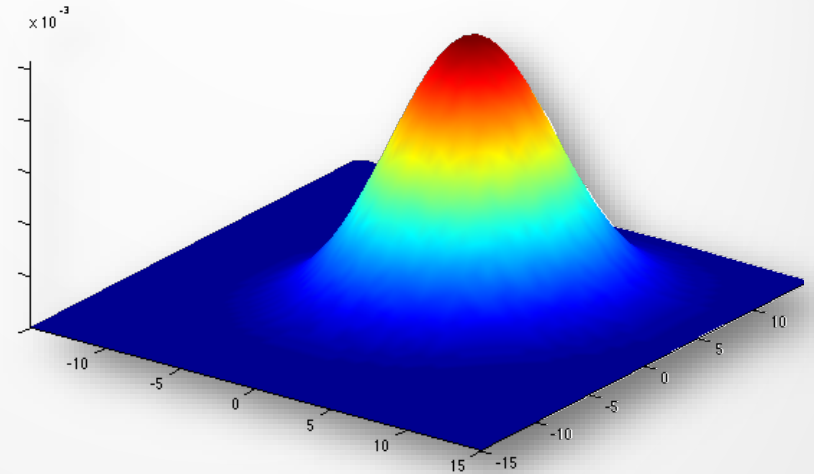
$$\vec{\mu} = E(\vec{x}) = \langle \mu[1], \mu[2], \dots, \mu[d] \rangle$$

the **covariance matrix** is

$$\Sigma[i, j] = E(x[i]x[j]) - \mu[i]\mu[j]$$

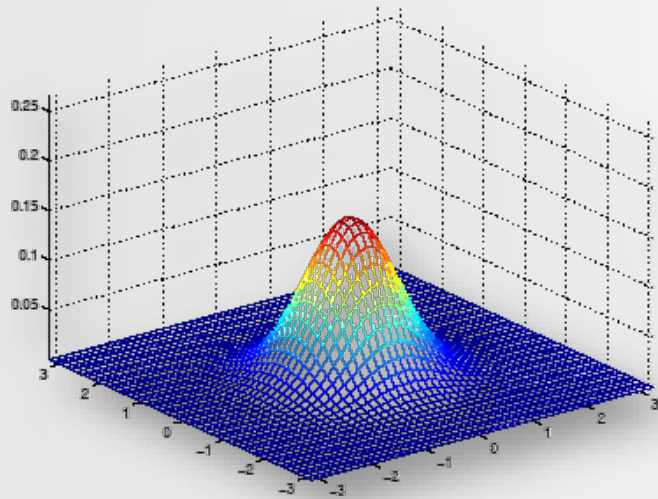
and

$$P(\vec{x}) = \frac{\exp\left(-\frac{(\vec{x} - \vec{\mu})^T \Sigma^{-1} (\vec{x} - \vec{\mu})}{2}\right)}{(2\pi)^{\frac{d}{2}} |\Sigma|^{\frac{1}{2}}}$$

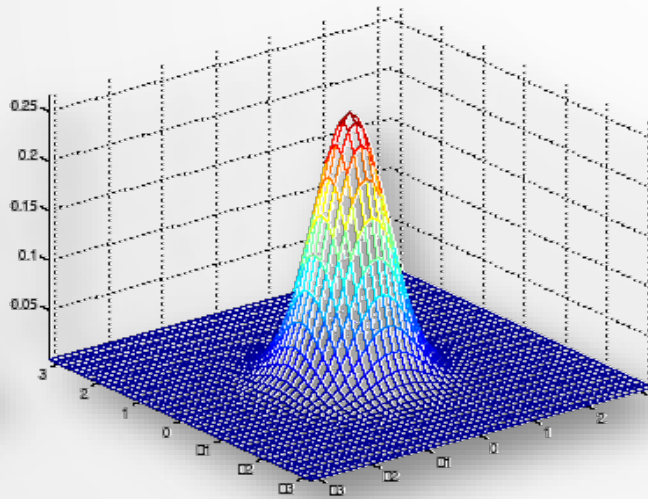


$A^T$  is the **transpose** of  $A$   
 $A^{-1}$  is the **inverse** of  $A$   
 $|A|$  is the **determinant** of  $A$

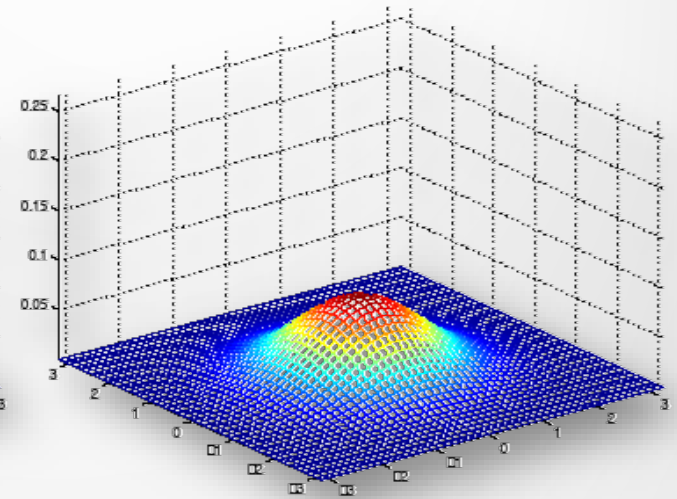
# Intuitions of covariance



$$\mu = [0 \ 0]$$
$$\Sigma = \mathbf{I}$$



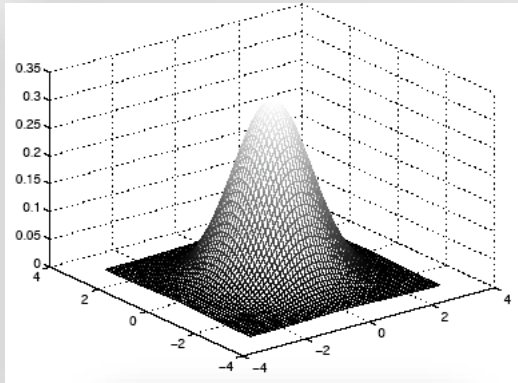
$$\mu = [0 \ 0]$$
$$\Sigma = 0.6\mathbf{I}$$



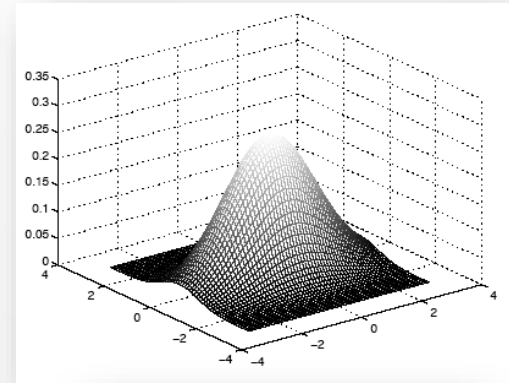
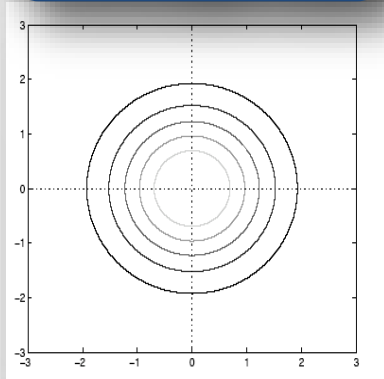
$$\mu = [0 \ 0]$$
$$\Sigma = 2.0\mathbf{I}$$

- As values in  $\Sigma$  become larger, the Gaussian spreads out.
- ( $\mathbf{I}$  is the identity matrix)

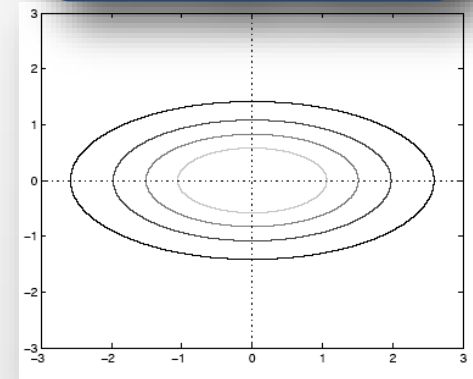
# Intuitions of covariance



$$\Sigma = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}$$



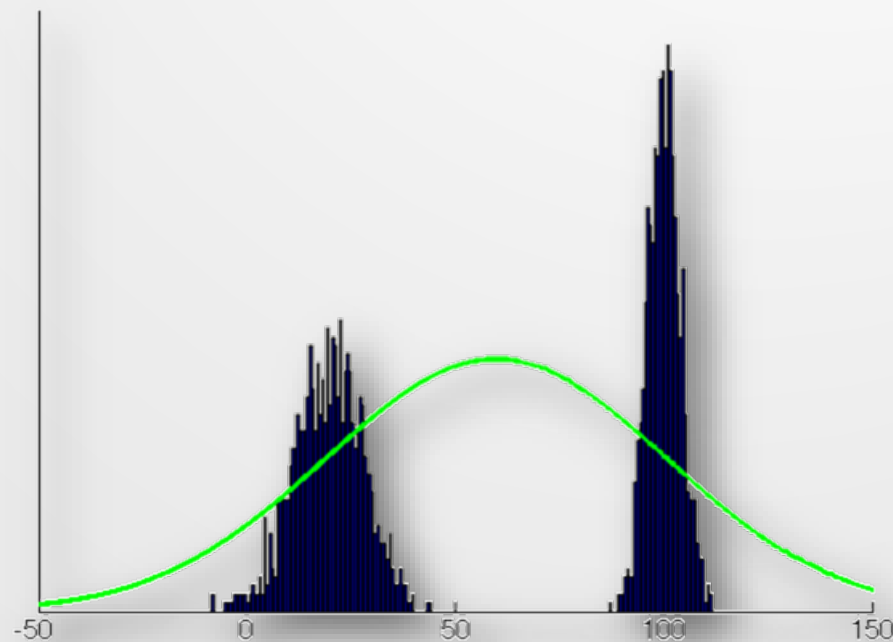
$$\Sigma = \begin{bmatrix} 2 & 0 \\ 0 & 0.6 \end{bmatrix}$$



- Different values on the diagonal result in different variances in their respective dimensions

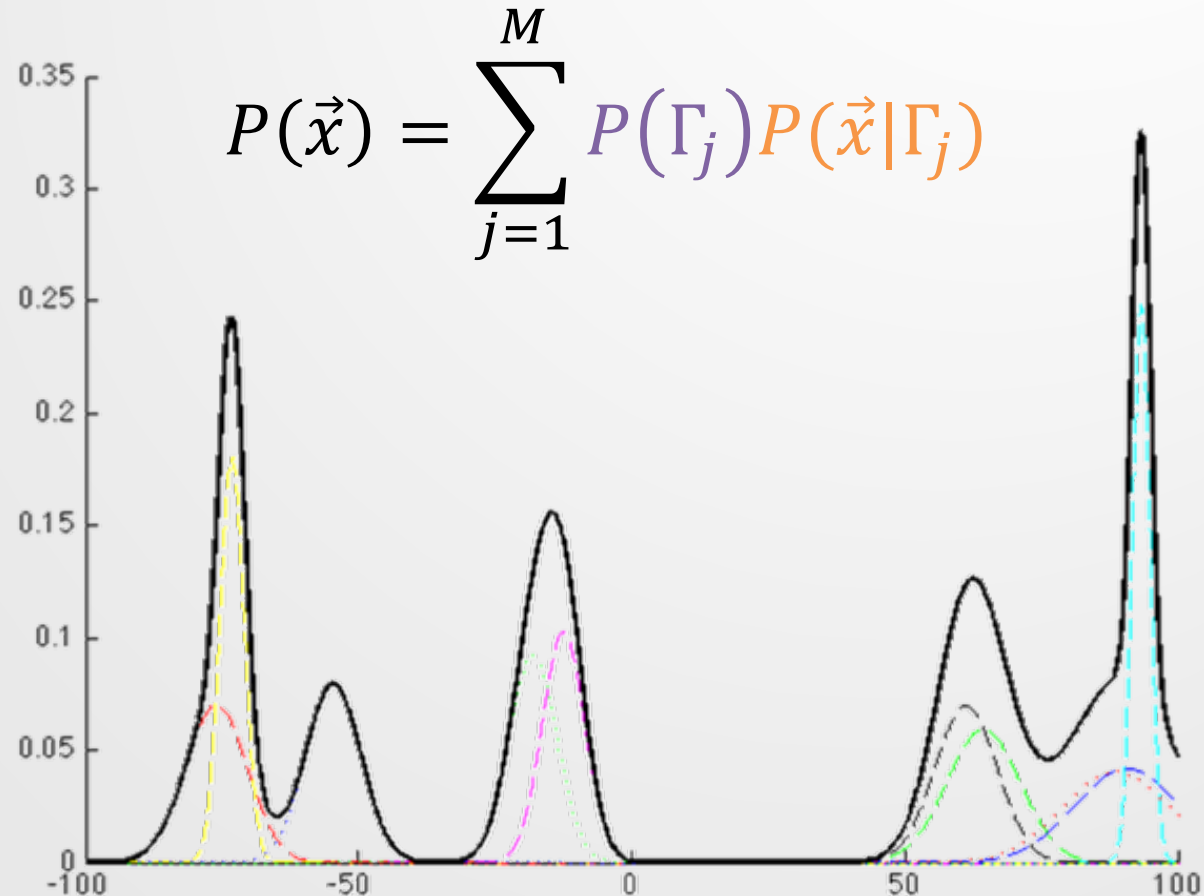
# Non-Gaussian observations

- Speech data are generally *not* unimodal.
- The observations below are **bimodal**, so fitting one Gaussian would not be representative.



# Mixtures of Gaussians

- **Gaussian mixture models (GMMs)** are a **weighted** linear combination of  $M$  component Gaussians,  $\langle \Gamma_1, \Gamma_2, \dots, \Gamma_M \rangle$ :



# Observation likelihoods

- Assuming MFCC dimensions are independent of one another, the **covariance matrix is diagonal** – i.e., 0 off the diagonal.
- Therefore, the probability of an observation vector given a Gaussian becomes

$$P(\vec{x}|\Gamma_m) = \frac{\exp\left(-\frac{1}{2} \sum_{i=1}^d \frac{(x[i] - \mu_m[i])^2}{\Sigma_m[i]}\right)}{(2\pi)^{\frac{d}{2}} \left(\prod_{i=1}^d \Sigma_m[i]\right)^{\frac{1}{2}}}$$

- Imagine that a GMM first chooses a Gaussian, then emits an observation from that Gaussian.*



# MLE for GMMs

- Let  $\omega_m = P(\Gamma_m)$  and  $b_m(\vec{x}_t) = P(\vec{x}_t | \Gamma_m)$ ,

'weight'

'component observation likelihood'

$$P_{\theta}(\vec{x}_t) = \sum_{m=1}^M \omega_m b_m(\vec{x}_t)$$


where  $\theta = \langle \omega_m, \vec{\mu}_m, \Sigma_m \rangle$  for  $m = 1..M$

- To estimate  $\theta$ , we solve  $\nabla_{\theta} \log L(X, \theta) = 0$  where

$$\log L(X, \theta) = \sum_{t=1}^T \log P_{\theta}(\vec{x}_t) = \sum_{t=1}^T \log \sum_{m=1}^M \omega_m b_m(\vec{x}_t)$$

# MLE for GMMs

- What happens when we try to find a maximum for  $\mu_m[n]$ ?

$$\begin{aligned}\frac{\delta \log L(X, \theta)}{\delta \mu_m[n]} &= \sum_{t=1}^T \frac{\delta}{\delta \mu_m[n]} \log \sum_{m'=1}^M \omega_{m'} b_{m'}(\vec{x}_t) = 0 \\ \sum_{t=1}^T \frac{1}{P_\theta(\vec{x}_t)} \frac{\delta}{\delta \mu_m[n]} \omega_m b_m(\vec{x}_t) &= \sum_{t=1}^T \frac{\omega_m b_m(\vec{x}_t)}{P_\theta(\vec{x}_t)} \left( \frac{x_t[n] - \mu_m[n]}{\Sigma_m[n]^2} \right) = 0 \\ \mu_m[n] &= \frac{\sum_{t=1}^T \frac{\omega_m b_m(\vec{x}_t)}{P_\theta(\vec{x}_t)} x_t[n]}{\sum_{t=1}^T \frac{\omega_m b_m(\vec{x}_t)}{P_\theta(\vec{x}_t)}} = \frac{\sum_{t=1}^T P_\theta(\Gamma_m | \vec{x}_t) x_t[n]}{\sum_{t=1}^T P_\theta(\Gamma_m | \vec{x}_t)}\end{aligned}$$


But this involves  $\mu_m[n]$ !

# Learning mixtures of gaussians

- If we knew *which* Gaussian generated each sample, then  $\langle \vec{\mu}_m, \Sigma_m \rangle$  can be learned by MLE.
- The MLE of  $P(\Gamma_j)$  would likewise be the count  $\frac{\#\vec{x}_t \text{ from } \Gamma_j}{T}$
- But we **don't** know this!
- Instead, we guess at “soft” mixture assignments  $P_\theta(\Gamma_m | \vec{x}_t)$  from another model...
- ...which we got from a previous round of maximization

# Expectation-Maximization for GMMs

- Overall idea:
  - First, initialize a set of model parameters.
  - “Expectation”: Compute the expected probabilities of observation, given these parameters.
  - “Maximization”: Update the parameters to maximize the aforementioned probabilities.
  - Repeat.

# Expectation-Maximization for GMMs

- The **expectation step** gives us:

$$P_{\theta}(\Gamma_m | \vec{x}_t) = \frac{\omega_m b_m(\vec{x}_t)}{P_{\theta}(\vec{x}_t)}$$

Proportion of overall probability contributed by  $m$

- The **maximization step** gives us:

$$\widehat{\mu}_m = \frac{\sum_t P_{\theta}(\Gamma_m | \vec{x}_t) \vec{x}_t}{\sum_t P_{\theta}(\Gamma_m | \vec{x}_t)}$$

$$\widehat{\Sigma}_m = \frac{\sum_t P_{\theta}(\Gamma_m | \vec{x}_t) \vec{x}_t^2}{\sum_t P_{\theta}(\Gamma_m | \vec{x}_t)} - \widehat{\mu}_m^2$$

$$\widehat{\omega}_m = \frac{1}{T} \sum_{t=1}^T P_{\theta}(\Gamma_m | \vec{x}_t)$$

Recall from slide 18, MLE wants:

$$\mu = \frac{\sum_i x_i}{n}$$
$$\sigma^2 = \frac{\sum_i (x_i - \mu)^2}{n}$$

# Recipe for GMM EM

- For each speaker, we learn a GMM given all  $T$  frames of their training data.

- 1. Initialize:** Guess  $\theta = \langle \omega_m, \overrightarrow{\mu_m}, \Sigma_m \rangle$  for  $m = 1..M$  either uniformly, randomly, or by  $k$ -means clustering.
- 2. E-step:** Compute  $P_\theta(\Gamma_m | \overrightarrow{x_t})$ .
- 3. M-step:** Update parameters for  $\langle \omega_m, \overrightarrow{\mu_m}, \Sigma_m \rangle$  with  $\langle \widehat{\omega_m}, \widehat{\overrightarrow{\mu_m}}, \widehat{\Sigma_m} \rangle$  as described on slide 29.