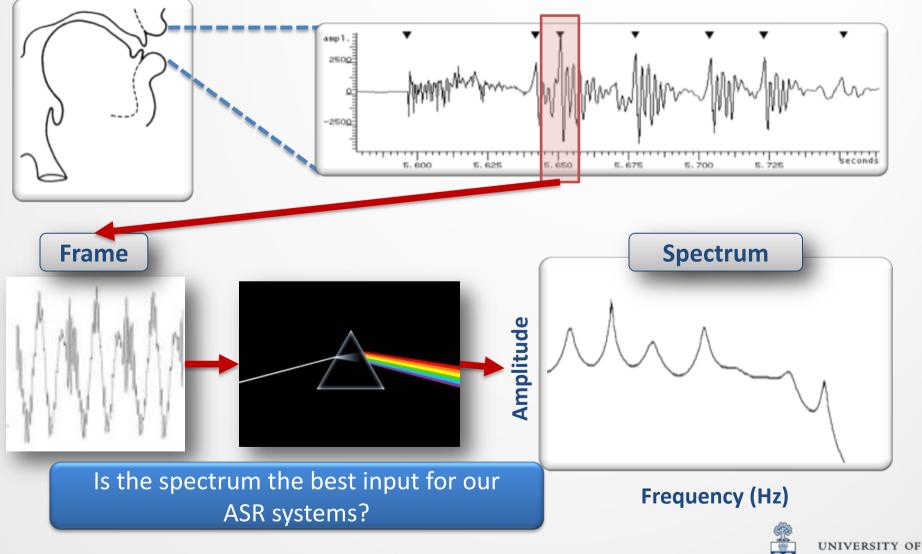
automatic speech recognition

CSC401/2511 – Natural Language Computing – Spring 2019 Lecture 9 Frank Rudzicz University of Toronto

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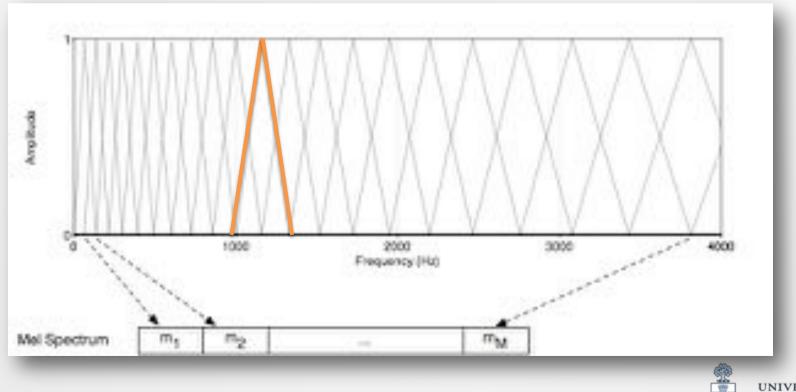
Recall our input to ASR



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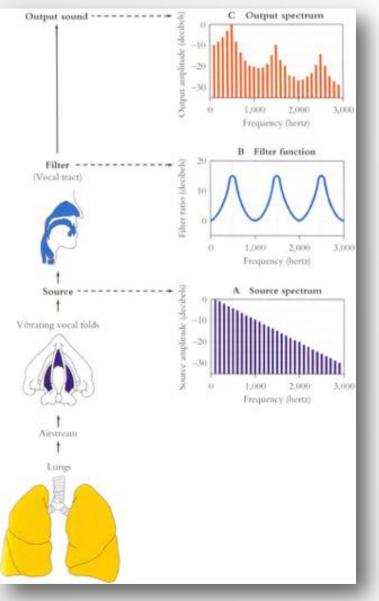
1. The Mel-scale filter bank

- To **mimic** the response of the **human ear** (and because it empirically improves speech recognition), we often discretize the spectrum using *M* triangular **filters**.
 - Uniform spacing before 1 kHz, logarithmic after 1 kHz



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2. Source and filter

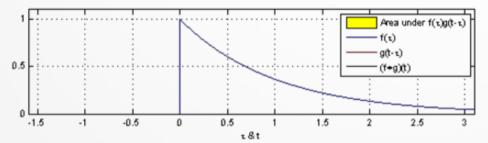


- The acoustics of speech are produced by a glottal pulse waveform (the source) passing through a vocal tract whose shape modifies that wave (the filter).
- The **shape** of the vocal tract is more important to phoneme recognition.
 - We to separate the source from the filter in the acoustics.



2. Source and filter (aside)

- Since speech is assumed to be the output of a linear time invariant system, it can be described as a convolution.
 - **Convolution**, *x* * *y*, is beyond the scope of this course, but can be conceived as the modification of one signal by another.



 For speech signal x[n], glottal signal g[n], and vocal tract transfer v[n] with spectra X[z], G[z], and V[z], respectively :

x[n] = g[n] * v[n]X[z] = G[z]V[z] $\log X[z] = \log G[z] + \log V[z]$

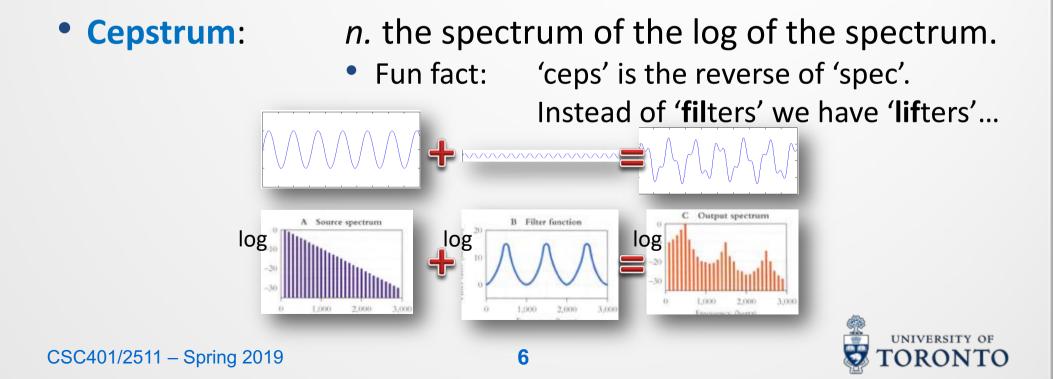
We've separated the source and filter into two terms!



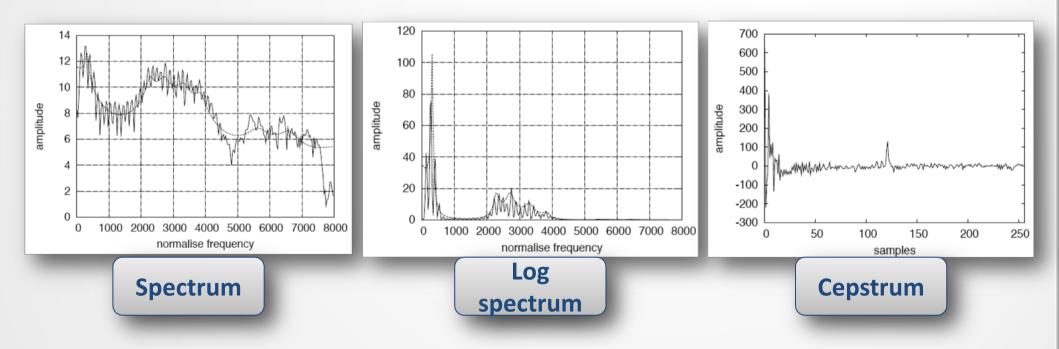
2. The cepstrum

- We separate the source and the filter by pretending the log of the spectrum is actually a time domain signal.
 - the log spectrum log X[z] is a sum of the log spectra of the source and filter, i.e., a superposition;

finding *its* spectrum will allow us to **isolate** these components.



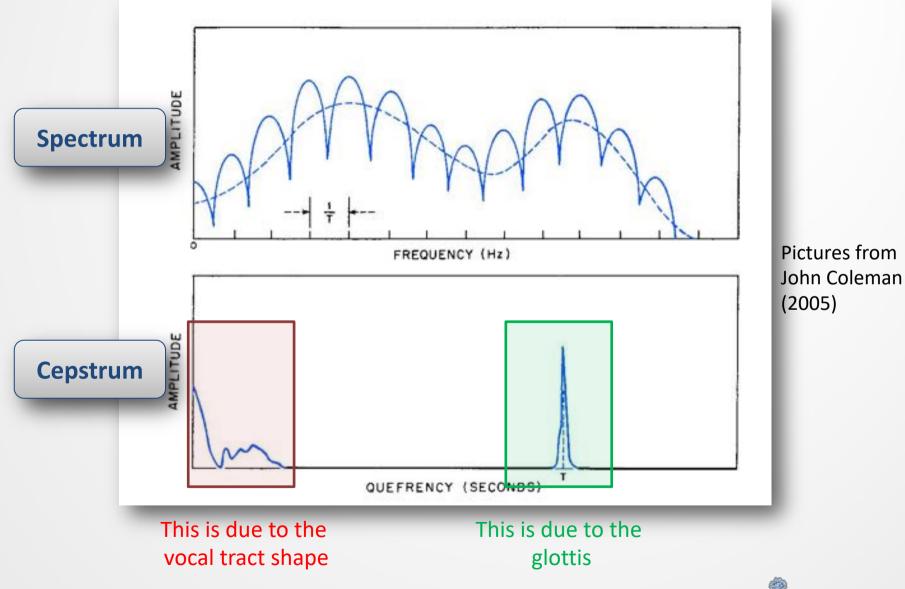
2. The cepstrum



 The domain of the cepstrum is quefrency (a play on the word 'frequency').



2. The cepstrum



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Mel-frequency cepstral coefficients

- Mel-frequency cepstral coefficients (MFCCs) are the most popular representation of speech used in ASR.
 - They are the spectra of the logarithms of the Mel-scaled filtered spectra of the windows of the waveform.





Advantages of MFCCs

- The cepstrum produces **highly uncorrelated features** (every dimension is useful).
 - This includes a **separation** of the **source** and **filter**.
- Historically, the cepstrum *has been* easier to learn than the spectrum for phoneme recognition.
- There is an efficient method to compute cepstra called the discrete cosine transform.



MFCCs in practice

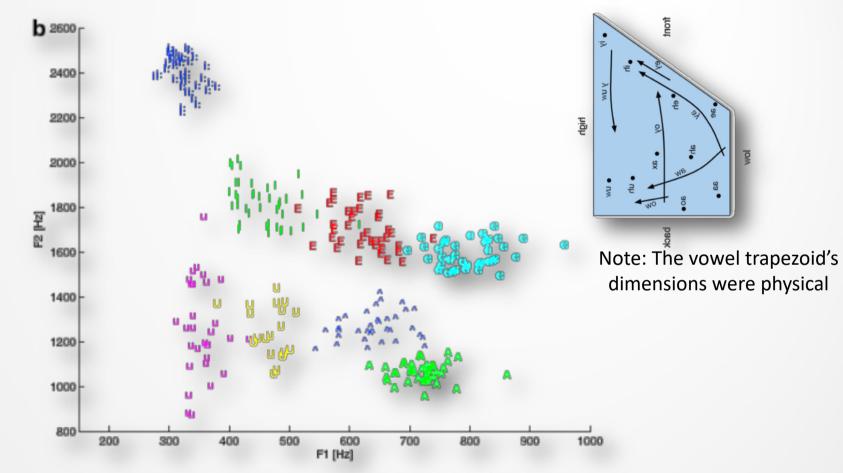
- An observation vector of MFCCs often consists of
 - The **first 13 cepstral coefficients** (i.e., the first 13 dimensions produced by this method),
 - An additional overall energy measure,
 - The **velocities** (δ) of each of those 14 dimensions,
 - i.e., the rate of change of each coefficient at a given time
 - The **accelerations** ($\delta\delta$) of each of original 14 dimensions.
- The result is that at a timeframe t we have an observation MFCC vector of (13+1)*3=42 dimensions.
 - This vector is what is used by our ASR systems...



GAUSSIAN CLUSTERS



Classifying speech sounds



 Speech sounds tend to cluster. This graph shows vowels, each in their own colour, according to the 1st two formants.



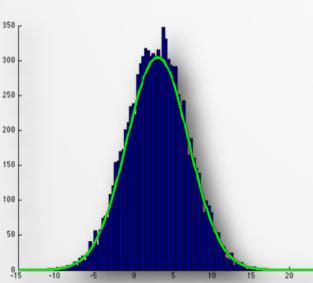
Classifying speakers

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



Fitting continuous distributions

 Since we are operating with continuous variables, we need to fit continuous probability functions to a discrete number of observations.

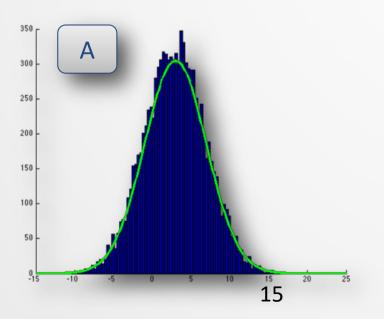


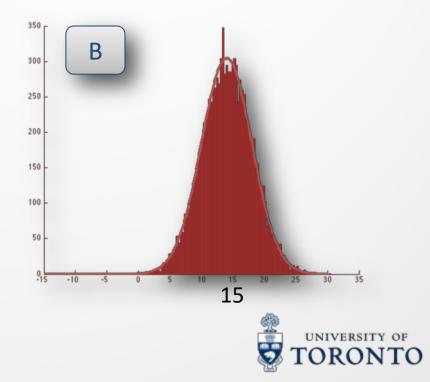
• If we assume the 1-dimensional data in **this histogram** is Normally distributed, we can fit a continuous Gaussian function simply in terms of the mean μ and variance σ^2 .



Comparing continuous distributions

- If we observe a particular value in this univariate space, e.g.,
 x = 15, we can say which of several distributions is most likely to have produced it.
 - Here, distribution B is more likely to have produced x = 15 because
 P(x; B) > P(x; A).

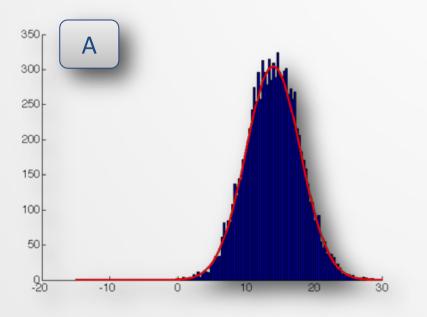


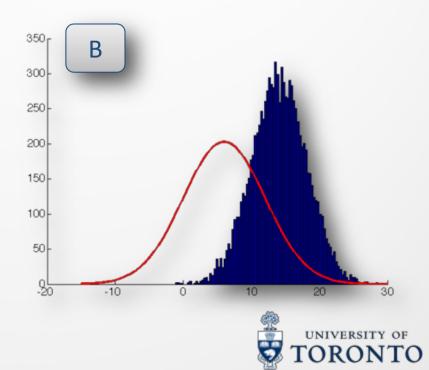


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Good fits

- Given some fixed training data, we want to be able to fit continuous probability functions that best match our observations.
 - The data in **this histogram** are **more likely** to have been produced from the parameterization on the **left**.





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)
• $\sigma^2 = E\left((x-\mu)^2\right) = \int (x-\mu)^2 P(x) dx$ (variance)

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



Maximum likelihood estimation

- Given data $X = \{x_1, x_2, ..., x_n\}$, 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, θ) provides a surface over all θ, 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 μ :

$$L(X,\mu) = P(X;\mu) = \prod_{i=1}^{n} P(x_i;\theta) = \prod_{i=1}^{n} \frac{\exp\left(-\frac{(x_i - \mu)^2}{2\sigma^2}\right)}{\sqrt{2\pi}\sigma}$$
$$\log L(X,\mu) = -\frac{\sum_i (x_i - \mu)^2}{2\sigma^2} - n\log\sqrt{2\pi}\sigma$$
$$\frac{\delta}{\delta\mu}\log L(X,\mu) = \frac{\sum_i (x_i - \mu)}{\sigma^2} = 0$$
$$\mu = \frac{\sum_i x_i}{n}$$
Similarly, $\sigma^2 = \frac{\sum_i (x_i - \mu)^2}{n}$



Multivariate Gaussians

• When data is *d*-dimensional, the input variable is $\vec{x} = \langle x[1], x[2], ..., x[d] \rangle$

the **mean** is

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

the **covariance matrix** is

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

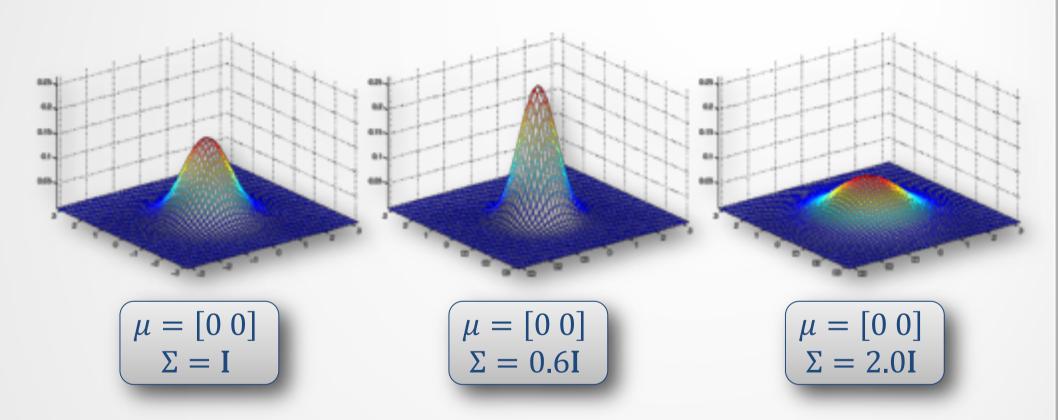
and

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

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



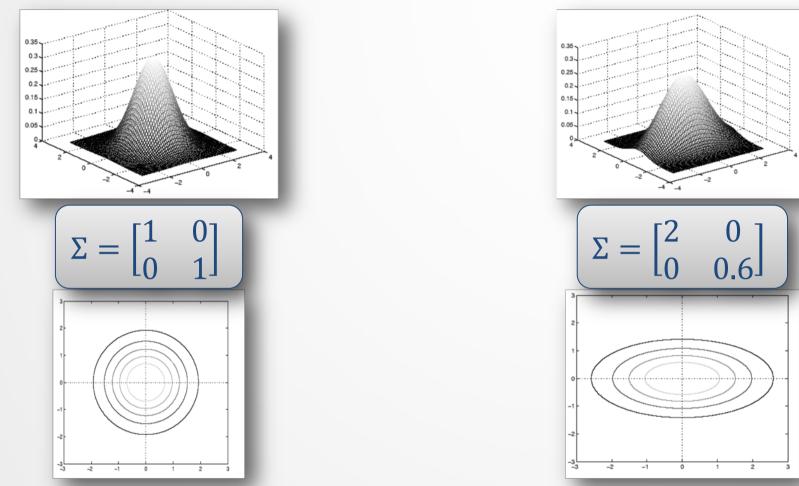
Intuitions of covariance



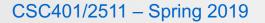
- As values in Σ become larger, the Gaussian spreads out.
- (I is the identity matrix 0 except for 1s on the diagonal)



Intuitions of covariance

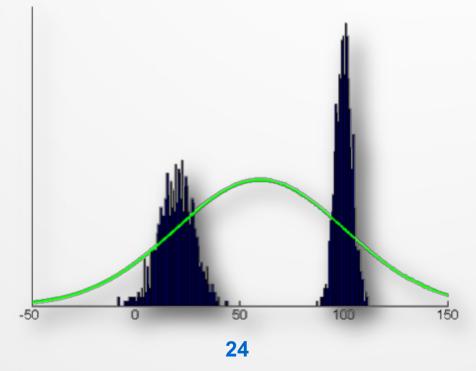


 Different values on the diagonal result in different variances in their respective dimensions



Non-Gaussian observations

- Speech data is generally *not* unimodal it's more complex.
- The observations below are **bimodal**, so fitting one Gaussian would not be representative.
 - E.g., if you usually keep your phone in your desk or on your table, it makes no sense looking for them floating in the air between them.

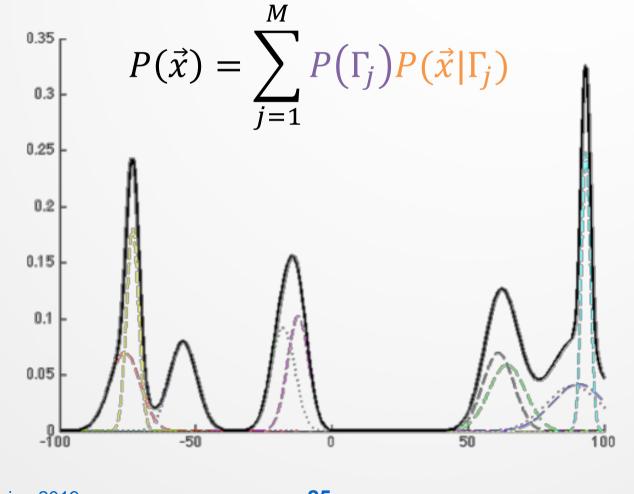




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Mixtures of Gaussians

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



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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 from slide 14 becomes

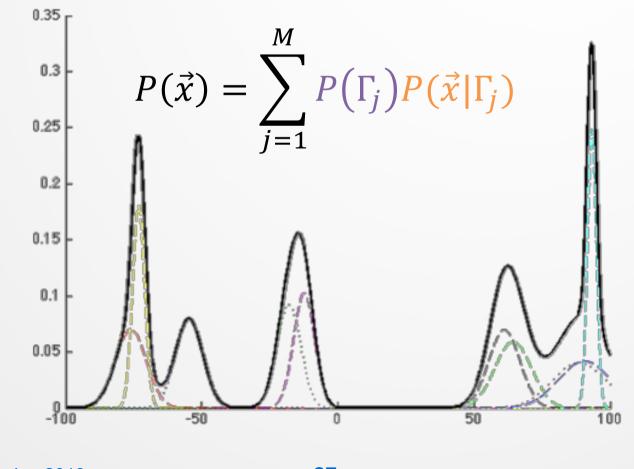
$$P(\vec{x}|\Gamma_m) = \frac{\exp\left(-\frac{1}{2}\sum_{i=1}^{d}\frac{(x[i] - \mu_m[i])^2}{\sum_m[i]}\right)}{(2\pi)^{\frac{d}{2}}\left(\prod_{i=1}^{d}\sum_m[i]\right)^{\frac{1}{2}}}$$

We imagine a GMM first chooses a Gaussian, then emits an observation from that Gaussian.



Mixtures of Gaussians

• If we knew which Gaussian generated each sample, we could learn $P(\Gamma_j)$ with MLE, but that data is **hidden**, so we must





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use...

Expectation-Maximization for GMMs

• If
$$\omega_m = P(\Gamma_m)$$
 and $b_m(\vec{x_t}) = P(\vec{x_t}|\Gamma_m)$, "component observation likelihood"
(weight'
 $P_{\theta}(\vec{x_t}) = \sum_{m=1}^{M} \omega_m b_m(\vec{x_t})$ "overall probability"

where $\theta = \langle \omega_m, \overrightarrow{\mu_m}, \Sigma_m \rangle$ for m = 1..M

• To estimate θ , we solve $\nabla_{\theta} \log L(X, \theta) = 0$ where $\log L(X, \theta) = \sum_{t=1}^{T} \log P_{\theta}(\vec{x_t}) = \sum_{t=1}^{T} \log \sum_{m=1}^{M} \omega_m b_m(\vec{x_t})$



Expectation-Maximization for GMMs

• We **differentiate** the log likelihood function w.r.t . $\mu_m[n]$ and set this to 0 to find the value of $\mu_m[n]$ at which the likelihood stops growing.

$$\frac{\delta \log L(X,\theta)}{\delta \mu_m[n]} = \sum_{t=1}^N \frac{1}{P_{\theta}(\vec{x_t})} \left[\frac{\delta}{\delta \mu_m[n]} \omega_m b_m(\vec{x_t}) \right] = 0$$



Expectation-Maximization for GMMs

• The **expectation step** gives us:

$$\boldsymbol{b}_{m}(\boldsymbol{x}_{t}) = P(\boldsymbol{x}_{t} | \boldsymbol{\Gamma}_{m})$$
$$\boldsymbol{P}(\boldsymbol{\Gamma}_{m} | \boldsymbol{x}_{t}; \boldsymbol{\theta}) = \frac{\omega_{m} b_{m}(\boldsymbol{x}_{t})}{P_{\theta}(\boldsymbol{x}_{t})} \mathbf{prod}$$

Proportion of overall orobability contributed by *m*

• The maximization step gives us:

$$\widehat{\mu_{m}} = \frac{\sum_{t} P(\Gamma_{m} | \overrightarrow{x_{t}}; \theta) \overrightarrow{x_{t}}}{\sum_{t} P(\Gamma_{m} | \overrightarrow{x_{t}}; \theta)}$$
$$\widehat{\Sigma_{m}} = \frac{\sum_{t} P(\Gamma_{m} | \overrightarrow{x_{t}}; \theta) \overrightarrow{x_{t}}^{2}}{\sum_{t} P(\Gamma_{m} | \overrightarrow{x_{t}}; \theta)} - \widehat{\overline{\mu_{m}}}^{2}$$
$$\widehat{\omega_{m}} = \frac{1}{T} \sum_{t=1}^{T} P(\Gamma_{m} | \overrightarrow{x_{t}}; \theta)$$

Recall from slide
20, MLE wants:
$$\mu = \frac{\sum_{i} x_{i}}{n}$$
$$\sigma^{2} = \frac{\sum_{i} (x_{i} - \mu)^{2}}{n}$$



Some notes...

- In the previous slide, the square of a vector, \$\vec{a}^2\$, is elementwise (i.e., numpy.multiply in Python)
 E.g., [2, 3, 4]² = [4, 9, 16]
- Since Σ is diagonal, it can be represented as a vector.

• Can
$$\widehat{\overline{\sigma_m^2}} = \frac{\sum_t P(\Gamma_m | \overrightarrow{x_t}; \theta) \overrightarrow{x_t^2}}{\sum_t P(\Gamma_m | \overrightarrow{x_t}; \theta)} - \widehat{\overline{\mu_m}}^2$$
 become negative?

• No.

• This is left as an exercise, but only if you're interested.



Speaker recognition

- Speaker recognition: *n*. the identification of a speaker among several speakers given only some acoustics.
- Each **speaker** will produce speech according to **different** probability distributions.
 - We train a **Gaussian mixture model** for each speaker, given annotated data (mapping utterances to speakers).
 - We choose the speaker whose model gives the highest probability for an observation.



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 $m = 1M$
	either uniformly, randomly, or by k-means
	clustering.

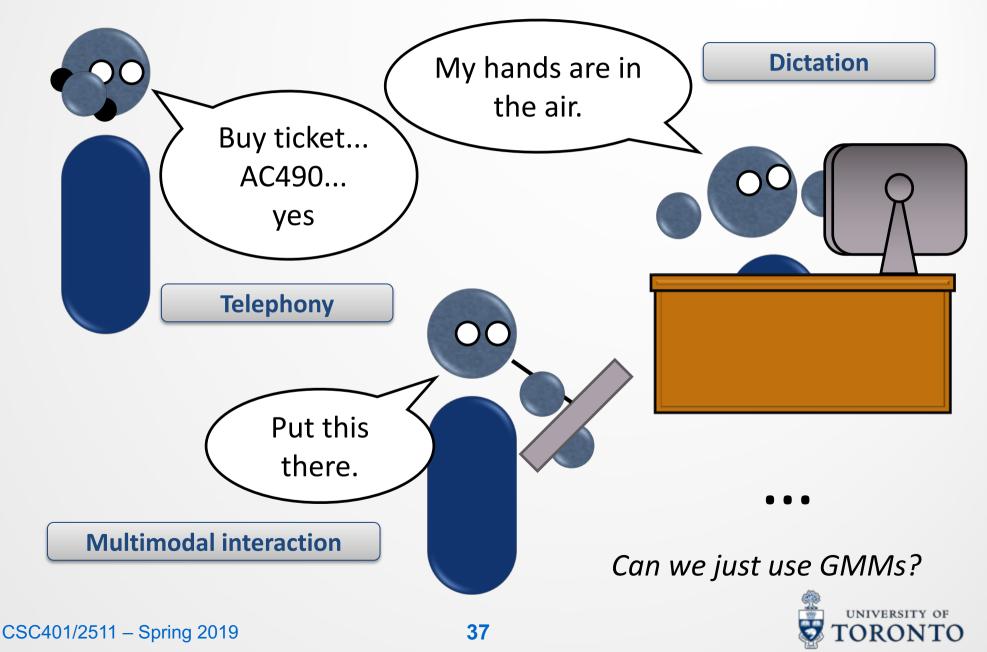
- **2. E-step**: Compute $b_m(\vec{x_t})$ and $P(\Gamma_m|\vec{x_t}; \theta)$.
- **3. M-step**: Update parameters for $\langle \omega_m, \overline{\mu_m}, \Sigma_m \rangle$ as described on slide 30.
- (see the Reynolds & Rose (1995) paper on the course webpage for details)



SPEECH RECOGNITION



Consider what we want speech to do



Speech databases

- Large-vocabulary continuous ASR is meant to encode full conversational speech, with a vocabulary of >64K words.
 - This requires *lots* of data to train our models.
- The Switchboard corpus contains 2430 conversations spread out over about 240 hours of data (~14 GB).
- The TIMIT database contains 63,000 sentences from 630 speakers.
 - Relatively small (~750 MB), but very popular.
- Speech data from conferences (e.g., TED) or from broadcast news tends to be between 3 GB and 30 GB.



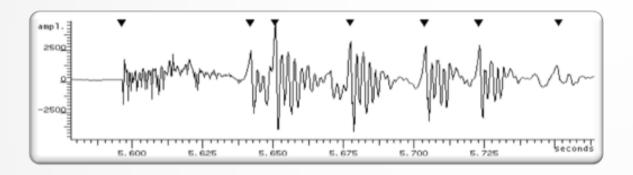
Aspects of ASR systems in the world

- Speaking mode: Isolated word (e.g., "yes") vs. continuous (e.g., "Siri, ask Cortana for the weather")
- Speaking style: Read speech vs. spontaneous speech; the latter contains many dysfluencies (e.g., stuttering, uh, like, ...)
- Enrolment: Speaker-dependent (all training data from one speaker) vs. speaker-independent (training data from many speakers).
 Vocabulary: Small (<20 words) or large (>50,000 words).
 Transducer: Cell phone? Noise-cancelling microphone? Teleconference microphone?



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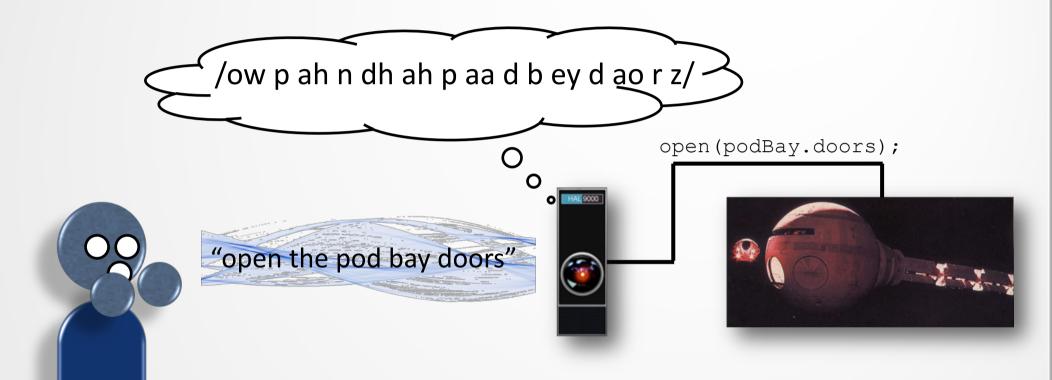
Speech is dynamic



- Speech changes over time.
 - GMMs are good for high-level clustering, but they encode no notion of order, sequence, or time.
- Speech is an expression of language.
 - We want to incorporate knowledge of how phonemes and words are ordered with **language models**.



Speech is sequences of phonemes^(*)



We want to convert a series of MFCC vectors into a sequence of phonemes.



Phoneme dictionaries

- There are many **phonemic dictionaries** that map words to pronunciations (i.e., lists of phoneme sequences).
- The CMU dictionary (<u>http://www.speech.cs.cmu.edu/cgi-bin/cmudict</u>) is popular.
 - 127K words transcribed with the ARPAbet.
 - Includes some rudimentary prosody markers.

I	EH2	V	AH0	L	UW1	SH	AH0	Ν				
1(2)	IY2	V	AH0	L	UW1	SH	AH0	Ν				
1(3)	EH2	V	0W0	L	UW1	SH	AH0	Ν				
1(4)	IY2	V	0W0	L	UW1	SH	AH0	Ν				
IARY	EH2	V	AH0	L	UW1	SH	AH0	Ν	EH2	R	IY0	



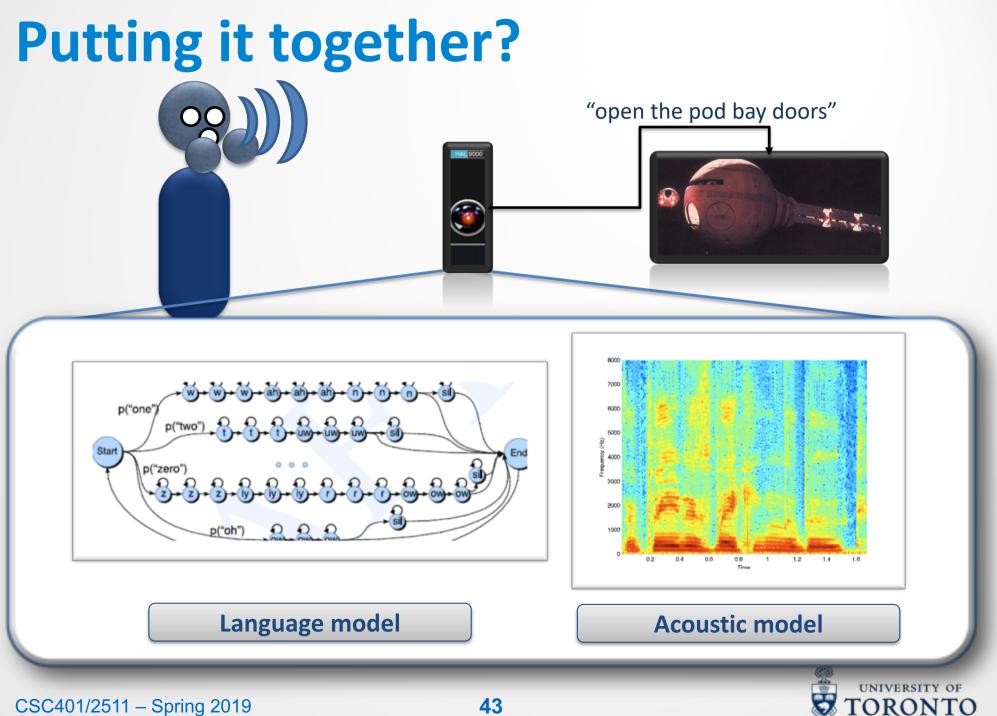
EVOLUTION

EVOLUTION

EVOLUTION

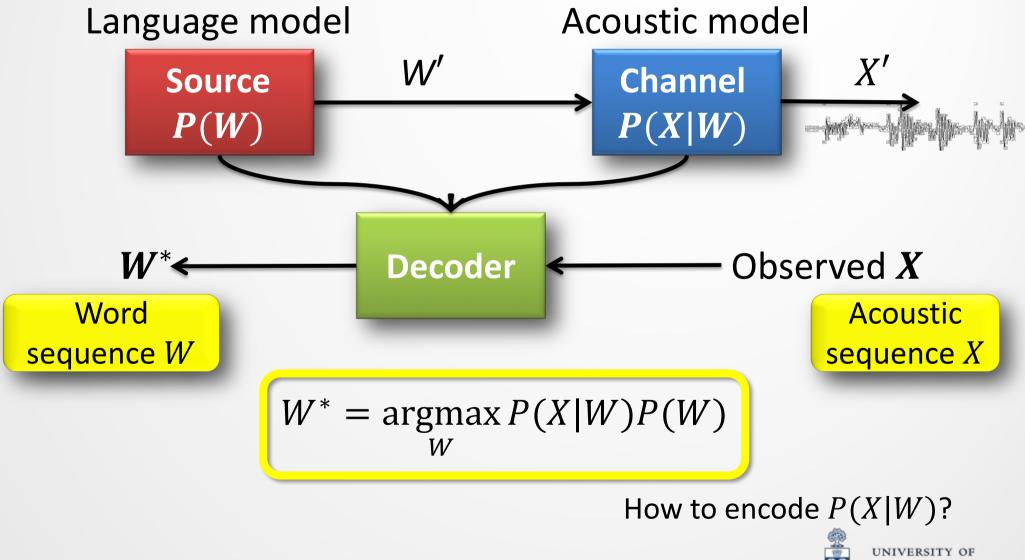
EVOLUTION

EVOLUTION



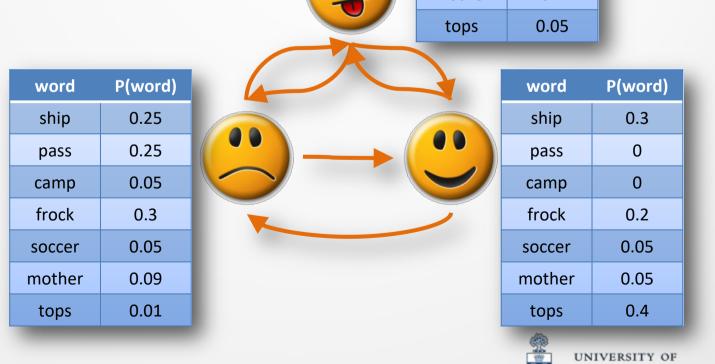
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The noisy channel model for ASR



Reminder – discrete HMMs

 Previously we saw discrete HMMs: at each state we observed a discrete symbol from a finite set of discrete symbols.



P(word)

0.1

0.05

0.05

0.6

0.05

0.1

word

ship

pass

camp

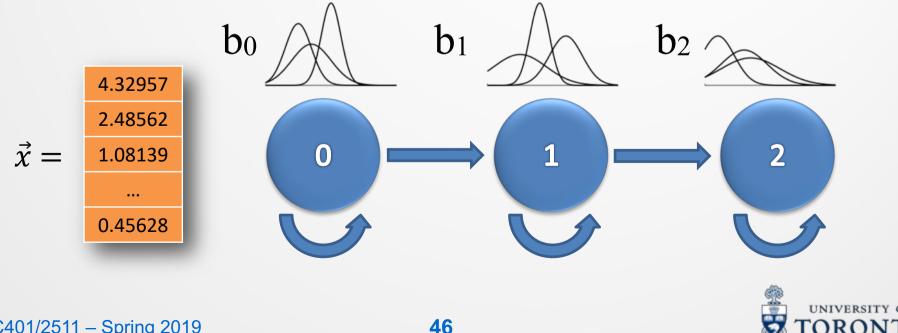
frock

soccer

mother

Continuous HMMs (CHMM)

- A continuous HMM has observations that are distributed over continuous variables.
 - Observation probabilities, b_i , are also continuous.
 - E.g., here $b_0(\vec{x})$ tells us the probability of seeing the (multivariate) continuous observation \vec{x} while in state 0.



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Defining CHMMs

- Continuous HMMs are very similar to discrete HMMs.
 - $S = \{s_1, \dots, s_N\}$
 - $X = \mathbb{R}^{42}$

$$\theta \begin{cases} \bullet \Pi = \{\pi_1, \dots, \pi_N\} \\ \bullet A = \{a_{ij}\}, i, j \in S \\ \bullet B = b_i(\vec{x}), i \in S, \vec{x} \in X \end{cases}$$

yielding

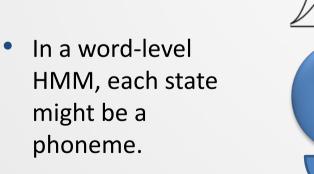
• $Q = \{q_0, \dots, q_T\}, q_i \in S$ • $\mathcal{O} = \{\sigma_0, \dots, \sigma_T\}, \sigma_i \in X$

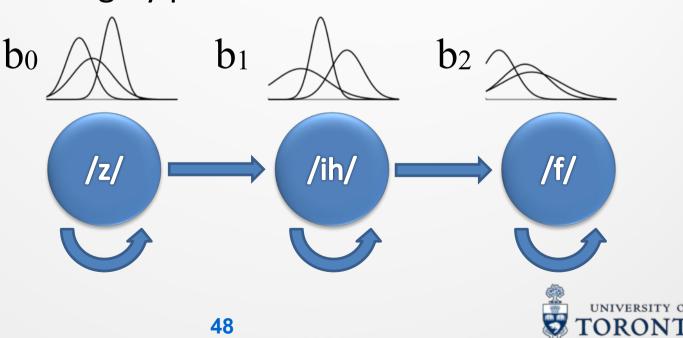
- : set of states (e.g., subphones)
- : continuous observation space
- : initial state probabilities
- : state transition probabilities
- : state output probabilities (i.e., Gaussian mixtures)
- : state sequence
- : observation sequence



Word-level HMMs?

- Imagine that we want to learn an HMM for each word in our lexicon (e.g., 60K words → 60K HMMs).
- No, thank you! Zipf's law tells us that many words occur very infrequently.
 - 1 (or a few) training examples of a word is *not* enough to train a model as highly parameterized as a CHMM.

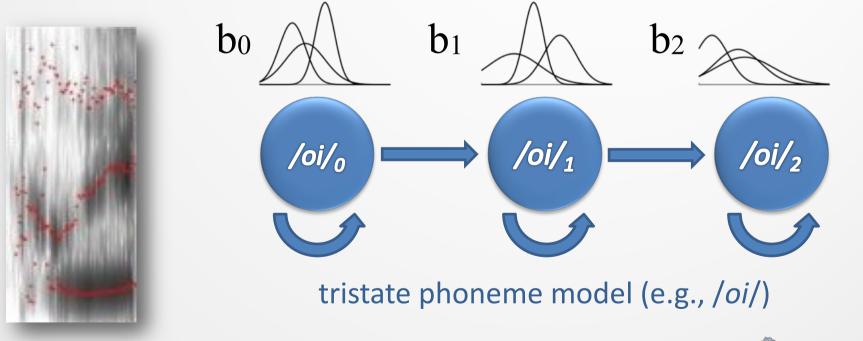




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Phoneme HMMs

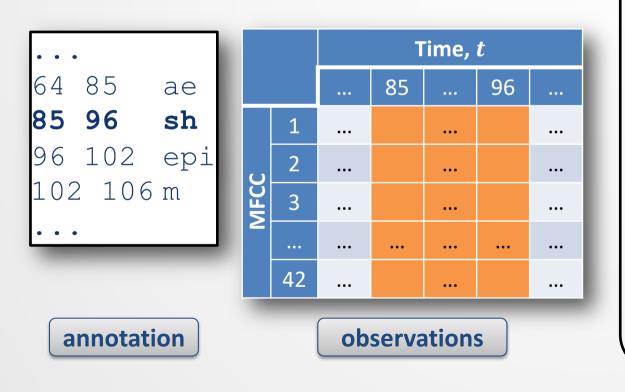
- Phonemes change over time we model these dynamics by building one HMM for each phoneme.
 - Tristate phoneme models are popular.
 - The centre state is often the 'steady' part.

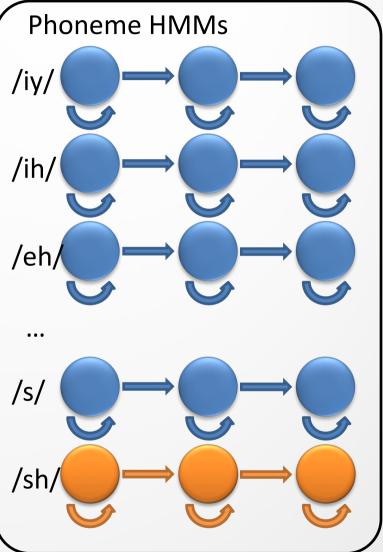




Phoneme HMMs

- We train each phoneme HMM using *all* sequences of that phoneme.
 - Even from different words.







Combining models

- We can learn an *N*-gram <u>language model</u> from word-level transcriptions of speech data.
 - These models are discrete and are trained using MLE.
- Our phoneme HMMs together constitute our <u>acoustic model</u>.
 - Each phoneme HMM tells us how a phoneme 'sounds'.
- We can combine these models by concatenating phoneme HMMs together according to a known lexicon.
 - We use a word-to-phoneme dictionary.



Combining models

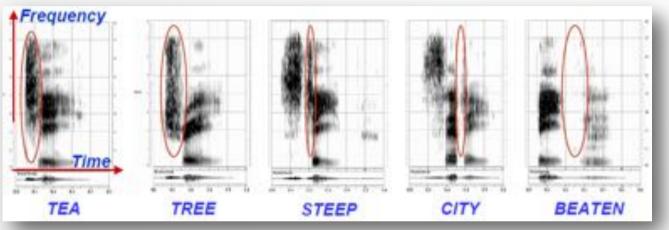
- If we know how phonemes combine to make words, we can simply concatenate together our phoneme models by inserting and adjusting transition weights.
 - e.g., Zipf is pronounced /z ih f/, so...



(It's a bit more complicated than this – normally phoneme HMMs have special 'handle' states at either end that connect to other HMMs)

Co-articulation and triphones

• **Co-articulation**: *n*. When a phoneme is influenced by adjacent phonemes.

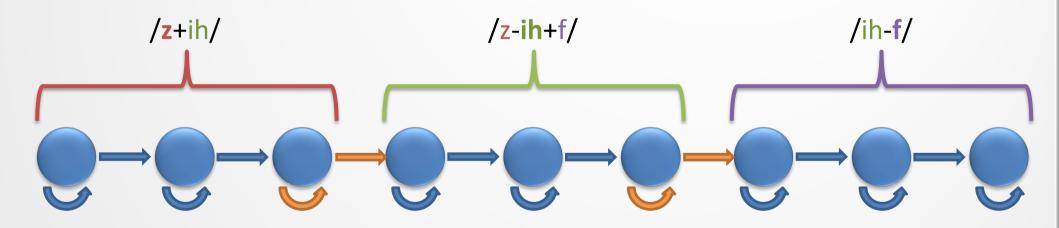


- A triphone HMM captures co-articulation.
 - Triphone model /a-b+c/ is phoneme b when preceded by a and followed by c.



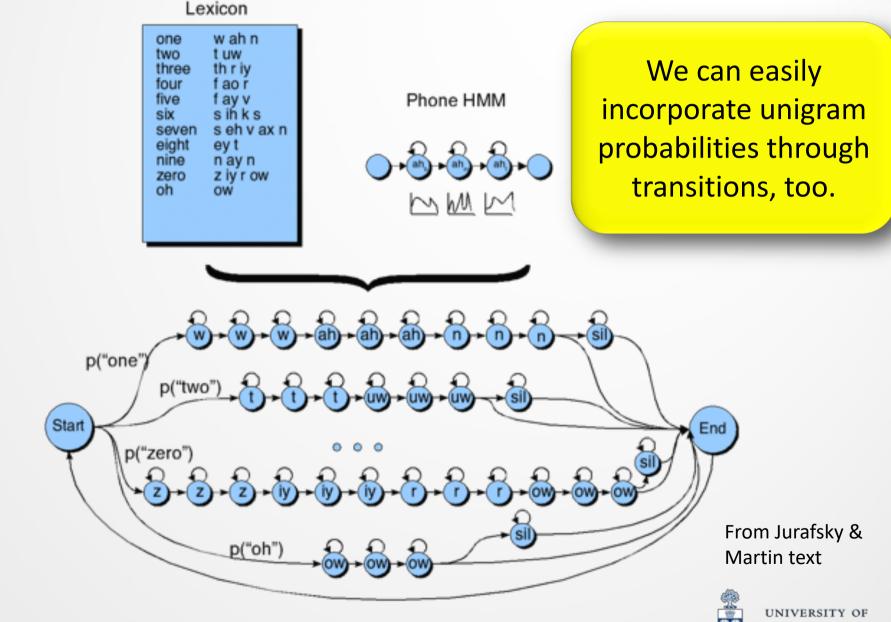
Combining triphone HMMs

 Triphone models can only connect to other triphone models that 'match'.



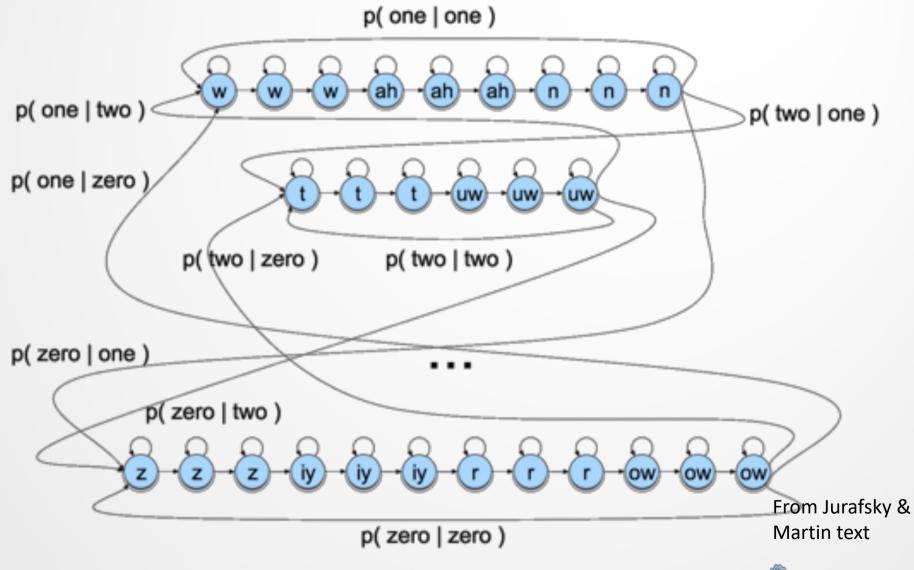


Concatenating phoneme models



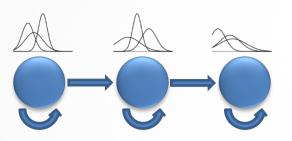
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Bigram models





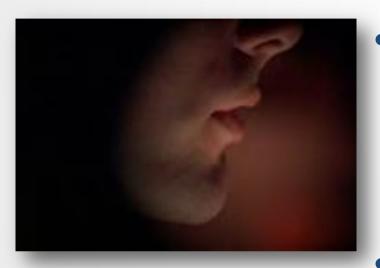
Using CHMMs



- As before, these HMMs are generative models that encode statistical knowledge of how output is generated.
- We train CHMMs with Baum-Welch (a type of Expectation-Maximization), as we did before with discrete HMMs.
 - Here, the observation parameters, $b_i(\vec{x})$, are adjusted using the GMM training 'recipe' from earlier.
- We find the best state sequences using Viterbi, as before.
 - Here, the best state sequence gives us a sequence of phonemes and words.



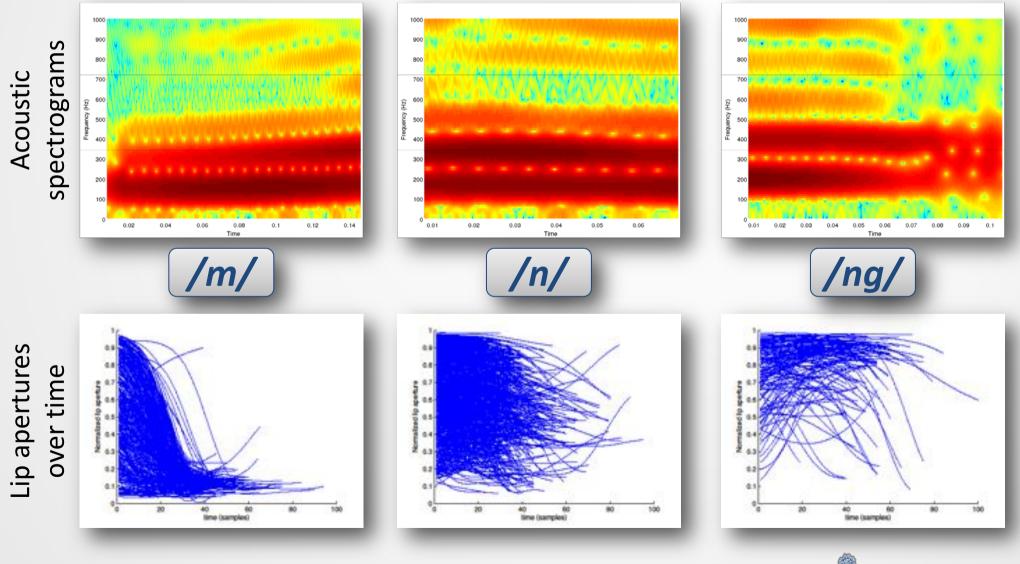
Audio-visual speech methods



- Observing the **vocal tract** directly, rather than through inference, can be very helpful in automatic speech recognition.
- The shape and aperture of the mouth gives some clues as to the phoneme being uttered.
 - Depending on the level of invasiveness, we can even measure the glottis and tongue directly.

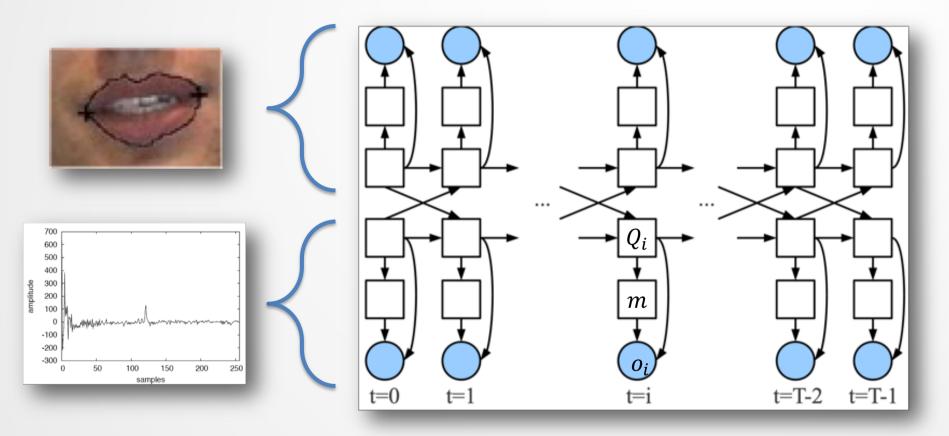


Example – Lip aperture and nasals





Coupled HMM



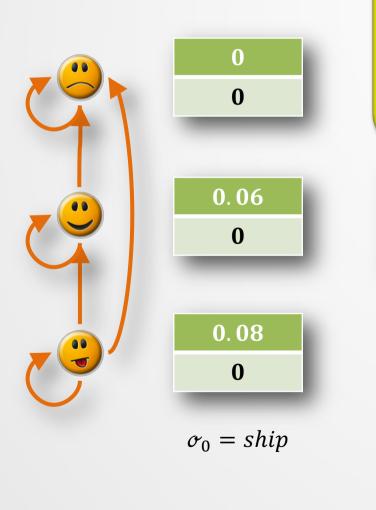
Where Q_i is the HMM state, m is the index into a GMM, and o_i is the observation at time i.

Nefian A V, Liang L, Pi X, et al. A coupled HMM for audio-visual speech recognition. In: International Conference on Acoustics, Speech and Signal Processing ICASSP'02. 2002. 2013–6. CSC401/2511 – Spring 2019 60

NEURAL SPEECH RECOGNITION



Remember Viterbi



The best path to state s_j at time t, $\delta_j(t)$, depends on the best path to each possible previous state, $\delta_i(t-1)$, and their transitions to j, a_{ij}

$$\delta_j(t) = \max_i \left[\delta_i(t-1)a_{ij} \ b_j(\sigma_t) \right]$$
$$\psi_j(t) = \operatorname*{argmax}_i \left[\delta_i(t-1)a_{ij} \right]$$

Do these probabilities need to be GMMs?

 $\sigma_1 = frock$

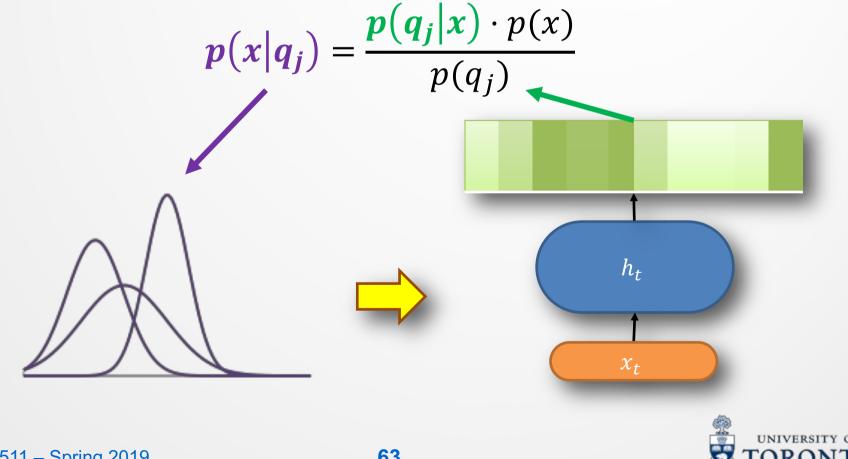
Observations, σ_t



 $\sigma_2 = tops$

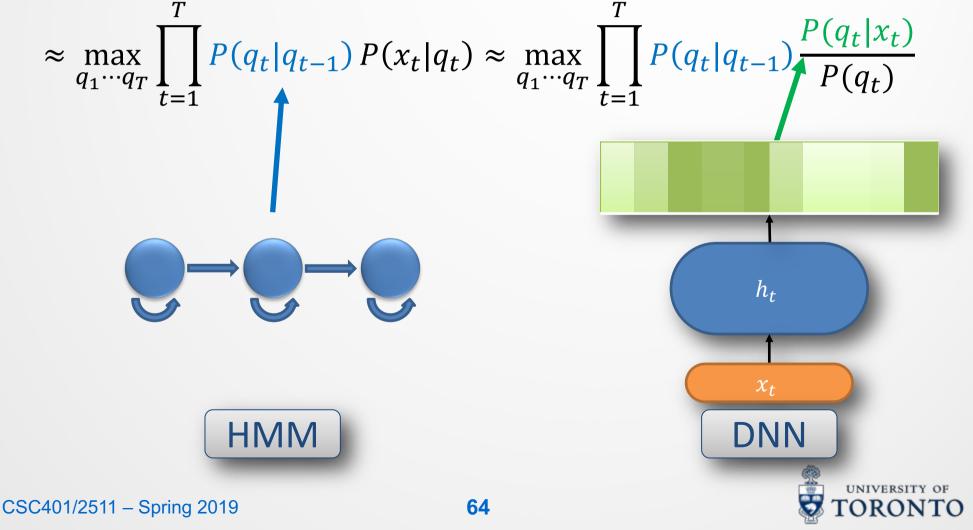
Replacing GMMs with DNNs

- Obtain $b_i(x) = p(x|q_i)$ with a neural network.
- We can't learn that continuous distribution directly, but we can use Bayes' rule:

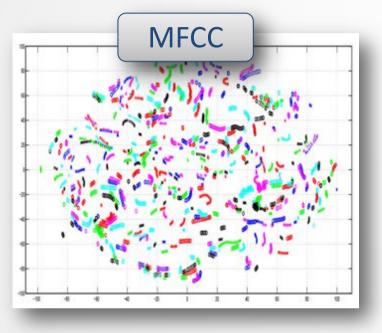


Replacing GMMs with DNNs

 The probability of a word sequence W comes *loosely* from P(X|W)



What are these DNNs learning?



8th layer

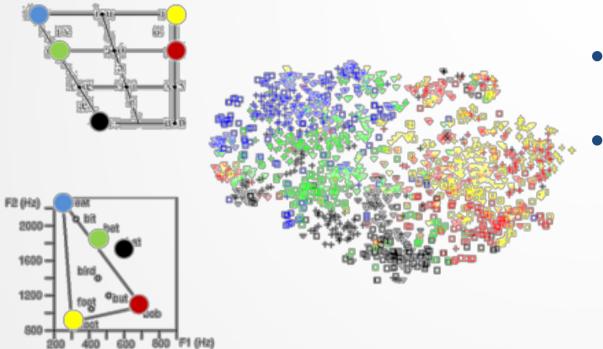
1st layer

- t-SNE visualizations in 2D.
- Deeper layers encode information about the speaker; more so given raw spectra than MFCCs (why?)

Mohamed, A., Hinton, G., & Penn, G. (2012). Understanding how deep belief networks perform acoustic modelling. In *ICASSP* (pp. 6–9).



What are these DNNs learning?



- t-SNE visualizations of hidden layer.
- Lower layers detect
 manner of articulation

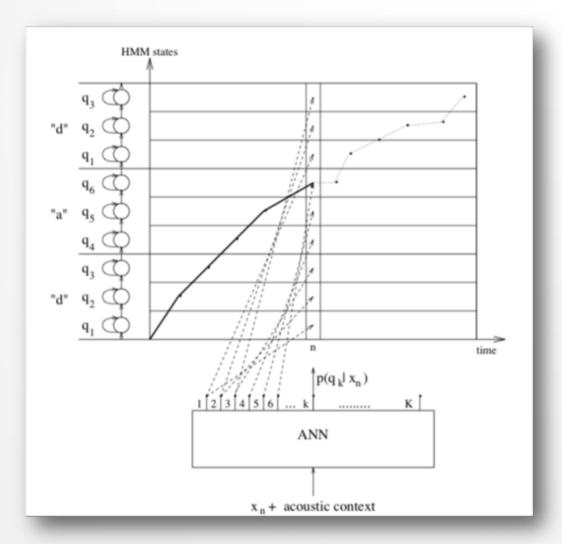
Figure 1: Multilingual BN features of five vowels from French (+), German (\Box) and Spanish (\bigtriangledown): /a/ (black), /i/ (blue), /e/ (green), /o/ (red), and /u/ (yellow)

Vu, N. T., Weiner, J., & Schultz, T. (2014). Investigating the learning effect of multilingual bottle-neck features for ASR. *Interspeech*, 825–829.



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Hybrid HMM and DNN

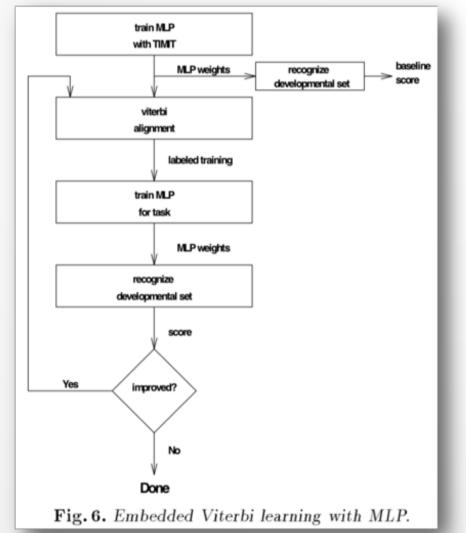


Bourlard H, Morgan. (1998) Hybrid HMM/ANN systems for speech recognition: Overview and new research directions. Adapt Process Seq Data Struct 1387:389–417. doi:10.1007/BFb0054006



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Hybrid HMM and DNN



Results: Often, it depends on the data and task.

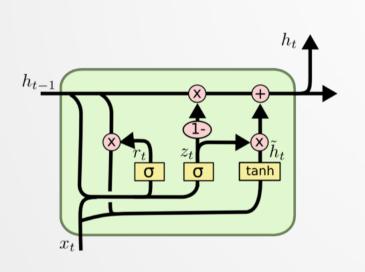
Sometimes, the hybrid approach is best. Sometimes, HMM-GMMs are still best, and sometimes...

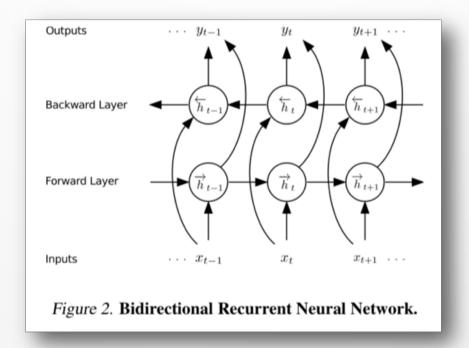
Bourlard H, Morgan. (1998) Hybrid HMM/ANN systems for speech recognition: Overview and new research directions. Adapt Process Seq Data Struct 1387:389–417. doi:10.1007/BFb0054006



End-to-end neural networks

- End-to-end neural network ASR often depends on two steps:
 - 1. A generalization of RNNs (e.g., GRUs) to be **bi-directional**. This allows us to use both Forward and Backward information, as in HMMs.





Graves A, Jaitly N. (2014) <u>Towards End-To-End Speech Recognition with Recurrent Neural Networks</u>. JMLR Workshop Conf Proc, 32:1764–1772.

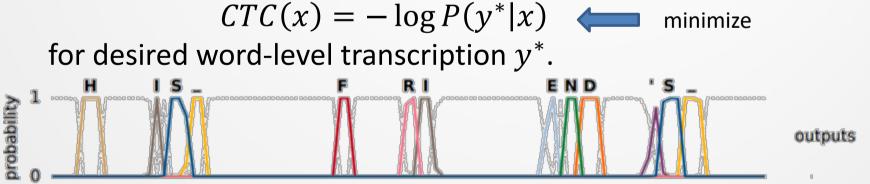


End-to-end neural networks

• Neural networks are typically trained at the **frame-level**.

- This requires a separate training target for every frame, which in turn requires the alignment between the audio and transcription sequences to be known.
- However, the alignment is only reliable once the classifier is trained.
- ..., the second step for end-to-end neural network ASR is:
 - 2. An objective function that allows sequence transcription *without* requiring prior alignment between the **input** and **target** sequences.

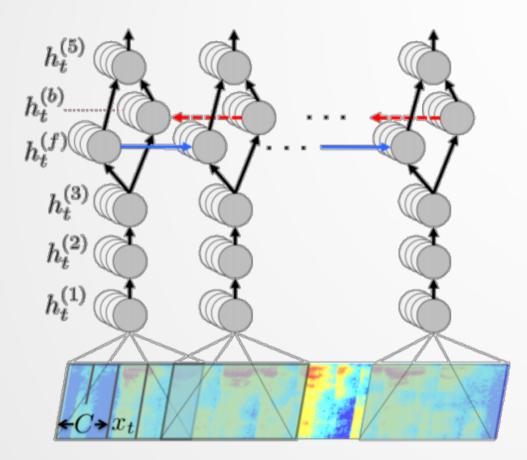
E.g., Connectionist Temporal Classification:



Graves A, Jaitly N. (2014) <u>Towards End-To-End Speech Recognition with Recurrent Neural Networks</u>. JMLR Workshop Conf Proc, 32:1764–1772.



End-to-end spectra-to-characters

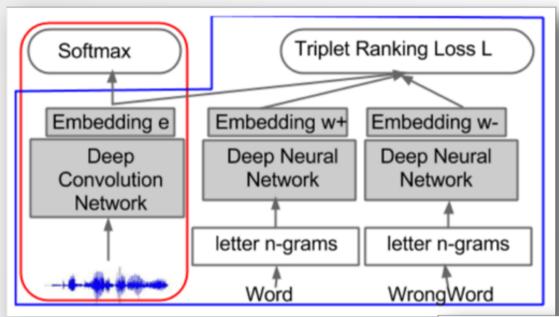


- Input: spectrograms
- Output: characters (incl. space and null characters)
- No phonemes or vocabulary means no OOV words.

A. Hannun, C. Case, J. Casper, B. Catanzaro, G. Diamos, E. Elsen, R. Prenger, S. Satheesh, S. Sengupta, A. Coates, A. Ng "Deep Speech: Scaling up end-to-end speech recognition", arXiv:1412.5567v2, 2014.



End-to-end acoustic/word hybrids



- Get word boundaries from some external tool.
- Train word/characters and acoustics simultaneously.
- Obtain up to 0.11% improvement in error rates

Table 2: Word Error Rates for the three compared models, with two different values of the beam search parameter.

Model	WER			
Woder	beam=11	beam=15		
Baseline	10.16	9.70		
Word embedding model	11.2	11.1		
Combination	10.07	9.59		

Bengio, S., & Heigold, G. (2014). Word Embeddings for Speech Recognition, Interspeech



End-to-end neural networks

Table 1. **Wall Street Journal Results.** All scores are word error rate/character error rate (where known) on the evaluation set. 'LM' is the Language model used for decoding. '14 Hr' and '81 Hr' refer to the amount of data used for training.

System	LM	14 HR	81 HR
RNN-CTC	None	74.2/30.9	30.1/9.2
RNN-CTC	DICTIONARY	69.2/30.0	24.0/8.0
RNN-CTC	Monogram	25.8	15.8
RNN-CTC	BIGRAM	15.5	10.4
RNN-CTC	TRIGRAM	13.5	8.7
RNN-WER	None	74.5/31.3	27.3/8.4
RNN-WER	DICTIONARY	69.7/31.0	21.9/7.3
RNN-WER	MONOGRAM	26.0	15.2
RNN-WER	BIGRAM	15.3	9.8
RNN-WER	TRIGRAM	13.5	8.2
BASELINE	None	—	—
BASELINE	DICTIONARY	56.1	51.1
BASELINE	MONOGRAM	23.4	19.9
BASELINE	BIGRAM	11.6	9.4
BASELINE	TRIGRAM	9.4	7.8
COMBINATION	TRIGRAM	—	6.7

Here, **lower** scores are **better**, because they are **error rates**.

But how to compute those error rates?

DNN/HMM

Graves A, Jaitly N. (2014) <u>Towards End-To-End Speech Recognition with Recurrent Neural Networks</u>. JMLR Workshop Conf Proc, 32:1764–1772.



EVALUATING SPEECH RECOGNITION



Evaluating ASR accuracy

- How can you tell how good an ASR system at recognizing speech?
 - E.g., if somebody said Reference: <u>how to recognize speech</u> but an ASR system heard Hypothesis: <u>how to wreck a nice beach</u> how do we quantify the error?
- One measure is word accuracy: #CorrectWords/#ReferenceWords
 - E.g., 2/4, above
 - This runs into problems similar to those we saw with SMT.
 - E.g., the hypothesis 'how to recognize speech boing boing boing boing 'has 100% accuracy by this measure.
 - Normalizing by #HypothesisWords also has problems...



Word-error rates (WER)

- ASR enthusiasts are often concerned with word-error rate (WER), which counts different kinds of errors that can be made by ASR at the word-level.
 - Substitution error: One word being mistook for another
 - Deletion error:
 - Insertion error:

e.g., 'shift' given 'ship'
An input word that is 'skipped'
e.g. 'I Torgo' given 'I am Torgo'
A 'hallucinated' word that was not in the input.
e.g., 'This Norwegian parrot is no more'

given 'This parrot is no more'



Evaluating ASR accuracy

- But how to decide which errors are of each type?
- E.g., Reference: how to recognize speech Hypothesis: how to wreck a nice beach,
- It's not so simple: 'speech' seems to be mistaken for 'beach', except the /s/ phoneme is incorporated into the preceding hypothesis word, 'nice' (/n ay s/).
 - Here, 'recognize' seems to be mistaken for 'wreck a nice'
 - Are each of 'wreck a nice' substitutions of 'recognize'?
 - Is 'wreck' a substitution for 'recognize'?
 - If so, the words 'a' and 'nice' must be insertions.
 - Is 'nice' a substitution for 'recognize'?
 - If so, the words 'wreck' and 'a' must be insertions.



- In practice, ASR people are often more concerned with overall WER, and don't care about how those errors are partitioned.
 - E.g., 3 substitution errors are 'equivalent' to 1 substitution plus 2 insertions.
- The Levenshtein distance is a straightforward algorithm based on dynamic programming that allows us to compute overall WER.



Allocate matrix R[n + 1, m + 1] // where *n* is the number of reference words // and *m* is the number of hypothesis words Initialize $R[0,0] \coloneqq 0$, and $R[i,j] \coloneqq \infty$ for all other i = 0 or j = 0for $i \coloneqq 1..n$ // #ReferenceWords for $j \coloneqq 1..m$ // #Hypothesis words $R[i,j] \coloneqq \min(\begin{array}{c} R[i-1,j]+1, \\ R[i-1,j-1], \\ (the j^{th} reference word and \\ (the j^{th} hypothesis word match \\ R[i,j-1]+1, \\ (the j^{th} hypothesis word match \\ R[i,j-1]+1, \\ (the j^{th} negret on match \\ ($



Levenshtein distance – initialization

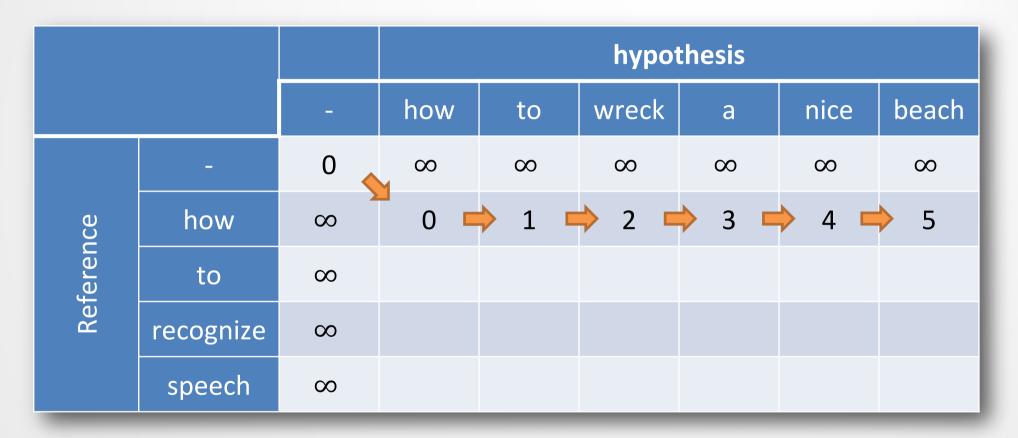
			hypothesis						
		-	how	to	wreck	а	nice	beach	
Reference	-	0	∞	∞	∞	∞	∞	∞	
	how	∞							
	to	∞							
	recognize	∞							
	speech	∞							

The value at cell (i, j) is the **minimum** number of **errors** necessary to align i with j.



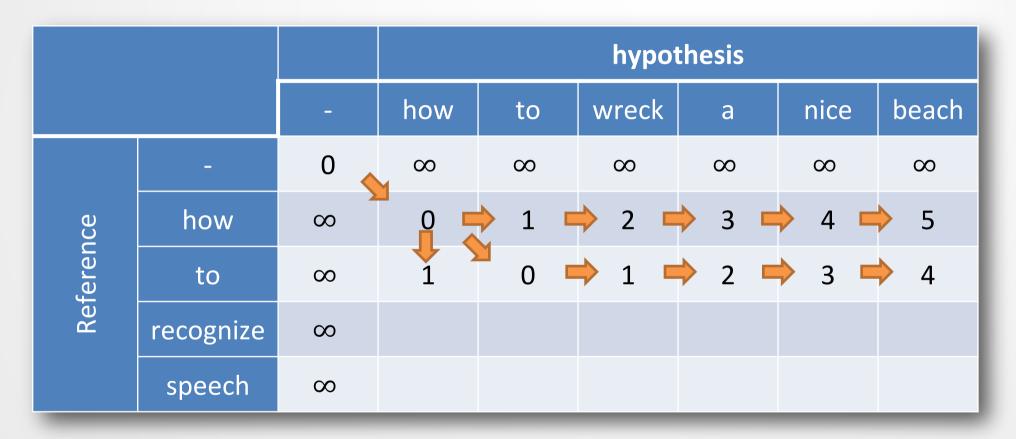
			hypothesis						
		-	how	to	wreck	а	nice	beach	
Reference	-	0	∞	∞	∞	∞	∞	∞	
	how	8	0						
	to	∞							
	recognize	∞							
	speech	∞							

- $R[1,1] = \min(\infty + 1, (0), \infty + 1) = 0$ (match)
- We put a little arrow in place to indicate the choice.
 - 'Arrows' are normally stored in a backtrace matrix.



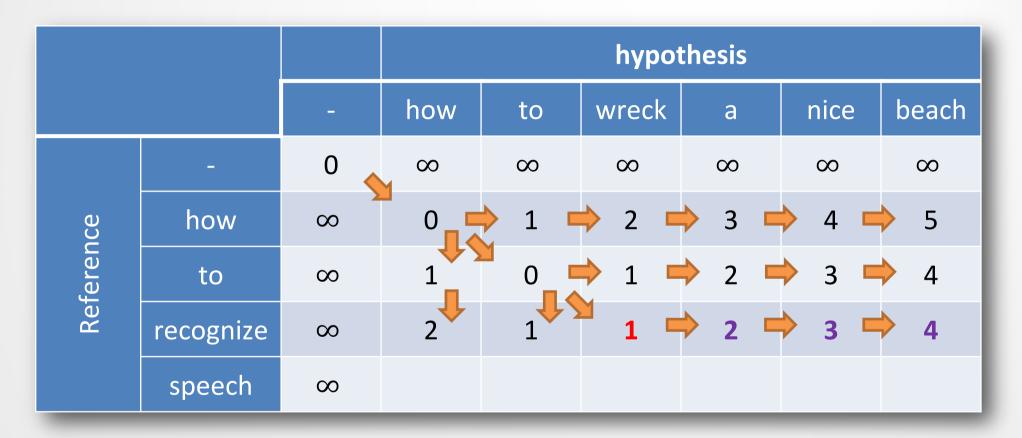
- We continue along for the first reference word...
 - These are all **insertion** errors





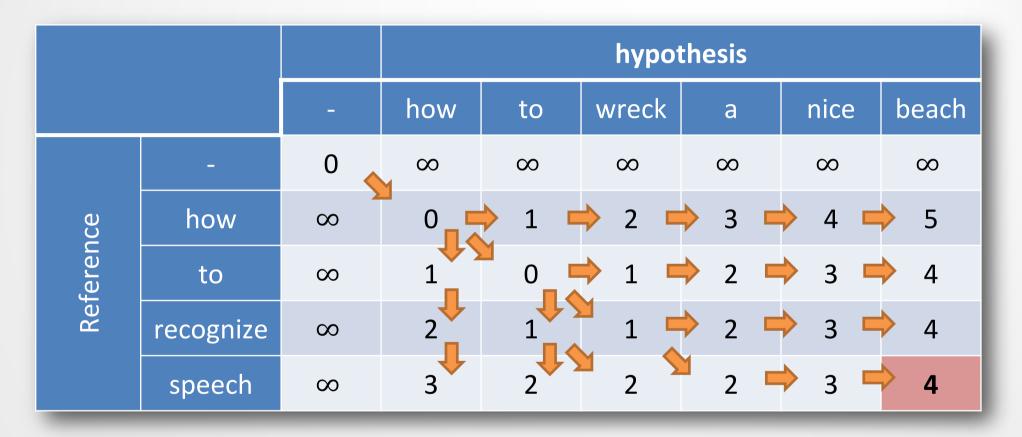
And onto the second reference word



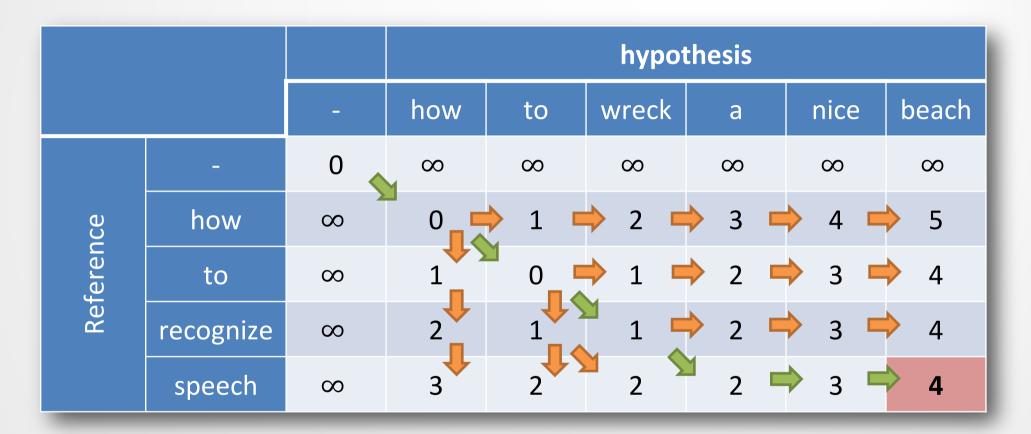


- Since *recognize* ≠ *wreck*, we have a **substitution** error.
- At some points, you have >1 possible path as indicated.
 - We can prioritize types of errors arbitrarily.



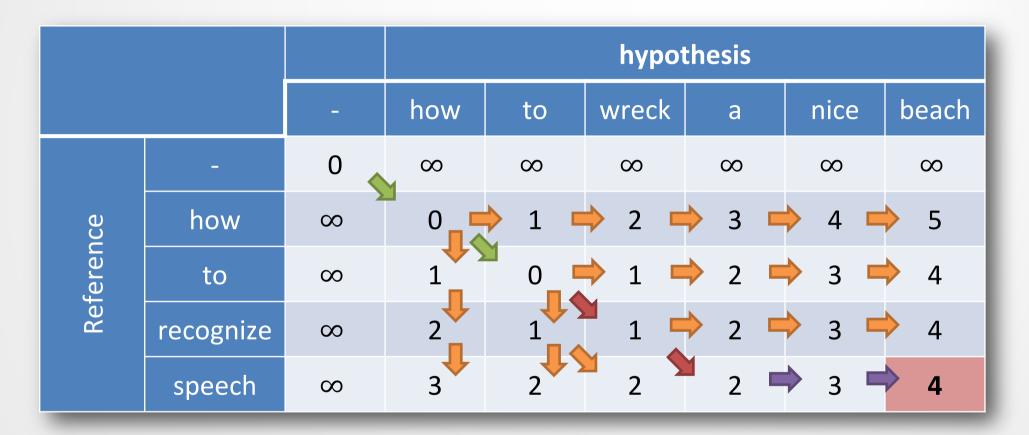


- And we finish the grid.
- There are R[n, m] = 4 word errors and a WER of 4/4 = 100%.
 - WER can be greater than 100% (relative to the reference).



• If we want, we can **backtrack** using our arrows to find the proportion of substitution, deletion, and insertion errors.





- Here, we estimate 2 substitution errors and 2 insertion errors.
- Arrows can be encoded within a special **backtrace** matrix.





NEURAL SPEECH RECOGNITION (SLIGHT RETURN)

Speaker adaptation

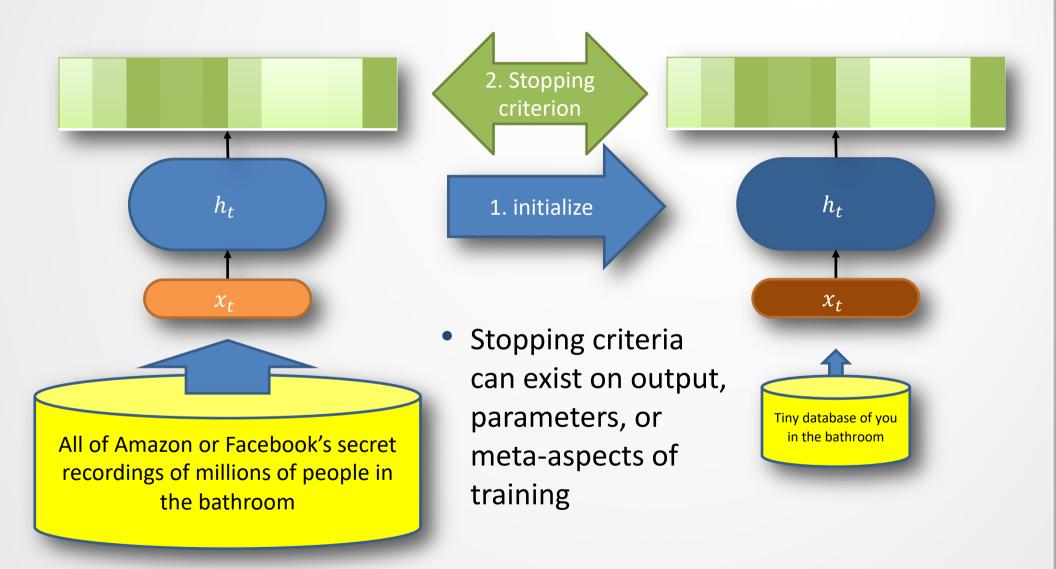
- Given a neural ASR system trained with many speakers, we want to adapt to the voice of a new individual.
- We know how to do this with HMMs
 - (e.g., with interpolation, or (aside) with MAP or MLLR training).
- DNNs need *lots* of data to be useful, though...
 - **Conservative**: re-train whole DNN, with some constraints
 - Transformative:

Speaker-aware:

only retrain one layer (or a few) do not really train the parameters

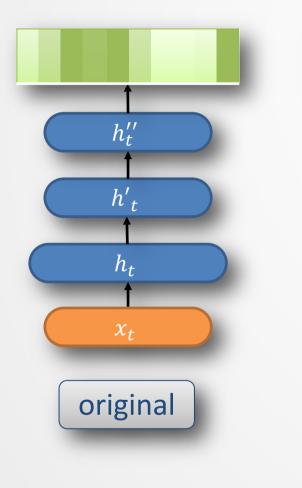


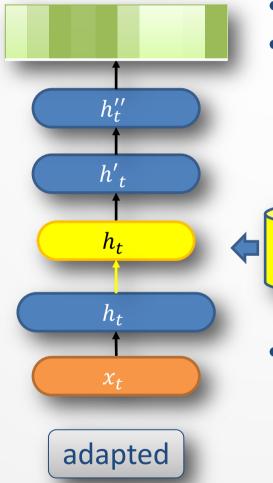
Conservative speaker adaptation





Transformative speaker adaptation





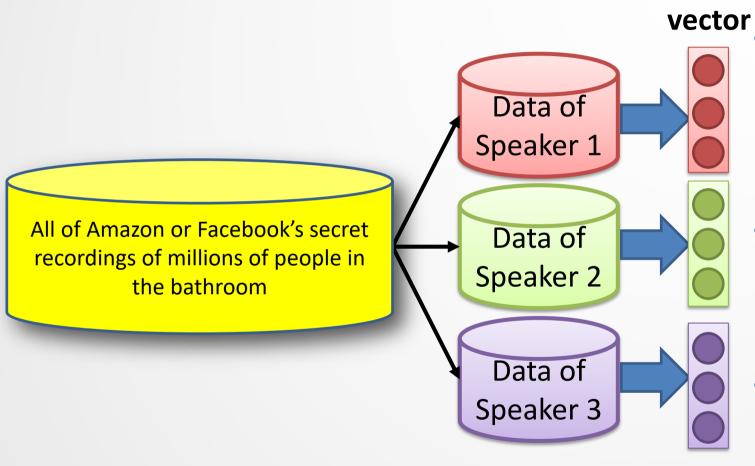
- Insert a new layer.
- Keeping all other parameters fixed, train the new ones to normalize speaker information.

 There are many alternatives...

Tiny database of you in the bathroom



Speaker-aware training



Speaker vector Fixed length low dimension vectors, obtained in a variety of ways.

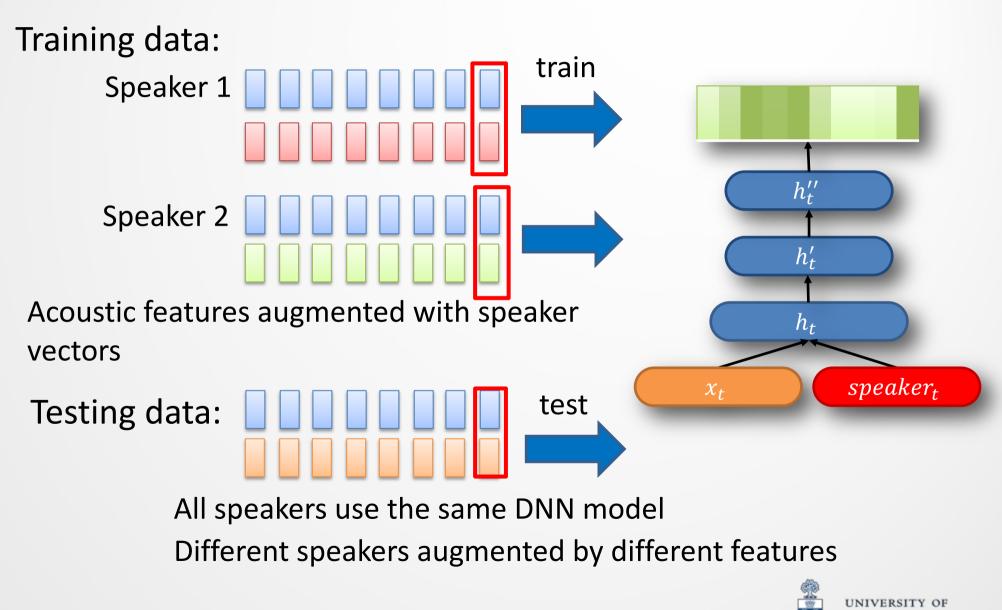
Note we can
 segment things by
 recording device,
 noise, etc.

This can be used to remove the channel effect.

Senior, A., & Lopez-Moreno, I. (2014). Improving DNN speaker independence with I-vector inputs. ICASSP, 225–229. <u>https://doi.org/10.1109/ICASSP.2014.6853591</u>



Speaker-aware training

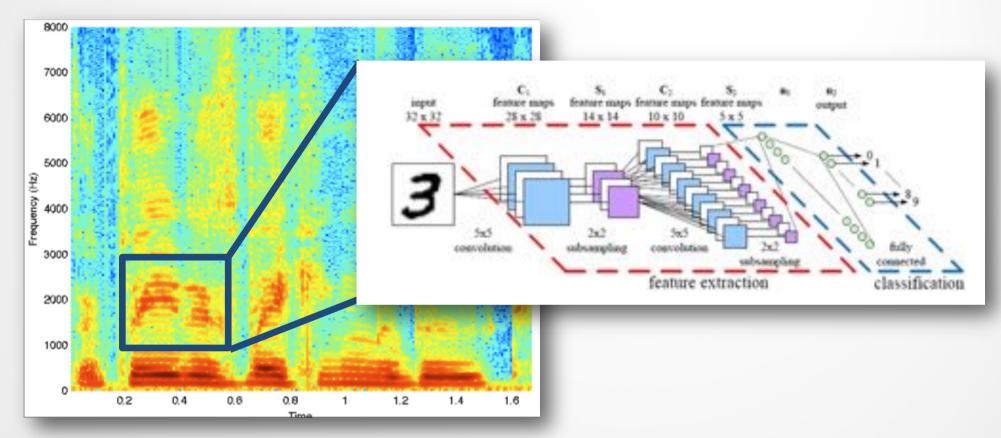


Aside – the open-source Kaldi ASR

- Kaldi is the *de-facto* open-source ASR toolkit: <u>http://kaldi-asr.org</u>
 - It has pretrained <u>models</u>, including the ASpIRE chain model trained on Fisher English, augmented with impulse responses and noises to create multi-condition training.
 - My favourite incarnation uses I-Vectors to account for the speaker.
 - It often (anecdotally) performs better than Google's <u>SpeechAPI</u>.
 - It is originally in C++, but a wrapper (<u>PyTorch-Kaldi</u>) exists in the much easier Python.



Aside – convolutional neural networks



• Spectrograms are kind of images, so lets use the kinds of neural networks used in computer vision.



Next...

- We've seen how to:
 - extract useful speech features with Mel-frequency cepstral coefficients.
 - cluster multi-modal speech data with Gaussian mixture models.
 - recognize speech with hidden Markov models and neural networks.
 - evaluate ASR performance with Levenshtein distance.
- Next, we'll see how to synthesize artificial speech.

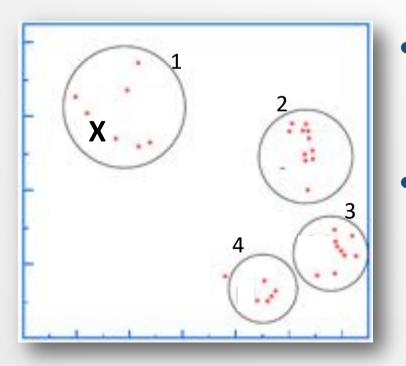


APPENDIX: CLUSTERING (EVERYTHING THAT FOLLOWS IS AN <u>ASIDE</u>. NOT ON THE EXAM. (THINGS NOT ON THE EXAM MAY BE USEFUL OR AT LEAST INTERESTING))



Clustering

- Quantization involves turning possibly multi-variate and continuous representations into univariate discrete symbols.
 - Reduced storage and computation costs.
 - Potentially tremendous loss of information.



- Observation **X** is in Cluster One, so we replace it with **1**.
- Clustering is unsupervised learning.
 - Number and form of clusters often unknown.



Aspects of clustering

- What **defines** a particular cluster?
 - Is there some **prototype** representing each cluster?
- What defines **membership** in a cluster?
 - Usually, some distance metric d(x, y) (e.g., Euclidean distance).
- How well do clusters represent unseen data?
 - How is a new point assigned to a cluster?
 - How do we modify that cluster as a result?



K-means clustering

- Used to group data into K clusters, $\{C_1, \dots, C_K\}$.
- Each cluster is represented by the mean of its assigned data.
 - (sometimes it's called the cluster's *centroid*).
- Iterative algorithm converges to local optimum:
 - **1.** Select *K* initial cluster means $\{\mu_1, ..., \mu_K\}$ from among data points.
 - **2. Until** (stopping criterion),
 - a) Assign each data sample to closest cluster

 $x \in C_i$ if $d(x, \mu_i) \le d(x, \mu_j), \quad \forall i \ne j$

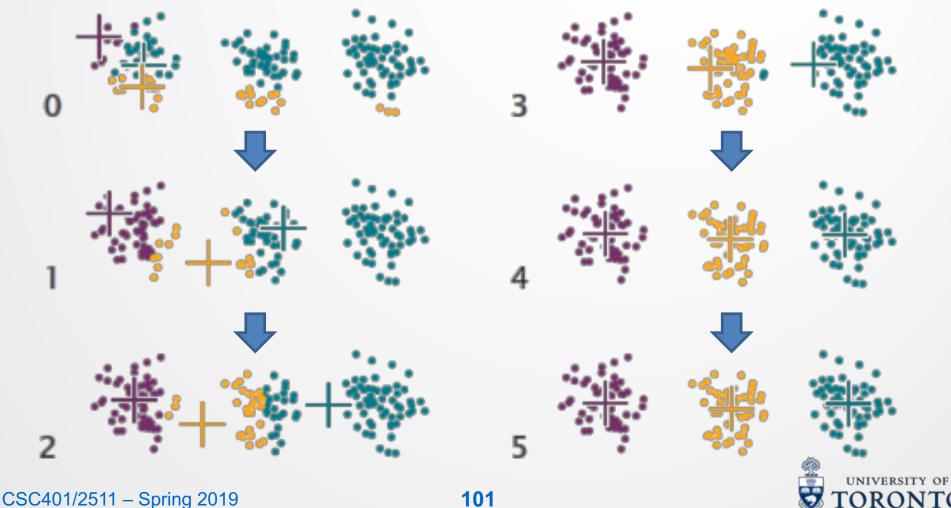
b) Update *K* means from assigned samples

$$\mu_i = E(x) \forall x \in C_i, \quad 1 \le i \le K$$



K-means example (K = 3)

- Initialize with a random selection of 3 data samples.
- Euclidean distance metric $d(x, \mu)$



K-means stopping condition

• The total distortion, \mathcal{D} , is the sum of squared error,

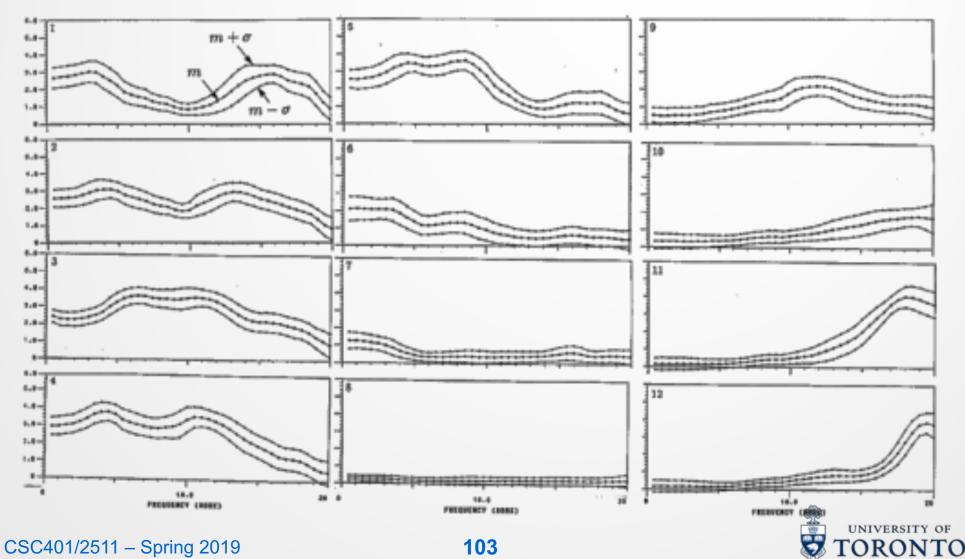
$$\mathcal{D} = \sum_{i=1}^{K} \sum_{x \in C_i} ||x - \mu_i||^2$$

- \mathcal{D} decreases between n^{th} and $(n+1)^{th}$ iteration.
- We can stop training when \mathcal{D} falls below some threshold \mathcal{T} . $1 - \frac{\mathcal{D}(n+1)}{\mathcal{D}(n)} < \mathcal{T}$



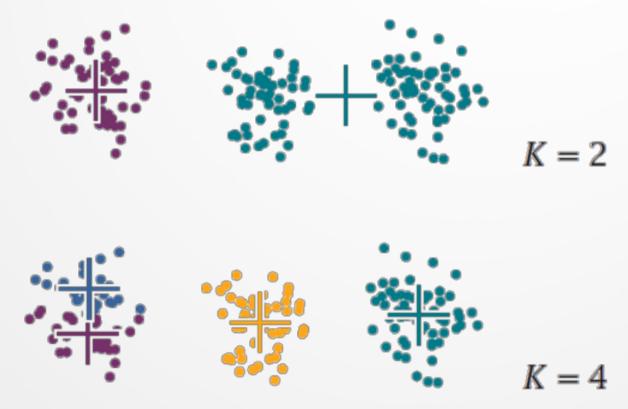
Acoustic clustering example

• 12 clusters of spectra, after training.



Number of clusters

- The number of true clusters is unknown.
- We can iterate through various values of *K*.
 - As K approaches the size of the data, \mathcal{D} approaches 0...





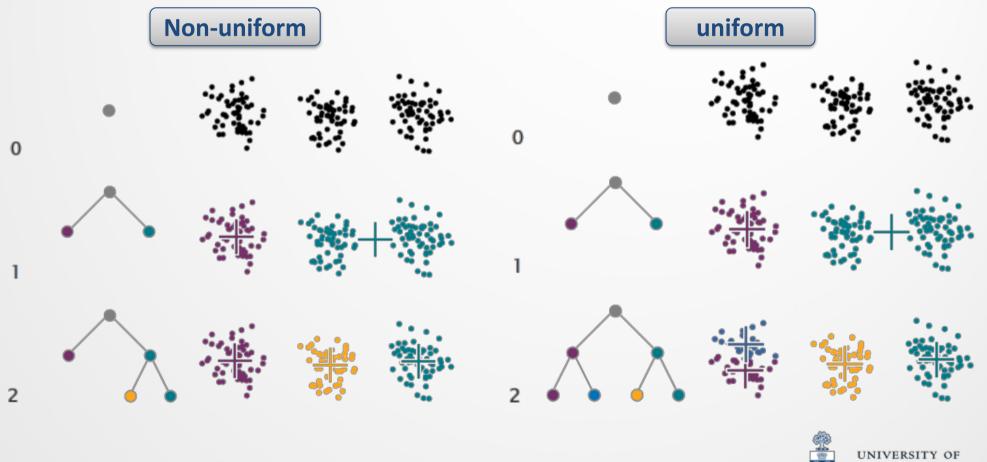
Hierarchical clustering

- Hierarchical clustering clusters data into hierarchical 'class' structures.
- Two types: top-down (divisive) or bottom-up (agglomerative).
- Often based on greedy formulations.
- Hierarchical structure can be used for hypothesizing classes.



Divisive clustering

 Creates hierarchy by successively splitting clusters into smaller groups.

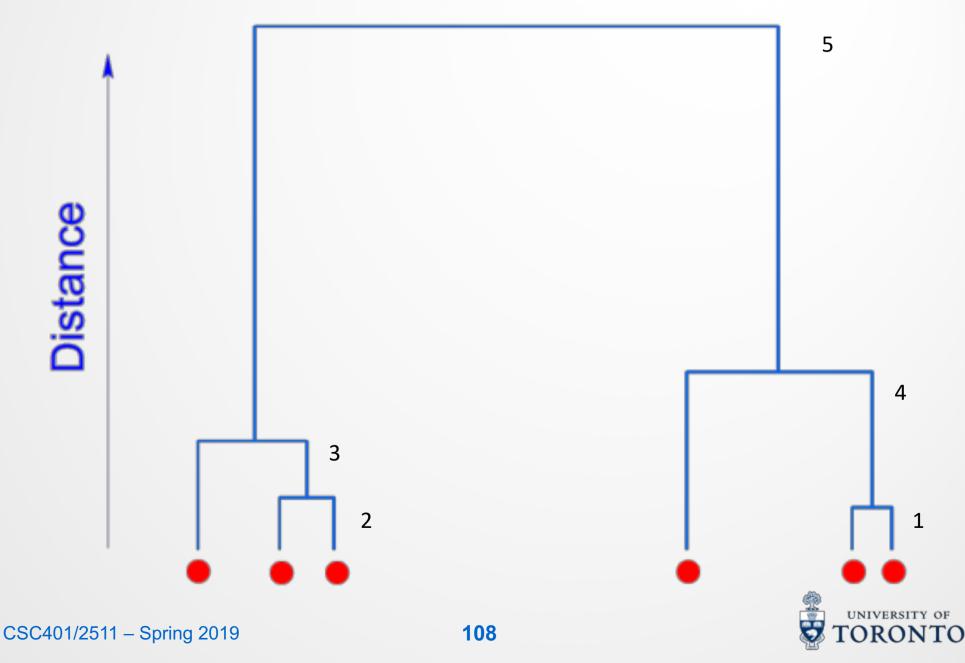


Agglomerative clustering

- Agglomerative clustering starts with N 'seed' clusters and iteratively combines these into a hierarchy.
- On each iteration, the **two most similar** clusters are **merged** together to form a new **meta-cluster**.
- After N 1 iterations, the hierarchy is complete.
- Often, when the similarity scores of new meta-clusters are tracked, the resulting graph (i.e., dendogram) can yield insight into the natural grouping of data.

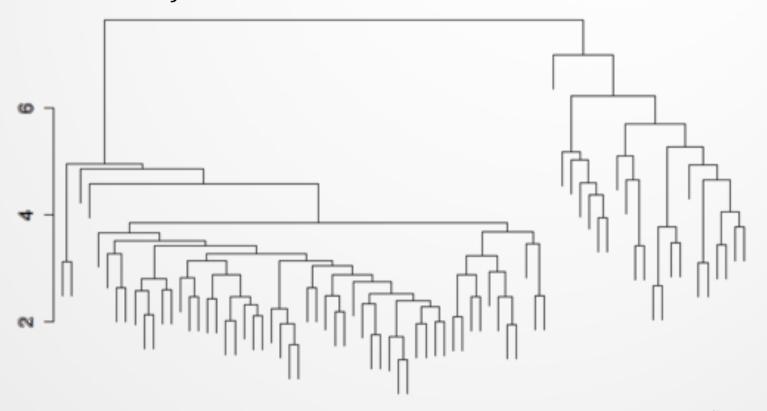


Dendogram example



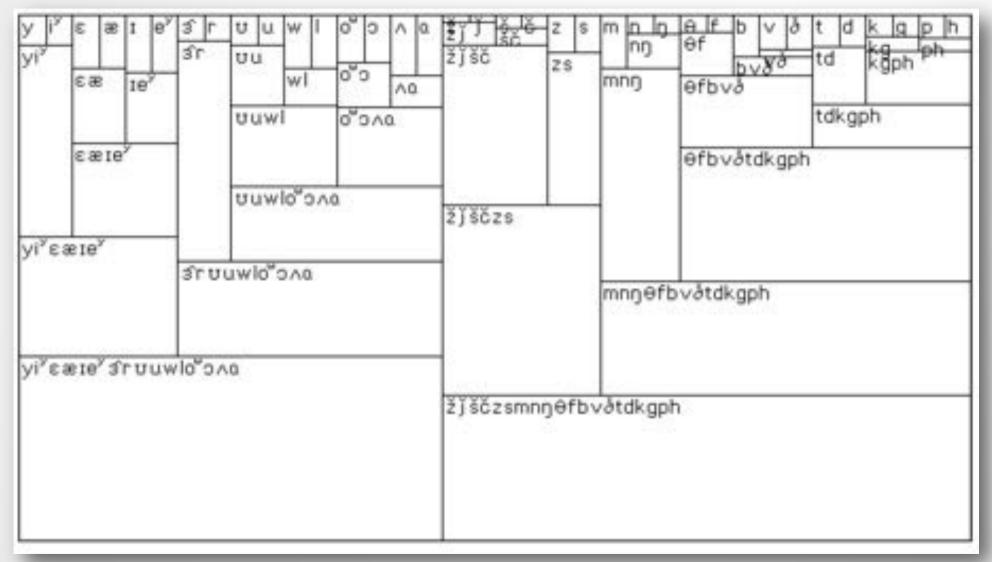
Speaker clustering

- 23 female and 53 male speakers from TIMIT.
- Data are vectors of average F1 and F2 for 9 vowels.
- Distance $d(C_i, C_j)$ is average of distances between members.





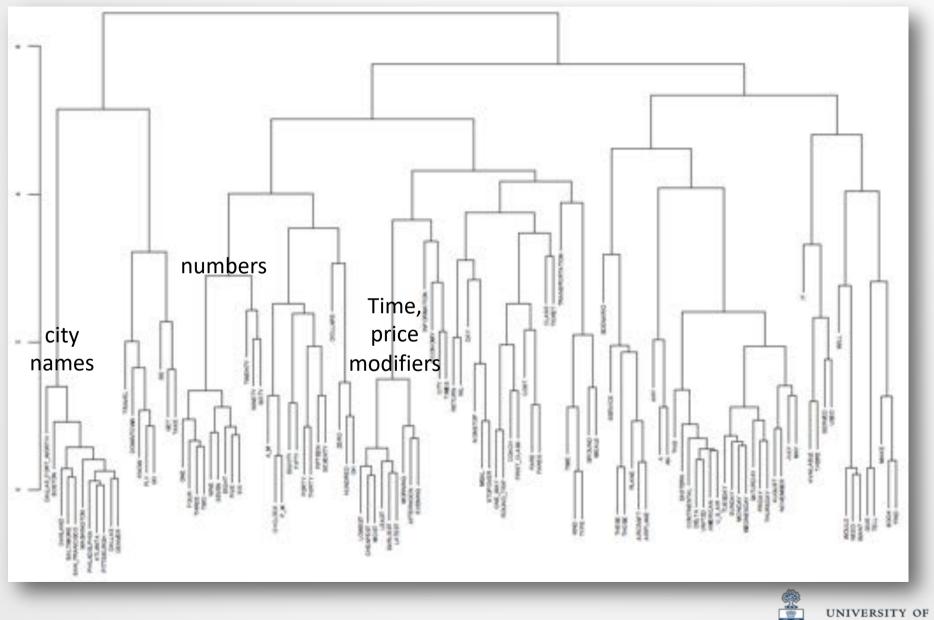
Acoustic-phonetic hierarchy



(this is basically an upside-down dendogram)



Word clustering



ORONTO