

## String Edit Distance

$D(i, j) = d(i, j) + \min [ D(i-1, j), D(i-1, j-1), D(i, j-1) ]$  for  $i=1.....M$  and  $j = 1.....N$   
 $Path(i, j) = \arg \min [ D(i-1, j), D(i-1, j-1), D(i, j-1) ]$

Initial condition path starts at  $D(1,1)$

Final condition path ends at  $D(M,N)$

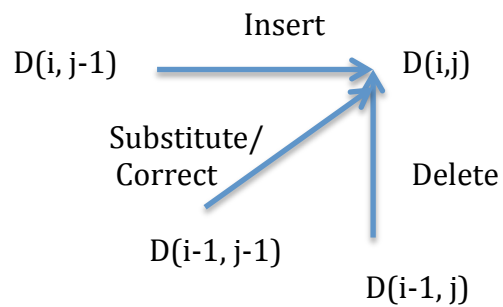
```
if str(i) = str(j)
```

```
    d(i,j) = 0
```

```
else
```

```
    d(i,j) = 1
```

```
end if
```



Backtrack the path to find the number of insertions, deletion, and substitution.

K	Inf					
R	Inf					
A	Inf					
P	Inf					
	0	Inf	Inf	Inf	Inf	Inf
		S	P	A	K	E

## Word Error Rate

Reference transcript: *she had a dark suit in greasy water all year*

ASR output: *he had a dark grey suit in water all there*

WER = (# Ins + # Del + # Sub)/(# of words in reference transcript)

<i>year</i>	Inf										
<i>all</i>	Inf										
<i>water</i>	Inf										
<i>greasy</i>	Inf										
<i>in</i>	Inf										
<i>suit</i>	Inf										
<i>dark</i>	Inf										
<i>a</i>	Inf										
<i>had</i>	Inf										
<i>she</i>	Inf										
	0	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf
		<i>he</i>	<i>had</i>	<i>a</i>	<i>dark</i>	<i>grey</i>	<i>suit</i>	<i>in</i>	<i>water</i>	<i>all</i>	<i>there</i>

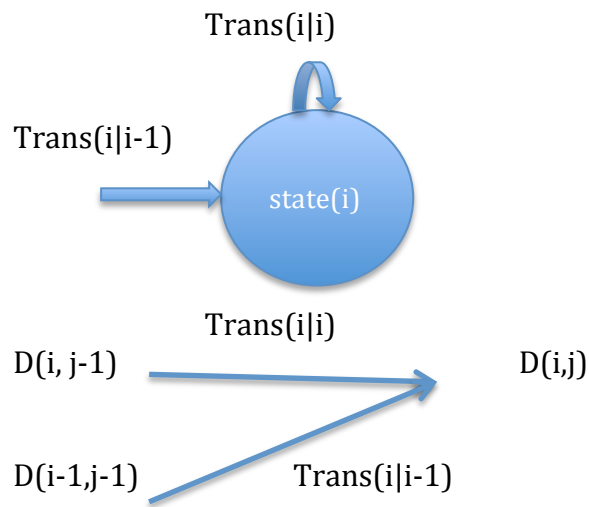
**Computation of log-likelihood of a HMM given an Observation Sequence**  
**using Viterbi algorithm**

$$D(i, j) = d(i, j) + \min [ (D(i-1, j-1) + \text{Trans}(i|i-1)), (D(i, j-1) + \text{Trans}(i|i)) ]$$
$$\text{Path}(i, j) = \arg \min [ (D(i-1, j-1) + \text{Trans}(i|i-1)), (D(i, j-1) + \text{Trans}(i|i)) ]$$

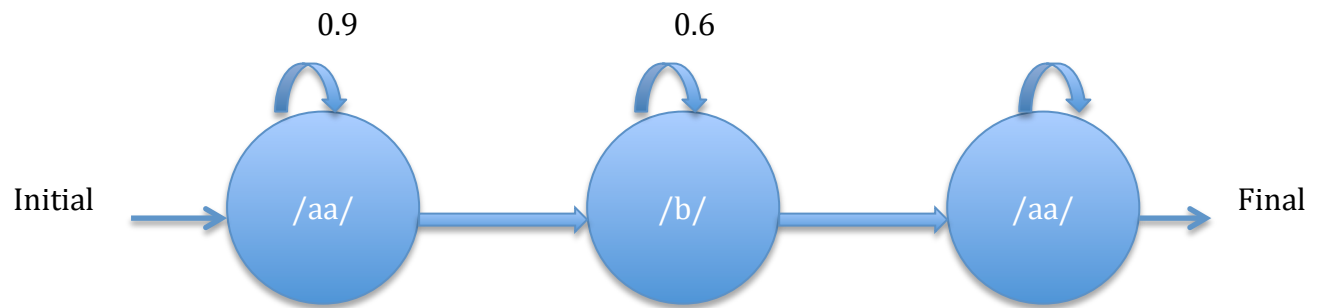
$$d(i, j) = -\log(P(\text{obs}(j)|\text{state}(i)))$$

$$\text{Trans}(i|i-1) = -\log(P(\text{state}(i)|\text{state}(i-1)))$$

NOTE: Local constraints are based on HMM topology



Example with single dimension observation



Initial and Final are non-emitting states

State /aa/:

mean = 15.1

standard deviation = 3.2

State /b/:

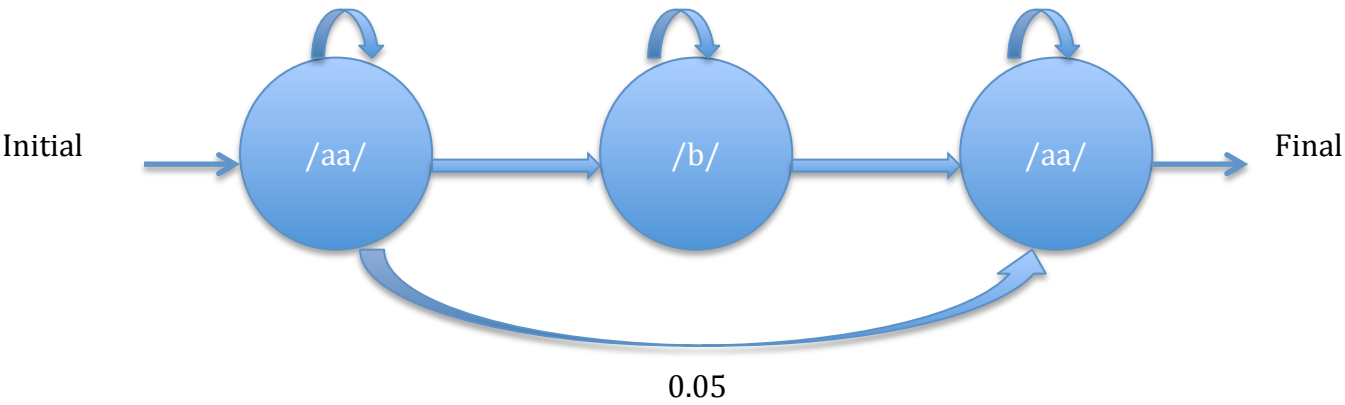
mean = 10.3

standard deviation = 4.4

Observation sequence: 16.0, 13.7, 12.0, 9.8, 10.0, 11.8, 14.2, 17.2, 12.5

/aa/	Inf									
/b/	Inf									
/aa/	Inf									
	0.0	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf
		16.0	13.7	12.0	9.8	10.0	11.8	14.2	17.2	12.5

Compute the log-likelihood for the following HMM topology. Note down the key changes in the computation that are to be made w.r.t the previous topology.



/aa/	Inf									
/b/	Inf									
/aa/	Inf									
	0.0	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf	Inf
		16.0	13.7	12.0	9.8	10.0	11.8	14.2	17.2	12.5