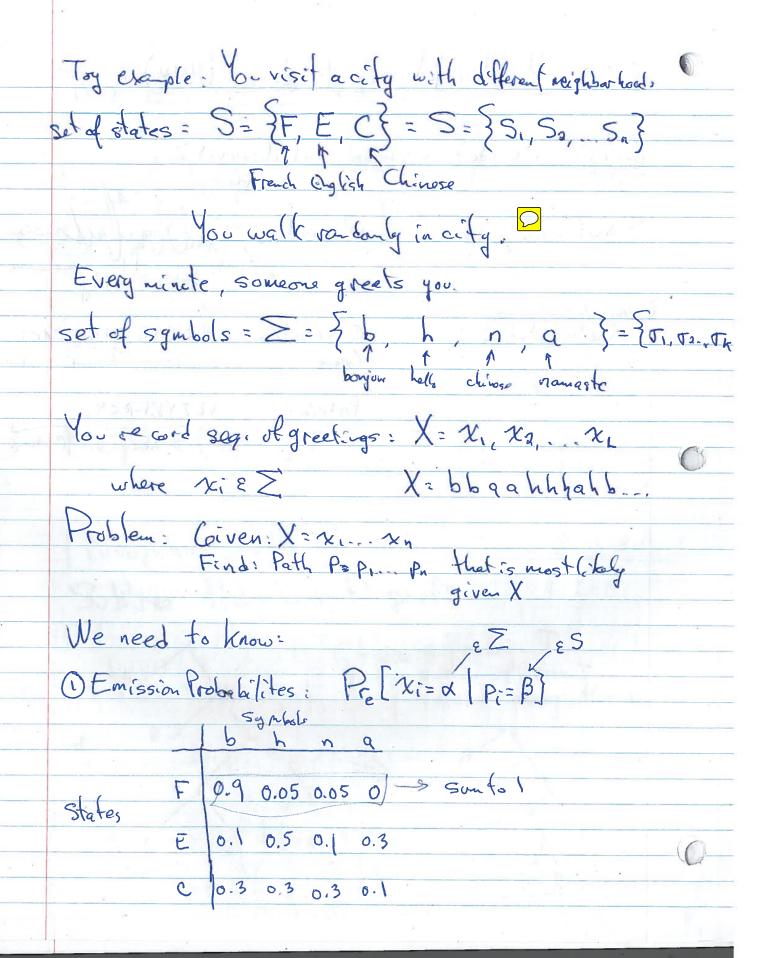
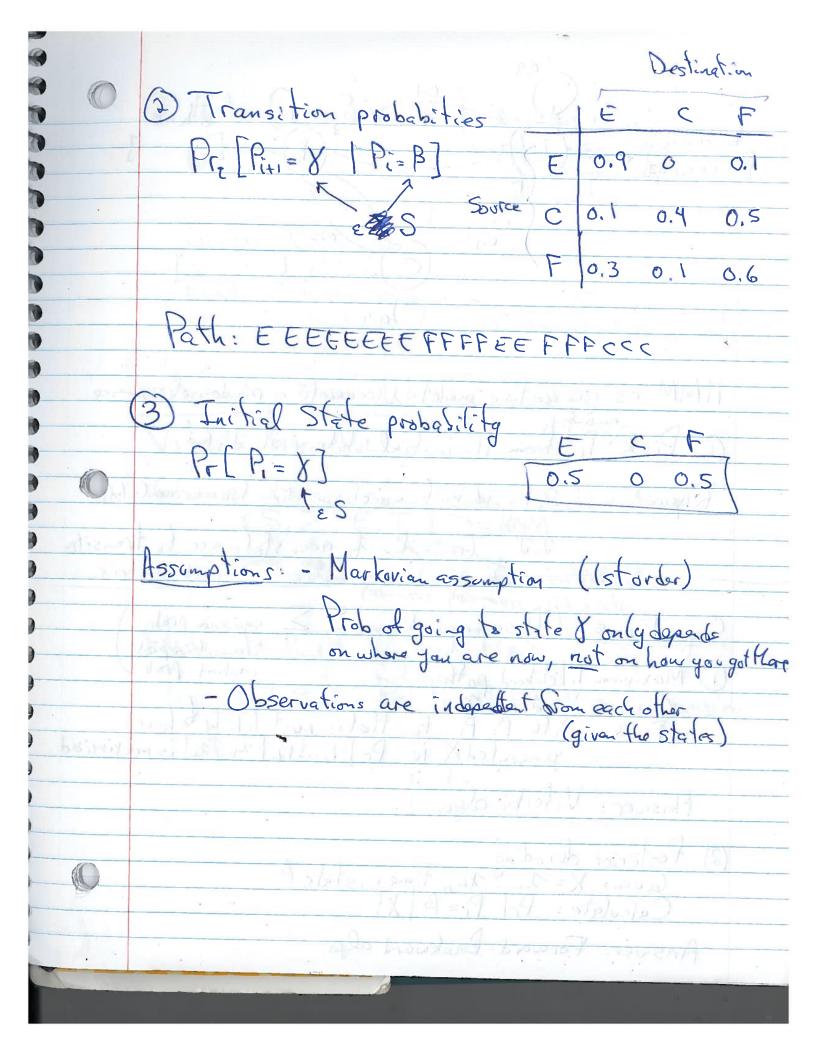
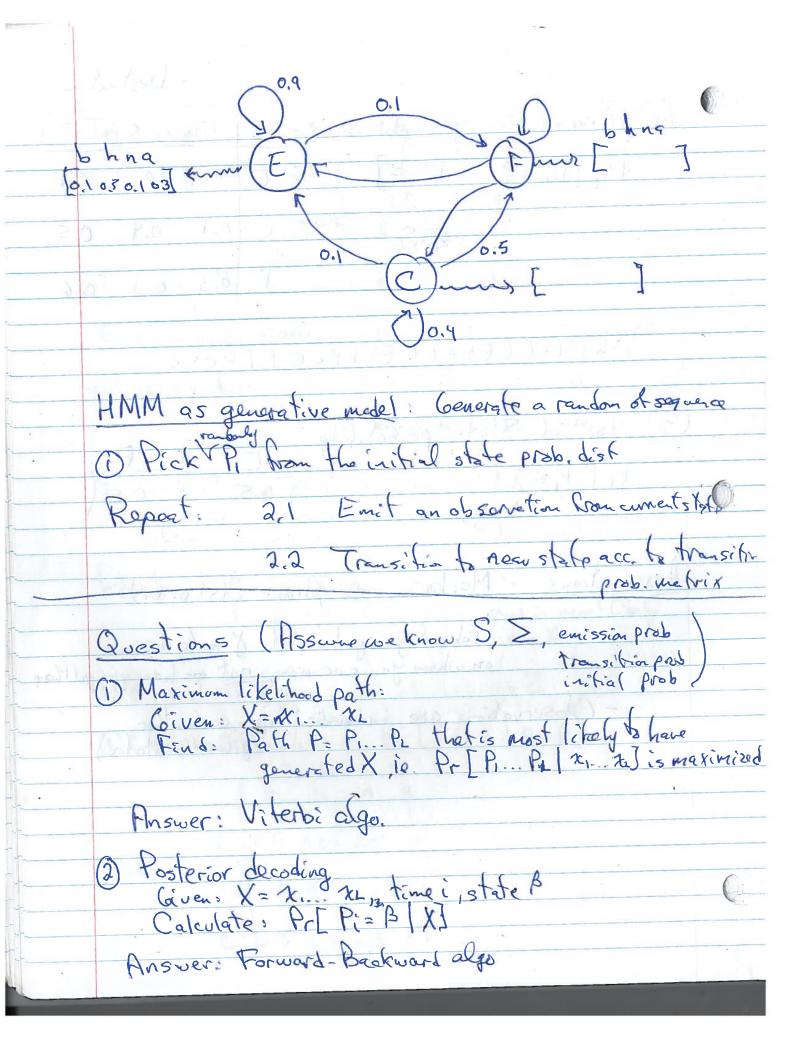
Hidden Markov Models (HMM) 10 Speech recognition: Observation: Hello Biological sequence Come samo fation Observation: ACTAGTAGGTAC Profeir: VLIYVPRCY, d-helin B-sh C







Estimation Problem: Assura we know S, E! but not emission prob transition prob Emission proils Secret Hat Pr[X E, T, I] is max Civen: X7 xi (seq. of Lobsenations) = HMM alphabet emission transition initial state probredix probretix probretix Find: P= P. Pe , where Pies stores such that PrePr. P. Xexi xn Jis maiximized P. (P. . P. . X. . X.) 10

Question 1: How to calculate Po[A. P. 1X=X1 XL] ? for a given path P=PiPi
for a given path P=Pu.P.
Pr[P. P. X=x1 = Pr[P1 - P. 1 X=x1 - x4]
Indopendent of Pr[X=x1xn] Path P
Pr[P P. 1 X=xxx] = Pr[X=xxx[PP.]. Pr[PP.]
Pr[P. P. 1 X=2 xx] = Pr[X=2xx[PP.]. Pr[PP.]. Pr[PP.]. Pr[PP.]. Pr[PP.].
Duestion 2: How to find Pst. Pr. R. M. m. J.
Viter bi algorithm
Define V(B,i) = Prob of the most lakely pathol
Define $V(\beta, i) = Prob of the most likely pathof length i, given observation XI Xi, ES EST If assuming path ends instate B$
= max { Pr[PPi, xxi]}
PiPi Whore Pi=B

Dyjana

395%

