

# CM224 HW 7 Solution

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**Problem 1.** Viterbi Algorithm

**Sub-Problem 1a.** Most likely population ancestry path with haplotype 10111?

**Answer 1a.** Most likely population ancestry path is 11122

**Solution 1a.** Since either population is equivalent and  $h_1 = 1$  we get,

$$C(1, 1) = 0.5 * \xi(h_1, e_{11}) = 0.5 * e_{11} = 0.5 * 0.3 = 0.15$$

$$C(1, 2) = 0.5 * \xi(h_1, e_{12}) = 0.5 * e_{12} = 0.5 * 0.2 = 0.1$$

For the second state  $s = 2$ , we have  $h_2 = 0$  we get,

$$C(s + 1, j) = \max(C(s, i)\delta_{ij}\xi(h_{s+1}, e_{s+1,j}))$$

$$C(2, 1) = \max(C(1, i)\delta_{i1}\xi(h_2, e_{21}))$$

$$C(2, 1) = \max(C(1, 1)\delta_{11}(1 - e_{21}), C(1, 2)\delta_{21}(1 - e_{21}))$$

$$C(2, 1) = \max(0.15 * 0.5 * 0.3, 0.1 * 0.4 * 0.3) = 0.0225$$

$$C(2, 2) = \max(C(1, i)\delta_{i2}\xi(h_2, e_{22}))$$

$$C(2, 2) = \max(C(1, 1)\delta_{12}(1 - e_{22}), C(1, 2)\delta_{22}(1 - e_{22}))$$

$$C(2, 2) = \max(0.15 * 0.5 * 0.2, 0.1 * 0.6 * 0.2) = 0.015$$

For the third state  $s = 3$ , we have  $h_3 = 1$  we get,

$$C(3, 1) = \max(C(2, i)\delta_{i1}\xi(h_3, e_{31}))$$

$$C(3, 1) = \max(C(2, 1)\delta_{11}e_{31}, C(2, 2)\delta_{21}e_{31})$$

$$C(3, 1) = \max(0.0225 * 0.5 * 0.3, 0.015 * 0.4 * 0.3) = 0.003375$$

$$C(3, 2) = \max(C(2, i)\delta_{i2}\xi(h_3, e_{32}))$$

$$C(3, 2) = \max(C(2, 1)\delta_{12}e_{32}, C(2, 2)\delta_{22}e_{32})$$

$$C(3, 2) = \max(0.0225 * 0.5 * 0.2, 0.015 * 0.6 * 0.2) = 0.00225$$

For the fourth state  $s = 4$ , we have  $h_4 = 1$  we get,

$$C(4, 1) = \max(C(3, i)\delta_{i1}\xi(h_2, e_{41}))$$

$$C(4, 1) = \max(C(3, 1)\delta_{11}e_{41}, C(3, 2)\delta_{21}e_{41})$$

$$C(4, 1) = \max(0.003375 * 0.5 * 0.2, 0.00225 * 0.4 * 0.2) = 0.0003375$$

$$C(4, 2) = \max(C(3, i)\delta_{i2}\xi(h_2, e_{42}))$$

$$C(4, 2) = \max(C(3, 1)\delta_{12}e_{42}, C(3, 2)\delta_{22}e_{42})$$

$$C(4, 2) = \max(0.003375 * 0.5 * 0.3, 0.00225 * 0.6 * 0.3) = 0.00050625$$

For the fifth state  $s = 5$ , we have  $h_5 = 1$  we get,

$$C(5, 1) = \max(C(4, i)\delta_{i1}\xi(h_2, e_{51}))$$

$$C(5, 1) = \max(C(4, 1)\delta_{11}e_{51}, C(4, 2)\delta_{21}e_{51})$$

$$C(5, 1) = \max(0.0003375 * 0.5 * 0.3, 0.00050625 * 0.4 * 0.3) = 0.00006075$$

$$C(5, 2) = \max(C(4, i)\delta_{i2}\xi(h_2, e_{52}))$$

$$C(5, 2) = \max(C(4, 1)\delta_{12}e_{52}, C(4, 2)\delta_{22}e_{52})$$

$$C(5, 2) = \max(0.0003375 * 0.5 * 0.25, 0.00050625 * 0.6 * 0.25) = 0.0000759375$$

On comparing above values, the most likely state is 11122

**Problem 2.** Similar to PCA

**Answer 2.** c)  $\sum_{i=1}^p x_i^t \hat{U} \hat{U}^t x_i = \lambda_1 + \lambda_2$

**Solution 2.**

$$\sum_{i=1}^p x_i^t U U^t x_i = \sum_{j=1}^2 u_j^t X X^t u_j$$

Let  $v_1, \dots, v_n$  be the eigenvectors of  $X X^t$

$$\sum_{i=1}^p x_i^t U U^t x_i = \sum_{j=1}^2 \sum_{k=1}^n (u_j^t v_k)^2 \lambda_k$$

Assuming orthogonality, the above maximizes when  $u_k = v_k$  and we get,

$$\sum_{i=1}^p x_i^t \hat{U} \hat{U}^t x_i = \sum_{k=1}^2 \lambda_k$$

$$\sum_{i=1}^p x_i^t \hat{U} \hat{U}^t x_i = \lambda_1 + \lambda_2$$

**Problem 3.** Methylation data PCA coding

**Sub-Problem 3a.** What is the proportion of variance explained by the first 10 principal components?

**Answer 3a.** 56.8808%

Explained variance ratio by first 10 components is 56.8808%.

Data normalization is done to make it mean 0 and variance 1. code is attached.

**Sub-Problem 3b.** Find component with highest squared correlation to the batch vector?

**Answer 3b.** Fourth PC

Fourth PC gives the highest squared correlation of 0.84 to the batch vector. code is attached

**Sub-Problem 3c.** Find component with highest squared correlation to the batch vector obtained using PCA with sparsity procedure?

**Answer 3c.** First PC

First PC in the sparse PCA analysis gives the highest squared correlation of 0.49 to the batch vector. code is attached.