## **Solutions for HW1:**

## Written part:

1.

(1). Predicting uniformly at random means scientists will label the fish in any one of the five labels with the same probabilities. So, for each species, scientists have 20% probability to do the right prediction. That is, the probability of incorrectly labeling a crab is:

$$25\% * 80\% + 5\% * 80\% + 20\% * 80\% + 35\% * 80\% + 15\% * 80\% = 0.8$$

(2). Scientists can predict either label 1 or 4 to improve their accuracy:

Label 1: error rate = 1 - 0.25 = 0.75Label 4: error rate = 1 - 0.35 = 0.65

2.

This question can be solved easily using the function from spicy:

```
>>> from scipy.stats import norm, multivariate_normal
|>>> import numpy as np
|>>>
|>>> x = np.array([11.1, 27.8])
|>>> y_1 = multivariate_normal.pdf(x, mean=[29.42, 33.98], cov=[[50.56, 57.49], [57.49, 65.54]])
|>>> y_2 = multivariate_normal.pdf(x, mean=[33.88, 37.80], cov=[[46.79, 52.19], [52.19, 58.59]])
|>>> print(y_1)
|2.1660889714243452e-277
|>>> print(y_2)
|2.1520877668415265e-141
|>>> |
```

As you can in this image, *multivariate\_normal.pdf()* function calculate the probability using the specified means and covariance. Obviously, y\_2 (orange) is larger than y\_1 (blue). So, the color of the crab is most likely to be orange.

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(1). 
$$P(HHTHH) = \theta^*\theta^*(1-\theta)^*\theta^*\theta = \theta^4(1-\theta)$$

(2). 
$$log P(HHTHH) = 4 * log \theta + log(1 - \theta)$$

(3). From the previous questions we know:

$$arg \max_{\theta} p(HHTHH|\theta) = arg \max_{\theta} \theta^{4}(1-\theta)$$
$$= arg \max_{\theta} (4 \log \theta + \log(1-\theta))$$

Using derivative we and make it equal to zero:

$$^{4}/_{\theta} = ^{1}/_{1-\theta}$$

So,  $\theta$  should be 0.8.

## Coding part:

\*\*No marks have been awarded to those who have used libraries other than the ones mentioned in the announcement or on piazza.

The estimated 
$$\mu_0$$
 and  $\Sigma_0$  of the Gaussian for the Alaskan salmon should be: 
$$\begin{cases} \mu_0 = [99.22, 428.64] \\ \Sigma_0 = \big[[264.35, -212.54], [-212.54, 1386.23] \big] \end{cases}$$

The estimated 
$$\mu_0$$
 and  $\Sigma_0$  of the Gaussian for the Canadian salmon should be: 
$$\begin{cases} \mu_0 = [136.93, 366.64] \\ \Sigma_0 = \big[[338.24, 162.82], [162.82, 712.85]\big] \end{cases}$$

The true label of the fish in the table:

The predicted label of the fish in the table:

So, the accuracy is 90%.

For the coding part, both biased and unbiased estimates are accepted (for  $\Sigma_0$ ).

$$\Sigma_0$$
 can be [[270.36, -217.37], [-217.37,1417.73]] for the Alaskan salmon

$$\Sigma_0$$
 can be [[345.93, 166.52], [166.52,729.05]] for the Canadian salmon

Using these slightly different  $\Sigma_0$  will lead to the same result for predictions. By the way, you can still use the *multivariate normal.pdf()* function to calculate the probabilities to decide which species fish belongs to.