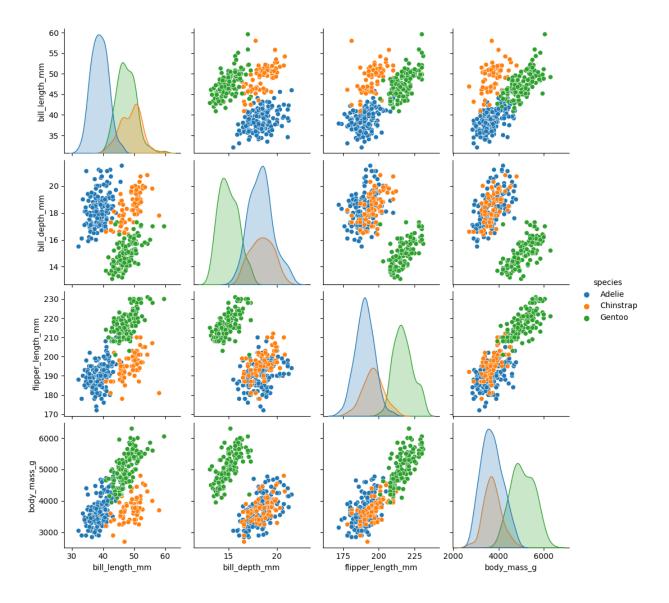
# Python Ex4 Danit Noa Yechezkel Keren Halpert



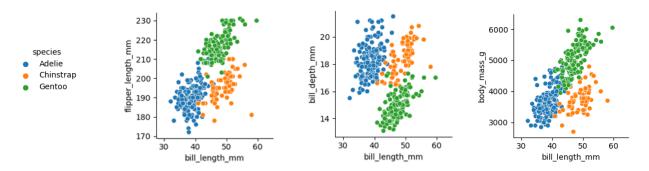
# **TASK 1:**

1. As depicted in this pair-plot, it is fairly easy to see that one of the best features is the "bill\_length\_mm" feature, the reason is that the three groups of penguin species are most segregated using this classifier as opposed to other features by comparison.



Now that the first feature was chosen, we need to decide which will be the second feature for the GNB 2-feature classifier.

In order to do so, we will look for the plot that is most segregated by color between the three plots using the first, chosen feature.



It is easy to see that the plot using the "body\_mass\_g" is not as good as the other two. But it is harder to tell between "flipper\_length\_mm" and "bill\_depth\_mm". So for this reason we will check both out using GNB and the accuracy score.

The accuracy score for the pair: 'bill\_length\_mm' and 'bill\_depth\_mm' is

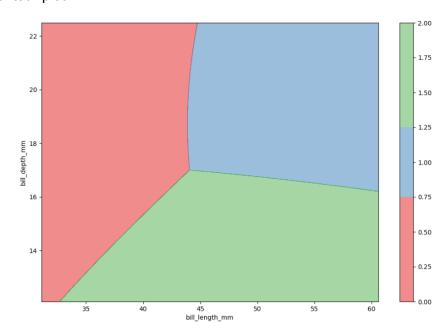
#### 0.9420289855072463

The accuracy score for the pair: 'bill\_length\_mm' and 'flipper\_length\_mm' is

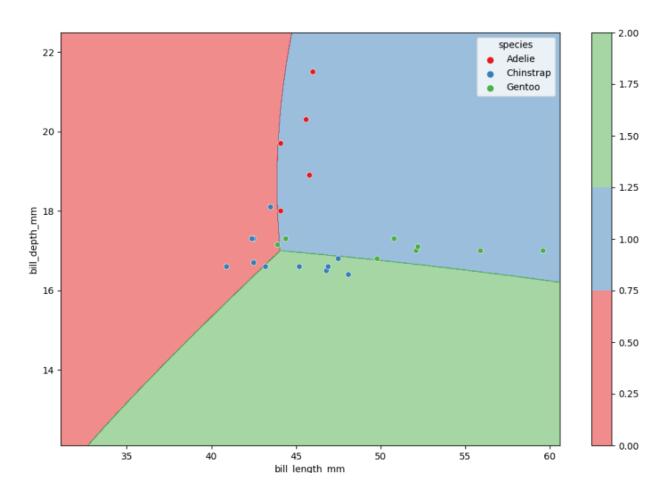
#### 0.927536231884058

Now we can see without a doubt that the winner of the two is 'bill\_depth\_mm'.

#### 3. Filled contour plot:



#### 4. Filled contour plot with scatter plot overlay of failed predictions:



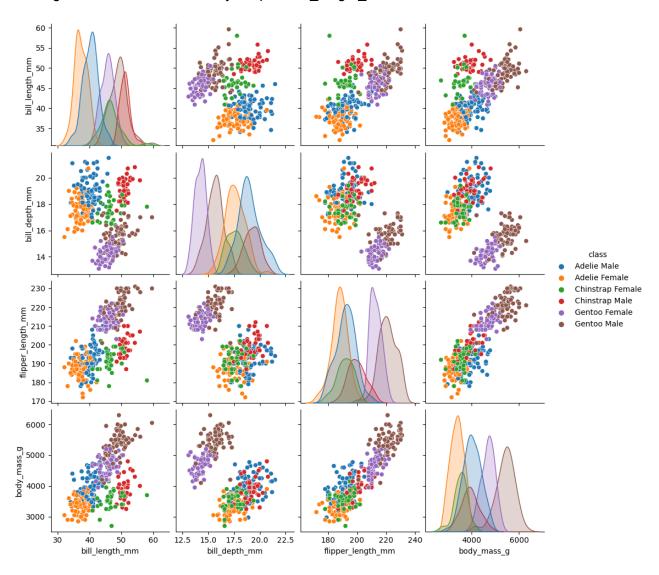
#### 5. Classification report:

	precision	recall	f1-score	support	
Adelie	1.00	0.94	0.97	34	
Chinstrap Gentoo	0.82 0.92	0.82 1.00	0.82 0.96	11 24	
accuracy			0.94	69	
macro avg weighted avg	0.91 0.94	0.92 0.94	0.92 0.94	69 69	

# **TASK 2:**

1. As done in task 1, we need to find the pair of features that provide the most segregated distribution of points.

Again, the first choice is fairly simple: "bill\_length\_mm".



But the second choice is a close call visually, therefore we will check out the GNB accuracy score.

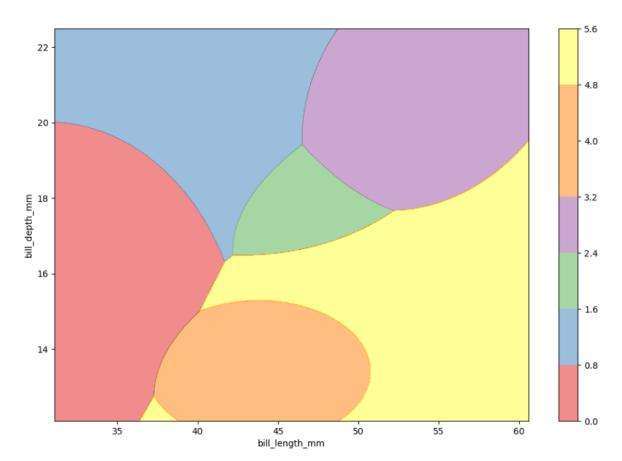
The accuracy score for the pair: 'bill\_length\_mm' and 'bill\_depth\_mm' is 0.8695652173913043

The accuracy score for the pair: 'bill\_length\_mm' and 'flipper\_length\_mm' is 0.782608695652174

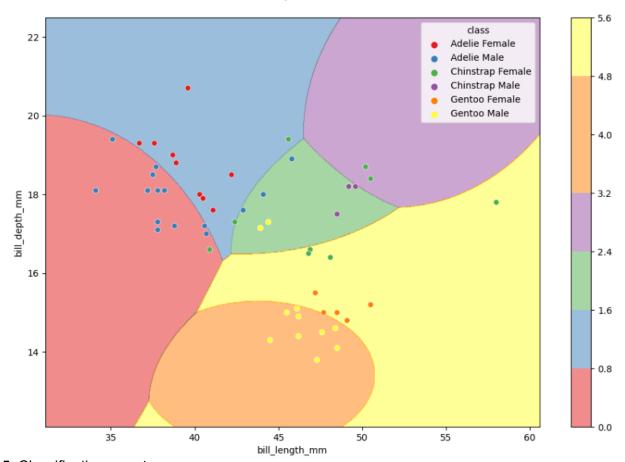
The accuracy score for the pair: 'bill\_length\_mm' and 'body\_mass\_g' is 0.8260869565217391

The clear winner is, again, 'bill\_depth\_mm'.

#### 3. Filled contour plot:



# 4. Filled contour plot with scatter plot overlay of failed predictions:



# 5. Classification report:

	precision	recall	f1-score	support	
Adelie Female	0.92	0.75	0.83	16	
Adelie Male	0.76	0.89	0.82	18	
Chinstrap Female	0.50	0.50	0.50	4	
Chinstrap Male	1.00	0.86	0.92	7	
Gentoo Female	1.00	1.00	1.00	7	
Gentoo Male	0.94	1.00	0.97	17	
accuracy			0.87	69	
macro avg	0.85	0.83	0.84	69	
weighted avg	0.88	0.87	0.87	69	