Gesture recognition Using the Kinect

Question 1: The joint angles, the hand positioning, and most commonly the speeds seem to vary among sequences of the same type. As the poses face towards the camera, you see the shoulders getting wider and arms seem to get longer. Since some poses are fairly similar, varying starting points would have probably confused any classification system. This would cause even more problems for nearest neighbor classifier due to it is heavy reliance on distances.

Question 2: By simply guessing the labels, there is $\frac{1}{9} = 11.11\%$ chance of correctly guessing any label which is in other words the expected accuracy of simply guessing. That's not the best possible accuracy you can get by simply guessing in a given run, but I assumed the questions is looking for the expected value.

Question 3: The L2 distance computation involves vector subtractions which you can only perform on vectors of the same length.

Upon implementing normalized frames normalize_frames(gesture_sets, num_frames), there're two situations to handle: 1) **Too long**: When a given frame length is greater then num_frames 2) **Too short**: When a given frame length is less than num_frames.

Too Long: I calculate step and offset as follows:

```
step = frames/num_remove; where frames= len(sequence.frames); and num_remove
= frames-num_frames;

offset = (step+frames%num_remove)/2;
```

I then compute the indices of the frames to be removed as follows.

```
to_remove = [offset+i*step for i in range(num_remove)]
```

It basically says, I want to remove num_remove frames spaced by step and starting from offset. The step parameter makes sure that the removed frames are equally spaced where the offset makes sure that the removed frames are around centered, about same distances from start(index = 0) and the end(index=n-1) of the array.

Too short: Again, compute step and offset as follows:

```
step = frames/missing; where frames= len(sequence.frames); and missing =
total-frames;

offset = (frames%missing)/2;
```

Finally, I compute indices of the frames to be duplicated as follows:

```
to_duplicate = [offset+i*step for i in range(missing)];
```

The logic behind the step and offset stays as in the case of **too long**. Note: my current implementation expects frames>=missing to hold. This is probably not a good idea and handling the situation where frames<missing to should be straightforward, but this seems hold for the given data so I didn't bother.

Question 4: Pros: The nearest neighbor classifier is very simple and fast. Cons: It relies on distances alone for information, which is not the only feature and probably not the most informative feature in this case. Features including relative angles, relative joint speeds, etc could more generally capture what's going in the gesture.

Question 5: The following table shows the results(accuracy) from different pairs of num_frames and ratio including the best among them(green):

| num_frames | ratio | accuracy | |
|------------|-------|--------------------|--|
| 20 | .3 | 0.582010582010582 | |
| 20 | .4 | 0.5987654320987654 | |
| 20 | .6 | 0.6203703703703703 | |
| 30 | .6 | 0.805555555555556 | |
| 30 | .7 | 0.8271604938271605 | |
| 40 | .7 | 0.8641975308641975 | |

Question 6: The test accuracy changes because train_test_split randomizes the test-train split.

Question 7: The following two for-loops create two dimensional python list of dimensions 30 * 9 by $33 * num_frames$ for the features and a 30 * 9 long array for the labels.

```
for gs in gesture_sets:
   for seq in gs.sequences:
```

```
sample = np.concatenate(list(map(lambda x: x.frame, seq.frames)))
samples.append(sample)
labels.append(int(gs.label))
```

The following line converts the label list to numpy array and the list of features to a Numpy matrix.

```
X, Y = np.vstack(samples), np.array(labels)
```

Of course you're doing this because the sklearn expects it but that's what's happening.

The following table shows the accuracy different values of train_size.

| train_size | lowest accuracy | highest accuracy | |
|------------|---------------------------------------|--------------------|--|
| 0.2 | 45.370370370370374 56.9444444444443 | | |
| 0.3 | 60.846560846560848 66.137566137566139 | | |
| 0.4 | 66.049382716049394 | | |
| 0.5 | 62.962962962962 | 71.11111111111114 | |
| 0.6 | 70.370370370370367 | 75.925925925925924 | |
| 0.7 | 70.370370370370367 | 75.308641975308646 | |
| 0.8 | 74.074074074074076 | 81.481481481481 | |
| 0.9 | 81.481481481481 | 81 92.592592592595 | |

Question 8: According to Wikipedia, the Gini impurity measures the frequency of incorrectly labeling a randomly chosen element from a distribution, if it was labeled according to a subset of the distribution. In other words, it has something to do with the likelihood of misclassifying a given element from a distribution of classes or **expected error rate**. It helps the algorithm decide the ordering of the features when building the tree. The following is the formula for the Gini Impurity.

$$I_G(p) = 1 - \sum_{i=1}^{J} p_i^2$$

If p_i is the probability of picking an item class i.

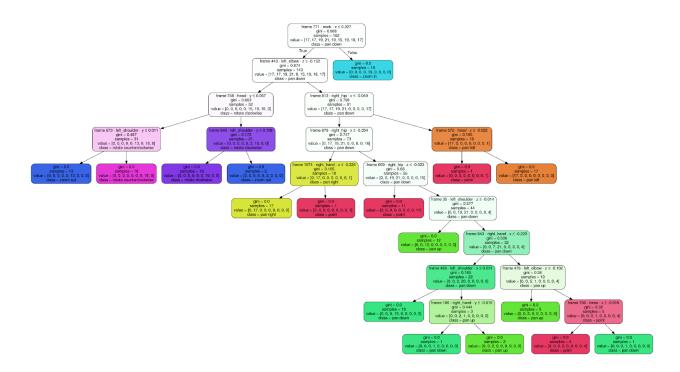
Question 9:

The training accuracy of the tree is **100%** and train accuracy was **70.37%**. Parameters.

- gini: Gini impurity at the node with the given split.

- **samples**: number of training samples passed to that node(all training samples for the root node).
- values: List of number of samples from each class at the current node.

The impurity is zero at the leaves because each leave classifies the elements of the same class. In other words, the **value** is zero everywhere except one index which makes the probability of misclassifying any element zero.



Question Bonus 1:

Approach 1: Added scaled-distance ratio between some joint positions as follows.

- 1. Added the ratio of the distances between the hand and shoulder on the same side to the corresponding arm length. $ratio = \frac{dist(shoulder\ pose,\ hand\ pose)}{arm\ length}$
- 2. Added the ratio of the distance between the hip and hand on the same side to the corresponding arm length. $ratio = \frac{dist(hip\ pose,\ hand\ pose)}{arm\ length}$.
- 3. This was done for both right and left side of the body.

Approach 2: Added the angles of the triangles created by the arms where the vertices are the shoulder, elbow, and hand. This was done for both sides of the body.

Approach 3: Add both.

The following table shows the results for each approach vs the original tree. As the table shows, the results are bit better for lower training ratio, but comparable for higher training ratios. This is because the whole idea of adding more features is counter productive. The problem is that the training accuracy is already 100% which means adding more features doesn't change match, in

fact it can help overfit even more. I have realized this pretty late so I didn't try to implement an approach that would limit the overfitting problem, but I have done what the staff have suggested so hope it was worth the effort in terms of grade.

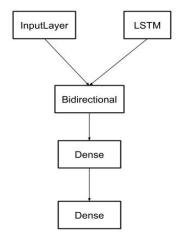
| | orig_acc | ratio | with_angs_acc | with_both_acc | with_dist_acc |
|---|-----------|-------|---------------|---------------|---------------|
| 0 | 39.917695 | 0.1 | 41.975309 | 45.679012 | 37.860082 |
| 1 | 55.55556 | 0.2 | 56.018519 | 55.555556 | 54.629630 |
| 2 | 62.962963 | 0.3 | 67.724868 | 64.021164 | 66.137566 |
| 3 | 68.518519 | 0.4 | 75.308642 | 73.456790 | 77.160494 |
| 4 | 73.333333 | 0.5 | 67.407407 | 71.111111 | 68.148148 |
| 5 | 75.925926 | 0.6 | 75.925926 | 83.333333 | 77.777778 |
| 6 | 76.543210 | 0.7 | 77.777778 | 74.074074 | 75.308642 |
| 7 | 75.925926 | 0.8 | 72.22222 | 74.074074 | 83.333333 |
| 8 | 88.88889 | 0.9 | 81.481481 | 77.777778 | 81.481481 |

Question 10: The input hasn't been changed from the Gesture_set representation. The features are arranged in 3D numpy array such that input layer of the network expects two dimensional numpy array(36 frames of 33 features each). This doesn't show that the LSTM is imposing strict requirements compare to the two other algorithms, it shows that the LSTM model architecture is more flexible or more generalized. An interesting observation is that the we don't actually split the data into test and validation for the LSTM beforehand, we only tell it what ratio to use with the validation_split parameter.

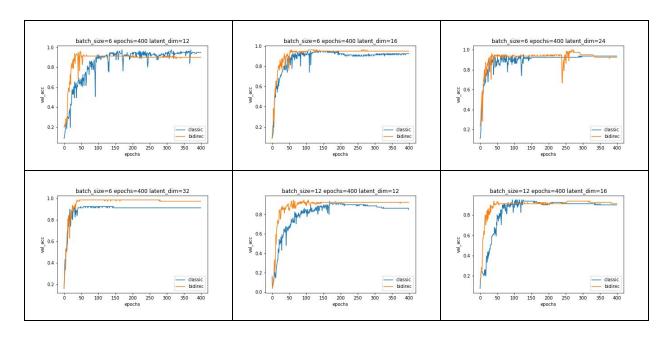
The best accuracy recorded was 95.06% which was found with multiple different parameter sets, but one of them was batch_size=8, epochs=200, and latent_dim=12.

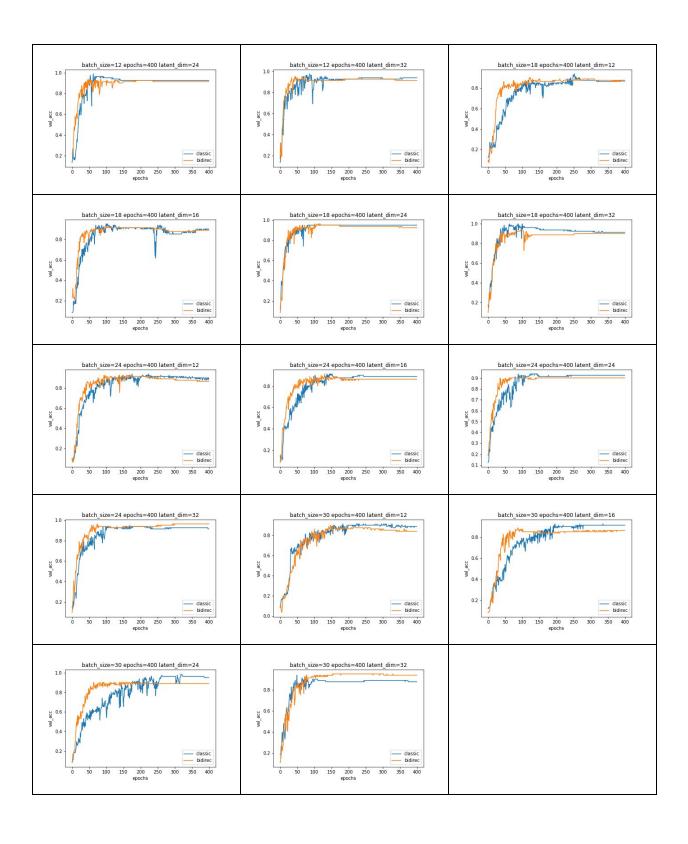
Question 11: I didn't use the hint, I used the **Bidirectional** object as shown by the line of code bilow.

lstm = Bidirectional(LSTM(latent_dim))(input_layer)

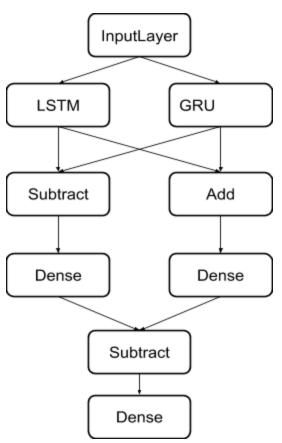


The following table of images show the validation accuracies of the normal LSTM and the Bidirectional LSTM for different values of the batch sizes and hidden layer dimensions as the epochs go from 0-400. Note the axis are label accordingly and the plot labels indicate which plot is which. The plots indicate that the two setups are very much comparable and their relative performances depend on the parameters(batch size and latent dim).





Question bonus 2:



The graph to the left shows the final architecture that produced the best results for the recurrent neural network bonus. The following are the module descriptions.

1. InputLayer: Inputlayer

2. LSTM: LSTM layer

3. GRU: Gated Recurrent Unit layer

4. Subtract: Merge by Subtraction layer.

5. Add: Merge by addition layer

6. **Dense**: Dense layer

The following plots show the performance of the selected bonus architecture against the initial(staff) set. Note: This architecture produce 100% final validation accuracies in some situations.

