```
In [32]:
          import numpy as np
          import pandas as pd
          import matplotlib.pyplot as plt
          import warnings
          from keras import layers
          from sklearn.model selection import train test split
          warnings.filterwarnings('ignore')
          import collections
          import keras
          import matplotlib.pyplot as plt
          import numpy as np
          import pandas as pd
          import seaborn as sns
          from keras import initializers
          from keras.datasets import mnist
          from tensorflow import keras
          import tensorflow as tf
          import itertools
          from keras import backend as K
          import time
          sed = 2022
          import random
```

Problem 1 ----- Perceptron

(1)

```
In [2]:
         # create function
         def generate data(length, seed):
             np.random.seed(seed)
             x 1 = np.expand dims(np.random.random(length),axis = 1)
             x 2 = np.expand dims(np.random.random(length),axis = 1)
             df 1 = pd.DataFrame(np.concatenate([x 1,x 2],axis = 1)).rename(\{0: x 1', 1: x 2'\},axis
             df 1['y'] = (df 1['x 1'] > df 1['x 2']).astype(int)
             df 1['y'][df 1['y'] == 0] = -1
             return df 1
         def train(df, seed, a, lr, epoch):
             # initialize weights
             np.random.seed(seed)
             w 1 = np.random.rand(1)
             w 2 = np.random.rand(1)
             for e in range(epoch):
                 for i in range(len(df)):
                      # forward
                     y \text{ hat} = df.iloc[i,2] * (df.iloc[i,0]*w 1 + df.iloc[i,1]*w 2)
                     loss = max(0,a-y hat)
                       # backward
                      if loss > 0:
                          gradient w1 = -1 * df.iloc[i,2] * df.iloc[i,0]
                          gradient w2 = -1 * df.iloc[i,2] * df.iloc[i,1]
                          w 1 -= lr * gradient w1
                          w 2 -= lr * gradient w2
                     else:
                          pass
             def predict func(x1, x2):
                 return 1 if w 1*x1 + w 2*x2 > 0 else -1
             return predict func, (w 1, w 2)
```

(array([0.09828205]), array([-0.09308899]))
Accuracy for this preceptron is: 0.978

(2) hinge loss

```
In [4]: # Train model
    a = 1
    df_2_train = generate_data(20, sed)
    preceptron_2, weights = train(df_2_train, sed, a, 0.1, 1000)
    print(weights)
    # get test accuracy
    df_2_test = generate_data(1000, sed)
    pred = []
    for i in range(len(df_2_test)):
        pred.append(preceptron_2(df_2_test.iloc[i,0], df_2_test.iloc[i,1]))
    df_2_result = pd.concat([df_2_test, pd.DataFrame(pred).rename({0:'y_pred'}, axis = 1)], axis
    df_2_result['AUC'] = df_2_result['y'] == df_2_result['y_pred']
    Auc_count = sum(df_2_result['AUC'])
    print('Accuracy for this preceptron is: '+ str(Auc_count/1000))
```

(array([12.80079885]), array([-11.97284392]))
Accuracy for this preceptron is: 0.971

(3)

- First case gives me a better accuracy.
- It's because in the first case, we only update the weights when we get a wrong prediction, so that our weights will only update towards a direction to minmize the classfication loss. However in the second case, when $0 < y(\bar{W} \cdot \bar{X}) < 1$ even though we get a correct prediction, we still have to adjust our weights and the gradient gained by the hindge loss is not always helpful in the classfication

(4)

• When the data set is evenly distributed along and close to the true decision boundary will we gain a stable result of classification of the same 1000 test instances.

Problem 2 ----- Weight Initialization, Dead Neurons, Leaky ReLU

(1)

In [5]:

copy the functions need from the github files

```
import keras
from keras.models import Sequential
from keras.layers import Conv2D, MaxPooling2D, Dense, Dropout, Flatten
from keras import backend as K
from matplotlib import pyplot as plt
from matplotlib import rcParamsDefault
def grid axes it(n plots, n cols=3, enumerate=False, fig=None):
        Iterate through Axes objects on a grid with n cols columns and as many
        rows as needed to accommodate n plots many plots.
                n plots: Number of plots to plot onto figure.
                n cols: Number of columns to divide the figure into.
                fig: Optional figure reference.
        Yields:
               n plots many Axes objects on a grid.
        n rows = n plots / n cols + int(n plots % n cols > 0)
        if not fig:
                default figsize = rcParamsDefault['figure.figsize']
                fig = plt.figure(figsize=(
                        default figsize[0] * n cols,
                        default figsize[1] * n rows
                ) )
        for i in range(1, n plots + 1):
                ax = plt.subplot(n rows, n cols, i)
                yield ax
def create mlp model(
       n hidden layers,
        dim layer,
        input shape,
        n classes,
        kernel initializer,
        bias initializer,
       activation,
):
        """Create Multi-Layer Perceptron with given parameters."""
        model = Sequential()
        model.add(Dense(dim layer, input shape=input shape, kernel initializer=kernel initiali
                                        bias initializer=bias initializer))
        for i in range(n hidden layers):
                model.add(Dense(dim layer, activation=activation, kernel initializer=kernel initializer=k
                                                 bias initializer=bias initializer))
        model.add(Dense(n classes, activation='softmax', kernel initializer=kernel initializer
                                         bias initializer=bias initializer))
        return model
def create cnn model(input shape, num classes, kernel initializer='glorot uniform',
                                          bias initializer='zeros'):
        """Create CNN model similar to
              https://github.com/keras-team/keras/blob/master/examples/mnist cnn.py."""
        model = Sequential()
        model.add(Conv2D(32, kernel size=(3, 3),
                                          activation='relu',
                                           input shape=input shape,
                                           kernel initializer=kernel initializer,
                                          bias initializer=bias initializer))
        model.add(Conv2D(64, (3, 3), activation='relu',
```

```
kernel initializer=kernel initializer,
                     bias initializer=bias initializer))
    model.add(MaxPooling2D(pool size=(2, 2)))
    model.add(Dropout(0.25))
    model.add(Flatten())
    model.add(Dense(128, activation='relu',
                    kernel initializer=kernel initializer,
                    bias initializer=bias initializer))
    model.add(Dropout(0.5))
    model.add(Dense(num classes, activation='softmax',
                    kernel initializer=kernel initializer,
                    bias initializer=bias initializer))
    return model
def compile model(model):
    model.compile(loss=keras.losses.categorical crossentropy,
                  optimizer=keras.optimizers.RMSprop(),
                  metrics=['accuracy'])
    return model
def get init id(init):
    Returns string ID summarizing initialization scheme and its parameters.
        init: Instance of some initializer from keras.initializers.
    try:
        init name = str(init).split('.')[2].split(' ')[0]
    except:
        init name = str(init).split(' ')[0].replace('.', ' ')
    param list = []
    config = init.get config()
    for k, v in config.items():
        if k == 'seed':
            continue
        param list.append('\{k\}-\{v\}'.format(k=k, v=v))
    init params = ' '.join(param list)
    return '|'.join([init name, init params])
def get activations(model, x, mode=0.0):
    """Extract activations with given model and input vector x."""
    outputs = [layer.output for layer in model.layers]
    activations = K.function([model.input], outputs)
    output elts = activations([x])
    return output elts
class LossHistory(keras.callbacks.Callback):
   """A custom keras callback for recording losses during network training."""
    def on train begin(self, logs={}):
        self.losses = []
        self.epoch losses = []
        self.epoch val losses = []
    def on batch end(self, batch, logs={}):
        self.losses.append(logs.get('loss'))
    def on epoch end(self, epoch, logs={}):
        self.epoch losses.append(logs.get('loss'))
        self.epoch val losses.append(logs.get('val loss'))
```

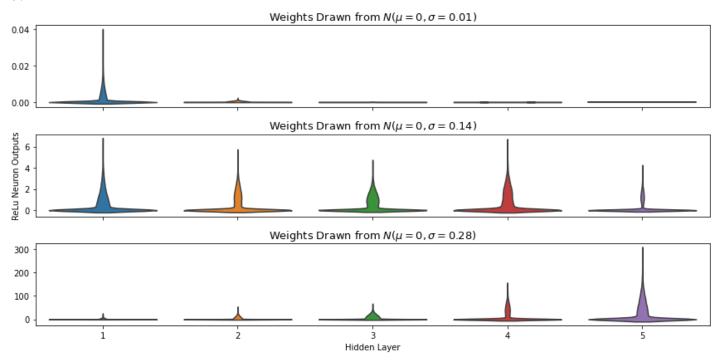
Relu Activation

```
In [6]:
         # copy code from plot activation layers.py file
         seed = 10
         # Number of points to plot
         n train = 1000
         n test = 100
         n classes = 10
         # Network params
         n hidden layers = 5
         dim layer = 100
         batch size = n train
         epochs = 12
         # Load and prepare MNIST dataset.
         n train = 60000
         n \text{ test} = 10000
         (x train, y train), (x test, y test) = mnist.load data()
         num classes = len(np.unique(y test))
         data dim = 28 * 28
         x train = x train.reshape(60000, 784).astype('float32')[:n train]
         x \text{ test} = x \text{ test.reshape}(10000, 784).astype('float32')[:n train]
         x train /= 255
         x_test /= 255
         y train = keras.utils.to categorical(y train, num classes)
         y test = keras.utils.to categorical(y test, num classes)
         # Run the data through a few MLP models and save the activations from
         # each layer into a Pandas DataFrame.
         rows = []
         sigmas = [0.01, 0.14, 0.28]
         for stddev in sigmas:
             init = initializers.RandomNormal(mean=0.0, stddev=stddev, seed=seed)
             activation = 'relu'
             model = create mlp model(
                 n hidden layers,
                 dim layer,
                 (data dim,),
                 n classes,
                 init,
                 'zeros',
                 activation
             compile model(model)
             output elts = get activations(model, x test)
             n layers = len(model.layers)
             i output layer = n layers - 1
             for i, out in enumerate(output elts[:-1]):
                 if i > 0 and i != i output layer:
                     for out i in out.ravel()[::20]:
                          rows.append([i, stddev, out i])
         df = pd.DataFrame(rows, columns=['Hidden Layer', 'Standard Deviation', 'Output'])
```

Plot previously saved activations from the 5 hidden layers

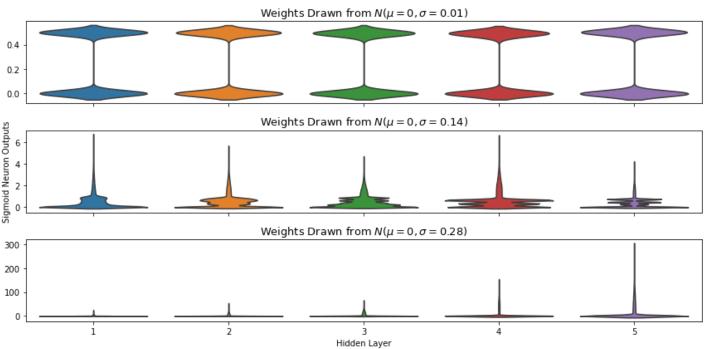
```
# using different initialization schemes.
fig = plt.figure(figsize=(12, 6))
axes = grid axes it(len(sigmas), 1, fig=fig)
for sig in sigmas:
    ax = next(axes)
    ddf = df[df['Standard Deviation'] == sig]
    sns.violinplot(x='Hidden Layer', y='Output', data=ddf, ax=ax, scale='count', inner=Nor
    ax.set xlabel('')
    ax.set ylabel('')
    ax.set title('Weights Drawn from $N(\mu = 0, \sigma = {\%.2f})$' \% sig, fontsize=13)
    if siq == sigmas[1]:
        ax.set ylabel("ReLu Neuron Outputs")
    if sig != sigmas[-1]:
        ax.set xticklabels(())
    else:
        ax.set xlabel("Hidden Layer")
plt.tight layout()
plt.show()
```

2022-10-11 23:17:21.761996: I tensorflow/core/platform/cpu_feature_guard.cc:151] This Tens orFlow binary is optimized with oneAPI Deep Neural Network Library (oneDNN) to use the fol lowing CPU instructions in performance-critical operations: AVX2 FMA To enable them in other operations, rebuild TensorFlow with the appropriate compiler flag s.



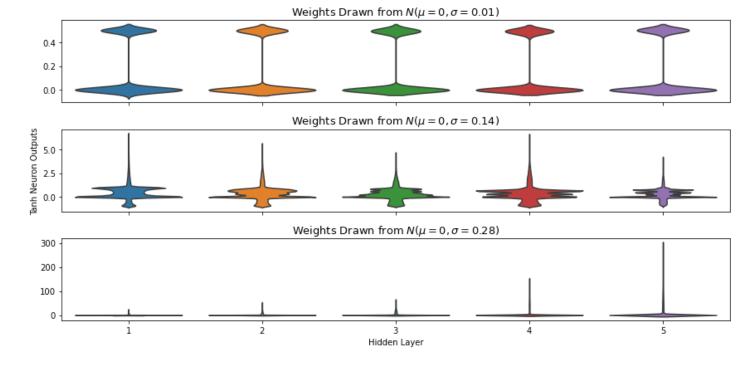
sigmoid activation

```
'zeros',
        activation
    compile model(model)
    output elts = get activations(model, x test)
    n layers = len(model.layers)
    i output layer = n layers - 1
    for i, out in enumerate(output elts[:-1]):
        if i > 0 and i != i output layer:
            for out i in out.ravel()[::20]:
                rows.append([i, stddev, out i])
df = pd.DataFrame(rows, columns=['Hidden Layer', 'Standard Deviation', 'Output'])
# Plot previously saved activations from the 5 hidden layers
# using different initialization schemes.
fig = plt.figure(figsize=(12, 6))
axes = grid axes it(len(sigmas), 1, fig=fig)
for sig in sigmas:
   ax = next(axes)
    ddf = df[df['Standard Deviation'] == sig]
    sns.violinplot(x='Hidden Layer', y='Output', data=ddf, ax=ax, scale='count', inner=Nor
    ax.set xlabel('')
    ax.set ylabel('')
    ax.set title('Weights Drawn from $N(\mu = 0, \sigma = {%.2f})$' % sig, fontsize=13)
    if sig == sigmas[1]:
        ax.set ylabel("Sigmoid Neuron Outputs")
    if sig != sigmas[-1]:
        ax.set xticklabels(())
    else:
        ax.set xlabel("Hidden Layer")
plt.tight layout()
plt.show()
                                 Weights Drawn from N(\mu = 0, \sigma = 0.01)
```



Tanh activation

```
sigmas = [0.01, 0.14, 0.28]
for stddev in sigmas:
   init = initializers.RandomNormal(mean=0.0, stddev=stddev, seed=seed)
    activation = 'tanh'
    model = create mlp model(
       n hidden layers,
        dim layer,
       (data dim,),
       n classes,
       init,
        'zeros',
        activation
    compile model(model)
    output elts = get activations(model, x test)
    n layers = len(model.layers)
    i output layer = n layers - 1
    for i, out in enumerate(output elts[:-1]):
        if i > 0 and i != i output layer:
            for out i in out.ravel()[::20]:
                rows.append([i, stddev, out i])
df = pd.DataFrame(rows, columns=['Hidden Layer', 'Standard Deviation', 'Output'])
# Plot previously saved activations from the 5 hidden layers
# using different initialization schemes.
fig = plt.figure(figsize=(12, 6))
axes = grid axes it(len(sigmas), 1, fig=fig)
for sig in sigmas:
   ax = next(axes)
    ddf = df[df['Standard Deviation'] == sig]
    sns.violinplot(x='Hidden Layer', y='Output', data=ddf, ax=ax, scale='count', inner=Nor
   ax.set xlabel('')
   ax.set ylabel('')
    ax.set title('Weights Drawn from $N(\mu = 0, \sigma = {\%.2f})$' \% sig, fontsize=13)
    if sig == sigmas[1]:
        ax.set ylabel("Tanh Neuron Outputs")
    if sig != sigmas[-1]:
        ax.set xticklabels(())
    else:
        ax.set xlabel("Hidden Layer")
plt.tight layout()
plt.show()
```

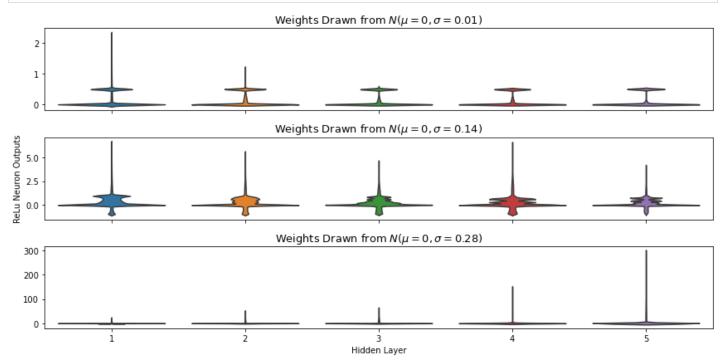


GlorotNormal initialization

```
In [9]:
         sigmas = [0.01, 0.14, 0.28]
         for stddev in sigmas:
             init = initializers.glorot normal(seed = seed)
             activation = 'relu'
             model = create mlp model(
                 n hidden layers,
                 dim layer,
                 (data dim,),
                 n classes,
                 init,
                 'zeros',
                 activation
             compile model(model)
             output elts = get activations(model, x test)
             n layers = len(model.layers)
             i output layer = n layers - 1
             for i, out in enumerate(output elts[:-1]):
                 if i > 0 and i != i output layer:
                     for out i in out.ravel()[::20]:
                          rows.append([i, stddev, out i])
         df = pd.DataFrame(rows, columns=['Hidden Layer', 'Standard Deviation', 'Output'])
         # Plot previously saved activations from the 5 hidden layers
         # using different initialization schemes.
         fig = plt.figure(figsize=(12, 6))
         axes = grid axes it(len(sigmas), 1, fig=fig)
         for sig in sigmas:
             ax = next(axes)
             ddf = df[df['Standard Deviation'] == sig]
             sns.violinplot(x='Hidden Layer', y='Output', data=ddf, ax=ax, scale='count', inner=Nor
             ax.set xlabel('')
             ax.set ylabel('')
             ax.set title('Weights Drawn from $N(\mu = 0, \sigma = {\%.2f})\$' \% sig, fontsize=13)
```

```
if sig == sigmas[1]:
    ax.set_ylabel("ReLu Neuron Outputs")
if sig != sigmas[-1]:
    ax.set_xticklabels(())
else:
    ax.set_xlabel("Hidden Layer")

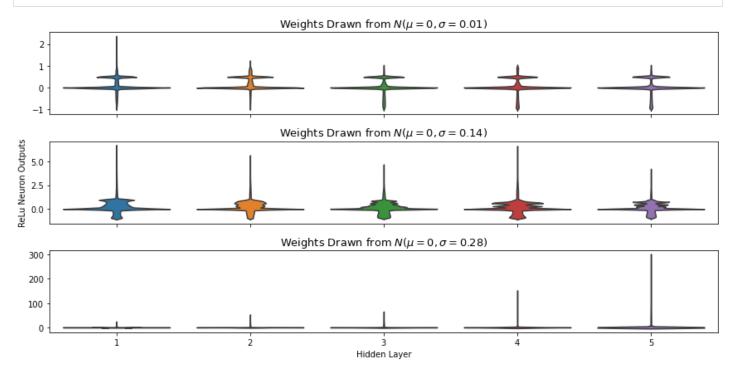
plt.tight_layout()
plt.show()
```



He initialization

```
In [10]:
          sigmas = [0.01, 0.14, 0.28]
          for stddev in sigmas:
              init = initializers.HeNormal(seed=seed)
              activation = 'tanh'
              model = create mlp model(
                  n hidden layers,
                  dim layer,
                  (data dim,),
                  n classes,
                  init,
                  'zeros',
                  activation
              compile model(model)
              output elts = get activations(model, x test)
              n layers = len(model.layers)
              i output layer = n layers - 1
              for i, out in enumerate(output elts[:-1]):
                  if i > 0 and i != i output layer:
                      for out i in out.ravel()[::20]:
                           rows.append([i, stddev, out i])
          df = pd.DataFrame(rows, columns=['Hidden Layer', 'Standard Deviation', 'Output'])
          # Plot previously saved activations from the 5 hidden layers
          # using different initialization schemes.
```

```
fig = plt.figure(figsize=(12, 6))
axes = grid axes it(len(sigmas), 1, fig=fig)
for sig in sigmas:
    ax = next(axes)
    ddf = df[df['Standard Deviation'] == sig]
    sns.violinplot(x='Hidden Layer', y='Output', data=ddf, ax=ax, scale='count', inner=Nor
    ax.set xlabel('')
    ax.set ylabel('')
    ax.set title('Weights Drawn from $N(\mu = 0, \sigma = {%.2f})$' % sig, fontsize=13)
    if sig == sigmas[1]:
        ax.set ylabel("ReLu Neuron Outputs")
    if sig != sigmas[-1]:
        ax.set xticklabels(())
    else:
        ax.set xlabel("Hidden Layer")
plt.tight layout()
plt.show()
```



- Here in order to see the effect clear I use the [0.01, 0.14, 0.28] As my sigmas, and as shown on the above plots, as the σ close to zero, we will face the Vanishing gradient since for front layers the gradient are close to 0 so that they're not updating.
- Comparied with relu, using simoid and Tanh is less likly to encounter Vanishing gradient.
- After change the initializer into GlorotNormal, the Vanishing gradient for relu activation is imporved.
- After change the initializer into Heinitialization, the Vanishing gradient for relu activation is imporved.

(2)

In [138...

```
print('The collapsed models are: ' + str(collapus_count)+' ,So the proportion is close to
```

The collapsed models are: 91 ,So the proportion is close to 90% as reported in the paper

(3)

```
In [139...
          leakyrelu = tf.keras.layers.LeakyReLU(alpha=0.01)
          collapus count leak relu = 0
          for i in range(1000):
              x = np.random.uniform(-np.sqrt(7), np.sqrt(7), 3000)
              y = np.abs(x)
              init = initializers.HeNormal(seed = sed)
              model 3 = keras.Sequential([layers.Dense(2, activation=leakyrelu,input shape = (1,),
                                                        kernel initializer=init, use bias=True),
                                        layers.Dense(2, activation=leakyrelu, kernel initializer=init,
                                        layers.Dense(1, use bias=False)])
              model 3.compile(loss='mean squared error', optimizer='adam', metrics=['MSE'])
              model 3.fit(x=x,y=y, epochs=1, batch size=64, verbose=0)
              pred = model 3.predict(x)
              if len(np.unique(pred)) == 1 and np.unique(pred)[0]==0:
                  collapus count leak relu+=1
```

```
In [140...
```

```
print('The collapsed models are: ' + str(collapus_count_leak_relu)+' , there is no collapsed
```

The collapsed models are: 0 , there is no collapse after using leakeyrelu!

Problem 3 ----- Batch Normalization, Dropout, MNIST

(1)

co-adaptation

• As is stated in Hinton's paper The ability of a set of genes to be able to work well with another random set of genes makes them more robust. Since a gene cannot rely on a large set of

partners to be present at all times, it must learn to do something useful on its own or in collaboration with a small number of other genes. In the neural network, neurons just works like gene in sexual reproduction. When adding dropout, the complex co-adaptation is reduced thus making neural network more robust and have better performance on test datasets.

covariance-shift

• Covariance-shift is defined as **the distribution of each layer's inputs changes during training, as the parameters of the previous layers change.** However by using the regularization like Batch
Normalization, we can actually let input of each layer transform into standard normal distribution, thus
let all hidden layers have the same distribution of input, and address the covariance-shift problem.

(2)

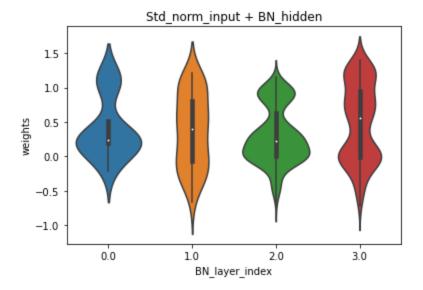
```
In [231...
       # preprocess data
       (x train, y train), (x val, y val) = mnist.load data()
       x train = np.expand dims(x train, axis = 3)
       x \text{ val} = \text{np.expand dims}(x \text{ val, axis} = 3)
       mean train = np.mean(x train)
       var train = np.var(x train)
       # define Lenet-5 and add BNlayer and train the model
       model 3 2 = tf.keras.models.Sequential([
             tf.keras.layers.Normalization(axis=-1, mean=mean train, variance=var train),
             tf.keras.layers.Conv2D(6, kernel size=5, strides = 1,input shape = (28,28,1),activ
             tf.keras.layers.BatchNormalization(),
             tf.keras.layers.AveragePooling2D(pool size = (2,2), strides = 2),
             tf.keras.layers.Conv2D(16, kernel size = 5, strides = 1, activation = 'tanh'),
             tf.keras.layers.BatchNormalization(),
             tf.keras.layers.AveragePooling2D(pool size = (2,2), strides = 2),
             tf.keras.layers.Conv2D(120, kernel size = 2, strides = 2, activation = 'tanh'),
             tf.keras.layers.BatchNormalization(),
             tf.keras.layers.Flatten(),
             tf.keras.layers.Dense(84, activation = 'tanh'),
             tf.keras.layers.BatchNormalization(),
             tf.keras.layers.Dense(10, activation = 'softmax'),
          ])
       model 3 2.compile(loss = 'sparse categorical crossentropy',optimizer = 'adam', metrics =
       history 3 2 = model 3 2.fit(x train, y train, epochs = 10, batch size = 64, validation dat
      Epoch 1/10
      - val loss: 0.0709 - val accuracy: 0.9774
      Epoch 2/10
      - val loss: 0.0531 - val accuracy: 0.9806
      Epoch 3/10
      - val loss: 0.0480 - val accuracy: 0.9851
      Epoch 4/10
      - val loss: 0.0412 - val accuracy: 0.9863
      Epoch 5/10
      - val loss: 0.0383 - val accuracy: 0.9871
      Epoch 6/10
      - val loss: 0.0347 - val accuracy: 0.9885
      Epoch 7/10
```

```
- val loss: 0.0460 - val accuracy: 0.9843
       Epoch 8/10
       - val loss: 0.0311 - val accuracy: 0.9898
       Epoch 9/10
       - val loss: 0.0347 - val accuracy: 0.9879
       Epoch 10/10
       - val loss: 0.0358 - val accuracy: 0.9893
In [232...
       # get parametres in BN layer
       BN index = [2, 5, 8, 11]
       BN parameters df = None
       BN index count = 0
       for i in BN index:
          weight = model 3 2.layers[i].get weights()
          weight = np.concatenate(weight)
          index = BN index count*np.ones like(weight)
          weight = np.expand dims(weight,axis = 1)
          index = np.expand dims(index,axis = 1)
          weight = np.concatenate([weight,index],axis = 1)
          weight = pd.DataFrame(weight)
          BN index count+=1
          if BN parameters df is not None:
             BN parameters df = pd.concat([BN parameters df,weight],axis = 0)
          else:
             BN parameters df = weight
```

BN parameters df = BN parameters df.rename({0:'weights',1:'BN layer index'},axis = 1)

y = BN parameters df['weights']).set(title = 'Std norm input + BN hidden')

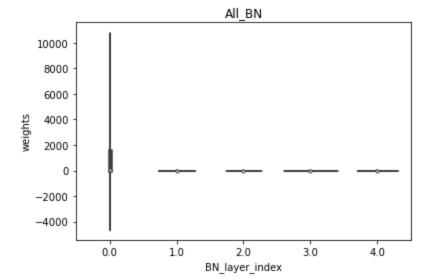
Out[232... [Text(0.5, 1.0, 'Std_norm_input + BN_hidden')]



sns.violinplot(x = BN parameters df['BN layer index'],

(3)

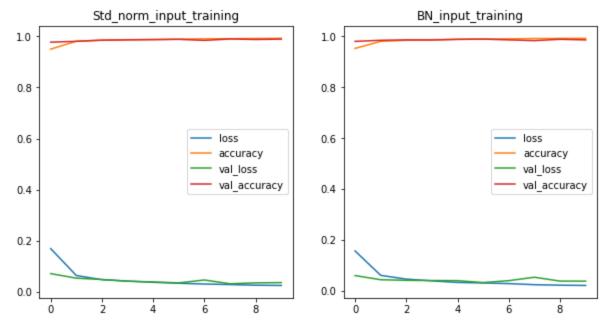
```
tf.keras.layers.Conv2D(120, kernel size = 2, strides = 2, activation = 'tanh'),
           tf.keras.layers.BatchNormalization(),
           tf.keras.layers.Flatten(),
           tf.keras.layers.Dense(84, activation = 'tanh'),
           tf.keras.layers.BatchNormalization(),
           tf.keras.layers.Dense(10, activation = 'softmax'),
         ])
      model 3 3.compile(loss = 'sparse categorical crossentropy',optimizer = 'adam', metrics =
      history 3 3 = model 3 3.fit(x train, y train, epochs = 10, batch size = 64, validation dat
      Epoch 1/10
      - val loss: 0.0611 - val accuracy: 0.9808
      Epoch 2/10
      - val loss: 0.0448 - val accuracy: 0.9850
      Epoch 3/10
      - val loss: 0.0425 - val accuracy: 0.9867
      Epoch 4/10
      - val loss: 0.0416 - val accuracy: 0.9861
      Epoch 5/10
      - val loss: 0.0411 - val accuracy: 0.9885
      Epoch 6/10
      - val loss: 0.0336 - val accuracy: 0.9898
      Epoch 7/10
      - val loss: 0.0411 - val accuracy: 0.9867
      Epoch 8/10
      - val loss: 0.0546 - val accuracy: 0.9837
      Epoch 9/10
      - val loss: 0.0393 - val accuracy: 0.9891
      Epoch 10/10
      - val loss: 0.0392 - val accuracy: 0.9866
In [290...
      BN index = [0, 2, 5, 8, 11]
      BN parameters df = None
      BN index count = 0
      for i in BN index:
         weight = model 3 3.layers[i].get weights()
         weight = np.concatenate(weight)
         index = BN index count*np.ones like(weight)
         weight = np.expand dims(weight,axis = 1)
         index = np.expand dims(index,axis = 1)
         weight = np.concatenate([weight,index],axis = 1)
         weight = pd.DataFrame(weight)
         BN index count+=1
         if BN parameters df is not None:
            BN parameters df = pd.concat([BN parameters df,weight],axis = 0)
         else:
            BN parameters df = weight
      BN parameters df = BN parameters df.rename((0:'weights',1:'BN layer index'),axis = 1)
      sns.violinplot(x = BN parameters df['BN layer index'],
                y = BN parameters df['weights']).set(title = 'All BN')
```



compare train/test loss

```
fig, ax = plt.subplots(1,2,figsize = (10,5))
    epoch_plot = range(10)
    for key in history_3_2.history.keys():
        ax[0].plot(epoch_plot,history_3_2.history[key],label = key)
        ax[0].legend()
        ax[0].set_title('Std_norm_input_training')

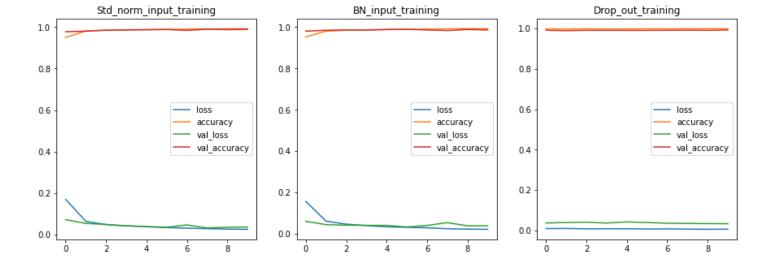
for key in history_3_3.history.keys():
        ax[1].plot(epoch_plot,history_3_3.history[key],label = key)
        ax[1].legend()
        ax[1].set_title('BN_input_training')
```



According to the plot, using the BN for input didn't improve the performace significantly, on the
contrary, it slightly increase the validation_loss a little bit on the 7th epoch. And also, according to the
violinplot the parameters in BN layer of input and hidden layer are not in the same scale.

(4)

```
tf.keras.layers.Conv2D(6, kernel size=5, strides = 1,activation = 'tanh'),
            tf.keras.layers.Dropout(0.5),
            tf.keras.layers.AveragePooling2D(pool size = (2,2), strides = 2),
            tf.keras.layers.Conv2D(16, kernel size = 5, strides = 1, activation = 'tanh'),
            tf.keras.layers.Dropout(0.5),
            tf.keras.layers.AveragePooling2D(pool size = (2,2), strides = 2),
            tf.keras.layers.Conv2D(120, kernel size = 2, strides = 2, activation = 'tanh'),
            tf.keras.layers.Dropout(0.5),
            tf.keras.layers.Flatten(),
            tf.keras.layers.Dense(84, activation = 'tanh'),
            tf.keras.layers.Dropout(0.5),
            tf.keras.layers.Dense(10, activation = 'softmax'),
         ])
      model 3 4.compile(loss = 'sparse categorical crossentropy',optimizer = 'adam', metrics =
      history 3 4 = model 3 3.fit(x train, y train, epochs = 10, batch size = 64, validation dat
      Epoch 1/10
      - val loss: 0.0379 - val accuracy: 0.9901
      Epoch 2/10
      - val loss: 0.0403 - val accuracy: 0.9881
      Epoch 3/10
      - val loss: 0.0420 - val accuracy: 0.9898
      Epoch 4/10
      - val loss: 0.0376 - val accuracy: 0.9896
      Epoch 5/10
      - val loss: 0.0434 - val accuracy: 0.9895
      Epoch 6/10
      - val loss: 0.0402 - val accuracy: 0.9894
      Epoch 7/10
      - val loss: 0.0367 - val accuracy: 0.9900
      Epoch 8/10
      - val loss: 0.0358 - val accuracy: 0.9903
      Epoch 9/10
      - val loss: 0.0350 - val accuracy: 0.9897
      Epoch 10/10
      - val loss: 0.0345 - val accuracy: 0.9913
In [257...
      fig, ax = plt.subplots(1, 3, figsize = (15, 5))
      epoch plot = range (10)
      for key in history 3 2.history.keys():
         ax[0].plot(epoch plot,history 3 2.history[key],label = key)
         ax[0].legend()
         ax[0].set title('Std norm input training')
      for key in history 3 3.history.keys():
         ax[1].plot(epoch plot,history 3 3.history[key],label = key)
         ax[1].legend()
         ax[1].set title('BN input training')
      for key in history 3 4.history.keys():
         ax[2].plot(epoch plot, history 3 4.history[key], label = key)
         ax[2].legend()
         ax[2].set title('Drop out training')
```



• After using Dropout layer instead of BN layer, the model converged within one epoch, it trained faster, but the final accuracy and loss seems don't have a significant difference.

(5)

```
In [258...
       model 3 5 = tf.keras.models.Sequential([
             tf.keras.layers.BatchNormalization(input shape = (28,28,1)),
             tf.keras.layers.Conv2D(6, kernel size=5, strides = 1,activation = 'tanh'),
             tf.keras.layers.BatchNormalization(),
             tf.keras.layers.Dropout(0.5),
             tf.keras.layers.AveragePooling2D(pool size =(2,2), strides =2),
             tf.keras.layers.Conv2D(16, kernel size = 5, strides = 1, activation = 'tanh'),
             tf.keras.layers.BatchNormalization(),
             tf.keras.layers.Dropout(0.5),
             tf.keras.layers.AveragePooling2D(pool size = (2,2), strides = 2),
             tf.keras.layers.Conv2D(120, kernel size = 2, strides = 2, activation = 'tanh'),
             tf.keras.layers.BatchNormalization(),
             tf.keras.layers.Dropout(0.5),
             tf.keras.layers.Flatten(),
             tf.keras.layers.Dense(84, activation = 'tanh'),
             tf.keras.layers.BatchNormalization(),
             tf.keras.layers.Dropout(0.5),
             tf.keras.layers.Dense(10, activation = 'softmax'),
          1)
       model 3 5.compile(loss = 'sparse categorical crossentropy',optimizer = 'adam', metrics =
       history 3 5 = model 3 5.fit(x train, y train, epochs = 10, batch size = 64, validation dat
      Epoch 1/10
      - val loss: 0.1129 - val accuracy: 0.9642
      Epoch 2/10
      - val loss: 0.0771 - val accuracy: 0.9751
      Epoch 3/10
      - val loss: 0.0607 - val accuracy: 0.9798
      Epoch 4/10
      - val loss: 0.0539 - val accuracy: 0.9828
      Epoch 5/10
       - val loss: 0.0552 - val accuracy: 0.9830
      Epoch 6/10
      - val loss: 0.0536 - val accuracy: 0.9821
```

```
Epoch 7/10
        - val loss: 0.0489 - val accuracy: 0.9848
        Epoch 8/10
        - val loss: 0.0492 - val accuracy: 0.9836
        Epoch 9/10
        - val loss: 0.0414 - val accuracy: 0.9861
        Epoch 10/10
        - val loss: 0.0436 - val accuracy: 0.9850
In [261...
        fig, ax = plt.subplots(1, 4, figsize = (20, 5))
        epoch plot = range (10)
        for key in history 3 2.history.keys():
            ax[0].plot(epoch plot, history 3 2.history[key], label = key)
            ax[0].legend()
            ax[0].set title('Std norm input training')
        for key in history 3 3.history.keys():
            ax[1].plot(epoch plot,history 3 3.history[key],label = key)
            ax[1].legend()
            ax[1].set title('BN input training')
        for key in history 3 4.history.keys():
            ax[2].plot(epoch plot,history_3_4.history[key],label = key)
            ax[2].legend()
            ax[2].set title('Drop out training')
        for key in history 3 5.history.keys():
            ax[3].plot(epoch plot, history 3 5.history[key], label = key)
            ax[3].legend()
            ax[3].set title('Mixed training')
             Std_norm_input_training
                                  BN input training
                                                      Drop out training
                                                                           Mixed training
                                                                    1.0
        1.0
                            1.0
                                                1.0
                                                                    0.8
        0.8
                            0.8
                                                0.8
                                         loss
                                                0.6
                                                                    0.6
                     accuracy
                                         accuracy
                                                             accuracy
                                                                                 accuracy
                     val loss
                                         val loss
                                                             val loss
                                                                                  val loss
                     val accuracy
                                         val_accuracy
                                                             val_accuracy
                                                                                 val accuracy
                            0.4
                                                0.4
        0.4
                                                                    0.4
        0.2
                            0.2
                                                                    0.2
                                                0.0
                            0.0
                                                                    0.0
```

 According to the above plot, After using BN and drop togather, model significantly reduces it's performance.

Problem 4

```
In [23]: # useful functions and generate data set

def eggholderfx(x_1,x_2):
    part_1 = -(x_2+47)*np.sin(np.sqrt(np.abs(x_1/2 + (x_2+47))))
    part_2 = - x_1*np.sin(np.sqrt(np.abs(x_1-(x_2+47))))
    return part_1 + part_2

def creat_y(x_1,x_2):
    return (eggholderfx(x_1,x_2) + np.random.normal(loc = 0, scale = np.sqrt(0.3)))
    np.random.seed(sed)
```

```
X = np.random.uniform(-512,512,(100000,2))
y = []
for i in X:
    x_i = creat_y(i[0],i[1])
    y.append(x_i)
y = np.array(y)

x_train, x_val, y_train, y_val = train_test_split(X,y,test_size=0.2, random_state=sed)
```

(1)

```
In [86]:
         unit selection = [16,32,64,128,256,512]
          my sgd = tf.keras.optimizers.SGD(nesterov=True)
          def root mean squared error(y true, y pred):
                  return K.sqrt(K.mean(K.square(y pred - y true)))
          def get layer and units(layer num, unit list):
              return list = []
              for i in itertools.product(unit list,repeat = layer num):
                  if sum(i) <= 512:
                      return list.append(i)
              return (return list)
          def my create model(unit list):
              model = keras.models.Sequential([keras.layers.Dense(unit list[0], input shape = (2,), &
              for i, unit in enumerate(unit list):
                  if i != 0:
                      model.add(keras.layers.Dense(unit, activation = 'relu'))
                      model.add(keras.layers.BatchNormalization())
              model.add(keras.layers.Dense(1))
              return model
          def problem4 experiment(layer, epoch num):
              num of units one layer = []
              rmses one layer = []
              num parameters one layer = []
              used time onelayer = []
              for i in range(len(layer)): # change layer parameters here
                  model = my create model(layer[i]) # change layer parameters here
                  model.compile(loss = root mean squared error,
                                optimizer = my sgd,metrics=[tf.keras.metrics.RootMeanSquaredError()]
                  start = time.monotonic()
                  model.fit(x train, y train, epochs=epoch num, batch size = 1000,
                            validation data=(x val, y val), verbose = 0)
                  end = time.monotonic()
                  total time = end-start
                  rmse = np.sqrt(np.mean(np.power((model.predict(x val).reshape(len(y val))-y val),2
                  used time onelayer.append(total time)
                  rmses one layer.append(rmse)
                  num of units one layer.append(sum(layer[i])) # change layer parameters here
                  num parameters one layer.append(model.count params())
                  print('Finished number: '+ str(i+1)+', total number of models: '+str(len(layer)))
              return num of units one layer, rmses one layer, num parameters one layer, used time onela
```

• Here if I want to train all combinations I need to train more than 10 hours on 3070 GPU, So for two layer model and three layer model, I only sample 10 combinations form each and train total 26 models.

```
In [87]:
    random.seed = sed
    one_layer = get_layer_and_units(1,unit_selection)
```

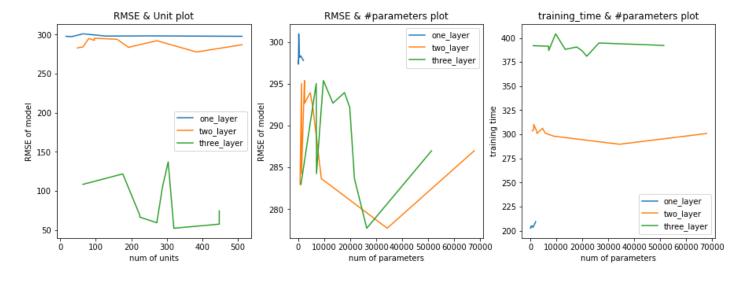
```
three layer = random.sample(get layer and units(3, unit selection), 10)
In [88]:
          # start training one layer model
          units one, rmse one, num parameters one, used time one = problem4 experiment(one layer,200
         Finished number: 1, total number of models: 6
         Finished number: 2, total number of models: 6
         Finished number: 3, total number of models: 6
         Finished number: 4, total number of models: 6
         Finished number: 5, total number of models: 6
         Finished number: 6, total number of models: 6
In [89]:
         # start training two layer model
          units two, rmse two, num parameters two, used time two = problem4 experiment(random.sample
         Finished number: 1, total number of models: 10
         Finished number: 2, total number of models: 10
         Finished number: 3, total number of models: 10
         Finished number: 4, total number of models: 10
         Finished number: 5, total number of models: 10
         Finished number: 6, total number of models: 10
         Finished number: 7, total number of models: 10
         Finished number: 8, total number of models: 10
         Finished number: 9, total number of models: 10
         Finished number: 10, total number of models: 10
In [90]:
         # start training three layer model
         units three, rmse three, num parameters three, used time three = problem4 experiment(rando
         Finished number: 1, total number of models: 10
         Finished number: 2, total number of models: 10
         Finished number: 3, total number of models: 10
         Finished number: 4, total number of models: 10
         Finished number: 5, total number of models: 10
         Finished number: 6, total number of models: 10
         Finished number: 7, total number of models: 10
         Finished number: 8, total number of models: 10
         Finished number: 9, total number of models: 10
         Finished number: 10, total number of models: 10
In [91]:
          units RMSE three sorted = np.array(sorted(zip(units three, rmse three), key = lambda x: x
          params RMSE two sorted = np.array(sorted(zip(num parameters two, rmse two), key = lambda >
          params RMSE three sorted = np.array(sorted(zip(num parameters three, rmse three), key = 1\epsilon
          params time two sorted = np.array(sorted(zip(num parameters two, used time two), key = <math>lar
          params time three sorted = np.array(sorted(zip(num parameters three, used time three), key
          fig, ax = plt.subplots(1, 3, figsize = (15, 5))
          ax[0].plot(units one, rmse one, label = 'one layer')
          ax[0].plot(units RMSE two sorted[:,0], units RMSE two sorted[:,1], label = 'two layer')
          ax[0].plot(units RMSE three sorted[:,0], units RMSE three sorted[:,1], label = 'three laye
          ax[0].set xlabel('num of units')
          ax[0].set ylabel('RMSE of model')
          ax[0].set title('RMSE & Unit plot')
          ax[0].legend()
          ax[1].plot(num parameters one, rmse one, label = 'one layer')
          ax[1].plot(params RMSE two sorted[:,0], params RMSE two sorted[:,1], label = 'two layer')
          ax[1].plot(params RMSE three sorted[:,0], params RMSE two sorted[:,1], label = 'three laye
          ax[1].set xlabel('num of parameters')
          ax[1].set ylabel('RMSE of model')
```

two layer = random.sample(get layer and units(2, unit selection), 10)

```
ax[1].set_title('RMSE & #parameters plot')
ax[1].legend()

ax[2].plot(num_parameters_one, used_time_one, label = 'one_layer')
ax[2].plot(params_time_two_sorted[:,0], params_time_two_sorted[:,1], label = 'two_layer')
ax[2].plot(params_time_three_sorted[:,0], params_time_three_sorted[:,1], label = 'three_lax[2].set_xlabel('num of parameters')
ax[2].set_ylabel('training time')
ax[2].set_title('training_time & #parameters plot')
ax[2].legend()
```

Out[91]: <matplotlib.legend.Legend at 0x1c02706f3a0>



- Sorry the second plot I use the wrong Y value, I should use params_RMSE_three_sorted instead of params_RMSE_two_sorted, however I don't have the time to rerun the result, Plz don't deduct score!
- Acording to the first plot, green line should below the orange line

(2)

- According to the plots, we are excepted to have a lower RMSE when we add more parameters or units to our model
- According to the plot in the middle hand side, going deeper don't seems to always have a lower RMSE.
 From 1 year to 2 layer, the RMSE decreased significantly, however from 2 layers to 3 layers, the RMSE doesn't drop as excepted.
- According to the plot on the right hand side, I don't see a similar trade off in training time. (with same layer, the training time don't have a huge variation as the num of parameters increase)