Fully connected neural network on MNIST dataset

- a) Open the notebook fcn_MNIST.ipynb. In this notebook we use a fully connected neural network to predict the handwritten digits of the MNIST dataset.
 - We have 4000 examples with 784 pixel values and 10 classes. Run the fist 3 cells.
- b) Write the missing TensorFlow code in cell 4 for the first hidden layer.
- c) Run the next two cells to store the graph and do a forward pass of the untrained network. Have a look at the network.
- d) Now lets train the model. Finish the code to calculate the loss and accuracy for the validation set.

Fully connected neural network on MNIST dataset (Tricks)

Note for docker users.

• In this notebook we create different runs so it might be beneficial to save them also outside the docker container. This is possible using the -v option when starting docker.

• If you experience crashes of the docker container do a two step procedure. First start docker in bash.

docker run -p 8888:8888 -p 6006:6006 -v /Users/oli/Documents/workspace/dl_course/:/notebooks

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Then start the jupyter notebook in the console with

jupyter notebook --NotebookApp.token=tensorchiefs

- a) Open the notebook fcn_MNIST_keras and run the first model (execute the cell after training) and visualize the result in TensorBoard (have a look at learning curves and the histograms / distributions of the weights)
- b) Remove the init='zero' argument of the dense layers, to have a proper internalization of your weights. Change the name from name = 'sigmoid_init0' to name = 'sigmoid'. Restart the kernel and repeat the training as in a). Compare the results in TensorBoard, describe your results.

- c) Change the activations / non-linearities from Activation('sigmoid') to Activation('relu') and change the name from name = 'sigmoid' to name = 'relu'. Continue as above, especially have a look at the validation loss do you observe overfitting.
- d) Add a dropout layer: Now add a dropout layer model.add(Dropout(0.3)) between the Dense-Layer and the Activation. Change the name from name = 'relu' to name = 'dropout'.
- e) Add a batch-normalization: Now add a batch-norm layer model.add(BatchNormalization())
 between the Dense-Layer and the Dropout. Change the name from name
 'dropout' to name = 'batch_dropout'. Continue as above

The network should look like:

Layer (type)	Output	Shape	Param #	Connected to
dense_1 (Dense)	(None,	500)	392500	dense_input_1[0][0]
batchnormalization_1 (BatchNorma	(None,	500)	2000	dense_1[0][0]
dropout_1 (Dropout)	(None,	500)	0	batchnormalization_1[0][
activation_1 (Activation)	(None,	500)	0	dropout_1[0][0]
dense_2 (Dense)	(None,	50)	25050	activation_1[0][0]
batchnormalization_2 (BatchNorma	(None,	50)	200	dense_2[0][0]
dropout_2 (Dropout)	(None,	50)	0	batchnormalization_2[0][
activation_2 (Activation)	(None,	50)	0	dropout_2[0][0]
dense_3 (Dense)	(None,	10)	510	activation_2[0][0]

Total params: 420,260 Trainable params: 419,160 Non-trainable params: 1,100