IST 718 Big Data Analytics

Lab 3 (week 9)

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Assignment: Identify Clothing Items

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1 Introduction

Can we use algorithms and compute to identify clothing items? Specifically, can we determine which algorithm and compute methodology provides us the most efficient approach for classifying simple fashion images

1.1 About the Data

Using the base samples available from Zalando Research:

Fashion-mnist

Fashion-MNIST is a dataset of Zalando's article images—consisting of a training set of 60,000 examples and a test set of 10,000 examples. Each example is a 28x28 grayscale image, associated with a label from 10 classes. It's intended that Fashion-MNIST serve as a direct **drop-in replacement** for the original MNIST dataset for benchmarking machine learning algorithms. It shares the same image size and structure of training and testing splits.

- Each row is a separate image
- Column 1 is the class label.
- Remaining columns are pixel numbers (784 total).
- Each value is the darkness of the pixel (1 to 255)

1.1.1 *Labels*

Each training and test example is assigned to one of the following labels:

Label	Description
0	T-shiert/top
1	Trouser
2	Pullover
3	Dress
4	Coat
5	Sandal
6	Shirt
7	Sneaker
8	Bag

Label	Description
9	Ankle boot

1.1.2 Dataset Info

- Full training dataset has 6000 rows and 784 columns
- Full testing dataset has 1000 rows and 784 columns

1.2 **Data Exploration & Cleaning**

X_Train Dataset Info:

class: ndarray shape: (48000, 784) strides: (784, 1) itemsize: 1 byteorder: little type: uint8

X_Test Dataset Info: class: ndarray shape: (10000, 784) strides: (784, 1) itemsize: 1 byteorder: little

type: uint8

Sample Set of Training Images:

Sample Class 0: T-shiert/top Images:

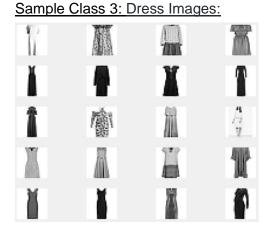




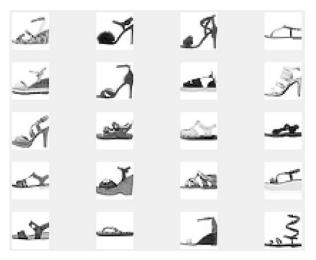
Sample Class 1: Trouser Images:



_ . _ . _ .



Sample Class 5: Sandle Images:



Sample Class 2: Pullover Images:



Sample Class 4: Coat Images:



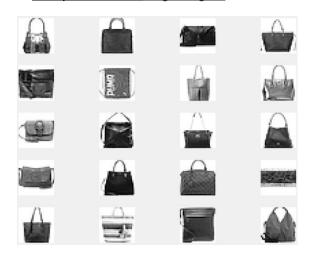


Sample Class 6: Shirt Images:

Sample Class 7: Sneaker Images:



Sample Class 8: Bag Images:



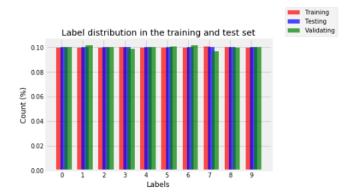
Sample Class 9: Ankle Boot Images:



1.3 Data Transformation

The training dataset of 6000 sample images was split into training/validation (80/20 split) sets for model validation and prediction accuracy measurement. The test dataset of 1000 sample images was held out for prediction accuracy measurements.

The training/validation/testing datasets have nearly equal number of each class label. These datasets are considered balanced.



1.4 Data Normalization

The 'label' attribute was encoded to a catagorical, nominal variable for class classification training/testing. This was done using the sklearn.preprocessing.LabelEncoder class. Encode labels with value between 0 and n_classes-1.

Training and test dataset's were converted to float32 objects to minimizing memory computation needs and normalized from RGB color to black and white (0-1). This was accomplished by dividing the training and test data by 255.

2 Model - Random Forest Classifier

Python Package: scikit-learn v0.21.3 sklearn.ensemble.RandomForestClassifier

A random forest is a meta estimator that fits a number of decision tree classifiers on various subsamples of the dataset and uses averaging to improve the predictive accuracy and control over-fitting. The sub-sample size is always the same as the original input sample size but the samples are drawn with replacement if bootstrap=True (default).

2.1 Analysis and Models

2.1.1 Data Preprocessing

Data preprocessing for this model is described above in section 1.4.

2.2 Model Baseline - Random Forest Classifier

This algorithm uses a perturb-and-combine techniques specifically designed for trees. A diverse set of classifieres are created by introducing randomness in the classifier construction. The prediction of the ensemble is given as the averaged prediction of the individual classifiers. Like decision trees, forests of trees also extend to multi-output problems. In random forests, each tree in the ensemble is built from a sample drawn with replacement. When splitting each node during the construction of a tree, the best split is found either from all input features or a random subset of size max features.

2.2.1 Model Details

{'bootstrap': True,
 'class_weight': None,
 'criterion': 'gini',
 'max_depth': None,
 'max_features': 'auto',
 'max_leaf_nodes': None,
 'min_impurity_decrease': 0.0,
 'min_impurity_split': None,
 'min_samples_leaf': 1,
 'min_samples_split': 2,
 'min_weight_fraction_leaf': 0.0,
 'n_estimators': 100,

'n_estimators': 100,
'n_jobs': None,
'oob_score': False,
'random_state': None,

'verbose': 2,

'warm_start': False}

2.2.2 Model Parameters

For this model, all default parameters were selected. Below is a listing of those parameters. For parameter tuning techniques, see <u>this</u> for more details.

Parameter and Value

Description

	-
n_estimators=100	The number of trees in the forest.
criterion="gini"	The function to measure the quality of a split. Supported criteria are "gini" for the Gini impurity and "entropy" for the information gain.
max_depth=None	The maximum depth of the tree. If None, then nodes are expanded until all leaves are pure or until all leaves contain less than min_samples_split samples
min_samples_split=2	The minimum number of samples required to split an internal node
min_samples_leaf=1	The minimum number of samples required to be at a leaf node. A split point at any depth will only be considered if it leaves at least min_samples_leaf training samples in each of the left and right branches
min_weight_fraction_leaf=0.0	The minimum weighted fraction of the sum total of weights (of all the input samples) required to be at a leaf node. Samples have equal weight when sample_weight is not provided.
max_features="auto"	The number of features to consider when looking for the best split: If "auto", then max_features=sqrt(n_features).
max_leaf_nodes=None	Grow trees with max_leaf_nodes in best-first fashion. Best nodes are defined as relative reduction in impurity. If None then unlimited number of leaf nodes.
min_impurity_decrease=0.0	A node will be split if this split induces a decrease of the impurity greater than or equal to this value.
min_impurity_split=None	Threshold for early stopping in tree growth. A node will split if its impurity is above the threshold, otherwise it is a leaf.
bootstrap=True	Whether bootstrap samples are used when building trees. If False, the whole datset is used to build each tree.
oob_score=False	Whether to use out-of-bag samples to estimate the generalization accuracy.
n_jobs=None	The number of jobs to run in parallel for both fit and predict. None means 1 unless in a joblib.parallel_backend context1 means using all processors.
random_state=None	if int, random_state is the seed used by the random number generator; If RandomState instance, random_state is the random number generator; If None, the random number generator is the RandomState instance used by np.random
verbose=2	Controls the verbosity when fitting and predicting.
warm_start=False	When set to True, reuse the solution of the previous call to fit and add more estimators to the

ensemble, otherwise, just fit a whole new forest.	
class_weight=None	Weights associated with classes in the form {class_label: weight}. If not given, all classes are supposed to have weight one. For multi-output problems, a list of dicts can be provided in the same order as the columns of y.

The main parameters to adjust when using these methods is n_estimators and max_features. The former is the number of trees in the forest. The larger the better, but also the longer it will take to compute. Also, the results will stop getting significantly better beyone a critical number of trees. The later is the size of the random subsets of features to consider when splitting a node. The lower the greater the reduction of variance, but also the greater the increase in bias.

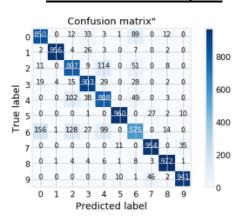
2.2.2.1 Model Results

l	Fit Score	% Accuracy	Fit Time	Score Time	Predict Time	Precision	Recal	F1-Score
	0.8806	87%	84.92204	0.0	0.5337	0.87	0.87	0.87

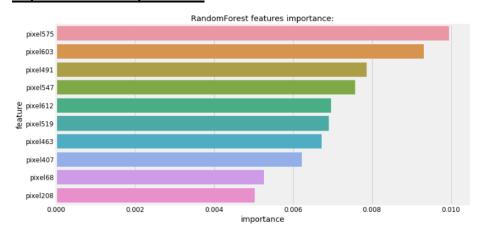
Classification Report:

		precision	recall	f1-score	support
	Class0	0.82	0.85	0.83	1000
	Class1	0.99	0.96	0.97	1000
	Class2	0.75	0.81	0.78	1000
	Class3	0.87	0.90	0.88	1000
	Class4	0.76	0.81	0.78	1000
	Class5	0.98	0.96	0.97	1000
	Class6	0.71	0.57	0.64	1000
	Class7	0.93	0.95	0.94	1000
	Class8	0.96	0.97	0.96	1000
	Class9	0.95	0.94	0.95	1000
a	ccuracy			0.87	10000
ma	cro avg	0.87	0.87	0.87	10000
weigh	ted avg	0.87	0.87	0.87	10000

Confusion Matrix Report:



Top 10 Feature Importance:



Model Tuned - Random Forest Classification 2.3

Model Details 2.3.1

{'bootstrap': True, 'class_weight': None, 'criterion': 'gini', 'max depth': 10, 'max_features': 'auto', 'max_leaf_nodes': None, 'min_impurity_decrease': 0.0, 'min_impurity_split': None, 'min_samples_leaf': 1, 'min_samples_split': 2,

'min_weight_fraction_leaf': 0.0,

'n_estimators': 200,

'n_jobs': 5,

'oob_score': False, 'random_state': 42,

'verbose': 0,

'warm_start': False}

2.3.2 **Model Parameters**

Parameter and Value

n_estimators=200	The number of trees in the forest.
criterion="gini"	The function to measure the quality of a split. Supported criteria are "gini" for the Gini impurity and "entropy" for the information gain.
max_depth=10	The maximum depth of the tree. If None, then nodes are expanded until all leaves are pure or until all leaves contain less than min_samples_split samples
min_samples_split=2	The minimum number of samples required to split an internal node
min_samples_leaf=1	The minimum number of samples required to be at a leaf node. A split point at any depth will only be considered if it leaves at least min_samples_leaf training samples in each of the left and right branches
min_weight_fraction_leaf=0.0	The minimum weighted fraction of the sum total of weights (of all the input samples) required to be at a leaf node. Samples have equal weight when sample_weight is not provided.
max_features="auto"	The number of features to consider when looking for the best split: If "auto", then max_features=sqrt(n_features).
max_leaf_nodes=None	Grow trees with max_leaf_nodes in best-first fashion. Best nodes are defined as relative reduction in impurity. If None then unlimited number of leaf nodes.
min_impurity_decrease=0.0	A node will be split if this split induces a decrease of the impurity greater than or equal to this value.

Description

min_impurity_split=None	Threshold for early stopping in tree growth. A node will split if its impurity is above the threshold, otherwise it is a leaf.
bootstrap=True	Whether bootstrap samples are used when building trees. If False, the whole datset is used to build each tree.
oob_score=False	Whether to use out-of-bag samples to estimate the generalization accuracy.
n_jobs=5	The number of jobs to run in parallel for both fit and predict. None means 1 unless in a joblib.parallel_backend context1 means using all processors.
random_state=42	if int, random_state is the seed used by the random number generator; If RandomState instance, random_state is the random number generator; If None, the random number generator is the RandomState instance used by np.random
verbose=0	Controls the verbosity when fitting and predicting.
warm_start=False	When set to True, reuse the solution of the previous call to fit and add more estimators to the ensemble, otherwise, just fit a whole new forest.
class_weight=None	Weights associated with classes in the form {class_label: weight}. If not given, all classes are supposed to have weight one. For multi-output problems, a list of dicts can be provided in the same order as the columns of y.

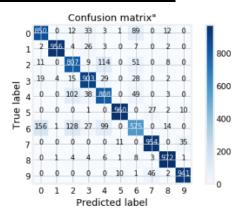
2.3.3 Model Results

Fit Score	% Accuracy	Fit Time	Score Time	Predict Time	Precision	Recal	F1-Score
0.851	84%	25.9838	0.0	0.4462	0.85	0.85	0.85

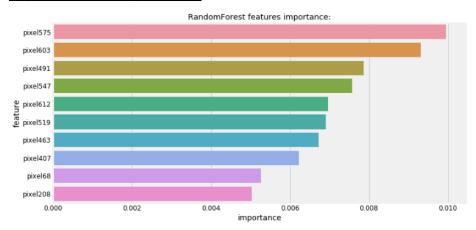
Classification Report:

		precision	recall	f1-score	support
	Class0	0.79	0.84	0.82	1202
	Class1	1.00	0.95	0.98	1219
	Class2	0.74	0.75	0.74	1205
	Class3	0.81	0.91	0.86	1184
	Class4	0.69	0.82	0.75	1202
	Class5	0.97	0.92	0.95	1211
	Class6	0.75	0.50	0.60	1218
	Class7	0.89	0.92	0.91	1159
	Class8	0.96	0.96	0.96	1197
	Class9	0.92	0.94	0.93	1203
a	ccuracy			0.85	12000
ma	cro avg	0.85	0.85	0.85	12000
weigh	ted avg	0.85	0.85	0.85	12000

Confusion Matrix Report:



Top 10 Feature Importance:



3 Model - XGBoosted Trees

XGBoost is an optimized distributed gradient boosting library designed to be highly efficient, flexible and portable. It implements machine learning algorithms under the Gradient Boosting framework. XGBoost provides a parallel tree boosting.

See documentation for more indepth details.

- XGBoost Documentation
- XGBoost Parameter details

3.1 Analysis and Models

3.1.1 Data Preprocessing

Data preprocessing for this model is described above in section 1.4.

3.2 Model Baseline - XGBoost

3.2.1 Model Details

```
{'base_score': 0.5,
'booster': 'gbtree',
'colsample_bylevel': 1,
'colsample_bynode': 1,
'colsample_bytree': 1,
'gamma': 0,
'learning_rate': 0.1,
'max delta step': 0,
'max_depth': 5,
'min_child_weight': 1,
'missing': nan,
'n_estimators': 50,
'nthread': 1,
'objective': 'multi:softprob',
'reg_alpha': 0,
'reg lambda': 1,
'scale pos weight': 1,
'seed': 0,
'subsample': 1,
'verbosity': 1}
```

3.2.2 Model Parameters

All other default parameters used for this model.

Parameter and Value Description

n_estimators=50	int Number of boosted trees to fit

max_depth=5	The maximum depth of the tree. If None, then nodes are expanded until all leaves are pure or
	until all leaves contain less than min_samples_split samples

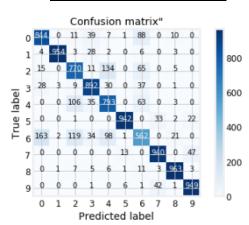
3.2.3 Model Results

Fit Score	% Accuracy	Fit Time	Score Time	Predict Time	Precision	Recal	F1-Score
0.8688	86%	654.5434	0.0	0.7612	0.86	0.86	0.86

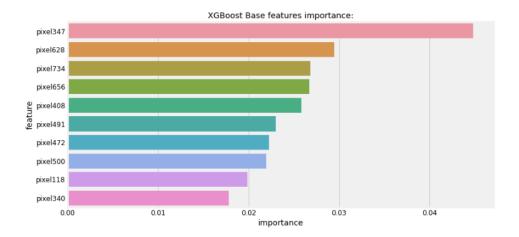
Classification Report:

	precision	recall	f1-score	support	
	•				
Class0	0.80	0.84	0.82	1000	
Class1	0.99	0.95	0.97	1000	
Class2	0.75	0.77	0.76	1000	
Class3	0.85	0.89	0.87	1000	
Class4	0.74	0.79	0.77	1000	
Class5	0.98	0.94	0.96	1000	
Class6	0.67	0.56	0.61	1000	
Class7	0.92	0.94	0.93	1000	
Class8	0.95	0.96	0.96	1000	
Class9	0.93	0.95	0.94	1000	
accuracy			0.86	10000	
macro avg	0.86	0.86	0.86	10000	
weighted avg	0.86	0.86	0.86	10000	

Confusion Matrix Report:



Top 10 Feature Importance



3.3 Model Tuned - XGBoost

3.3.1 Model Details

XGBoost with GridSearch, an exhaustive search over specified parameter values for an estimator. GridSearchCV implements a 'fit' a 'score', and 'predict' method. The parameters of the estimator used to apply these methods are optimized by cross-validated grid-search over a parameter grid.

Python package: scikit-learn v0.21.3 sklearn.model_selection.GridSearchCV

```
{'base_score': 0.5,
'booster': 'gbtree',
'colsample_bylevel': 1,
'colsample bynode': 1,
'colsample_bytree': 0.8,
'gamma': 0,
'learning_rate': 0.1,
'max_delta_step': 0,
'max_depth': 10,
'min_child_weight': 1,
'missing': nan,
'n_estimators': 100,
'nthread': 1,
'objective': 'multi:softprob',
'reg_alpha': 0,
'reg_lambda': 1,
'scale_pos_weight': 1,
'seed': 42,
'subsample': 1,
'verbosity': 1}
```

3.3.2 Model Parameters

Using GridSearch parameters specified below, all other parameters are set to default values.

```
param_grid = [{
    'max_depth':[5,10],
    'n_estimators':[100],
    'learning_rate':[0.05, 0.1],
    'colsample_bytree':[0.8, 0.95]
}]
```

Parameter and Value

Description

estimator	This is assumed to implement the scikit-learn estimator interface. Either estimator needs to provide a score function, or scoring must be passed.
param_grid	Dictionary with parameters names (string) as keys and lists of parameter settings to try as values, or a list of such dictionaries, in which case the grids spanned by each dictionary in the list are explored. This enables searching over any sequence of parameter settings.

scoring	A single string or a callable to evaluate the predictions on the test set.
n_jobs	Number of jobs to run in parallel. None means 11means using all processers
pre_dispatch	Controls the number of jobs that get dispatched during parallel execution. Reducing this number can be useful to avoid an explosion of memory consumption when more jobs get dispatched than CPUs can process. This parameter can be: None, in which case all the jobs are immediately created and spawned. Use this for lightweight and fast-running jobs, to avoid delays due to on-demand spawning of the jobs. An int, giving the exact number of total jobs that are spawned. A string, giving an expression as a function of n_jobs, as in '2*n_jobs'.
iid	If True, return the average score across folds, weighted by the number of samples in each test set. If False, return the average score across folds. Default is True
cv	Determines the cross-validation splitting strategy. Possible inputs for cv are: None, to use the default 3-fold cross validation, integer, to specify the number of folds in a (Stratified)KFold
refit	Refit an estimator using the best found parameters on the whole dataset. For multiple metric evaluation, this needs to be a string denoting the scorer that would be used to find the best parameters for refitting the estimator at the end.
verbose	Controls the verbosity; the higher, the more messages
error_score	Value to assign to the score if an error in estimator fitting. If set to 'raise', the error is raised. If a numeric value is given, FitFailedWanring is raised.
return_train_score	If False, the cv_results_ attribute will not include training scores. Computing training scores is used to get insights on how different parameter settings impact the overfitting/underfitting trade-off. However computing the scores on the training set can be computationally expensive and is not strictly required to select the parameters that yield the best generalization performance.

3.3.3 Model Results

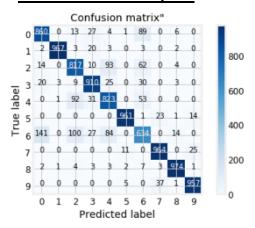
Fit Score	% Accuracy	Fit Time	Score Time	Predict Time	Precision	Recal	F1-Score
0.8961	89%	25462.3460	0.0	2.6526	0.89	0.89	0.89

Top 10

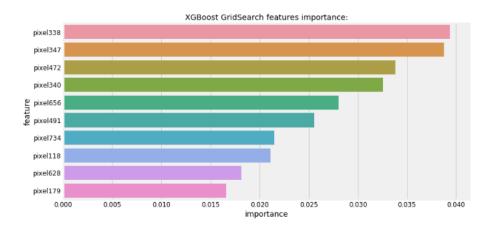
Classification Report:

precision recall f1-score support Class0 0.83 0.86 0.84 Class1 0.99 0.97 0.98 1000 0.80 Class2 0.79 0.82 1000 Class3 0.89 0.91 0.90 1000 Class4 0.80 0.82 0.81 1000 0.98 1000 Class5 0.96 0.97 0.63 0.67 1000 Class6 0.72 Class7 0.94 0.96 0.95 1000 Class8 0.97 0.97 0.97 1000 0.96 0.96 1000 Class9 0.96 accuracy 0.89 10000 macro avg 0.89 0.89 0.89 10000 weighted avg 0.89 10000 0.89 0.89

Confusion Matrix Report:



Feature Importance:



4 Neural Networks with Keras

Python package: keras

Keras is a high-level neural networks API, written in Python and capable of running on top of TensorFlow, CNTK, or Theano. It was developed with a focus on enabling fast experimentation.

- Allows for easy and fast prototyping (through user friendliness, modularity, and extensibility).
- Supports both convolutional networks and recurrent networks, as well as combinations of the two.
- Runs seamlessly on CPU and GPU.

4.1 Analysis and Models

4.1.1 Data Preprocessing

Data preprocessing for this model is described above in section 1.3 and 1.4.

Additionally, one hot encoding was performed to convert class vectores to binary class matrices. This means that a column will be created for each output category and a binary variable is inputted for each category.

4.2 Model Baseline - Single Layer Neural Network

4.2.1 Model Details

This is a Keras Sequential model, with two Dense layers. The Sequential model is a linear stack of layers. The input_shape for this model is the number of pixels for this dataset of images which is 784.

The compile method, which configures the model for training, uses an 'adam' optimizer and for loss the 'categorical_crossentropy' objective function.

This models most notable training (fit) parameters are:

- epochs=300
- batch size=1000
- verbose=2
- shuffle=True

4.2.1.1 Parameters - Input Dense Layer

```
{'batch_size': 1000,
  'epochs': 300,
  'steps': None,
  'samples': 48000,
  'verbose': 2,
  'do_validation': True,
  'metrics': ['loss', 'acc', 'val loss', 'val acc']}
```

Parameter and Value

Description

units=784	Positive integer, dimensionality of the output space
input_dim=784	Input shape, nD tensor
kernel_initializer='normal'	Initializer for the kernel weights matrix
activation='relu'	Activation function to use. If you don't specify anything, no activation is applied (i.e., 'linear' activation)

4.2.1.2 Parameters - Output Dense Layer

Parameter and Value

Description

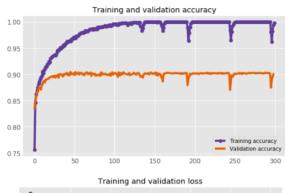
unit=10	Positive integer, dimensionality of the output space			
kernel_initializer='normal'	Initializer for the kernel weights matrix			
activation='softmax'	Activation function to use.			

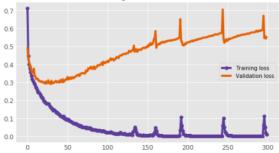
4.2.1.3 Model Summary

Layer (type)	Output Shape	Param #
dense 1 (Dense)	(None, 784)	615440
dense_1 (bense)	(NONE, 704)	013440
dense_2 (Dense)	(None, 10)	7850

Total params: 623,290 Trainable params: 623,290 Non-trainable params: 0

Training and Validation Accuracy and Validation Loss





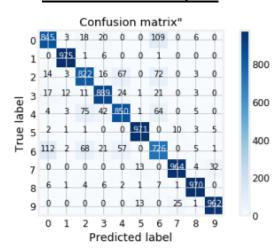
4.2.2 Model Results

Fit Score	% Accuracy	Fit Time	Score Time	Predict Time	Precision	Recal	F1-Score
0.897	90%	676.8362	0.6127	0.0	0.90	0.90	0.90

Classification Report:

	precision	recall	f1-score	support
Class0	0.84	0.84	0.84	1000
Class1	0.99	0.97	0.98	1000
Class2	0.82	0.82	0.82	1000
Class3	0.91	0.89	0.90	1000
Class4	0.81	0.85	0.83	1000
Class5	0.98	0.97	0.97	1000
Class6	0.73	0.73	0.73	1000
Class7	0.95	0.96	0.96	1000
Class8	0.97	0.97	0.97	1000
Class9	0.96	0.96	0.96	1000
accuracy			0.90	10000
macro avg	0.90	0.90	0.90	10000
weighted avg	0.90	0.90	0.90	10000

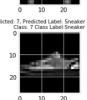
Confusion Matrix Report:



Sample Correct Predicted Images:

Predicted: 1, Predicted Label: Trouser
Class: 1 Class Label: Trouser
0
10
20
0 10 20
Predicted: 2, Predicted Label: Pullover

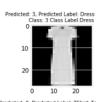
edicted: 2, Predicted Label: Pullover Class: 2 Class Label: Pullover 0













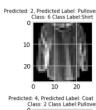


Sample Incorrect Predicted Images:















4.3 Model - Convelutional Neural Network

Deciding on 3x3 as the optimal kernel size for CNN, see here for details.

4.3.1 Model Details

This is a Keras Sequential model, with three 2D convelutional layers. The Sequential model is a linear stack of layers. The input_shape for this model is a 3x3x1. The 2D convolution filter like 3x3 will always have a third dimension in size. The third dimension is equal to the number of channels of the input image. For gray-scale images such as this dataset, that has 1 black and white channel, the third dimension is a 1. Our input_shape is then, (28, 28, 1)

The compile method, which configures the model for training, uses an <u>'adam'</u> optimizer and for loss the <u>'categorical crossentropy'</u> objective function.

This models most notable training (fit) parameters are:

- epochs=300
- batch size=1000
- verbose=2
- shuffle=True

4.3.1.1 Parameters - Layer 1: Conv2D

Parameter and Value	Description

filters=28	Integer, the dimensionality of the output space (i.e. the number of output filters in the convolution)
kernel_size=3	An integer or tuple/list of 2 integers, specifying the height and width of the 2D convolution window. Can be a single integer to specify the same value for all spatial dimensions.
padding='same'	One of 'valid' or 'same'
activation='relu'	Activation function to use.
kernel_initializer='he_normal'	Initializer for the kernel weights matrix
input_shape=(28, 28, 1)	When using this layer as the first layer in a model, this keyword argument is required. Tuble of integers specifing the image shape.

4.3.1.2 Parameters - Layer 2: MaxPooling2D

Parameter and value	Description
pool_size=(2,2)	Integer or tuple of 2 integers, factors by which to downscale (vertical, horizontal). (2,2) will halve the input in both spatial dimension. If only one integer is specified, the same window length will be used for both dimesions.

4.3.1.3 Parameters - Layer 3: Dropout

Doromotor and Value	Description
Parameter and Value	Description

rate=0.25	Float between 0 and 1. Fraction of the input units to drop.

4.3.1.4 Parameters - Layer 4: Conv2D

Parameter and Value

Description

filters=64	Integer, the dimensionality of the output space (i.e. the number of output filters in the convolution)
kernel_size=3	An integer or tuple/list of 2 integers, specifying the height and width of the 2D convolution window. Can be a single integer to specify the same value for all spatial dimensions.
padding='same'	One of 'valid' or 'same'
activation='relu'	Activation function to use.

4.3.1.5 Parameters - Layer 5: MaxPooling2D

Parameter and Value

Description

pool_size=(2,2)	Integer or tuple of 2 integers, factors by which to downscale (vertical, horizontal). (2,2) will halve the input in both spatial dimension. If only one integer is specified, the same window length will be used for both dimensions.
	be used for both dimesions.

4.3.1.6 Parameters - Layer 6: Dropout

Parameter and Value

Description

rate=0.25	Float between 0 and 1. Fraction of the input units to drop.
rate=0.25	Float between 0 and 1. Fraction of the input units to drop.

4.3.1.7 Parameters - Layer 7: Conv2D

Parameter and Value

Description

filters=128	Integer, the dimensionality of the output space (i.e. the number of output filters in the convolution)
kernel_size=3	An integer or tuple/list of 2 integers, specifying the height and width of the 2D convolution window. Can be a single integer to specify the same value for all spatial dimensions.
activation='relu'	Activation function to use.

4.3.1.8 Parameters - Layer 8: Dropout

Parameter and Value

Description

rate=0.4	Float between 0 and 1. Fraction of the input units to drop.

4.3.1.9 Parameters - Layer 9: Flatten

Parameter and Value Description

4.3.1.10 Parameters - Layer 10: Dense

Parameter and Value Description

unit=748	Positive integer, dimensionality of the output space
activation='relu'	Activation function to use.

4.3.1.11 Parameters - Layer 11: Dropout

Parameter and Value Description

rate=0.3	Float between 0 and 1. Fraction of the input units to drop.

4.3.1.12 Parameters - Layer 12: Dense

Parameter and Value Description

unit=10	Positive integer, dimensionality of the output space
activation='softmax'	Activation function to use.

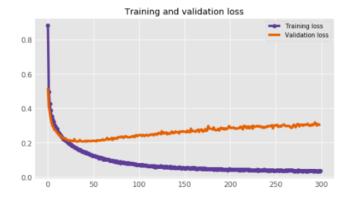
4.3.1.13 Model Summary

Layer (type)	Output	Shape	Param #
conv2d_4 (Conv2D)	(None,	28, 28, 28)	280
max_pooling2d_3 (MaxPooling2	(None,	14, 14, 28)	0
dropout_5 (Dropout)	(None,	14, 14, 28)	0
conv2d_5 (Conv2D)	(None,	12, 12, 64)	16192
max_pooling2d_4 (MaxPooling2	(None,	6, 6, 64)	0
dropout_6 (Dropout)	(None,	6, 6, 64)	0
conv2d_6 (Conv2D)	(None,	4, 4, 128)	73856
dropout_7 (Dropout)	(None,	4, 4, 128)	0
flatten_2 (Flatten)	(None,	2048)	0
dense_7 (Dense)	(None,	784)	1606416
dropout_8 (Dropout)	(None,	784)	0
dense_8 (Dense)	(None,	10)	7850

Total params: 1,704,594 Trainable params: 1,704,594 Non-trainable params: 0

Training and Validation Accuracy and Validation Loss





4.3.2 Model Results

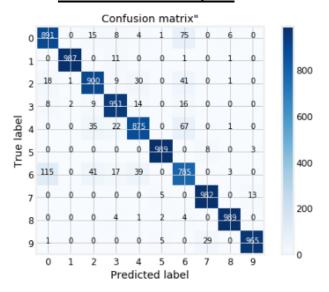
Fit Score	% Accuracy	Fit Time	Score Time	Predict Time	Precision	Recal	F1-Score
0.9314	93%	24986.7392	4.5842	0.0	0.93	0.93	0.93

Classification Report:

Each of the classes highlighted in green below represent significant improvements in predicting these classes correctly over all other models evaluated in this report.

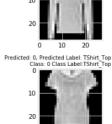
		precision	recall	f1-score	support
	Class0	0.86	0.89	0.88	1000
	Class1	1.00	0.99	0.99	1000
	Class2	0.90	0.90	0.90	1000
	Class3	0.93	0.95	0.94	1000
	Class4	0.91	0.88	0.89	1000
	Class5	0.99	0.99	0.99	1000
	Class6	0.79	0.79	0.79	1000
	Class7	0.96	0.98	0.97	1000
	Class8	0.99	0.99	0.99	1000
	Class9	0.98	0.96	0.97	1000
	ccuracy cro avg	0.93	0.93	0.93 0.93	10000 10000
weigh	ted avg	0.93	0.93	0.93	10000

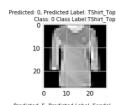
Confusion Matrix Report:

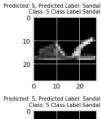


Sample Correct Predicted Images:

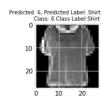
10 Predicted: 2, Predicted Label: Pullover Class: 2 Class Label:Pullover 0

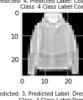






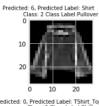






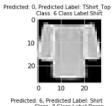


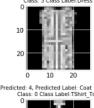
Sample Incorrect Predicted Images:

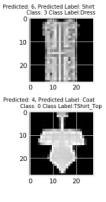


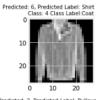


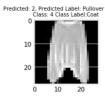












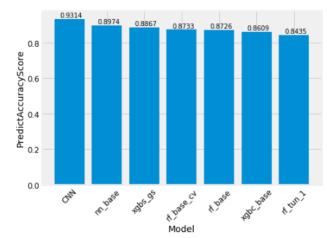


5 Algorithm Performance Comparison

Model Performance Ranking Bar Graph, sorted by PredictAccuracyScore descending:

Overall, the Convelutional Neural Network had the best predictor accuracy score, however it's compute time was second to last.

The CNN performed significantly better than all other models in classifing the more challenging images (Class6-Shirt, Class4-Coat, Class2-Pullover). Below is a breakdown of each models f1-score per class prediction.



	Model	Class0	Class1	Class2	Class3	Class4	Class5	Class6	Class7	Class8	Class9
0	CNN	0.88	0.99	0.90	0.94	0.89	0.99	0.79	0.97	0.99	0.97
1	nn_base	0.84	0.98	0.82	0.90	0.83	0.97	0.73	0.96	0.97	0.96
2	xgbs_gs	0.84	0.98	0.80	0.90	0.81	0.97	0.67	0.95	0.97	0.96
3	xgbs_base	0.82	0.97	0.76	0.86	0.77	0.96	0.61	0.93	0.96	0.94
4	rf_tun_1	0.82	0.98	0.74	0.86	0.75	0.95	0.60	0.91	0.96	0.93
5	rf_base	0.83	0.97	0.78	0.88	0.78	0.97	0.64	0.94	0.96	0.95

<u>Model Performance Ranking Bar Graph, sorted by log of Total Compute Time ascending:</u>

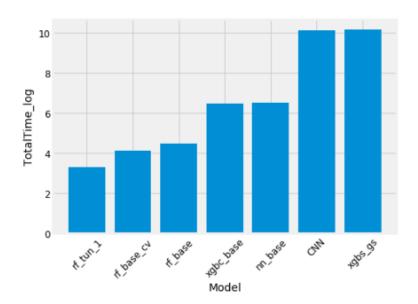
	ModelName	TestAccuracy Score	PredictAccuracy Score	TrainTime	TestTime	ScoreTime	PredictTime	TotalTime
1	CNN	0.931400	0.9314	24986.739215	0.000000	4.584205	0.000000	24991.323420
0	nn_base	0.897400	0.8974	676.836291	0.000000	0.612764	0.000000	677.449054
4	xgbs_gs	0.896167	0.8867	25462.346012	1.900499	0.000000	2.652688	25466.899199
1	rf_base_cv	0.883280	0.8733	59.931356	0.000000	0.987812	0.520665	61.439834
0	rf_base	0.880667	0.8726	84.922047	0.641522	0.000000	0.533718	86.097286
3	xgbc_base	0.868833	0.8609	654.543446	0.609808	0.000000	0.761276	655.914530
2	rf_tun_1	0.851000	0.8435	25.983807	0.433044	0.000000	0.446281	26.863133

Model Performance Ranking Table, sorted by Total Compute Time ascending:

**Note - The compute time for these models are based on the processing power of the system. These were ran locally, on an Intel Core i7-8650U CPU@1.90 GHz (8 CPUs)

The tuned Random Forest had the fastest compute times, however it was last in prediction accuracy scoring at 84%.

The tradeoff between these models is accuracy versus compute time. It would depend on the use case in chosing which model to select.



	ModelName	TestAccuracyScore	PredictAccuracy Score	TrainTime	TestTime	ScoreTime	PredictTime	TotalTime
2	rf_tun_1	0.851000	0.8435	25.983807	0.433044	0.000000	0.446281	26.863133
1	rf_base_cv	0.883280	0.8733	59.931356	0.000000	0.987812	0.520665	61.439834
0	rf_base	0.880667	0.8726	84.922047	0.641522	0.000000	0.533718	86.097286
3	xgbc_base	0.868833	0.8609	654.543446	0.609808	0.000000	0.761276	655.914530
0	nn_base	0.897400	0.8974	676.836291	0.000000	0.612764	0.000000	677.449054
1	CNN	0.931400	0.9314	24986.739215	0.000000	4.584205	0.000000	24991.323420
4	xgbs_gs	0.896167	0.8867	25462.346012	1.900499	0.000000	2.652688	25466.899199

6 Appendix - Submission Items

- Report with graphics
- Supporting notebook for the report
- Description of your tool and methodology if you used another statistical/software package

7 Appendix B - Behind Fashion-MNIST

The original MNIST dataset contains a lot of handwritten digits. Members of the AI/ML/Data Science community love this dataset and use it as a benchmark to validate their algorithms. In fact, MNIST is often the first dataset researchers try. "If it doesn't work on MNIST, it won't work at all", they said. "Well, if it does work on MNIST, it may still fail on others."

- MNIST is too easy. Convolutional nets can achieve 99.7% on MNIST. Classic machine learning algorithms can also achieve 97% easily. Check out <u>our side-by-side benchmark for Fashion-MNIST vs. MNIST</u>, and read "Most pairs of MNIST digits can be distinguished pretty well by just one pixel."
- **MNIST is overused.** In this April 2017 Twitter thread, Google Brain research scientist and deep learning expert Ian Goodfellow calls for people to move away from MNIST.
- MNIST can not represent modern CV tasks, as noted in this April 2017 Twitter thread, deep learning expert/Keras author François Chollet.