

finding_donors

April 17, 2020

0.1 Supervised Learning

0.2 Project: Finding Donors for *CharityML*

In this notebook, some template code has already been provided for you, and it will be your job to implement the additional functionality necessary to successfully complete this project. Sections that begin with '**Implementation**' in the header indicate that the following block of code will require additional functionality which you must provide. Instructions will be provided for each section and the specifics of the implementation are marked in the code block with a '`TODO`' statement. Please be sure to read the instructions carefully!

In addition to implementing code, there will be questions that you must answer which relate to the project and your implementation. Each section where you will answer a question is preceded by a '**Question X**' header. Carefully read each question and provide thorough answers in the following text boxes that begin with '**Answer:**'. Your project submission will be evaluated based on your answers to each of the questions and the implementation you provide.

Note: Please specify WHICH VERSION OF PYTHON you are using when submitting this notebook. Code and Markdown cells can be executed using the **Shift + Enter** keyboard shortcut. In addition, Markdown cells can be edited by typically double-clicking the cell to enter edit mode.

0.3 Getting Started

In this project, you will employ several supervised algorithms of your choice to accurately model individuals' income using data collected from the 1994 U.S. Census. You will then choose the best candidate algorithm from preliminary results and further optimize this algorithm to best model the data. Your goal with this implementation is to construct a model that accurately predicts whether an individual makes more than \$50,000. This sort of task can arise in a non-profit setting, where organizations survive on donations. Understanding an individual's income can help a non-profit better understand how large of a donation to request, or whether or not they should reach out to begin with. While it can be difficult to determine an individual's general income bracket directly from public sources, we can (as we will see) infer this value from other publically available features.

The dataset for this project originates from the [UCI Machine Learning Repository](#). The dataset was donated by Ron Kohavi and Barry Becker, after being published in the article "*Scaling Up the Accuracy of Naive-Bayes Classifiers: A Decision-Tree Hybrid*". You can find the article by Ron Kohavi [online](#). The data we investigate here consists of small changes to the original dataset, such as removing the '`fnlwgt`' feature and records with missing or ill-formatted entries.

0.4 Exploring the Data

Run the code cell below to load necessary Python libraries and load the census data. Note that the last column from this dataset, 'income', will be our target label (whether an individual makes more than, or at most, \$50,000 annually). All other columns are features about each individual in the census database.

```
In [1]: # Import libraries necessary for this project
import numpy as np
import pandas as pd
from time import time
from IPython.display import display # Allows the use of display() for DataFrames

# Import supplementary visualization code visuals.py
import visuals as vs

# Pretty display for notebooks
%matplotlib inline

# Load the Census dataset
data = pd.read_csv("census.csv")

# Success - Display the first record
print(data.shape[0])
display(data.head())
```

45222

	age	workclass	education_level	education-num	marital-status	\
0	39	State-gov	Bachelors	13.0	Never-married	
1	50	Self-emp-not-inc	Bachelors	13.0	Married-civ-spouse	
2	38	Private	HS-grad	9.0	Divorced	
3	53	Private	11th	7.0	Married-civ-spouse	
4	28	Private	Bachelors	13.0	Married-civ-spouse	

	occupation	relationship	race	sex	capital-gain	\
0	Adm-clerical	Not-in-family	White	Male	2174.0	
1	Exec-managerial	Husband	White	Male	0.0	
2	Handlers-cleaners	Not-in-family	White	Male	0.0	
3	Handlers-cleaners	Husband	Black	Male	0.0	
4	Prof-specialty	Wife	Black	Female	0.0	

	capital-loss	hours-per-week	native-country	income
0	0.0	40.0	United-States	<=50K
1	0.0	13.0	United-States	<=50K
2	0.0	40.0	United-States	<=50K

3	0.0	40.0	United-States	<=50K
4	0.0	40.0	Cuba	<=50K

0.4.1 Implementation: Data Exploration

A cursory investigation of the dataset will determine how many individuals fit into either group, and will tell us about the percentage of these individuals making more than \$50,000. In the code cell below, you will need to compute the following: - The total number of records, 'n_records' - The number of individuals making more than \$50,000 annually, 'n_greater_50k'. - The number of individuals making at most \$50,000 annually, 'n_at_most_50k'. - The percentage of individuals making more than \$50,000 annually, 'greater_percent'.

**** HINT: **** You may need to look at the table above to understand how the 'income' entries are formatted.

```
In [2]: data.describe()
```

```
Out[2]:
```

	age	education-num	capital-gain	capital-loss	hours-per-week
count	45222.000000	45222.000000	45222.000000	45222.000000	45222.000000
mean	38.547941	10.118460	1101.430344	88.595418	40.938017
std	13.217870	2.552881	7506.430084	404.956092	12.007508
min	17.000000	1.000000	0.000000	0.000000	1.000000
25%	28.000000	9.000000	0.000000	0.000000	40.000000
50%	37.000000	10.000000	0.000000	0.000000	40.000000
75%	47.000000	13.000000	0.000000	0.000000	45.000000
max	90.000000	16.000000	99999.000000	4356.000000	99.000000

```
In [3]: print(data["income"].unique())
print(">50K", len(data[data["income"] == ">50K"]))
print("<=50K", len(data[data["income"] == "<=50K"]))
```

```
['<=50K' '>50K']
>50K 11208
<=50K 34014
```

```
In [4]: # TODO: Total number of records
n_records = data.shape[0]

# TODO: Number of records where individual's income is more than $50,000
n_greater_50k = len(data[data["income"] == ">50K"])

# TODO: Number of records where individual's income is at most $50,000
n_at_most_50k = len(data[data["income"] == "<=50K"])

# TODO: Percentage of individuals whose income is more than $50,000
greater_percent = n_greater_50k / n_records * 100

# Print the results
```

```

print("Total number of records: {}".format(n_records))
print("Individuals making more than $50,000: {}".format(n_greater_50k))
print("Individuals making at most $50,000: {}".format(n_at_most_50k))
print("Percentage of individuals making more than $50,000: {}%".format(greater_percent))

```

Total number of records: 45222

Individuals making more than \$50,000: 11208

Individuals making at most \$50,000: 34014

Percentage of individuals making more than \$50,000: 24.78439697492371%

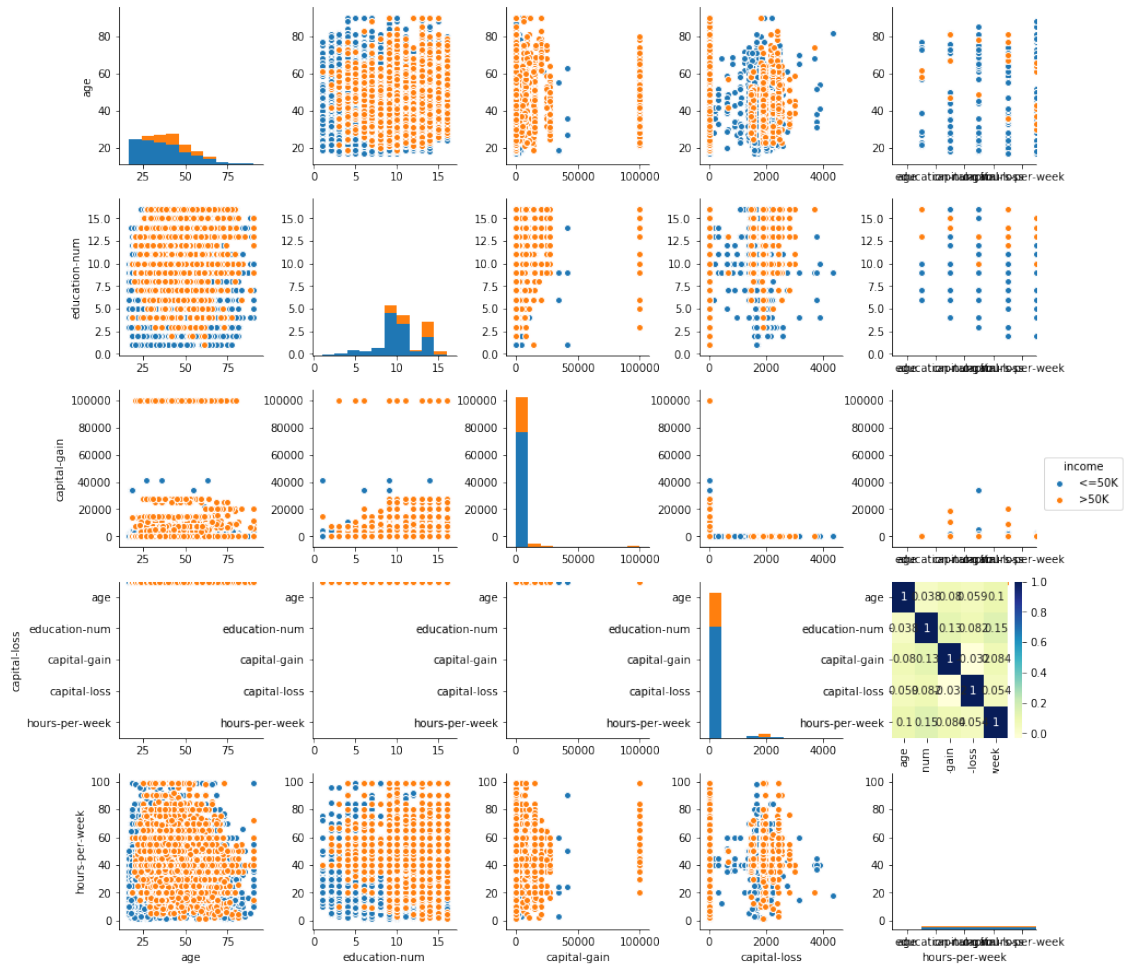
**** Featureset Exploration ****

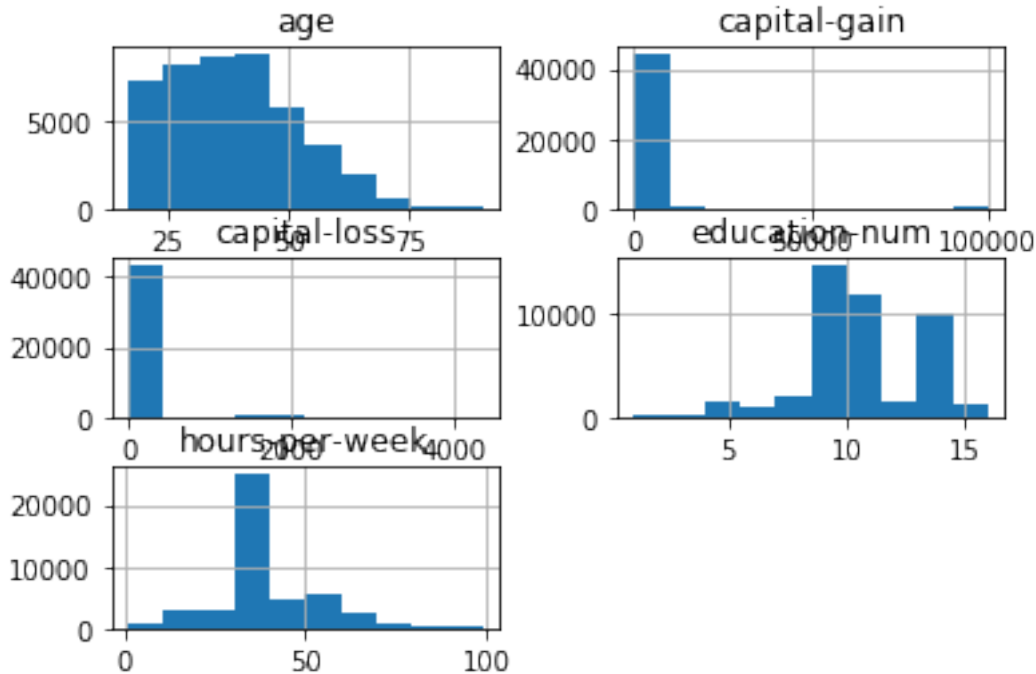
- **age**: continuous.
- **workclass**: Private, Self-emp-not-inc, Self-emp-inc, Federal-gov, Local-gov, State-gov, Without-pay, Never-worked.
- **education**: Bachelors, Some-college, 11th, HS-grad, Prof-school, Assoc-acdm, Assoc-voc, 9th, 7th-8th, 12th, Masters, 1st-4th, 10th, Doctorate, 5th-6th, Preschool.
- **education-num**: continuous.
- **marital-status**: Married-civ-spouse, Divorced, Never-married, Separated, Widowed, Married-spouse-absent, Married-AF-spouse.
- **occupation**: Tech-support, Craft-repair, Other-service, Sales, Exec-managerial, Prof-specialty, Handlers-cleaners, Machine-op-inspct, Adm-clerical, Farming-fishing, Transport-moving, Priv-house-serv, Protective-serv, Armed-Forces.
- **relationship**: Wife, Own-child, Husband, Not-in-family, Other-relative, Unmarried.
- **race**: Black, White, Asian-Pac-Islander, Amer-Indian-Eskimo, Other.
- **sex**: Female, Male.
- **capital-gain**: continuous.
- **capital-loss**: continuous.
- **hours-per-week**: continuous.
- **native-country**: United-States, Cambodia, England, Puerto-Rico, Canada, Germany, Outlying-US(Guam-USVI-etc), India, Japan, Greece, South, China, Cuba, Iran, Honduras, Philippines, Italy, Poland, Jamaica, Vietnam, Mexico, Portugal, Ireland, France, Dominican-Republic, Laos, Ecuador, Taiwan, Haiti, Columbia, Hungary, Guatemala, Nicaragua, Scotland, Thailand, Yugoslavia, El-Salvador, Trinidad&Tobago, Peru, Hong, Holand-Netherlands.

```

In [5]: import seaborn as sns
        sns.pairplot(data, hue="income");
        sns.heatmap(data.corr(), annot=True, cmap="YlGnBu");
        data.hist();

```





0.5 Preparing the Data

Before data can be used as input for machine learning algorithms, it often must be cleaned, formatted, and restructured — this is typically known as **preprocessing**. Fortunately, for this dataset, there are no invalid or missing entries we must deal with, however, there are some qualities about certain features that must be adjusted. This preprocessing can help tremendously with the outcome and predictive power of nearly all learning algorithms.

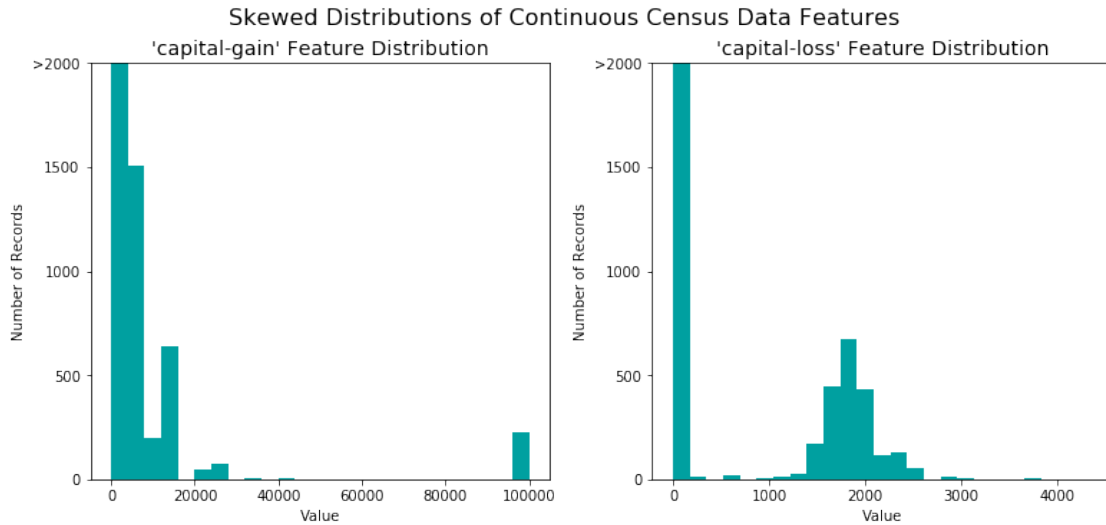
0.5.1 Transforming Skewed Continuous Features

A dataset may sometimes contain at least one feature whose values tend to lie near a single number, but will also have a non-trivial number of vastly larger or smaller values than that single number. Algorithms can be sensitive to such distributions of values and can underperform if the range is not properly normalized. With the census dataset two features fit this description: 'capital-gain' and 'capital-loss'.

Run the code cell below to plot a histogram of these two features. Note the range of the values present and how they are distributed.

```
In [6]: # Split the data into features and target label
income_raw = data['income']
features_raw = data.drop('income', axis = 1)

# Visualize skewed continuous features of original data
vs.distribution(data)
```

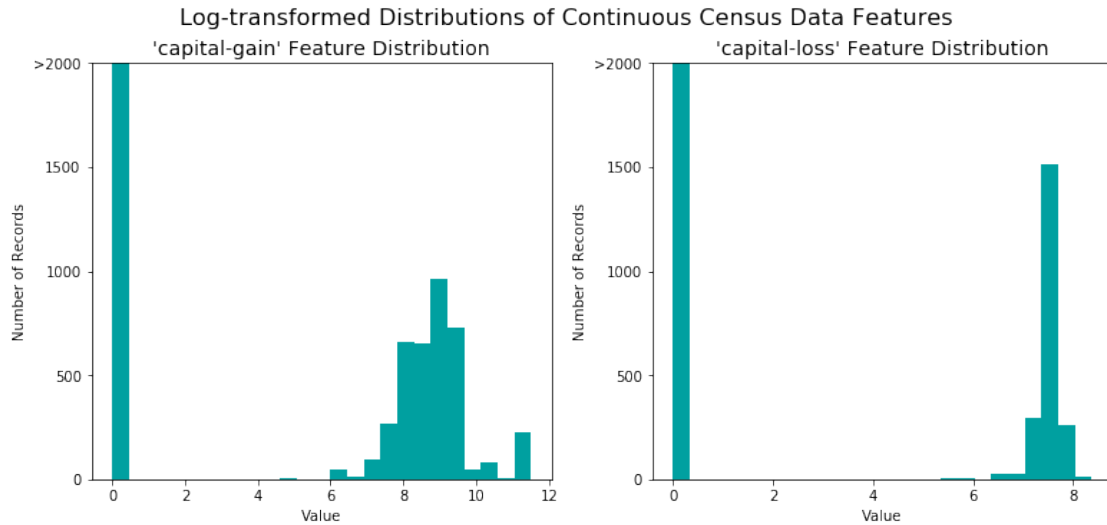


For highly-skewed feature distributions such as 'capital-gain' and 'capital-loss', it is common practice to apply a logarithmic transformation on the data so that the very large and very small values do not negatively affect the performance of a learning algorithm. Using a logarithmic transformation significantly reduces the range of values caused by outliers. Care must be taken when applying this transformation however: The logarithm of 0 is undefined, so we must translate the values by a small amount above 0 to apply the the logarithm successfully.

Run the code cell below to perform a transformation on the data and visualize the results. Again, note the range of values and how they are distributed.

```
In [7]: # Log-transform the skewed features
skewed = ['capital-gain', 'capital-loss']
features_log_transformed = pd.DataFrame(data = features_raw)
features_log_transformed[skewed] = features_raw[skewed].apply(lambda x: np.log(x + 1))

# Visualize the new log distributions
vs.distribution(features_log_transformed, transformed = True)
```



0.5.2 Normalizing Numerical Features

In addition to performing transformations on features that are highly skewed, it is often good practice to perform some type of scaling on numerical features. Applying a scaling to the data does not change the shape of each feature's distribution (such as 'capital-gain' or 'capital-loss' above); however, normalization ensures that each feature is treated equally when applying supervised learners. Note that once scaling is applied, observing the data in its raw form will no longer have the same original meaning, as exemplified below.

Run the code cell below to normalize each numerical feature. We will use `sklearn.preprocessing.MinMaxScaler` for this.

```
In [8]: # Import sklearn.preprocessing.StandardScaler
        from sklearn.preprocessing import MinMaxScaler

        # Initialize a scaler, then apply it to the features
        scaler = MinMaxScaler() # default=(0, 1)
        numerical = ['age', 'education-num', 'capital-gain', 'capital-loss', 'hours-per-week']

        features_log_minmax_transform = pd.DataFrame(data = features_log_transformed)
        features_log_minmax_transform[numerical] = scaler.fit_transform(features_log_transformed[numerical])

        # Show an example of a record with scaling applied
        display(features_log_minmax_transform.head(n = 5))
```

	age	workclass	education_level	education-num \
0	0.301370	State-gov	Bachelors	0.800000
1	0.452055	Self-emp-not-inc	Bachelors	0.800000
2	0.287671	Private	HS-grad	0.533333
3	0.493151	Private	11th	0.400000
4	0.150685	Private	Bachelors	0.800000

	marital-status	occupation	relationship	race	sex \
0	Never-married	Adm-clerical	Not-in-family	White	Male
1	Married-civ-spouse	Exec-managerial	Husband	White	Male
2	Divorced	Handlers-cleaners	Not-in-family	White	Male
3	Married-civ-spouse	Handlers-cleaners	Husband	Black	Male
4	Married-civ-spouse	Prof-specialty	Wife	Black	Female

	capital-gain	capital-loss	hours-per-week	native-country
0	0.667492	0.0	0.397959	United-States
1	0.000000	0.0	0.122449	United-States
2	0.000000	0.0	0.397959	United-States
3	0.000000	0.0	0.397959	United-States
4	0.000000	0.0	0.397959	Cuba

0.5.3 Implementation: Data Preprocessing

From the table in **Exploring the Data** above, we can see there are several features for each record that are non-numeric. Typically, learning algorithms expect input to be numeric, which requires that non-numeric features (called *categorical variables*) be converted. One popular way to convert categorical variables is by using the **one-hot encoding** scheme. One-hot encoding creates a "dummy" variable for each possible category of each non-numeric feature. For example, assume someFeature has three possible entries: A, B, or C. We then encode this feature into someFeature_A, someFeature_B and someFeature_C.

```
| someFeature | | someFeature_A | someFeature_B | someFeature_C |
:- | :- | | :- | :- | :- |
0 | B | | 0 | 1 | 0 |
1 | C | ----> one-hot encode ----> | 0 | 0 | 1 |
2 | A | | 1 | 0 | 0 |
```

Additionally, as with the non-numeric features, we need to convert the non-numeric target label, 'income' to numerical values for the learning algorithm to work. Since there are only two possible categories for this label (" $\leq 50K$ " and " $> 50K$ "), we can avoid using one-hot encoding and simply encode these two categories as 0 and 1, respectively. In code cell below, you will need to implement the following: - Use `pandas.get_dummies()` to perform one-hot encoding on the 'features_log_minmax_transform' data. - Convert the target label 'income_raw' to numerical entries. - Set records with " $\leq 50K$ " to 0 and records with " $> 50K$ " to 1.

```
In [9]: # TODO: One-hot encode the 'features_log_minmax_transform' data using pandas.get_dummies
features_final = pd.get_dummies(features_log_minmax_transform)

# TODO: Encode the 'income_raw' data to numerical values
income = income_raw.replace({"<=50K":0, ">50K":1})

# Print the number of features after one-hot encoding
encoded = list(features_final.columns)
print("{} total features after one-hot encoding.".format(len(encoded)))
```

```
# Uncomment the following line to see the encoded feature names
print(encoded)
```

103 total features after one-hot encoding.

```
['age', 'education-num', 'capital-gain', 'capital-loss', 'hours-per-week', 'workclass_ Federal-g
```

0.5.4 Shuffle and Split Data

Now all *categorical variables* have been converted into numerical features, and all numerical features have been normalized. As always, we will now split the data (both features and their labels) into training and test sets. 80% of the data will be used for training and 20% for testing.

Run the code cell below to perform this split.

```
In [10]: # Import train_test_split
         from sklearn.cross_validation import train_test_split

         # Split the 'features' and 'income' data into training and testing sets
         X_train, X_test, y_train, y_test = train_test_split(features_final,
                                                             income,
                                                             test_size = 0.2,
                                                             random_state = 0)

         # Show the results of the split
         print("Training set has {} samples.".format(X_train.shape[0]))
         print("Testing set has {} samples.".format(X_test.shape[0]))
```

Training set has 36177 samples.

Testing set has 9045 samples.

```
/opt/conda/lib/python3.6/site-packages/sklearn/cross_validation.py:41: DeprecationWarning: This
    "This module will be removed in 0.20.", DeprecationWarning)
```

0.6 Evaluating Model Performance

In this section, we will investigate four different algorithms, and determine which is best at modeling the data. Three of these algorithms will be supervised learners of your choice, and the fourth algorithm is known as a *naive predictor*.

0.6.1 Metrics and the Naive Predictor

CharityML, equipped with their research, knows individuals that make more than \$50,000 are most likely to donate to their charity. Because of this, *CharityML* is particularly interested in predicting who makes more than \$50,000 accurately. It would seem that using **accuracy** as a metric for evaluating a particular model's performance would be appropriate. Additionally, identifying someone

that *does not* make more than \$50,000 as someone who does would be detrimental to *CharityML*, since they are looking to find individuals willing to donate. Therefore, a model's ability to precisely predict those that make more than \$50,000 is *more important* than the model's ability to **recall** those individuals. We can use **F-beta score** as a metric that considers both precision and recall:

$$F_{\beta} = (1 + \beta^2) \cdot \frac{\text{precision} \cdot \text{recall}}{(\beta^2 \cdot \text{precision}) + \text{recall}}$$

In particular, when $\beta = 0.5$, more emphasis is placed on precision. This is called the **F_{0.5} score** (or F-score for simplicity).

Looking at the distribution of classes (those who make at most \$50,000, and those who make more), it's clear most individuals do not make more than \$50,000. This can greatly affect **accuracy**, since we could simply say "*this person does not make more than \$50,000*" and generally be right, without ever looking at the data! Making such a statement would be called **naïve**, since we have not considered any information to substantiate the claim. It is always important to consider the *naïve prediction* for your data, to help establish a benchmark for whether a model is performing well. That been said, using that prediction would be pointless: If we predicted all people made less than \$50,000, *CharityML* would identify no one as donors.

Note: Recap of accuracy, precision, recall ** Accuracy ** measures how often the classifier makes the correct prediction. It's the ratio of the number of correct predictions to the total number of predictions (the number of test data points).

** Precision ** tells us what proportion of messages we classified as spam, actually were spam. It is a ratio of true positives(words classified as spam, and which are actually spam) to all positives(all words classified as spam, irrespective of whether that was the correct classificatio), in other words it is the ratio of

[True Positives/(True Positives + False Positives)]

** Recall(sensitivity)** tells us what proportion of messages that actually were spam were classified by us as spam. It is a ratio of true positives(words classified as spam, and which are actually spam) to all the words that were actually spam, in other words it is the ratio of

[True Positives/(True Positives + False Negatives)]

For classification problems that are skewed in their classification distributions like in our case, for example if we had a 100 text messages and only 2 were spam and the rest 98 weren't, accuracy by itself is not a very good metric. We could classify 90 messages as not spam(including the 2 that were spam but we classify them as not spam, hence they would be false negatives) and 10 as spam(all 10 false positives) and still get a reasonably good accuracy score. For such cases, precision and recall come in very handy. These two metrics can be combined to get the F1 score, which is weighted average(harmonic mean) of the precision and recall scores. This score can range from 0 to 1, with 1 being the best possible F1 score(we take the harmonic mean as we are dealing with ratios).

0.6.2 Question 1 - Naïve Predictor Performace

- If we chose a model that always predicted an individual made more than \$50,000, what would that model's accuracy and F-score be on this dataset? You must use the code cell below and assign your results to 'accuracy' and 'fscore' to be used later.

** Please note ** that the the purpose of generating a naïve predictor is simply to show what a base model without any intelligence would look like. In the real world, ideally your base model

would be either the results of a previous model or could be based on a research paper upon which you are looking to improve. When there is no benchmark model set, getting a result better than random choice is a place you could start from.

**** HINT: ****

- When we have a model that always predicts '1' (i.e. the individual makes more than 50k) then our model will have no True Negatives(TN) or False Negatives(FN) as we are not making any negative('0' value) predictions. Therefore our Accuracy in this case becomes the same as our Precision(True Positives/(True Positives + False Positives)) as every prediction that we have made with value '1' that should have '0' becomes a False Positive; therefore our denominator in this case is the total number of records we have in total.
- Our Recall score(True Positives/(True Positives + False Negatives)) in this setting becomes 1 as we have no False Negatives.

```
In [11]: TP = np.sum(income)
print("TP: ",TP)
FP = income.count() - TP
print("FP", FP)
TN = 0
FN = 0
```

```
TP: 11208
FP 34014
```

```
In [29]: '''
TP = np.sum(income) # Counting the ones as this is the naive case. Note that 'income' is
encoded to numerical values done in the data preprocessing step.
FP = income.count() - TP # Specific to the naive case

TN = 0 # No predicted negatives in the naive case
FN = 0 # No predicted negatives in the naive case
'''

# TODO: Calculate accuracy, precision and recall
accuracy = TP / (income.count())
recall = TP / (TP + FN)
precision = TP / (TP + FP)

# TODO: Calculate F-score using the formula above for beta = 0.5 and correct values for
fscore = 2 * (1+0.5**2) * precision * recall / (0.5**2*precision + recall)

# Print the results
print("Naive Predictor: [Accuracy score: {:.4f}, F-score: {:.4f}]"
      .format(accuracy, fscore))
```

```
Naive Predictor: [Accuracy score: 0.2478, F-score: 0.5835]
```

0.6.3 Supervised Learning Models

The following are some of the supervised learning models that are currently available in [scikit-learn](#) that you may choose from: - Gaussian Naive Bayes (GaussianNB) - Decision Trees

- Ensemble Methods (Bagging, AdaBoost, Random Forest, Gradient Boosting) - K-Nearest Neighbors (KNeighbors) - Stochastic Gradient Descent Classifier (SGDC) - Support Vector Machines (SVM) - Logistic Regression

0.6.4 Question 2 - Model Application

List three of the supervised learning models above that are appropriate for this problem that you will test on the census data. For each model chosen

- Describe one real-world application in industry where the model can be applied.
- What are the strengths of the model; when does it perform well?
- What are the weaknesses of the model; when does it perform poorly?
- What makes this model a good candidate for the problem, given what you know about the data?

**** HINT: ****

Structure your answer in the same format as above, with 4 parts for each of the three models you pick. Please include references with your answer.

Answer:

Ensemble Methods - Random Forests

- Real-world application: Random forests are used to predict Alzheimer's disease: <https://www.sciencedirect.com/science/article/pii/S2213158214001326>
- Strengths: this model is robust with missing data. It can be easily explained. It has a low number of hyperparameters.
- Weaknesses: it can overfit quickly.
- Good candidate? Given the problem, it is a good candidate because we'll need to explain how it works to CharityML and experience shows it performs well on this kind of tasks.

AdaBoosting

- Real-world application: AdaBoost can be used to predict NFL performances: <https://towardsdatascience.com/understanding-adaboost-2f94f22d5bfe>
- Strengths: It's a very computationally efficient model and easy to setup.
- Weaknesses: It's complex, can overfit and it can be hard to explain how it works.
- Good candidate? It's ease to implement makes it a good candidate for CharityML. Moreover, the data is pretty clean and we could iterate over the learners the model needs.

K-Nearest Neighbors

- Real-world application: KNN can be used to detect intrusion in Network Data: <https://research.ijcaonline.org/volume97/number7/pxc3897306.pdf>
- Strengths: It has very few hyperparameters.
- Weaknesses: It can have a high computation cost in high dimensions. And it's sensitive to noise and outliers.
- Good candidate? Given the data, we don't have too many dimensions and the data is pretty clean without noise and outlier. This is said, KNN is a good candidate.

Sources: * <http://phirilytics.blogspot.com/2017/07/ensemble-model-strengthsand-some.html> * <https://towardsdatascience.com/ensemble-methods-bagging-boosting-and-stacking-c9214a10a205> * <https://elitedatascience.com/machine-learning-algorithms>

0.6.5 Implementation - Creating a Training and Predicting Pipeline

To properly evaluate the performance of each model you've chosen, it's important that you create a training and predicting pipeline that allows you to quickly and effectively train models using various sizes of training data and perform predictions on the testing data. Your implementation here will be used in the following section. In the code block below, you will need to implement the following: - Import `fbeta_score` and `accuracy_score` from `sklearn.metrics`. - Fit the learner to the sampled training data and record the training time. - Perform predictions on the test data `X_test`, and also on the first 300 training points `X_train[:300]`. - Record the total prediction time. - Calculate the accuracy score for both the training subset and testing set. - Calculate the F-score for both the training subset and testing set. - Make sure that you set the `beta` parameter!

```
In [13]: # TODO: Import two metrics from sklearn - fbeta_score and accuracy_score
         from sklearn.metrics import fbeta_score, accuracy_score

         def train_predict(learner, sample_size, X_train, y_train, X_test, y_test):
             """
             inputs:
                 - learner: the learning algorithm to be trained and predicted on
                 - sample_size: the size of samples (number) to be drawn from training set
                 - X_train: features training set
                 - y_train: income training set
                 - X_test: features testing set
                 - y_test: income testing set
             """

             results = {}

             # TODO: Fit the learner to the training data using slicing with 'sample_size' using
             start = time() # Get start time
             learner.fit(X_train[:sample_size], y_train[:sample_size])
             end = time() # Get end time

             # TODO: Calculate the training time
             results['train_time'] = end - start

             # TODO: Get the predictions on the test set(X_test),
             #         then get predictions on the first 300 training samples(X_train) using .predict
             start = time() # Get start time
             predictions_test = learner.predict(X_test)
             predictions_train = learner.predict(X_train[:300])
             end = time() # Get end time

             # TODO: Calculate the total prediction time
```

```

results['pred_time'] = end - start

# TODO: Compute accuracy on the first 300 training samples which is y_train[:300]
results['acc_train'] = accuracy_score(y_train[:300], predictions_train)

# TODO: Compute accuracy on test set using accuracy_score()
results['acc_test'] = accuracy_score(y_test, predictions_test)

# TODO: Compute F-score on the the first 300 training samples using fbeta_score()
results['f_train'] = fbeta_score(y_train[:300], predictions_train, beta=0.5)

# TODO: Compute F-score on the test set which is y_test
results['f_test'] = fbeta_score(y_test, predictions_test, beta=0.5)

# Success
print("{} trained on {} samples.".format(learner.__class__.__name__, sample_size))

# Return the results
return results

```

0.6.6 Implementation: Initial Model Evaluation

In the code cell, you will need to implement the following: - Import the three supervised learning models you've discussed in the previous section. - Initialize the three models and store them in 'clf_A', 'clf_B', and 'clf_C'. - Use a 'random_state' for each model you use, if provided. - **Note:** Use the default settings for each model — you will tune one specific model in a later section. - Calculate the number of records equal to 1%, 10%, and 100% of the training data. - Store those values in 'samples_1', 'samples_10', and 'samples_100' respectively.

Note: Depending on which algorithms you chose, the following implementation may take some time to run!

```

In [14]: # TODO: Import the three supervised learning models from sklearn
from sklearn.ensemble import AdaBoostClassifier, RandomForestClassifier
from sklearn.neighbors import KNeighborsClassifier

# TODO: Initialize the three models
clf_A = AdaBoostClassifier(random_state=42)
clf_B = RandomForestClassifier(random_state=42)
clf_C = KNeighborsClassifier()

# TODO: Calculate the number of samples for 1%, 10%, and 100% of the training data
# HINT: samples_100 is the entire training set i.e. len(y_train)
# HINT: samples_10 is 10% of samples_100 (ensure to set the count of the values to be `
# HINT: samples_1 is 1% of samples_100 (ensure to set the count of the values to be `in
samples_100 = y_train.count()
samples_10 = int(round(y_train.count() * 0.1))
samples_1 = int(round(y_train.count() * 0.01))

```

```

# Collect results on the learners
results = {}
for clf in [clf_A, clf_B, clf_C]:
    clf_name = clf.__class__.__name__
    results[clf_name] = {}
    for i, samples in enumerate([samples_1, samples_10, samples_100]):
        results[clf_name][i] = \
            train_predict(clf, samples, X_train, y_train, X_test, y_test)

# Run metrics visualization for the three supervised learning models chosen
vs.evaluate(results, accuracy, fscore)

```

AdaBoostClassifier trained on 362 samples.

AdaBoostClassifier trained on 3618 samples.

AdaBoostClassifier trained on 36177 samples.

RandomForestClassifier trained on 362 samples.

RandomForestClassifier trained on 3618 samples.

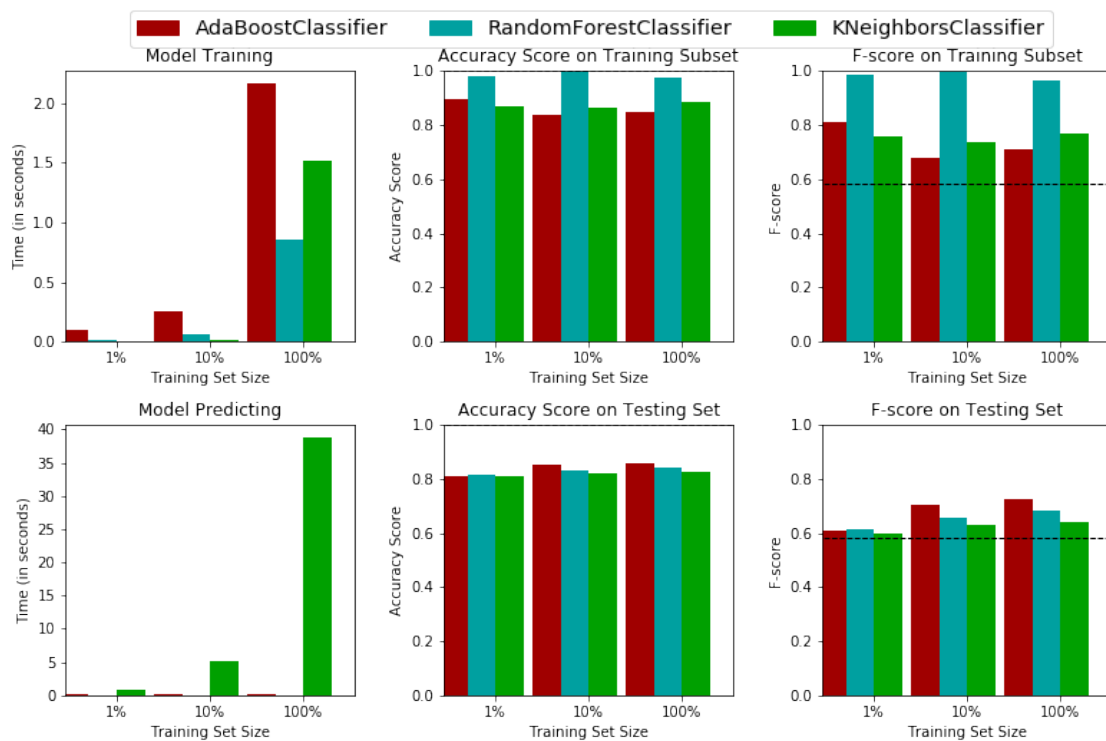
RandomForestClassifier trained on 36177 samples.

KNeighborsClassifier trained on 362 samples.

KNeighborsClassifier trained on 3618 samples.

KNeighborsClassifier trained on 36177 samples.

Performance Metrics for Three Supervised Learning Models



0.7 Improving Results

In this final section, you will choose from the three supervised learning models the *best* model to use on the student data. You will then perform a grid search optimization for the model over the entire training set (`X_train` and `y_train`) by tuning at least one parameter to improve upon the untuned model's F-score.

0.7.1 Question 3 - Choosing the Best Model

- Based on the evaluation you performed earlier, in one to two paragraphs, explain to *CharityML* which of the three models you believe to be most appropriate for the task of identifying individuals that make more than \$50,000.

**** HINT: **** Look at the graph at the bottom left from the cell above (the visualization created by `vs.evaluate(results, accuracy, fscore)`) and check the F score for the testing set when 100% of the training set is used. Which model has the highest score? Your answer should include discussion of the: * metrics - F score on the testing when 100% of the training data is used, * prediction/training time * the algorithm's suitability for the data.

Answer:

I believe the Random Forest Classifier is the most appropriate for this case. First, training and prediction times are very low. We can barely see the prediction time in the graph above.

Then, this model clearly performs overfitting. Accuracy and F score are almost equal to 1 on training set. Nevertheless, with testing set, its score is equivalent to the others (between 65% and 70%). I can expect that hypertuning this model will reduce overfitting and help its generalization to perform better on new data.

Finally, the explainability of the model is one important point for *CharityML*. Random Forest will be clearly the simpler model to explain.

0.7.2 Question 4 - Describing the Model in Layman's Terms

- In one to two paragraphs, explain to *CharityML*, in layman's terms, how the final model chosen is supposed to work. Be sure that you are describing the major qualities of the model, such as how the model is trained and how the model makes a prediction. Avoid using advanced mathematical jargon, such as describing equations.

**** HINT: ****

When explaining your model, if using external resources please include all citations.

Answer:

The model is a Random Forest Classifier. The idea behind is very basic as it looks like the way we, as humans, make decisions.

A forest is an ensemble of trees. Each tree splits the data under specific circumstances: is the age greater than 50 ? Is capital gain lower than 20k ? ... So each tree has its own algorithm and parameters.

To predict if a person will donate or not, each tree will make its own prediction. And to decide which decision we keep, it's like an election : we keep the most common prediction.

By default, a forest has 100 trees. But we will tune this parameter to find the best number of trees of *CharityML*.

0.7.3 Implementation: Model Tuning

Fine tune the chosen model. Use grid search (GridSearchCV) with at least one important parameter tuned with at least 3 different values. You will need to use the entire training set for this. In the code cell below, you will need to implement the following: - Import `sklearn.grid_search.GridSearchCV` and `sklearn.metrics.make_scorer`. - Initialize the classifier you've chosen and store it in `clf`. - Set a `random_state` if one is available to the same state you set before. - Create a dictionary of parameters you wish to tune for the chosen model. - Example: `parameters = {'parameter' : [list of values]}`. - **Note:** Avoid tuning the `max_features` parameter of your learner if that parameter is available! - Use `make_scorer` to create an `fbeta_score` scoring object (with $\beta = 0.5$). - Perform grid search on the classifier `clf` using the '`scorer`', and store it in `grid_obj`. - Fit the grid search object to the training data (`X_train`, `y_train`), and store it in `grid_fit`.

Note: Depending on the algorithm chosen and the parameter list, the following implementation may take some time to run!

```
In [26]: # TODO: Import 'GridSearchCV', 'make_scorer', and any other necessary libraries
from sklearn.grid_search import GridSearchCV
from sklearn.metrics import make_scorer, fbeta_score
from sklearn.ensemble import RandomForestClassifier

# TODO: Initialize the classifier
clf = RandomForestClassifier(random_state=42)

# TODO: Create the parameters list you wish to tune, using a dictionary if needed.
# HINT: parameters = {'parameter_1': [value1, value2], 'parameter_2': [value1, value2]}
parameters = {
    'n_estimators': list(range(100,250,25)),
    #'n_estimators': [20, 50, 100],
    #'max_depth': [5,10, 20 , 50],
    'max_depth': list(range(10,25,2)),
    #'min_samples_split': [2,5,10],
    #'min_samples_leaf': [1,2,5]
}

# TODO: Make an fbeta_score scoring object using make_scorer()
scorer = make_scorer(fbeta_score, beta=0.5)

# TODO: Perform grid search on the classifier using 'scorer' as the scoring method using
grid_obj = GridSearchCV(clf, param_grid=parameters, scoring=scorer)

# TODO: Fit the grid search object to the training data and find the optimal parameters
grid_fit = grid_obj.fit(X_train, y_train)

# Get the estimator
best_clf = grid_fit.best_estimator_

# Make predictions using the unoptimized and model
```

```

predictions = (clf.fit(X_train, y_train)).predict(X_test)
best_predictions = best_clf.predict(X_test)

# Report the before-and-afterscores
print("Unoptimized model\n-----")
print("Accuracy score on testing data: {:.4f}".format(accuracy_score(y_test, predictions)))
print("F-score on testing data: {:.4f}".format(fbeta_score(y_test, predictions, beta = 0.5)))
print("\nOptimized Model\n-----")
print("Final accuracy score on the testing data: {:.4f}".format(accuracy_score(y_test, best_predictions)))
print("Final F-score on the testing data: {:.4f}".format(fbeta_score(y_test, best_predictions, beta = 0.5)))
print(best_clf)

```

Unoptimized model

Accuracy score on testing data: 0.8431

F-score on testing data: 0.6842

Optimized Model

Final accuracy score on the testing data: 0.8586

Final F-score on the testing data: 0.7313

```

RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                        max_depth=16, 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=1,
                        oob_score=False, random_state=42, verbose=0, warm_start=False)

```

0.7.4 Question 5 - Final Model Evaluation

- What is your optimized model's accuracy and F-score on the testing data?
- Are these scores better or worse than the unoptimized model?
- How do the results from your optimized model compare to the naive predictor benchmarks you found earlier in **Question 1**?

Note: Fill in the table below with your results, and then provide discussion in the **Answer** box.

Metric	Unoptimized Model	Optimized Model
Accuracy Score	0.84	0.86
F-score	0.68	0.73

Results: Answer:

The scores from the optimized model are better than the unoptimized one.

In comparison with the naive predictor in Question 1, (accuracy: 0.2478, F-score: 0.2917), we have much higher scores: we have + 0.61 in accuracy and +0.44 in the F-score. This is a huge

increase.

0.8 Feature Importance

An important task when performing supervised learning on a dataset like the census data we study here is determining which features provide the most predictive power. By focusing on the relationship between only a few crucial features and the target label we simplify our understanding of the phenomenon, which is most always a useful thing to do. In the case of this project, that means we wish to identify a small number of features that most strongly predict whether an individual makes at most or more than \$50,000.

Choose a scikit-learn classifier (e.g., adaboost, random forests) that has a `feature_importance_` attribute, which is a function that ranks the importance of features according to the chosen classifier. In the next python cell fit this classifier to training set and use this attribute to determine the top 5 most important features for the census dataset.

0.8.1 Question 6 - Feature Relevance Observation

When **Exploring the Data**, it was shown there are thirteen available features for each individual on record in the census data. Of these thirteen records, which five features do you believe to be most important for prediction, and in what order would you rank them and why?

Answer:

I think these five features will be the most important : * age: you basically have a better job and so a higher income when you're older and get more experience * education: the best education you have, the higher is your income when you begin and higher it will be along your career-life * sex: unfortunately, statistics show there is a 30% gap in salary between men and women * occupation: your income is tightly linked to what kind of occupation you have * marital-status: I think that when you're married, you're on a more stable situation

0.8.2 Implementation - Extracting Feature Importance

Choose a scikit-learn supervised learning algorithm that has a `feature_importance_` attribute available for it. This attribute is a function that ranks the importance of each feature when making predictions based on the chosen algorithm.

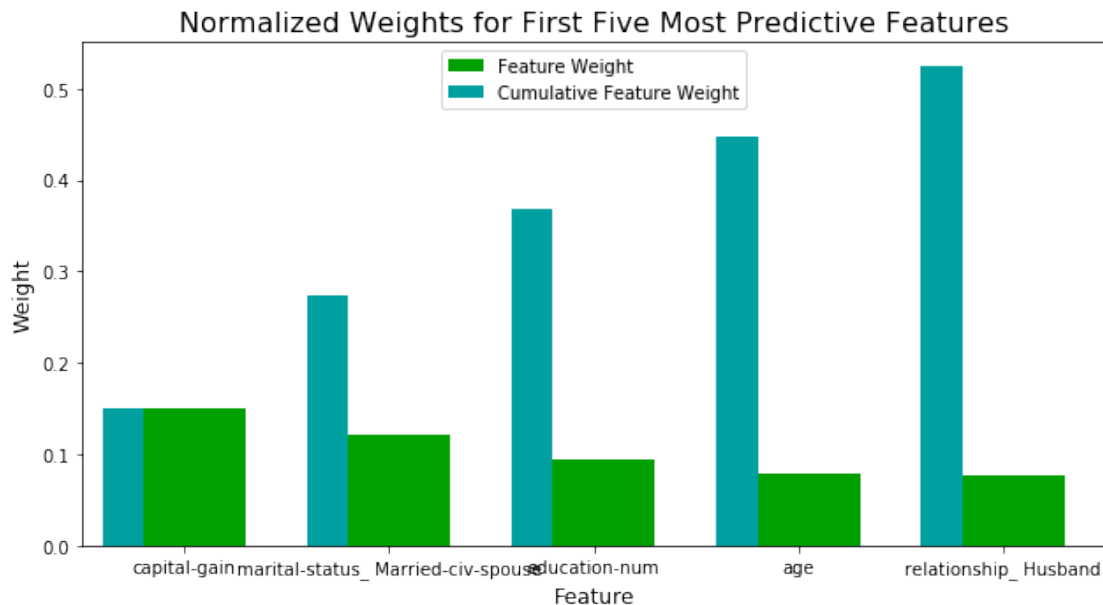
In the code cell below, you will need to implement the following: - Import a supervised learning model from sklearn if it is different from the three used earlier. - Train the supervised model on the entire training set. - Extract the feature importances using `'.feature_importances_'`.

```
In [27]: # TODO: Import a supervised learning model that has 'feature_importances_'
         from sklearn.ensemble import RandomForestClassifier

         # TODO: Train the supervised model on the training set using .fit(X_train, y_train)
         model = RandomForestClassifier(random_state=42, n_estimators=200, max_depth=16)
         model.fit(X_train, y_train)

         # TODO: Extract the feature importances using .feature_importances_
         importances = model.feature_importances_
```

```
# Plot
vs.feature_plot(importances, X_train, y_train)
```



0.8.3 Question 7 - Extracting Feature Importance

Observe the visualization created above which displays the five most relevant features for predicting if an individual makes at most or above \$50,000.

* How do these five features compare to the five features you discussed in **Question 6**? * If you were close to the same answer, how does this visualization confirm your thoughts? * If you were not close, why do you think these features are more relevant?

Answer:

The main ideas I believed are shown in the graph: * age is the number 4 most important feature * "education idea", is present but through the feature education-num, not with the feature I thought * sex, but with the feature relationship_Husband which tells us two things: Man and Married * occupation is not present * marital-status: being married is the 2nd most important feature

I missed the capital-gain. But it makes totally sense. I don't how I could miss it.

0.8.4 Feature Selection

How does a model perform if we only use a subset of all the available features in the data? With less features required to train, the expectation is that training and prediction time is much lower — at the cost of performance metrics. From the visualization above, we see that the top five most important features contribute more than half of the importance of **all** features present in the data. This hints that we can attempt to *reduce the feature space* and simplify the information required for the model to learn. The code cell below will use the same optimized model you found earlier, and train it on the same training set *with only the top five important features*.

```

In [28]: # Import functionality for cloning a model
         from sklearn.base import clone

         # Reduce the feature space
         X_train_reduced = X_train[X_train.columns.values[(np.argsort(importances)[:,-1])[:5]]]
         X_test_reduced = X_test[X_test.columns.values[(np.argsort(importances)[:,-1])[:5]]]

         # Train on the "best" model found from grid search earlier
         clf = (clone(best_clf)).fit(X_train_reduced, y_train)

         # Make new predictions
         reduced_predictions = clf.predict(X_test_reduced)

         # Report scores from the final model using both versions of data
         print("Final Model trained on full data\n-----")
         print("Accuracy on testing data: {:.4f}".format(accuracy_score(y_test, best_predictions)))
         print("F-score on testing data: {:.4f}".format(fbeta_score(y_test, best_predictions, beta=2)))
         print("\nFinal Model trained on reduced data\n-----")
         print("Accuracy on testing data: {:.4f}".format(accuracy_score(y_test, reduced_predictions)))
         print("F-score on testing data: {:.4f}".format(fbeta_score(y_test, reduced_predictions, beta=2)))

```

Final Model trained on full data

Accuracy on testing data: 0.8586

F-score on testing data: 0.7313

Final Model trained on reduced data

Accuracy on testing data: 0.8425

F-score on testing data: 0.6865

0.8.5 Question 8 - Effects of Feature Selection

- How does the final model's F-score and accuracy score on the reduced data using only five features compare to those same scores when all features are used?
- If training time was a factor, would you consider using the reduced data as your training set?

Answer:

The accuracy is almost the same on reduced data (-0.01) but the F-score loses 0.05 points, corresponding to the score of the unoptimized model.

If training time was a factor, I will definitely reduce the training set. But I'll also fine-tune the model specifically for this new set of features.

Note: Once you have completed all of the code implementations and successfully answered each question above, you may finalize your work by exporting the iPython Notebook as an HTML document. You can do this by using the menu above and navigating to

File -> Download as -> HTML (.html). Include the finished document along with this notebook as your submission.

0.9 Before You Submit

You will also need run the following in order to convert the Jupyter notebook into HTML, so that your submission will include both files.

```
In [30]: !!jupyter nbconvert *.ipynb
```

```
Out[30]: ['[NbConvertApp] Converting notebook finding_donors.ipynb to html',  
          '[NbConvertApp] Writing 795542 bytes to finding_donors.html']
```

```
In [ ]:
```