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0.30 -	Feature importances
0.15 -	lead_ti lead_ti lead_ti n al_date_mo day_of_mo parking_spa arket_segm reekend_nig ival_date_y ad customer_t ecial_reque oution_chan deposit_ti is_cance is_cance is_cance is_cancellati epeated_gu in_waiting_ bak
Method The algoprocedur	earn that the feature "adr" is significantly more clearly to the prediction in comparison to the other features, for instance, times bigger (2.877) than the second most important feature.  d 2 - Recursive feature elimination (RFE) with Random Forest Classification rithm assign weights to each of features. Whose absolute weights are the smallest are pruned from the current set feature is recursively repeated on the pruned set until the desired number of features.  = RandomForestClassifier (n_estimators=20)
rfe = Free rfe = r	RFE (estimator=clf_rf_, n_features_to_select=11, step=1)  rfe.fit(X_train, y_train)  Chosen best 11 feature by RFE:',X_train.columns[rfe.support_])  best 11 feature by RFE: Index(['lead_time', 'arrival_date_year', 'arrival_date_month', 'arrival_date_day_of_month', 'stays_in_weekend_nights',  'stays_in_week_nights', 'meal', 'market_segment',  'distribution_channel', 'adr', 'required_car_parking_spaces'],  type='object')  mod brought me the same 11 most important features as the first method, therefore I will use only those 11 features and thm once again.
<pre>'reserv n_waiti  X = nev y = nev  X_train  model =</pre>	<pre>cel_supervised = hotel_supervised.drop(columns = ['deposit_type', 'children', 'booking_claration_status', 'is_canceled', 'previous_bookings_not_canceled', 'previous_cancellations.org_list','is_repeated_guest','babies','distribution_channel', 'deposit_type'])  or_hotel_supervised.drop(columns = ['hotel']).copy()  or_hotel_supervised["hotel"].copy()  or_hotel_supervised["hote</pre>
0.72889	ed the accuracy by 0.008 points - unfortunately not so significantly.  atmap(confusion_matrix(y_test, y_pred), annot=True, fmt="d")  atlib.axessubplots.AxesSubplot at 0x20edbdc0d30>  -20000 -17500 -15000
<pre>print(' print(' print('</pre>	12500 -12500 -10000 -7500 -5000 -2500 fn, tp = confusion_matrix(y_test, y_pred).ravel() TP:", tp) FN:", fn) FP:", fp) TN:", tn)
hotel_t	confusion matrix, we can see that the algorithm predicted better the city hotel. However, the prediction of the resulting has before.  Expe = hotel_supervised.groupby('hotel')[['lead_time', 'arrival_date_year', 'arrival_date' 'arrival_date_day_of_month', 'stays_in_weekend_nights', 'stays_in_week_nights', 'adults', 'meal', 'adr', 'required_car_parking_spaces', 'total_of_special_requests']].mean()
In concluis the "ace Except the weekend hotel (on Finally, a the number of the second	ead_time arrival_date_year arrival_date_month arrival_date_day_of_month stays_in_weekend_nights stays_in_week_nights arrival_date_year arrival_date_day_of_month stays_in_weekend_nights arrival_date_year arrival_date_day_of_month stays_in_weekend_nights arrival_date_year arrival_date_month arrival_date_day_of_month stays_in_weekend_nights arrival_date_year.  15.787094 0.795187 2.182  1.189815 3.128  2.182  2.182  2.182  3.198  3.128  2.182  3.128  2.182  3.128  2.182  3.128  2.182  3.128  2.182  3.128  3.12
Can we so To answer partition of cluster.  from slater sla	move one to the next the unsupervised question:  split our data to N clusters such that every cluster is including similar booking features?  er that, I will use KMeans algorithm. This algorithm is a method of vector quantization, originally from signal processing, nobservations into k clusters in which each observation belongs to the cluster with the nearest mean, serving as a protein observation into k clusters in which each observation belongs to the cluster with the nearest mean, serving as a protein elearn.cluster import KMeans  sclearn.metrics import silhouette_score  need to find out how many clusters do I need to divide into each data I used. In order to deal with this dilemma, I will builted 'number_of_clusters()'. This function will return us two graphs. The first graph is called "Sum of square error ", in this what is the sum of square error as long as the number of clusters increases. Now comes the question of what is the number are "suspicious" to be optimal. For answering this question, I will use the "Elbow Method". In this method, when we see that are "suspicious" to be optimal. For answering this question, I will use the "Elbow Method". In this method, when we see that are "suspicious" to be optimal.
will use is value is of got. I need that has the	number clusters, we will take that number as an option about how many clusters should we use in this data set. The sector called "Silhouette", the silhouette is a measure about how good our algorithm work, his range is between -1 to 1. As lot close to 1 we can say that the clustering was good. In our graph, I will wish to choose the number of clusters by the best do to find the best cluster by a combination of those two graphs, a number of clusters which has a good "Elbow break" a the best silhouette.    Description of the clusters (data_frame):
	<pre>x1 = (range(2, 11)) x2 = (range(2, 11)) y1 = sum_squared y2 = silhouette plt.subplot(2, 1, 1) plt.plot(x1, y1) plt.title(r'Sum of Squred Error \${R_2}\$', fontsize=15) plt.grid() plt.ylabel(r'\${R_2}\$') plt.subplot(2, 1, 2) plt.plot(x2, y2) plt.title('Silhouette', fontsize=15) plt.xlabel('No. of Clusters') plt.ylabel('Silhouette') plt.grid() plt.show()</pre>
number_ 10000 8000	2 3 4 Silhosette 8 9 10
the higher data into will choose Below is	n see, in the first plot, the sharper elbow break is when I divide the data into 3 clusters. However, when we look at the set of the silhouette is when I divide the data into 4 clusters. In the second plot, we see that although the best silhouette is to divide the silhouette of 3 clusters is not much lower than the 4. Moreover, the silhouette score is low in those both one to divide the data into 3 clusters.
• clus  def cre try exc	<pre>return KMeans(n_clusters=k, init='k-means++') sept:     print("Something got wrong - create_kmeans_calssifier")  usters_information(data_frame):</pre>
kmeans kmeans silhous 0.26888 hotel_ucluster	<pre>print(group)</pre>
0 0 1 2 3 4  40055 40056 40057 40058 40059	hotel is_canceled lead_time arrival_date_year arrival_date_month \ 1.0
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	0.5 2.0 0.5 2.0 0.5 2.0 0.5 2.0  x 26 columns]  In see here, the silhouette score that we got is very low. An explanation for this is because the values of the features are at the algorithm had difficulties to handle those samples and to separate them into groups that have the same behavior.