0 Resort Hotel 1 Resort Hotel	0	_ got wrong	g")		hotel_bookings.  al_date_week_number  27	
2 Resort Hotel 3 Resort Hotel 4 Resort Hotel 119385 City Hotel 119386 City Hotel	0	7 13 14  23	2015 2015 2015 2017	July July July August	27 27 27  35	
119387 City Hotel  119388 City Hotel  119389 City Hotel  119390 rows ×  The data is rea	0 0 32 columns	34 109 205	2017 2017 2017	August  August  August	35 35 35	
print("The The number:	number number number of columber of row of columns in	lumns in ho	otel data are" L data are 119	are", hotel_dat , hotel_data.sh 390		
'arr 'arr 'sta 'cou: 'is_ 'pre 'ass 'com 'req 'res	ival_date_monival_date_monival_date_dayys_in_week_nintry', 'markerepeated_guesvious_bookingigned_room_typany', 'days_uired_car_par	nth', 'arri y_of_month' ghts', 'ac et_segment' st', 'previ gs_not_canc ype', 'book _in_waiting cking_space	ival_date_week_ ', 'stays_in_w dults', 'child ', 'distributi ious_cancellat celed', 'reser king_changes', g_list', 'cust	eekend_nights', ren', 'babies', on_channel', ions', ved_room_type',  'deposit_type' omer_type', 'ac _special_reques	'meal', , 'agent',	
hotel_data.  hotel is_canceled lead_time arrival_data arrival_data arrival_data stays_in_we	dtypes  e_year e_month e_week_number e_day_of_mont ekend_nights		object int64 int64 object int64 int64			
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177 hotel_data[ 5	'country'].dr	n_channel']	ates().count() ].drop_duplica es()	tes().count()		
40060 Name: hotel There are only	sort Hotel City Hotel , dtype: obje two types of hote 'hotel'].valu Number of Res	els. ue_counts()		ar", title="Nur	wher of Resort a	nd City Hotels"
70000 60000 50000 40000 30000 20000 10000						
hotel_kinds for name, k	= hotel_data ind in hotel_ The average r	a.groupby(' _kinds:			"are:", kind['	children'].mean
6 The average 15 We can learn fr	number of ch	nildren and at children and of the number	d babies in Re	sort Hotel are:	.09136979048483 0.128681977034 hotels. ering date of the book	44834 0.0139041
<pre>plt.ylabel( plt.title("  Text(0.5, 1</pre>	The Distribut	cribution (	Of The Days Be	The Vacation") fore The Vacati	on')	
0.006 0.004 0.002 0.000	100 200 30	00 400 500 Lead Time	0 600 700 800			
<pre>Let's check whi  countries = countries.p plt.axhline plt.xlabel( plt.ylabel()</pre>	hotel_data.glot.bar() (32,color = "Countries") "Count")	groupby('co	ountry') ['coun	et. try'].value_cou	O days from their arri	ival date.
350 - 300 - 250 - 150 -	.0, 'The numb		els in each co	untry')		
100 - 50 - (MgA, MgA)  We can see that	at "AGO" and "AR	ountries  (ARC, ARE ARC		than the other	iis data set. Moreove	r, The average
each country in  Checking the n  hotel_data. plt.grid(co plt.title('  Text(0.5, 1	umbers of repeat groupby('hotelor='black', The number of	ed guests in e	each type of hotel.  epeated_guest'  e'', linewid guests in each		.nd='bar',color= alpha=0.9)	
The nur 2000 1750 1500 1250 1000 750 250	or repeated	scs in each	, re of hotel			
The city hotels the gap of the r	consists of a bit r number of the hot	els in each ty	pe. we can infer th		r, the gap between th ikely visit again in res	
Checking the n  hotel_data. plt.grid(co plt.title('  Text(0.5, 1	umber of adults in groupby('hote lor='black', The number of	n each type of el')['adult linestyle= f adults ir per of adul	fhotel.  ts'].sum().plo ='', linewid n each type of  tts in each ty	t(kind='bar',co th=1, axis='y', hotel')	olor='royalblue'	
120000 100000 80000 60000 40000	itel –		lotel –			
Now let's see if graph = sns		nce between to	the lead time in city  col='hotel')	nber of hotels consisy hotels to resort ho		
graph.map(p <seaborn.ax ho 20000 - 15000 - 10000 -</seaborn.ax 		ad_time', k	oins=20)			
Both are pretty Checking NaN			200 400 600 lead_time			
arrival_date stays_in_wee stays_in_wee adults children	e_year e_month e_week_number e_day_of_mont ekend_nights		0 0 0 0 0 0 0 0 0			
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Now let's check	(hotel_data.c Heatmap', for s['figure.dpi	ation between  corr(), anr  ntsize = 20  gsize'] =	missing as well.  In the features.  In the features.			
arriva arrival_ arrival_date_da	is_canceled - 1 0. lead_time - 0.29 al_date_year - 0.017 0. date_month - 0.011 0. ay_of_month0.0061 0.0 kend_nights0.0018 0.0	1 0.04 0.13 04 1 -0.53 .13 -0.53 1	0.0022 0.086 0.17 ( -0.00028 0.021 0.031 ( -0.026 0.018 0.019 0  1 -0.016 -0.028 -0.	0.12 -0.038 -0.021 -0.12 0.03 0.055 -0.013 0.01 0.029 0.0054 0.01 -0.031 0015 0.015 -0.00023-0.0062	0.11 -0.057 -0.14 0.054 0.086 -0.074 0.00014 0.17 -0.12 0.029 0.031 -0.056 0.037 -0.022 0.0048 0.019 -0.027 -0.0003 0.011 0.023	-0.063 -0.12 -0.096 7 0.2 -0.014 0.11
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In [43]:	<pre>print("Feature ranking:") for f in range(X_train.shape[1]):     print("%d. feature %d (%f)" % (f + 1, indices[f], importances[indices[f]]))  Feature ranking: 1. feature 20 (0.305477) 2. feature 6 (0.093916) 3. feature 1 (0.088147) 4. feature 3 (0.070110) 5. feature 10 (0.069752)</pre>
	6. feature 4 (0.060717) 7. feature 11 (0.040345) 8. feature 5 (0.038731) 9. feature 21 (0.029089) 11. feature 12 (0.025174) 12. feature 7 (0.023088) 13. feature 19 (0.021813) 14. feature 22 (0.021399) 15. feature 8 (0.014975) 16. feature 16 (0.013429)
In [44]:	17. feature 17 (0.011353) 18. feature 23 (0.009392) 19. feature 0 (0.006285) 20. feature 18 (0.004780) 21. feature 15 (0.004295) 22. feature 14 (0.004279) 23. feature 13 (0.003171) 24. feature 9 (0.001630)  plt.figure(1, figsize=(8, 8)) plt.title("Feature importances")
	<pre>plt.title("Feature importances") plt.bar(range(X_train.shape[1]), importances[indices], color="g", yerr=std[indices], align="center") plt.xticks(range(X_train.shape[1]), X_train.columns[indices], rotation=90) plt.xlim([-1, X_train.shape[1]]) plt.show()</pre> Feature importances
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	We can learn that the feature "adr" is significantly more clearly to the prediction in comparison to the other features, for instance, this feature almost 3 times bigger (2.877) than the second most important feature.  Method 2 - Recursive feature elimination (RFE) with Random Forest Classification  The algorithm assign weights to each of features. Whose absolute weights are the smallest are pruned from the current set features. That procedure is recursively repeated on the pruned set until the desired number of features.
	<pre>clf_rf_ = RandomForestClassifier(n_estimators=20)     rfe = RFE(estimator=clf_rf_, n_features_to_select=11, step=1)     rfe = rfe.fit(X_train, y_train)  print('Chosen best 11 feature by RFE:',X_train.columns[rfe.support_])  Chosen best 11 feature by RFE: Index(['lead_time', 'arrival_date_year', 'arrival_date_month',</pre>
In [47]:	<pre>'required_car_parking_spaces'],     dtype='object')  This method brought me the same 11 most important features as the first method, therefore I will use only those 11 features and will activate the algorithm once again.  new_hotel_supervised = hotel_supervised.drop(columns = ['deposit_type', 'children', 'booking_changes', 'reservation_status', 'is_canceled', 'previous_bookings_not_canceled', 'previous_cancellations','days_i n_waiting_list','is_repeated_guest','babies','distribution_channel', 'deposit_type'])</pre>
In [49]: In [50]:	<pre>X = new_hotel_supervised.drop(columns = ['hotel']).copy() y = new_hotel_supervised["hotel"].copy()  X_train, X_test, y_train, y_test = split_test_train(X, y, 0.3)  model = create_naive_bayes_classifier(X_train, y_train)  y_pred = model.predict(X_test)</pre>
Out[52]: In [53]:	accuracy_score(y_test, y_pred)  0.7288921152557516  I increased the accuracy by 0.008 points - unfortunately not so significantly.  sns.heatmap(confusion_matrix(y_test, y_pred), annot=True, fmt="d") <matplotlib.axessubplots.axessubplot 0x2aldoffd790="" at=""></matplotlib.axessubplots.axessubplot>
	- 22213 1682 - 20000 - 17500 - 12500 - 10000
In [54]:	tn, fp, fn, tp = confusion_matrix(y_test, y_pred).ravel() print("TP:", tp) print("FN:", fn) print("FP:", fp) print("TN:", tn)
	TP: 3893 FN: 8028 FP: 1682 TN: 22213  From the confusion matrix, we can see that the algorithm predicted better the city hotel. However, the prediction of the resulting hotel was less good than before.
In [55]: Out[55]:	hotel_type = hotel_supervised.groupby('hotel')[['lead_time', 'arrival_date_year', 'arrival_date_month',
	In conclusion, in both tests, the accuracy of the algorithm was around 0.72. The feature that has the most clarity about the type of the hotel is the "adr" which is the daily rate, the "adr" of the city hotel is significantly higher than the resort hotel (on average of 10.348 higher). Except this feature, I can say that people book for a longer vacation in resort hotel since the average of the stays in the week and the weekend night is longer in resort hotel than the city hotel. Moreover, the lead time of the booking for the city hotel is longer than the resort hotel (on average of 17.07 longer). Finally, although the accuracy of the algorithm was above 0.5 (our starting point) the algorithm got difficulties to predict the resort hotel since the number of the city hotel is much higher in this data set than the resort hotel, and the range of almost every feature was very alike to the
	Now let's move one to the next the unsupervised question:  Can we split our data to N clusters such that every cluster is including similar booking features?  To answer that, I will use KMeans algorithm. This algorithm is a method of vector quantization, originally from signal processing, that aims to partition n observations into k clusters in which each observation belongs to the cluster with the nearest mean, serving as a prototype of the cluster.
In [56]:	Firstly, I 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 build a function which called 'number_of_clusters()'. This function will return us two graphs. The first graph is called "Sum of square error ", in this graph we will see what is the sum of square error as long as the number of clusters increases. Now comes the question of what is the number of clusters that are "suspicious" to be optimal. For answering this question, I will use the "Elbow Method". In this method, when we see a break in some number clusters, we will take that number as an option about how many clusters should we use in this data set. The second graph I will use is called "Silhouette", the silhouette is a measure about how good our algorithm work, his range is between -1 to 1. As long as the
In [57]:	<pre>value is 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 silhouette I got. I need to find the best cluster by a combination of those two graphs, a number of clusters which has a good "Elbow break" and ones that has the best silhouette.  def number_of_clusters(data_frame):     try:         sum_squared = []         silhouette = []         for i in range(2, 11):              kmeans = KMeans(n_clusters=i, init='k-means++')</pre>
	<pre>kmeans.fit(data_frame) sum_squared.append(kmeans.inertia_) silhouette.append(silhouette_score(data_frame, kmeans.labels_))  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)</pre>
	<pre>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()  except:     print("Something got wrong - number_of_clusters")</pre>
In [58]:	number_of_clusters(hotel_unsupervised)  Sum of Squred Error R <sub>2</sub> 300000 200000 200000
	As we can see, in the first plot, the sharper elbow break is when I divide the data into 6 clusters. However, when we look at the second plot,
	the highest silhouette is when I divide the data into 10 clusters although the silhouette score in each number of clusters is not high. Therefore, I will divide the data into 6 clusters.  Below is two functions:  • create_kmeans_classifier - creating KMeans algorithm.  • clusters_information - display each cluster and his statistic details.
In [59]: In [60]:	<pre>try:     return KMeans(n_clusters=k, init='k-means++') except:     print("Something got wrong - create_kmeans_calssifier")  def clusters_information(data_frame):     try:         clusters = data_frame.groupby("label")         for name, group in clusters:</pre>
In [61]:	<pre>print(name)     print(group)     print(group.describe())  except:     print("Something got wrong - clusters_information")  Activating the functions:</pre>
	<pre>kmeans = create_kmeans_calssifier(6) kmeans.fit(hotel_unsupervised) silhouette_score(hotel_unsupervised, kmeans.labels_)  0.2557500178616348</pre>

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As we car		houette score that we got is	1 . 00 s very low. An explanation		e the values of the features are s that have the same behavior.

119364

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