```
In [3]: import pandas as pd
import numpy as np
from sklearn import preprocessing
import matplotlib.pyplot as plt
import seaborn as sns
sns.set(style="white")#white background
sns.set(style="whitegrid",color_codes=True)
import warnings
warnings.simplefilter(action='ignore')
```

In [4]: df=pd.read_csv(r"C:\Users\DELL E5490\Downloads\heart disease.csv")
 df

Out[4]:

	male	age	education	currentSmoker	cigsPerDay	BPMeds	prevalentStroke	prevalentHyp	diabetes	totChol	sysBP	diaBP	BMI	heartR
0	1	39	4.0	0	0.0	0.0	0	0	0	195.0	106.0	70.0	26.97	8
1	0	46	2.0	0	0.0	0.0	0	0	0	250.0	121.0	81.0	28.73	9
2	1	48	1.0	1	20.0	0.0	0	0	0	245.0	127.5	80.0	25.34	7
3	0	61	3.0	1	30.0	0.0	0	1	0	225.0	150.0	95.0	28.58	6
4	0	46	3.0	1	23.0	0.0	0	0	0	285.0	130.0	84.0	23.10	8
4233	1	50	1.0	1	1.0	0.0	0	1	0	313.0	179.0	92.0	25.97	6
4234	1	51	3.0	1	43.0	0.0	0	0	0	207.0	126.5	80.0	19.71	6
4235	0	48	2.0	1	20.0	NaN	0	0	0	248.0	131.0	72.0	22.00	8
4236	0	44	1.0	1	15.0	0.0	0	0	0	210.0	126.5	87.0	19.16	8
4237	0	52	2.0	0	0.0	0.0	0	0	0	269.0	133.5	83.0	21.47	8

4238 rows × 16 columns

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In [5]: df.head()

Out[5]:

	male	age	education	currentSmoker	cigsPerDay	BPMeds	prevalentStroke	prevalentHyp	diabetes	totChol	sysBP	diaBP	вмі	heartRate
0	1	39	4.0	0	0.0	0.0	0	0	0	195.0	106.0	70.0	26.97	80.0
1	0	46	2.0	0	0.0	0.0	0	0	0	250.0	121.0	81.0	28.73	95.0
2	1	48	1.0	1	20.0	0.0	0	0	0	245.0	127.5	80.0	25.34	75.0
3	0	61	3.0	1	30.0	0.0	0	1	0	225.0	150.0	95.0	28.58	65.0
4	0	46	3.0	1	23.0	0.0	0	0	0	285.0	130.0	84.0	23.10	85.0
4														>

In [6]: df.shape

Out[6]: (4238, 16)

In [7]: df.describe

Out[7]:	<box< th=""><th>d metho</th><th>od ND</th><th>Frame.d</th><th>escribe</th><th>of</th><th>male</th><th>age educ</th><th>ation</th><th>current</th><th>Smoker</th><th>cigsPerDay</th><th>BPMeds</th><th>\</th></box<>	d metho	od ND	Frame.d	escribe	of	male	age educ	ation	current	Smoker	cigsPerDay	BPMeds	\
	0	1	39		4.0		0	0.0	0.0					
	1	0	46		2.0		0	0.0	0.0					
	2	1	48		1.0		1	20.0	0.0					
	3	0	61		3.0		1	30.0	0.0					
	4	0	46		3.0		1	23.0	0.0					
	• • •	• • •			• • •		• • •	• • •	• • •					
	4233	1	50		1.0		1	1.0	0.0					
	4234	1	51		3.0		1	43.0	0.0					
	4235	0	48		2.0		1	20.0	NaN					
	4236	0	44		1.0		1	15.0	0.0					
	4237	0	52		2.0		0	0.0	0.0					
		preva	lentS	troke	prevale	ntHvn	diahetes	totChol	sysBP	diaBP	BMI	\		
	0	p. c.u.		0	pi cvaze.	0	0		106.0	70.0	26.97	•		
	1			0		0	0		121.0	81.0	28.73			
	2			0		0	0		127.5	80.0	25.34			
	3			0		1	0		150.0	95.0				
	4			0		0	0		130.0	84.0	23.10			
	4233			0		1	0	313.0	179.0	92.0	25.97			
	4234			0		0	0	207.0	126.5	80.0	19.71			
	4235			0		0	0	248.0	131.0	72.0	22.00			
	4236			0		0	0	210.0	126.5	87.0	19.16			
	4237			0		0	0	269.0	133.5	83.0	21.47			
		heart	22+0	glucos	a TanV	earCHD								
	0		30.0	77.		0								
	1		95.0	76.		0								
	2		75.0	70.		0								
	3		55.0	103.		1								
	4		35.0	85.		0								
	4233	(56.0	86.		1								
	4234		55.0	68.		0								
	4235		34.0	86.		0								
	4236		36.0	Na		0								
	4237		30.0	107.		0								

localhost:8888/notebooks/heart disease.ipynb

[4238 rows x 16 columns]>

In [8]: df.info

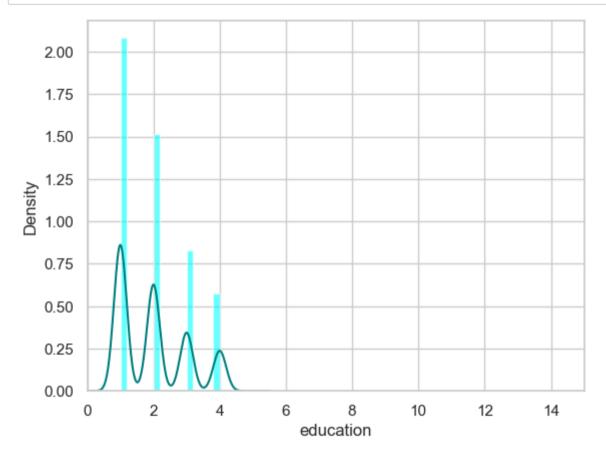
Out[8]:	<box< th=""><th>d metho</th><th>od Da</th><th>taFrame.</th><th>info of</th><th>male</th><th>age</th><th>educat</th><th>ion cu</th><th>ırrentSm</th><th>oker</th><th>cigsPerDay</th><th>BPMeds</th><th>\</th></box<>	d metho	od Da	taFrame.	info of	male	age	educat	ion cu	ırrentSm	oker	cigsPerDay	BPMeds	\
	0	1	39	۷	1.0	0	•	0.0	0.0					
	1	0	46	2	2.0	0		0.0	0.0					
	2	1	48	1	L.0	1		20.0	0.0					
	3	0	61	3	3.0	1		30.0	0.0					
	4	0	46		3.0	1		23.0	0.0					
					• •	• • •								
	4233	1	50	1	1.0	1		1.0	0.0					
	4234	1	51	3	3.0	1		43.0	0.0					
	4235	0	48	2	2.0	1		20.0	NaN					
	4236	0	44	1	1.0	1		15.0	0.0					
	4237	0	52	2	2.0	0		0.0	0.0					
		nnova	lon+C	tnoko r	prevalentHyp	diaboto	c +	o+Chol	sysBP	diaBP	BMI	\		
	0	pi eva.	Telles	0 0	0 evalencing		3 C	195.0	106.0	70.0	26.97			
	1			0	0		0	250.0	121.0	81.0	28.73			
	2			0	0		0	245.0	127.5	80.0	25.34			
							0	225.0	150.0					
	3 4			0 0	1 0		0	285.0		95.0 84.0	28.58 23.10			
				0					130.0					
	 4233			0	1	• •	0	313.0	 179.0	92.0	25.97			
	4234			0	0		0	207.0	126.5	80.0	19.71			
	4234			0	0		0	248.0	131.0	72.0	22.00			
	4235			0	0		0	210.0	126.5	87.0	19.16			
	4237			0	0		0	269.0	133.5	83.0	21.47			
	7237			O	O	•	O	203.0	100.0	03.0	21,7/			
		heart		glucose)								
	0		80.0	77.0										
	1		95.0	76.0										
	2		75.0	70.6	9									
	3	(65.0	103.6) 1									
	4	;	85.0	85.6	9 0									
	• • •		• • •	• • •										
	4233		66.0	86.6										
	4234		65.0	68.6										
	4235		84.0	86.6										
	4236		86.0	NaN										
	4237	;	80.0	107.6	9 0									
	_			_										

localhost:8888/notebooks/heart disease.ipynb

[4238 rows x 16 columns]>

In [9]:	<pre>df.isnull().sum()</pre>	
Out[9]:	male	0
	age	0
	education	105
	currentSmoker	0
	cigsPerDay	29
	BPMeds	53
	prevalentStroke	0
	prevalentHyp	0
	diabetes	0
	totChol	50
	sysBP	0
	diaBP	0
	BMI	19
	heartRate	1
	glucose	388
	TenYearCHD	0
	dtype: int64	

```
In [10]: ax=df["education"].hist(bins=15,density=True,stacked=True,color='cyan',alpha=0.6)
    df["education"].plot(kind='density',color='teal')
    ax.set(xlabel='education')
    plt.xlim(-0,15)
    plt.show()
```

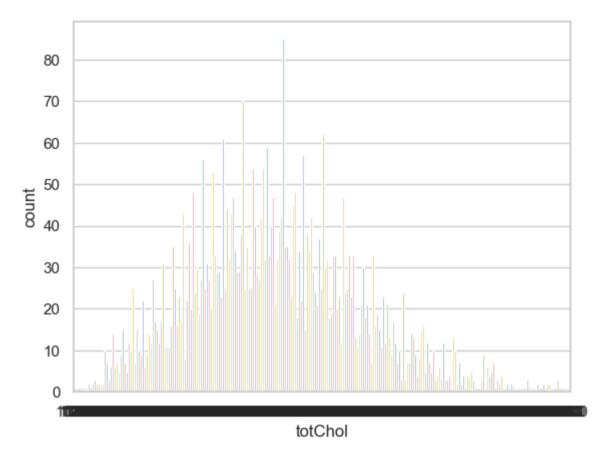


```
In [11]: print(df["education"].mean(skipna=True))
print(df["education"].median(skipna=True))
```

1.9789499153157513

2.0

```
In [14]: print(df['totChol'].value_counts())
    sns.countplot(x='totChol',data=df,palette='Set2')
           plt.show()
           240.0
                      85
           220.0
                      70
           260.0
                      62
           210.0
                      61
           232.0
                      59
           392.0
                       1
           405.0
                       1
           359.0
                       1
           398.0
                       1
           119.0
                       1
           Name: totChol, Length: 248, dtype: int64
```

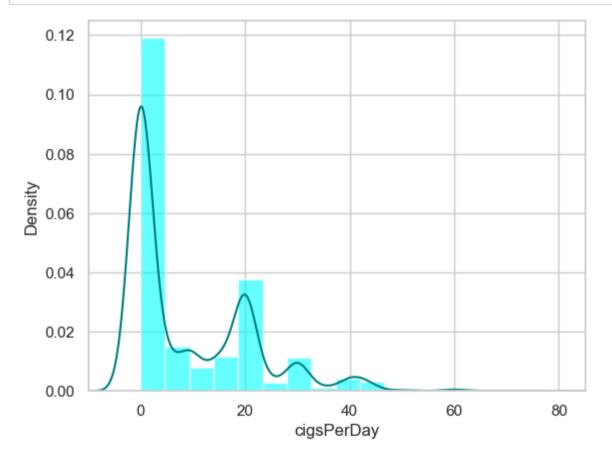


```
In [15]: print(df['totChol'].value_counts().idxmax())
240.0

In [16]: data=df.copy()
    data["education"].fillna(df["education"].median(skipna=True),inplace=True)
    data["totChol"].fillna(df["totChol"].median(skipna=True),inplace=True)
    data.drop('glucose',axis=1,inplace=True)
```

In [17]:	data.isnull().sum	()
Out[17]:	male	0
	age	0
	education	0
	currentSmoker	0
	cigsPerDay	29
	BPMeds	53
	prevalentStroke	0
	prevalentHyp	0
	diabetes	0
	totChol	0
	sysBP	0
	diaBP	0
	BMI	19
	heartRate	1
	TenYearCHD	0
	dtype: int64	

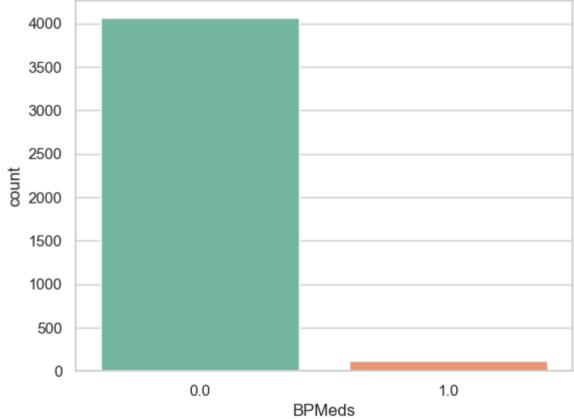
```
In [18]: ax=df["cigsPerDay"].hist(bins=15, density=True, stacked=True, color='cyan', alpha=0.6)
    df["cigsPerDay"].plot(kind='density',color='teal')
    ax.set(xlabel='cigsPerDay')
    plt.xlim(-10,85)
    plt.show()
```



```
In [19]: print(df["cigsPerDay"].mean(skipna=True))
print(df["cigsPerDay"].median(skipna=True))
```

9.003088619624615

0.0



```
In [24]: print(df['heartRate'].value_counts().idxmax())
```

75.0

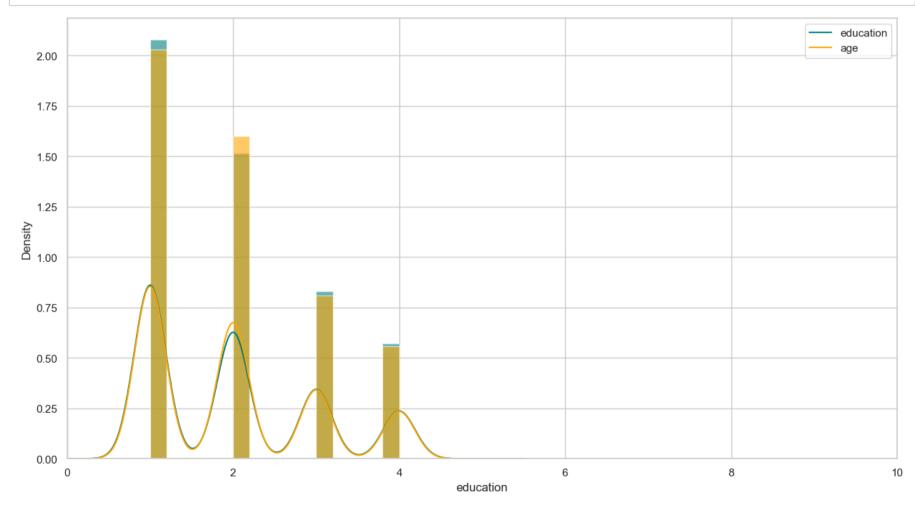
```
In [25]: data=df.copy()
         data["cigsPerDay"].fillna(df["cigsPerDay"].median(skipna=True),inplace=True)
         data["BPMeds"].fillna(df["BPMeds"].value_counts().idxmax(),inplace=True)
         data["education"].fillna(df["education"].median(skipna=True),inplace=True)
         data["totChol"].fillna(df["totChol"].value counts().idxmax(),inplace=True)
         data.drop('glucose',axis=1,inplace=True)
         data.drop('BMI', axis=1, inplace=True)
         data.drop('heartRate',axis=1,inplace=True)
          data.isnull().sum()
In [26]:
Out[26]: male
                             0
                             0
         age
         education
                             0
         currentSmoker
         cigsPerDay
         BPMeds
                             0
         prevalentStroke
         prevalentHyp
                             0
         diabetes
         totChol
                             0
         sysBP
         diaBP
                             0
         TenYearCHD
         dtype: int64
```

In [27]: data.head()

Out[27]:

	male	age	education	currentSmoker	cigsPerDay	BPMeds	prevalentStroke	prevalentHyp	diabetes	totChol	sysBP	diaBP	TenYearCHD	
0	1	39	4.0	0	0.0	0.0	0	0	0	195.0	106.0	70.0	0	
1	0	46	2.0	0	0.0	0.0	0	0	0	250.0	121.0	81.0	0	
2	1	48	1.0	1	20.0	0.0	0	0	0	245.0	127.5	80.0	0	
3	0	61	3.0	1	30.0	0.0	0	1	0	225.0	150.0	95.0	1	
4	0	46	3.0	1	23.0	0.0	0	0	0	285.0	130.0	84.0	0	

```
In [28]: plt.figure(figsize=(15,8))
    ax=df["education"].hist(bins=15, density=True, stacked=True, color='teal', alpha=0.6)
    df["education"].plot(kind='density',color='teal')
    ax=data["education"].hist (bins=15, density=True, stacked=True, color='orange', alpha=0.6)
    data["education"].plot(kind='density',color='orange')
    ax.legend(["education","age"])
    ax.set(xlabel='education')
    plt.xlim(-0,10)
    plt.show()
```



```
In [29]: data['Disease']=np.where((data["prevalentHyp"]+data[ "prevalentStroke"])>0,0,1)
    data.drop('prevalentHyp', axis=1,inplace=True)
    data.drop('prevalentStroke', axis=1, inplace=True)
```

```
In [30]: training=pd.get_dummies (data, columns=["currentSmoker", "totChol", "sysBP"])
    training.drop('TenYearCHD', axis=1, inplace=True)
    training.drop('male',axis=1, inplace=True)
    training.drop('diaBP',axis=1, inplace=True)
    final_train=training
    final_train.head()
```

Out[30]:

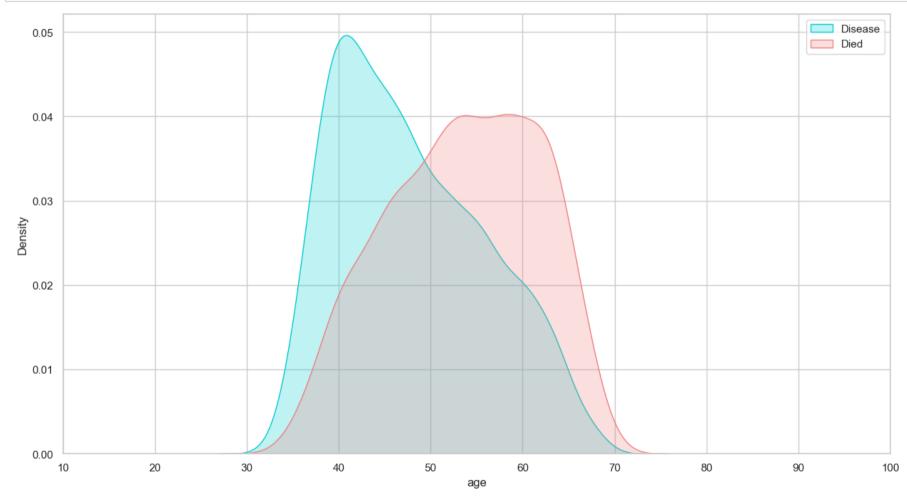
	age	education	cigsPerDay	BPMeds	diabetes	Disease	currentSmoker_0	currentSmoker_1	totChol_107.0	totChol_113.0	 sysBP_215.0 sy
0	39	4.0	0.0	0.0	0	1	1	0	0	0	 0
1	46	2.0	0.0	0.0	0	1	1	0	0	0	 0
2	48	1.0	20.0	0.0	0	1	0	1	0	0	 0
3	61	3.0	30.0	0.0	0	0	0	1	0	0	 0
4	46	3.0	23.0	0.0	0	1	0	1	0	0	 0

5 rows × 490 columns

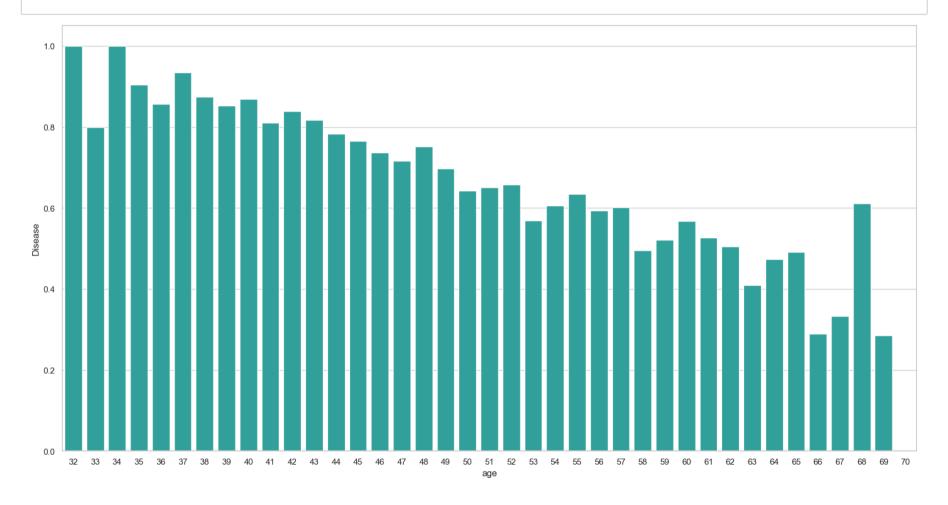
→

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```
In [31]: plt.figure(figsize=(15,8))
    ax = sns.kdeplot(final_train["age"][final_train.Disease==1], color="darkturquoise", shade=True)
    sns.kdeplot(final_train["age"][final_train.Disease == 0], color="lightcoral", shade=True)
    plt.legend(['Disease', 'Died'])
    ax.set(xlabel='age')
    plt.xlim(10, 100)
    plt.show()
```

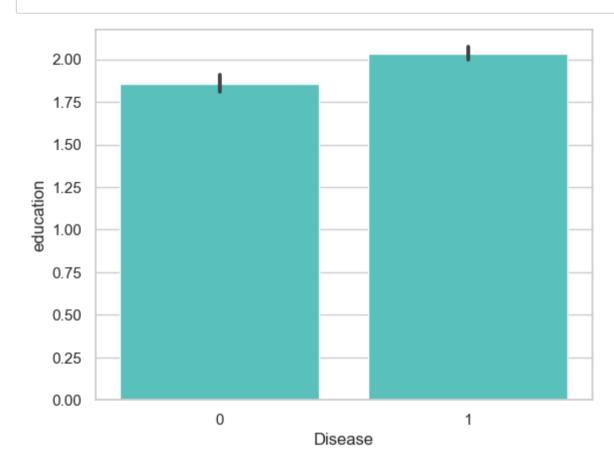


```
In [39]: plt.figure(figsize=(20,10))
    avg_survival_byage = final_train[["age", "Disease"]].groupby(['age'], as_index=False).mean()
    g=sns.barplot(x='age', y='Disease', data=avg_survival_byage, color="LightSeaGreen")
    plt.show()
```

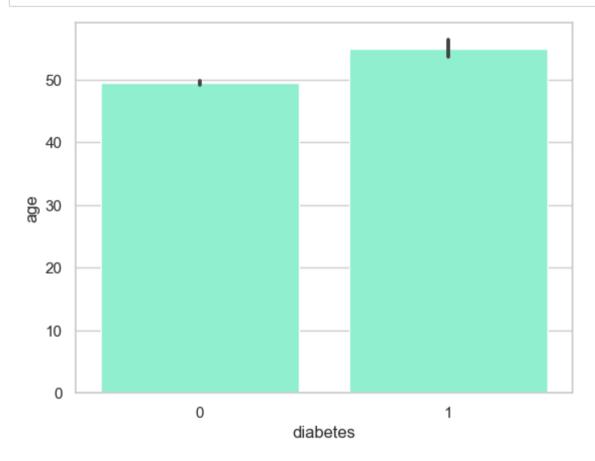


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```
In [37]: | sns.barplot(x='Disease', y='education', data=final_train, color="mediumturquoise")
    plt.show()
```



```
In [38]: import seaborn as sns
import matplotlib.pyplot as plt
# Assuming 'train_df' is your DataFrame containing the data
sns.barplot(x='diabetes', y='age', data=df, color='aquamarine')
plt.show()
```



In []: