Team ID	PNT2022TMID52707
Project Name	Statistical Machine Learning Approaches to Liver Disease Prediction.

## **Data Visualization**





## data.info()

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1636 entries, 0 to 1635
Data columns (total 11 columns):
```

#	Column	Non-Null Count	Dtype
0	Age	1636 non-null	int64
1	Gender	1636 non-null	object
2	Total_Bilirubin	1636 non-null	float64
3	Direct_Bilirubin	1636 non-null	float64
4	Alkaline_Phosphotase	1636 non-null	int64
5	Alamine_Aminotransferase	1636 non-null	int64
6	Aspartate_Aminotransferase	1636 non-null	int64
7	Total_Protiens	1636 non-null	float64
8	Albumin	1636 non-null	float64
9	Albumin_and_Globulin_Ratio	1624 non-null	float64
16	ð Dataset	1636 non-null	int64

dtypes: float64(5), int64(5), object(1)

memory usage: 140.7+ KB

	Age	Gender	Total_Bilirubin	Direct_Bilirubin	Alkaline_Phosphotase	Alamine_Aminotransferase	Aspartate_
1626	22	Female	2.2	1.0	215	159	
1627	28	Female	0.8	0.2	309	55	
1628	38	Male	0.7	0.2	110	22	
1629	25	Male	0.8	0.1	130	23	
1630	45	Female	0.7	0.2	164	21	
1631	45	Female	0.6	0.1	270	23	
1632	28	Female	0.6	0.1	137	22	
1633	28	Female	1.0	0.3	90	18	
1634	66	Male	1.0	0.3	190	30	
1635	66	Male	0.8	0.2	165	22	

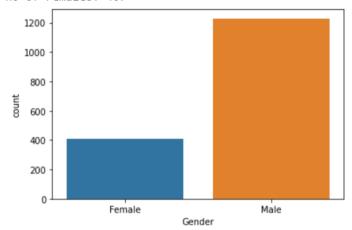
_	[8]	<pre>data.describe()</pre>

Total\_Bilirubin Direct\_Bilirubin Alkaline\_Phosphotase Alamine\_Aminotransferase Aspartate count 1636.000000 1636.000000 1636.000000 1636.000000 1636.000000 mean 44.727995 3.114792 1.387286 293.103912 80.944377 16.295775 5.955451 2.631630 248.412910 186.409237 std 10.000000 min 4.000000 0.400000 0.100000 63.000000 33.000000 23.000000 25% 0.800000 0.200000 175.000000 50% 45.000000 1.000000 0.300000 208.000000 35.000000 75% 58.000000 2.400000 1.200000 298.000000 60.000000 90.000000 75.000000 19.700000 2110.000000 2000.000000 max



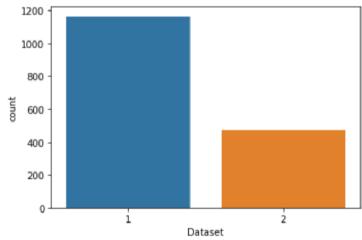
```
[13] sns.countplot(data=data,x='Gender',label='Count')
    m,f=data['Gender'].value_counts()
    print("No of Males:",m)
    print("no of Females:",f)
```

No of Males: 1229 no of Females: 407





liver disease patients: 1164 non-liver disease patients: 472



## [16] sns.heatmap(data.isnull(),yticklabels=False,cmap='viridis')



