

## Data Collection and Preprocessing Phase

Date	15 July 2024
Team ID	739716
Project Title	Predicting Baseline Histological staging in HCV patients using ML
Maximum Marks	6 Marks

Section	Description
Data Overview	<p>Now that we know the nature of the data, let's preprocess the information that was gathered.</p> <p>The downloaded data set may contain too much randomness to be used for training a machine learning model, so to get good results, the dataset must be carefully cleaned. The following steps are involved in this activity.</p> <ul style="list-style-type: none"> <li>? Handling missing values</li> <li>? Handling categorical data</li> </ul> <p>There are no categorical variables in our datasets</p> <p>The general procedures for pre-processing data before applying it to machine learning are as follows. Your dataset's state will determine whether or not you need to follow each of these stages.</p> <p>The first step will be to find the shape i.e. dimensions of the dataset. To find the shape of our data, the df. shape method is used. To find the data type, the df.info() function is used.</p> <pre>df.shape</pre> <p>(1385, 29)</p> <p>Thus, our dataset contains 1385 rows and 29 columns.</p>



```
: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1385 entries, 0 to 1384
Data columns (total 29 columns):
#   Column                                     Non-Null Count  Dtype
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0   Age                                         1385 non-null   int64
1   Gender                                     1385 non-null   int64
2   BMI                                         1385 non-null   int64
3   Fever                                       1385 non-null   int64
4   Nausea/Vomting                             1385 non-null   int64
5   Headache                                    1385 non-null   int64
6   Diarrhea                                    1385 non-null   int64
7   Fatigue & generalized bone ache           1385 non-null   int64
8   Jaundice                                    1385 non-null   int64
9   Epigastric pain                           1385 non-null   int64
10  WBC                                         1385 non-null   int64
11  RBC                                         1385 non-null   float64
12  HGB                                         1385 non-null   int64
13  Plat                                        1385 non-null   float64
14  AST 1                                       1385 non-null   int64
15  ALT 1                                       1385 non-null   int64
16  ALT4                                        1385 non-null   float64
17  ALT 12                                      1385 non-null   int64
18  ALT 24                                      1385 non-null   int64
19  ALT 36                                      1385 non-null   int64
20  ALT 48                                      1385 non-null   int64
21  ALT after 24 w                             1385 non-null   int64
22  RNA Base                                    1385 non-null   int64
23  RNA 4                                       1385 non-null   int64
24  RNA 12                                      1385 non-null   int64
25  RNA EOT                                    1385 non-null   int64
26  RNA EF                                     1385 non-null   int64
27  Baseline histological Grading              1385 non-null   int64
28  Baselinehistological staging               1385 non-null   int64
dtypes: float64(3), int64(26)
memory usage: 313.9 KB
```

From the above, we can see that there are no null values in this dataset.

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## Data exploration and preprocessing report

Data set variables will be statically analyzed to identify patterns and outliers with python employed for preprocessing tasks like normalization and feature engineering. data cleaning will address missing values and outliers, ensuring quality for subsequent analysis and modelling and forming a strong foundation for insights and predictions.

images will be

