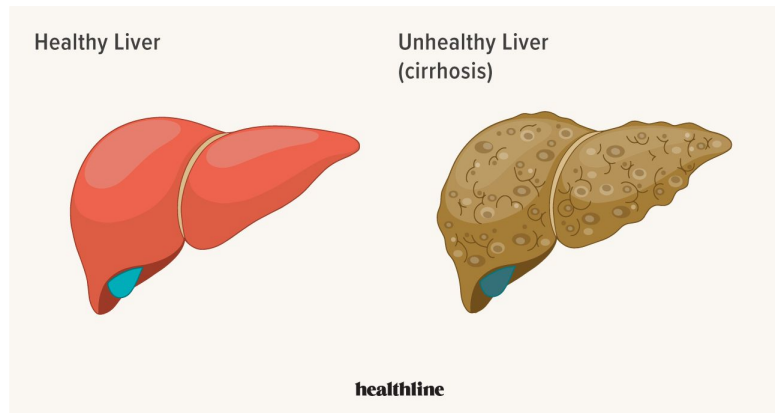


LIVER DISEASE PREDICTIONS



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DSP 556

DATA OVERVIEW

- Data gathered from South Korea spanning 2002–2021 through the South Korean National Health Insurance Service
- An annual random sampling of ~ 1 million individuals who underwent basic yearly check-ups covered by the South Korean National Health Insurance
- Dataset acquired via Kaggle

LITERATURE

- 4.5 million adults in the US are diagnosed with Liver disease each year
- Link between lipid profile (LDL, HDL, triglycerides) and liver disease observed in patients with cirrhosis
- Serum lipid levels serve as crucial indicators of liver damage
- GGT (glutamyltransferase) is a powerful predictor of liver disease
- Alcohol consumption and obesity as significant risk factors for developing cirrhosis

2009 vs 2019

- Analyzed data from 2009 and 2019
- The selection of 2009 and 2019 was deliberate to ensure a significant gap between the two sets of collected data
- Additionally, the exclusion of 2020 was intentional due to the disruptive impact of the COVID-19 pandemic.

DATA

- Variable Explanations

- Sex - M/F
- Age - groups 1-18 for every 5 years
- Height - centimeters
- Weight - kilograms
- BP_High & BP_LWST - Systolic and Diastolic Blood Pressure (N/mm Hg)
- BLDS - Blood Glucose level per 100ml of blood (N/mg/dL)
- TOT_CHOLE - Sum of ester and non-ester cholesterol (mg/dL)
 - Normal values 150-250 mg/dL
- HDL_CHOLE - Cholesterol level in HDL (mg/dL)
 - Normal values 30-65 mg/dL
- Triglyceride - amount of simple or neutral lipids (mg/dL)
 - Normal values 30-135 mg/dL
- HMG - level of pigment protein in blood, helps carry oxygen (g/dL)
- SGOT_AST - blood test level that indicates liver function (IU/L)
 - Normal values 0-40 IU/L, high concentration can indicate organ damage

GOAL

Generate machine learning models that utilize different health indicators to predict a patient's liver enzyme levels and thus their chance of possible liver disease

HCHK_YEAR	IDV_ID	SEX	AGE_GROUP	SIDO	HEIGHT	WEIGHT
Min. :2009	Min. : 1	Min. :1.000	Min. : 1.00	Min. :11.00	Min. :125.0	Min. : 25.00
1st Qu.:2009	1st Qu.: 250001	1st Qu.:1.000	1st Qu.: 4.00	1st Qu.:26.00	1st Qu.:155.0	1st Qu.: 55.00
Median :2009	Median : 500001	Median :1.000	Median : 6.00	Median :41.00	Median :160.0	Median : 60.00
Mean :2009	Mean : 500001	Mean :1.451	Mean : 6.02	Mean :33.15	Mean :161.9	Mean : 61.86
3rd Qu.:2009	3rd Qu.: 750000	3rd Qu.:2.000	3rd Qu.: 8.00	3rd Qu.:43.00	3rd Qu.:170.0	3rd Qu.: 70.00
Max. :2009	Max. :1000000	Max. :2.000	Max. :14.00	Max. :49.00	Max. :190.0	Max. :125.00

WAIST	SIGHT_LEFT	SIGHT_RIGHT	HEAR_LEFT	HEAR_RIGHT	BP_HIGH	BP_LWST
Min. : 51.00	Min. :0.1000	Min. :0.1000	Min. :1.000	Min. :1.000	Min. : 60.0	Min. : 30.00
1st Qu.: 74.00	1st Qu.:0.7000	1st Qu.:0.7000	1st Qu.:1.000	1st Qu.:1.000	1st Qu.:110.0	1st Qu.: 70.00
Median : 80.00	Median :1.0000	Median :1.0000	Median :1.000	Median :1.000	Median :120.0	Median : 78.00
Mean : 80.16	Mean :0.9732	Mean :0.9707	Mean :1.027	Mean :1.026	Mean :122.4	Mean : 76.29
3rd Qu.: 86.00	3rd Qu.:1.2000	3rd Qu.:1.2000	3rd Qu.:1.000	3rd Qu.:1.000	3rd Qu.:130.0	3rd Qu.: 80.00
Max. :129.00	Max. :9.9000	Max. :9.9000	Max. :2.000	Max. :2.000	Max. :310.0	Max. :190.00
	NA's :154	NA's :177	NA's :301	NA's :297		

BLDS	TOT_CHOLE	TRIGLYCERIDE	HDL_CHOLE	LDL_CHOLE	HMG	OLIG_PROTE_CD
Min. : 27.00	Min. : 43.0	Min. : 1.0	Min. : 1.00	Min. : 0.0	Min. : 0.10	Min. :1.000
1st Qu.: 85.00	1st Qu.:169.0	1st Qu.: 75.0	1st Qu.: 45.00	1st Qu.: 90.0	1st Qu.:12.90	1st Qu.:1.000
Median : 93.00	Median :192.0	Median :109.0	Median : 53.00	Median :111.0	Median :14.00	Median :1.000
Mean : 97.25	Mean :195.1	Mean :134.7	Mean : 56.39	Mean :120.9	Mean :13.96	Mean :1.083
3rd Qu.:102.00	3rd Qu.:217.0	3rd Qu.:163.0	3rd Qu.: 63.00	3rd Qu.:134.0	3rd Qu.:15.10	3rd Qu.:1.000
Max. :999.00	Max. :999.0	Max. :999.0	Max. :991.00	Max. :9998.0	Max. :23.80	Max. :6.000
				NA's :79		NA's :3410

CREATININE	SGOT_AST	SGPT_ALT	GAMMA_GTP	SMK_STAT_TYPE_CD	DRK_YN
Min. : 0.100	Min. : 1.00	Min. : 1.00	Min. : 1.00	Min. :1.000	Length:1000000
1st Qu.: 0.800	1st Qu.: 19.00	1st Qu.: 15.00	1st Qu.: 16.00	1st Qu.:1.000	Class :character
Median : 0.900	Median : 23.00	Median : 20.00	Median : 23.00	Median :1.000	Mode :character
Mean : 1.121	Mean : 25.36	Mean : 25.35	Mean : 36.63	Mean :1.666	
3rd Qu.: 1.100	3rd Qu.: 28.00	3rd Qu.: 29.00	3rd Qu.: 39.00	3rd Qu.:3.000	
Max. :99.900	Max. :999.00	Max. :999.00	Max. :999.00	Max. :3.000	
				NA's :3932	

HCHK_OE_INSPEC_YN	CRS_YN	TTH_MSS_YN	ODT_TRB_YN	WSDM_DIS_YN	TTR_YN
Length:1000000	Length:1000000	Length:1000000	Length:1000000	Length:1000000	Length:1000000
Class :character	Class :character	Class :character	Class :character	Class :character	Class :character
Mode :character	Mode :character	Mode :character	Mode :character	Mode :character	Mode :character

2009 SUMMARY STATS

DATA_STD_DT
Length:1000000
Class :character
Mode :character

> summary(Checkup_2)

	YEAR	IDV_ID	AREA_CODE	SEX	AGE_GROUP
Min.	:2019	Min. : 1	Min. :11.00	Min. :1.000	Min. : 5.0
1st Qu.:	:2019	1st Qu.: 202287	1st Qu.:27.00	1st Qu.:1.000	1st Qu.: 8.0
Median :	:2019	Median : 468191	Median :41.00	Median :1.000	Median :11.0
Mean :	:2019	Mean : 478699	Mean :33.71	Mean :1.481	Mean :10.5
3rd Qu.:	:2019	3rd Qu.: 734096	3rd Qu.:43.00	3rd Qu.:2.000	3rd Qu.:13.0
Max.	:2019	Max. :1000000	Max. :50.00	Max. :2.000	Max. :18.0

	HEIGHT	WEIGHT	WAIST	SIGHT_LEFT	SIGHT_RIGHT
Min.	:130.0	Min. : 30.00	Min. : 3.00	Min. :0.1000	Min. :0.1000
1st Qu.:	:155.0	1st Qu.: 55.00	1st Qu.: 74.00	1st Qu.:0.7000	1st Qu.:0.7000
Median :	:160.0	Median : 60.00	Median : 81.00	Median :1.0000	Median :1.0000
Mean :	:162.3	Mean : 63.56	Mean : 81.32	Mean :0.9719	Mean :0.9707
3rd Qu.:	:170.0	3rd Qu.: 70.00	3rd Qu.: 88.00	3rd Qu.:1.2000	3rd Qu.:1.2000
Max.	:195.0	Max. :145.00	Max. :999.00	Max. :9.9000	Max. :9.9000
			NA's :423	NA's :207	NA's :200

	HEAR_LEFT	HEAR_RIGHT	BP_HIGH	BP_LWST	BLDS
Min.	:1.000	Min. :1.000	Min. : 57.0	Min. : 28.00	Min. : 2.0
1st Qu.:	:1.000	1st Qu.:1.000	1st Qu.:112.0	1st Qu.: 70.00	1st Qu.: 89.0
Median :	:1.000	Median :1.000	Median :121.0	Median : 76.00	Median : 96.0
Mean :	:1.033	Mean :1.032	Mean :122.5	Mean : 75.76	Mean :100.9
3rd Qu.:	:1.000	3rd Qu.:1.000	3rd Qu.:131.0	3rd Qu.: 82.00	3rd Qu.:105.0
Max.	:2.000	Max. :2.000	Max. :260.0	Max. :200.00	Max. :960.0
NA's	:191	NA's :188	NA's :5799	NA's :5800	NA's :5904

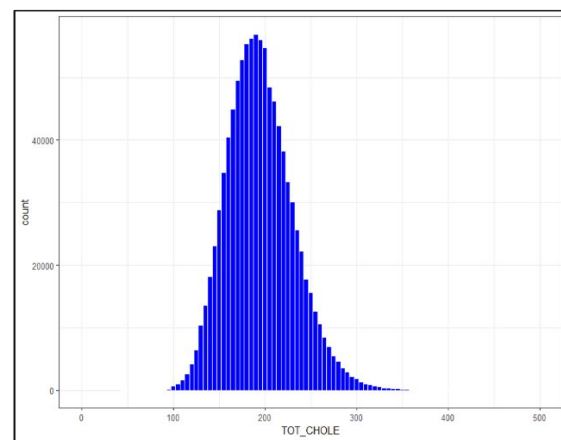
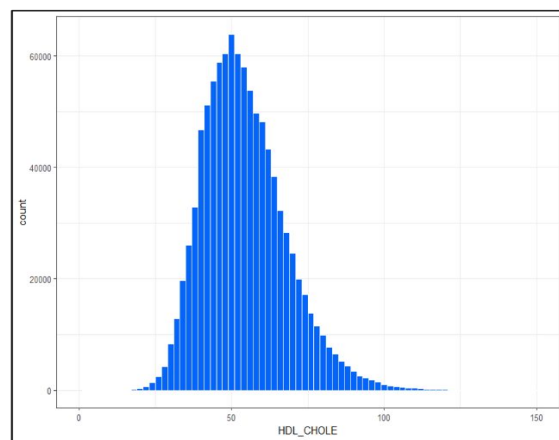
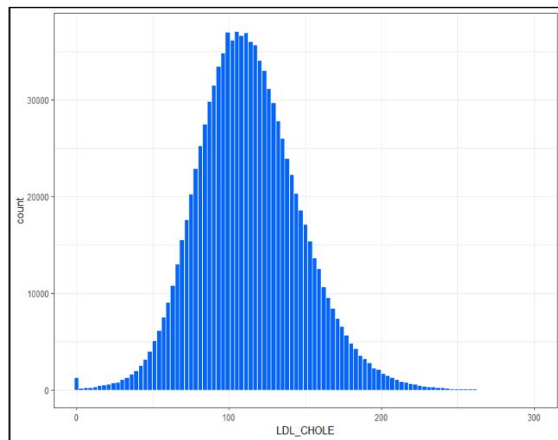
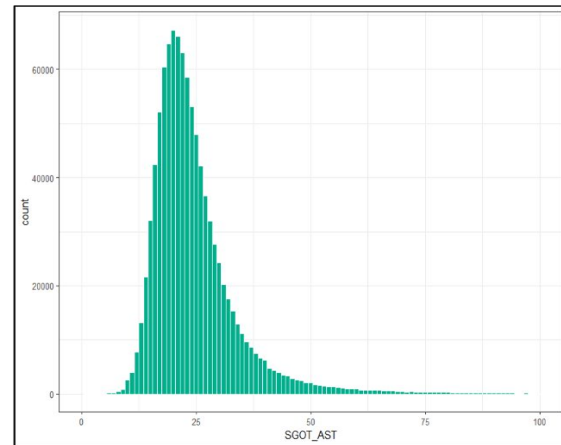
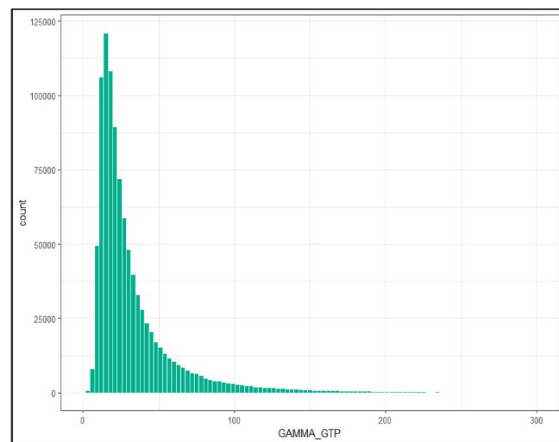
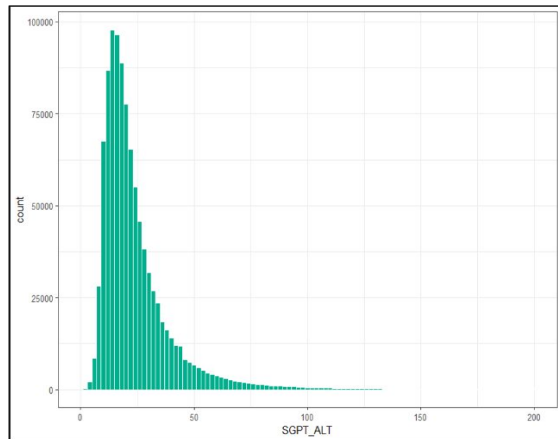
	TOT_CHOLE	TRIGLYCERIDE	HDL_CHOLE	LDL_CHOLE	HMG
Min.	: 50.0	Min. : 4.0	Min. : 1.0	Min. : 1.0	Min. : 0.80
1st Qu.:	:169.0	1st Qu.: 76.0	1st Qu.:46.0	1st Qu.: 89.0	1st Qu.:13.20
Median :	:195.0	Median :109.0	Median : 55.0	Median :112.0	Median :14.30
Mean :	:196.4	Mean :133.9	Mean : 56.6	Mean :113.9	Mean :14.25
3rd Qu.:	:221.0	3rd Qu.:161.0	3rd Qu.: 65.0	3rd Qu.:137.0	3rd Qu.:15.40
Max.	:2389.0	Max. :4879.0	Max. :588.0	Max. :2278.0	Max. :25.00
NA's	:708726	NA's :708731	NA's :708734	NA's :715543	NA's :5915

2019 SUMMARY STATS

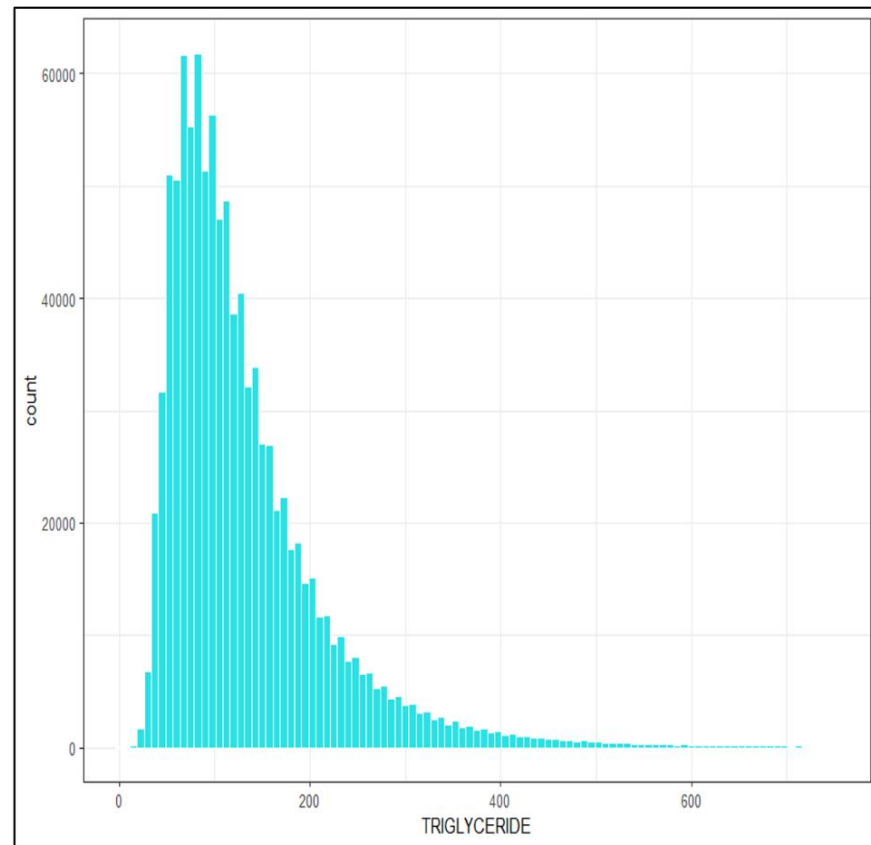
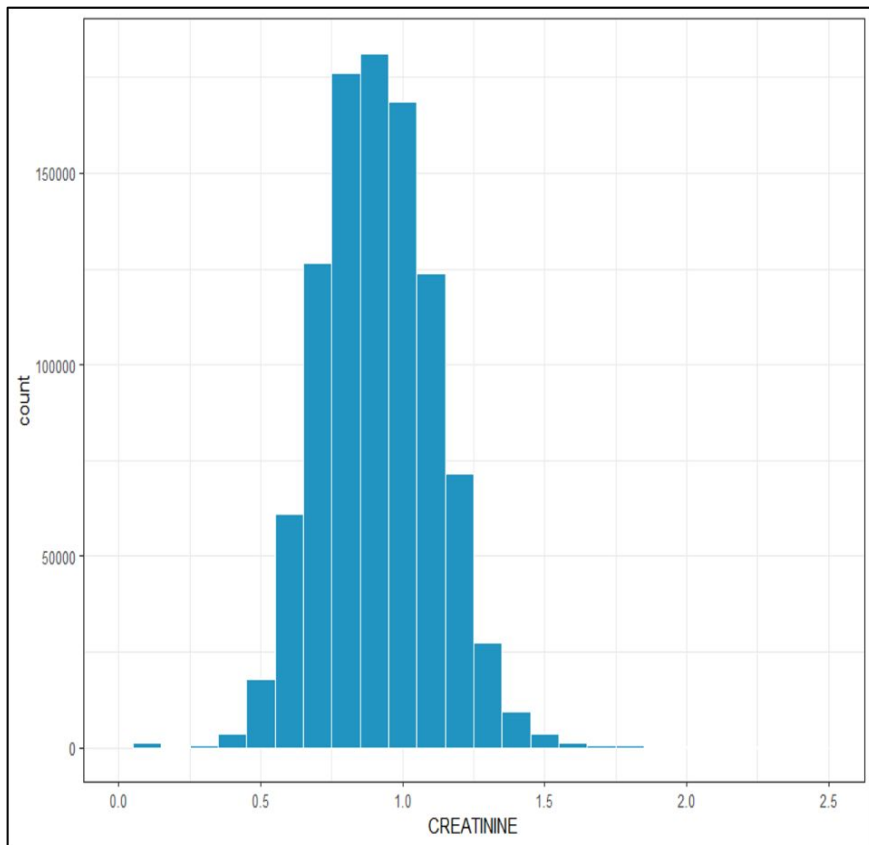
2019 SUMMARY STATS

TOT_CHOLE	TRIGLYCERIDE	HDL_CHOLE	LDL_CHOLE	HMG
Min. : 50.0	Min. : 4.0	Min. : 1.0	Min. : 1.0	Min. : 0.80
1st Qu.: 169.0	1st Qu.: 76.0	1st Qu.: 46.0	1st Qu.: 89.0	1st Qu.:13.20
Median : 195.0	Median : 109.0	Median : 55.0	Median : 112.0	Median :14.30
Mean : 196.4	Mean : 133.9	Mean : 56.6	Mean : 113.9	Mean :14.25
3rd Qu.: 221.0	3rd Qu.: 161.0	3rd Qu.: 65.0	3rd Qu.: 137.0	3rd Qu.:15.40
Max. :2389.0	Max. :4879.0	Max. :588.0	Max. :2278.0	Max. :25.00
NA's :708726	NA's :708731	NA's :708734	NA's :715543	NA's :5915
OLIG_PROTE_CD	CREATININE	SGOT_AST	SGPT_ALT	GAMMA_GTP
Min. :1.000	Min. : 0.06	Min. : 1.00	Min. : 1	Min. : 1.00
1st Qu.:1.000	1st Qu.: 0.70	1st Qu.: 19.00	1st Qu.: 15	1st Qu.: 16.00
Median :1.000	Median : 0.80	Median : 23.00	Median : 20	Median : 23.00
Mean :1.104	Mean : 0.86	Mean : 26.23	Mean : 26	Mean : 36.33
3rd Qu.:1.000	3rd Qu.: 1.00	3rd Qu.: 29.00	3rd Qu.: 30	3rd Qu.: 39.00
Max. :6.000	Max. :98.00	Max. :7362.00	Max. :6435	Max. :999.00
NA's :10976	NA's :5907	NA's :5903	NA's :5904	NA's :5911
SMK_STAT	DRK_YN	HCHK_CE_IN	CRS_YN	TTR_YN
Min. :1.000	Min. :0	Min. :0.0000	Min. :0.0	Min. :0.0
1st Qu.:1.000	1st Qu.:1	1st Qu.:0.0000	1st Qu.:0.0	1st Qu.:0.0
Median :1.000	Median :1	Median :0.0000	Median :0.0	Median :1.0
Mean :1.369	Mean :1	Mean :0.3977	Mean :0.2	Mean :0.6
3rd Qu.:2.000	3rd Qu.:1	3rd Qu.:1.0000	3rd Qu.:0.0	3rd Qu.:1.0
Max. :2.000	Max. :1	Max. :1.0000	Max. :1.0	Max. :2.0
NA's :174	NA's :377809		NA's :640647	NA's :640647
DATE	Alcohol			
Length:1063619	Min. :1			
Class :character	1st Qu.:1			
Mode :character	Median :1			
	Mean :1			
	3rd Qu.:1			
	Max. :1			
	NA's :377809			

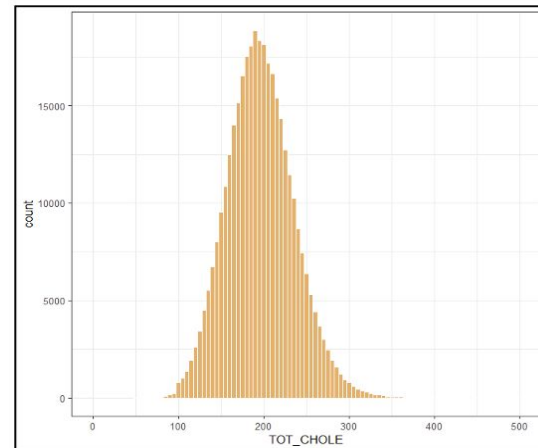
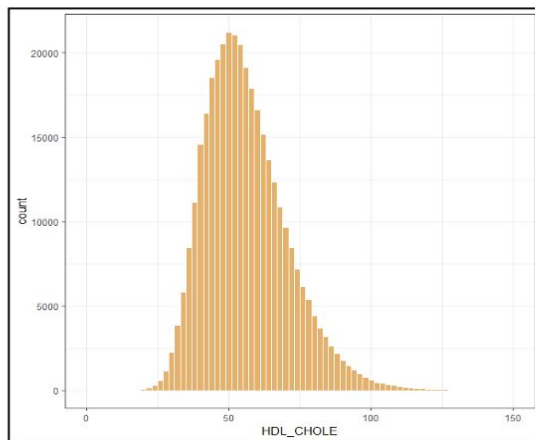
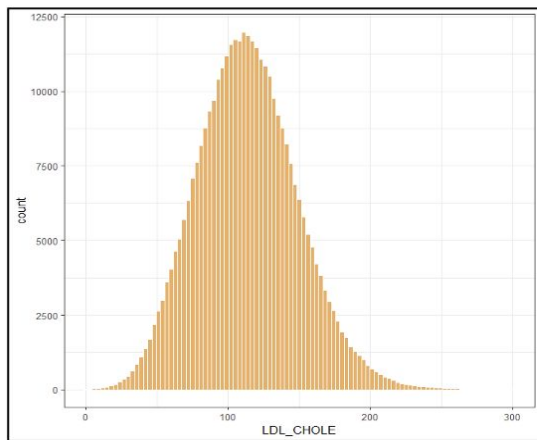
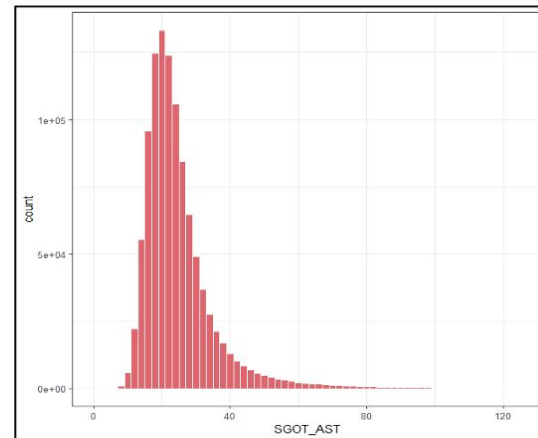
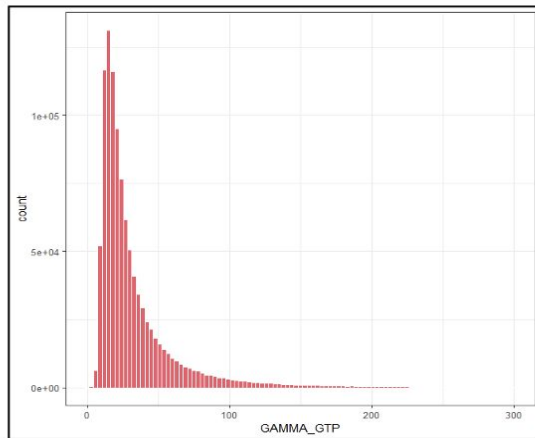
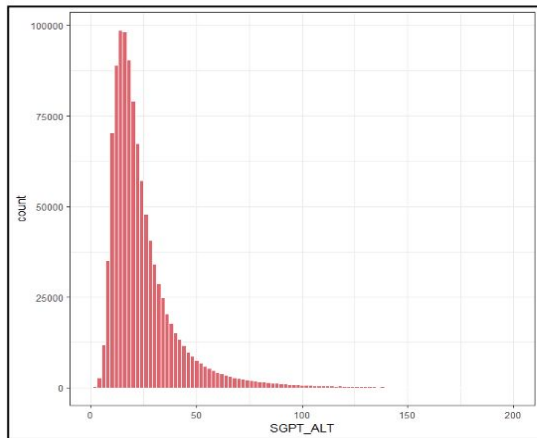
2009 KEY VARIABLES



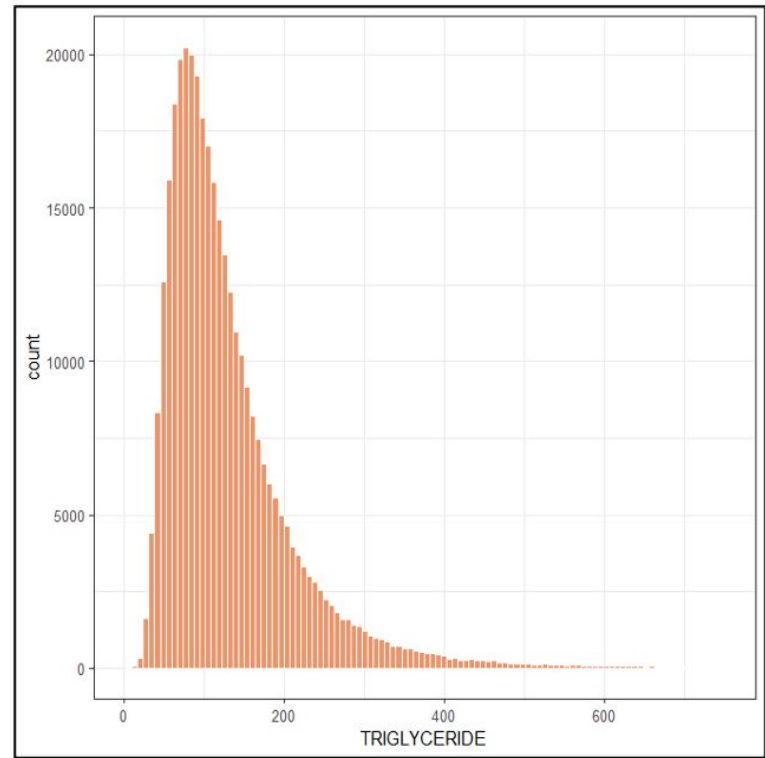
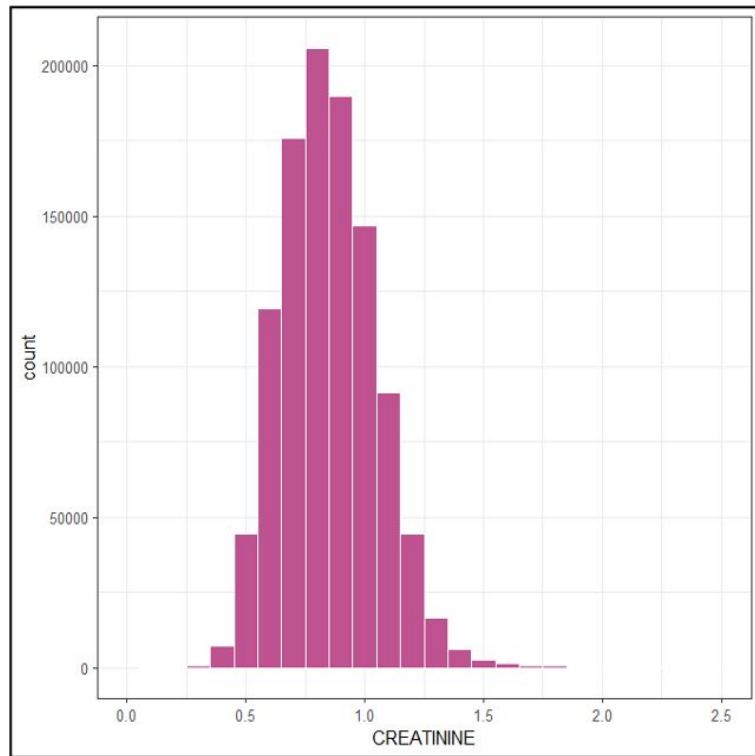
2009 KEY VARIABLES



2019 KEY VARIABLES



2019 KEY VARIABLES



METHODS AND DATA PREPROCESSING

- Utilized the following models from the SKLearn library
 - Logistic Regression
 - Decision Tree Classifier
 - Random Forest Classifier
- Numeric variables scaled using StandardScaler
- Our Response Variable, SGOT_AST, was encoded from a numeric to yes/no depending on the value being above or below 40(0-40 being the normal level expected)
- Hyperparameter Tuning with GridSearchCV
 - Tol and C in Logistic Regression Model
 - Max_Depth and Criterion in Decision Tree Classifier Model
 - Max_Depth and N_estimators in Random Forest Classifier Model
- Training/Test Splits created with ratio of 75:25

RESULTS

- 2009

```
[0] Logistic Regression Accuracy: 0.8914666666666666
[1] Decision Tree Accuracy: 0.9466666666666667
[2] Random Forest Classifier Accuracy: 0.9373333333333334
```

```
model 0
[[2311 189]
 [ 389 211]]
Testing Accuracy = 0.8122157244964262
```

```
model 1
[[2392 108]
 [ 326 2174]]
Testing Accuracy = 0.8520790729379687
```

```
model 2
[[2407 93]
 [ 357 2143]]
Testing Accuracy = 0.847457627118644
```

2019

```
[0] Logistic Regression Accuracy: 0.8916
[1] Decision Tree Accuracy: 0.9366666666666666
[2] Random Forest Classifier Accuracy: 0.93
```

```
model 0
[[1125 132]
 [ 163 1080]]
Testing Accuracy = 0.8099226804123711
```

```
model 1
[[1151 106]
 [ 75 1168]]
Testing Accuracy = 0.8741307371349096
```

```
model 2
[[1137 120]
 [ 75 1168]]
Testing Accuracy = 0.8657024793388429
```

ENSEMBLE METHOD RESULTS

- 2009

Accuracy Score of Bagging Classifier: 0.92

Accuracy Score of Stacking Classifier: 0.94

Accuracy Score of GradientBoost: 0.91

2019

Accuracy Score of Bagging Classifier: 0.90

Accuracy Score of Stacking Classifier: 0.94

Accuracy Score of GradientBoost: 0.91

RESULTS

- Able to build several models that predicted the correct liver enzyme levels at an accuracy rate between 81% and 94%
 - We are able to accurately predict elevated SGOT_AST liver enzyme levels and thus possible liver problems at a high rate
- Utilized subsets of the data for each model to avoid long computation times with 1 million observations

FUTURE EFFORTS

- Utilize more observations to create a more accurate predictive model
- Incorporate other indicators into the model
- Analyze liver enzyme levels for people of different countries
- Work to implement these models into a clinical setting to help notify doctors when a patient may be at risk of a liver problem due to heightened liver enzyme levels



THANK YOU!