In [1]:	import numpy as numport pandas as	ries	ON DETECTIO	N=====================================	
	<pre>import matplotlib import seaborn as  from sklearn.prep  from sklearn.mode from sklearn.mode from sklearn.mode from sklearn.mode from sklearn.mode from sklearn.mode  from sklearn.mode  from sklearn.neig from sklearn.line from sklearn.tree from sklearn.tree from sklearn.metr</pre>	.pyplot as plt sns  rocessing import  L_selection impor L_selection impor L_selection impor L_selection impor  hbors import KNei ar_model import L import DecisionT import SVC  ics import accura	<pre>t train_test_split t KFold t cross_val_score t GridSearchCV ghborsClassifier ogisticRegression</pre>		
<pre>In [2]: Out[2]:</pre>	dataset.head()  Age Gender Total_Bilirubi  0 65 Female 0  1 62 Male 10  2 62 Male 7  3 58 Male 1	csv( a/KUBUNTU/DATASET  in Direct_Bilirubin Alkaline_I  7 0.1  9 5.5	S/CSV_DATASETS/ind:  Phosphotase Alamine_Aminotra  187  699  490  182  195		
In [3]:	<ul><li>5 Alamine_Aminotransf</li><li>6 Aspartate_Aminotran</li><li>7 Total_Protiens</li><li>8 Albumin</li><li>9 Albumin_and_Globuli</li></ul>	ne.DataFrame'> 0 to 582 columns):  Non-Null Count 583 non-null 583 non-null 583 non-null 6e 583 non-null ierase 583 non-null sferase 583 non-null 583 non-null 583 non-null 583 non-null 583 non-null	Dtype int64 object float64 float64 int64 int64 int64 float64 float64 float64		
In [4]:	memory usage: 50.2+ KB  preprocessing the  # filling the nul  dataset['Albumin_	and_Globulin_Ratimin_and_Globulin_  Non-Null Count  1	Ratio'].mean(), inpose interview of the content of		its mean
In [5]:	4 Alkaline_Phosphotas 5 Alamine_Aminotrans 6 Aspartate_Aminotran 7 Total_Protiens 8 Albumin 9 Albumin_and_Globuli 10 Dataset dtypes: float64(5), int6 memory usage: 50.2+ KB   Converting each n  dataset = pd.get_  visualzing each at  def draw_hist(dat	se 583 non-null serase 583 non-null serase 583 non-null 583 non-null 583 non-null 583 non-null 583 non-null 583 non-null 584(5), object(1)  con-numeric data dummies (dataset) ctribute  aset, rows, cols)	int64 int64 float64 float64 float64 int64  to numeric for vis		
In [7]:	<pre>counter = 0 for i in rang for j in     if co     b     name     sns.d     count  draw_hist(dataset</pre>	<pre>range(cols): unter &gt;= len(name reak = names[counter] istplot(a=dataset er += 1 =dataset, rows=3,</pre>	s): [name], ax=axes[i, cols=4)		
	function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7 function and will be rem l function with similar warnings.warn(msg, Fut /usr/local/lib/python3.7	doved in a future version flexibility) or histple flexibility histple flexibility histple flexibility histple flexibility hist	distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619: Fon. Please adapt your code of (an axes-level funct distributions.py:2619	le to use either `displosion for histograms).  SutureWarning: `distplosion for histograms).	is a deprecated of a figure-leve is a figure-leve is a figure-leve is a figure-leve is a figure-lev
In [8]:	0.015 0.000 0.005 0.000 0.005 0.0010 0.0012 0.0008 0.0004 0.0002 0.0000 0.0004 0.0002 0.0000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.75 0.50 0.25 0.000 0.001 0.001 0.002 0.0001 0.0	0.3 - 0.2 - 0.1 - 0.008 - 0.008 - 0.006 - 0.005 - 0.004 - 0.003 - 0.002 - 0.001 - 0.000 - 0.00	0.4 - 0.3 - 0.2 - 0.1 - 0.0 2 4 Total Tota	t_Bilirubin	Gender_Male
Out[8]:	1	=dataset[selected]  id at 0x7fda1182a5d0>	_pair_cols], hue="	Dataset", kind='r	Dataset 1 2
<pre>In [9]: Out[9]:</pre>	using the correlati  corr_matrix = dat  fig, ax = plt.sub	on matrix  aset.corr() plots(figsize=(12 matrix, annot=Tru	, 12)) e, linewidths=.5, a		6 8 10 Protiens - 1.00
	Direct_Bilirubin - 0.007  Alkaline_Phosphotase - 0.08  Alamine_Aminotransferase - 0.02  Aspartate_Aminotransferase - 0.02  Albumin - 0.27  Albumin_and_Globulin_Ratio - 0.22  Dataset - 0.14  Gender_Female - 0.057  Gender_Male - 0.057	0.21 0.23 1 0.13  0.21 0.23 0.13 1  0.24 0.26 0.17 0.79  0 0.0081 0.00014 0.029 0.043  0 0.22 0.23 0.17 0.03  1 0.22 0.23 0.17 0.03  1 0.089 0.1 0.027 0.082  1 0.089 0.1 0.027 0.082	0.17	-0.15	- 0.50 - 0.25 - 0.00 0.25 0.50 0.75
In [10]:	creating the mode craete train and test of X = dataset.drop( y = dataset['Data X_train, X_test,	dataset 'Dataset', axis=1 set'] # target	) # data  train_test_split()	y, stratify=y, test_size=0.3,	
In [12]:	models.append(("L models.append(("K models.append(("D models.append(("S training and fillting	R", LogisticRegre NN", KNeighborsCl T", DecisionTreeC VM", SVC()))	ssion())) assifier(n_neighboo lassifier()))		
	<pre>print(10 * "= print("Accura print("Confus print("Classi</pre>	<pre>predict(X_test) ", "{} Results fo cy Score:{:0.2f}" ion Matrix:". fication Report:\ ation_report(y_te ")</pre>		core(y_test, y_pr	
	to converge (status=1): STOP: TOTAL NO. of ITERA  Increase the number of i     https://scikit-learn Please also refer to the     https://scikit-learn     extra_warning_msg=_LOG	0.90 0.82 0.26 0.34  0.71 0.58 0.58 0.71 0.68	125 50 175 175 175 175 175 175 175 175 175 175	in: regression	
In [14]:	on and F-score are ill-orameter to control this _warn_prf(average, model image).  Ising the cross variable to cross variable the cross variable to converge (status=1):  STOP: TOTAL NO. of ITERAL Increase the number of in https://scikit-learn.	defined and being set to behavior.  lifier, msg_start, len(rediction)  n models: (n_splits=5) ross_val_score(moders)  oss_validation_rediction_redictions  result.mean())))  dist-packages/sklearn/ attions_reactions_tier() ord lorg/stable/modules/predictions_reactions	del, X, y, cv=kfolosult of model:{:0.2	d, scoring="accurate".format(	cy")
	https://scikit-learn extra_warning_msg=_LOG /usr/local/lib/python3.7 to converge (status=1): STOP: TOTAL NO. of ITERA  Increase the number of i     https://scikit-learn Please also refer to the     https://scikit-learn extra_warning_msg=_LOG /usr/local/lib/python3.7 to converge (status=1): STOP: TOTAL NO. of ITERA  Increase the number of i     https://scikit-learn Please also refer to the     https://scikit-learn extra_warning_msg=_LOG /usr/local/lib/python3.7 to converge (status=1): STOP: TOTAL NO. of ITERA  Increase the number of i     https://scikit-learn Please also refer to the     https://scikit-learn extra_warning_msg=_LOG /usr/local/lib/python3.7 to converge (status=1): STOP: TOTAL NO. of ITERA  Increase the number of i     https://scikit-learn Please also refer to the     https://scikit-learn Please also refer to the     https://scikit-learn Please also refer to the     https://scikit-learn	i.org/stable/modules/linisTIC_SOLVER_CONVERGENCE/dist-packages/sklearn/modules/productions (max_iter) or i.org/stable/modules/productions (max_iter) or i.org/stable/modules/linisTIC_SOLVER_CONVERGENCE/dist-packages/sklearn/modules/productions (max_iter) or i.org/stable/modules/productions (max_iter) or i.org/stable/modules/productions (max_iter) or i.org/stable/modules/linisTIC_SOLVER_CONVERGENCE/dist-packages/sklearn/modules/productions (max_iter) or i.org/stable/modules/productions (max_iter) or i.org/stable/modules/productions (max_iter) or i.org/stable/modules/productions (max_iter) or i.org/stable/modules/linisTIC_SOLVER_CONVERGENCE/dist-packages/sklearn/modules/productions (max_iter) or i.org/stable/modules/productions (max_iter) or i.org/stable/modules/productions (max_iter) or i.org/stable/modules/linisTIC_SOLVER_CONVERGENCE/dist-packages/sklearn/modules/linisTIC_SO	r scale the data as shown eprocessing.html ernative solver options: near_model.html#logistic.p r scale the data as shown eprocessing.html ernative solver options: near_model.html#logistic.p r scale the data as shown eprocessing.html ernative solver options: near_model.html#logistic.p r scale the data as shown eprocessing.html ernative solver options: near_model.html#logistic.p r scale the data as shown eprocessing.html ernative solver options: near_model.html#logistic.p r scale the data as shown eprocessing.html ernative solver options: near_model.html#logistic.p	oy:940: ConvergenceWarn:  in in:  regression  oy:940: ConvergenceWarn:  orin:  regression  oy:940: ConvergenceWarn:  orin:  regression  oy:940: ConvergenceWarn:  orin:  orin:  orin:  orin:  orin:  orin:  orin:  orin:	ing: lbfgs failed
In [15]:	<pre>params_clfs = lis  svm_params = [{         'kernel': ['r         'gamma': [1e-] }, {         'kernel': ['l         'C': [1, 10, }]  params_clfs.appen  lr_params = {'pen params_clfs.appen  clf = DecisionTre dt_params = {         'max_features         'min_samples_         'random_state }</pre>	<pre>t() bf'], 3, 1e-4] inear'], 100, 1000] d((SVC(), svm_para alty': ['l1', 'l2 d((LogisticRegres eClassifier() ': ['auto', 'sqrt split': [2, 3, 4, leaf': [1], ': [123]</pre>	ams)) '], 'C': np.logspac sion(), lr_params)		15],
In [16]:	grid_search = grid_search.f print((%0 * " print(("{} sk	GridSearchCV(clf it(X_train, y_tra *")) learn result for ssname))) params:{}".format grid_search.cv_re ge test result:{: _search.predict(X_  rameter result:{: sion Matrix:" gsize=(12, 8)) ata=confusion_mat nnot=True, map="coolwarm", ticklabels=['Heal ticklabels=['Heal ticklabels=['Heal redicted values') ctual values') nfusion Matrix by  ification report(y_te	in)  GridSearch for svc  (grid_search.best_  sults_['mean_test_! .2f}".format(np.mea_ test)  .2f}".format(accura .format(confusion_r rix(y_test, y_pred  thy', 'Unhealthy'] thy', 'Unhealthy']  Grid Search')  ".format(	<pre>params_))) score'] an(test_means)))) acy_score(y_test,</pre>	

