Multimedia Appendix 1. Hyperparameter tuning.

The hyperparameters of the ML classifiers obtained through the grid search cross-validation were as follows:

Model	Hyperparameters	Best combination	
		All factors	Oxidative stress factors only
		assification	
	C_ values: [0.001, 0.01, 0.1, 1, 10, 100]	C_value: 1	C_value: 0.001
LR	Penalty: $[l1, l2]$	Penalty: 12	Penalty: 12
	Solvers: liblinear, lbfgs, newton-cg, sag, saga]	Solver: liblinear	Solver: liblinear
RF	n_estimators: [50, 100, 150, 200, 300] max_depth: [None, 5, 10, 15, 20, 25, 30]	n_estimators: 50	n_estimators: 50
	min_samples_split: [2, 5, 10]	$\max_depth: 15$	$\max_depth:$ None
	min_samples_leaf': [1, 2, 4]	$\min_samples_split: 2$	$\min_samples_split: 5$
	bootstrap: [True, False]	min_samples_leaf: 1	min_samples_leaf: 1
	max features: [auto, sqrt, log2]	bootstrap: False	bootstrap: False
	max_leaf_nodes: [None, 5, 10, 20, 50, 100]	max_leaf_nodes: None	max_leaf_nodes: None
KNN	n_neighbors : [3, 5,7,9,11,13,15]	n_neighbors: 11	n_neighbors: 15
	weights: [uniform, distance]	weights: distance	weights: distance
	metric: [minkowski, euclidean, manhattan]	Metric: manhattan	Metric: manhattan
	C: [0.1, 1, 10, 100]	C: 1	C: 1
SVM	gamma: $[1, 0.1, 0.01, 0.001]$	gamma: 0.1	gamma: 1
	kernel: [linear, rbf]	kernel: rbf	kernel:rbf
NB	Naive Bayes has almost no hyperpar	rameters to tune, so it usua	ally generalizes well
	solvers = ['newton-cg', 'liblinear']	solver: newton-cg	solver: liblinear
WLR	penalty = ['l1', 'l2']	penalty: l2	penalty: 12
	$c_values = [100, 10, 1.0, 0.1, 0.01]$	C: 1	C: 0.1
	n_estimators: [50, 100, 150, 200, 300] max_depth: [None, 5, 10, 15, 20, 25, 30]	n_estimators: 50	n_estimators: 100
WRF	min samples split: [2, 5, 10]	$\max_depth: None$	$\max_depth: 25$
	min_samples_leaf': [1, 2, 4]	min_samples_split: 2	min_samples_split: 2
	bootstrap: [True, False]	min_samples_leaf: 1	min_samples_leaf: 1
	max_features: [auto, sqrt, log2]	bootstrap: True	bootstrap: False
	max_leaf_nodes: [None, 5, 10, 20, 50, 100]	max_leaf_nodes: None	max_leaf_nodes: None
	Multi-class	Classification	

Table 1: Hyperparameter value range and optimal values for all ML methods.

	solvers = ['newton-cg', 'lbfgs', 'liblinear']	Solver: liblinear	Solver: liblinear
LR	penalty = ['11', '12']	Penalty: 12	Penalty: 12
	$c_values = [100, 10, 1.0, 0.1, 0.01]$	C: 10	C: 0.001
	n_estimators: [50, 100, 150, 200, 300] max_depth: [None, 5, 10, 15, 20, 25, 30]	n_estimators: 100	n_estimators: 100
	min samples split: $[2, 5, 10]$	max_depth: 15	$\max_depth: 15$
RF	$min_samples_leaf': [1, 2, 4]$	min_samples_split: 2	$\min_samples_split: 2$
	bootstrap: [True, False]	min_samples_leaf: 1	min_samples_leaf: 1
	max_features: [auto, sqrt, log2]	bootstrap: False	bootstrap: False
	max_leaf_nodes: [None, 5, 10, 20, 50, 100]	max_leaf_nodes: None	max_leaf_nodes: None
	$n_{neighbors'} : [5,7,9,11,13,15]$	n_neighbors: 13	n_neighbors: 15
KNN	weights: ['uniform','distance']	weights: distance	weights: distance
	metric:['minkowski','euclidean', 'manhattan']	Metric: Manhattan	Metric: Manhattan
	C: [1, 10, 100]	C: 1	C: 100
SVM	gamma: [1,0.1,0.01,0.001]	gamma: 0.1	gamma: 1
	kernel: ['rbf']	kernel: rbf	kernel: rbf
NB	naive Bayes has almost no hyperpar	ameters to tune, so it usua	ally generalizes well
	solvers = ['newton-cg', 'liblinear']	solver: liblinear	solver: liblinear
WLR	penalty = ['l1', 'l2']	penalty: 12	penalty: 12
	$c_values = [100, 10, 1.0, 0.1, 0.01]$	C: 0.001	C: 0.1
	n_estimators: [50, 100, 150, 200, 300]	n_estimators: 50	n_estimators: 300
	<pre>max_depth: [None, 5, 10, 15, 20, 25, 30] min_samples_split: [2, 5, 10]</pre>	max_depth: None	$\max_depth: 15$
WRF	min_samples_leaf': [1, 2, 4]	min_samples_split: 2	min_samples_split: 2
	bootstrap: [True, False]	min_samples_leaf: 1	min_samples_leaf: 1
	max features: [auto, sqrt, log2]	bootstrap: True	bootstrap: True
	max_leaf_nodes: [None, 5, 10, 20, 50, 100]	max_leaf_nodes: None	max_leaf_nodes: None

Multimedia Appendix 2. Confusion Matrices

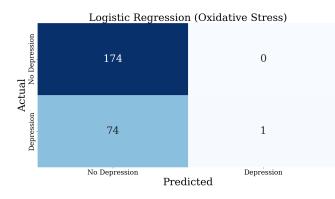


Figure 1: Confusion matrix of LR with oxidative stress biomarkers as the main features.

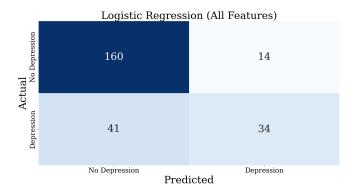


Figure 2: Confusion matrix of LR with all features.

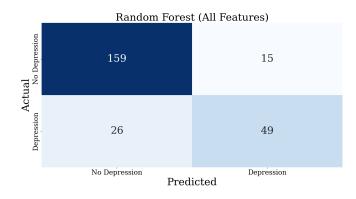


Figure 3: Confusion matrix of RF with all features.

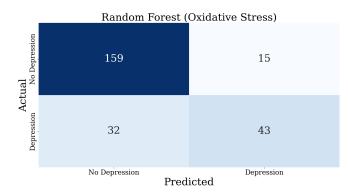


Figure 4: Confusion matrix of RF with oxidative stress biomarkers as the main features.

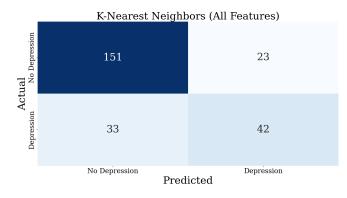


Figure 5: Confusion matrix of KNN with all features.

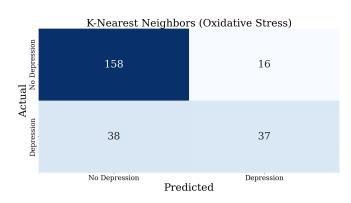


Figure 6: Confusion matrix of KNN with oxidative stress biomarkers as the main features.

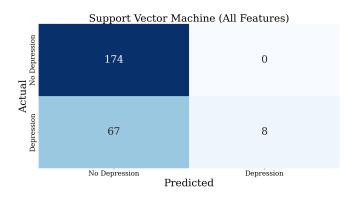


Figure 7: Confusion matrix of SVM with all features.

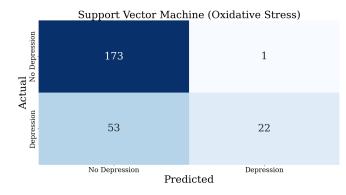


Figure 8: Confusion matrix of SVM with oxidative stress biomarkers as the main features.

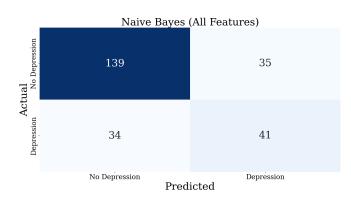


Figure 9: Confusion matrix of NB with all features.

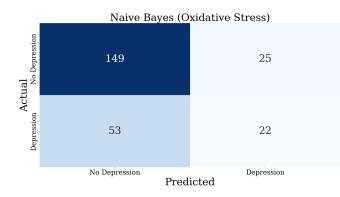


Figure 10: Confusion matrix of NB with oxidative stress biomarkers as the main features.

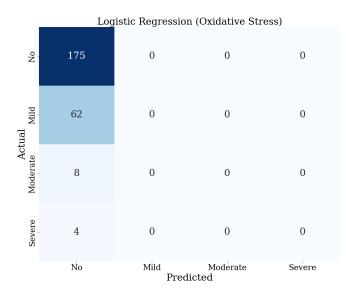


Figure 11: Confusion matrix of LR with oxidative stress biomarkers as the main features.

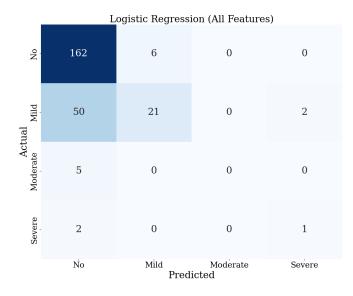


Figure 12: Confusion matrix of LR with all features.

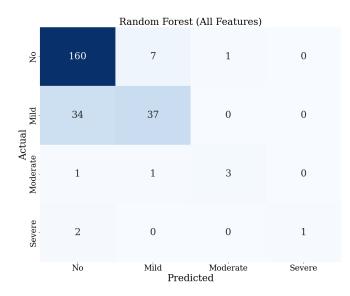


Figure 13: Confusion matrix of RF with all features.

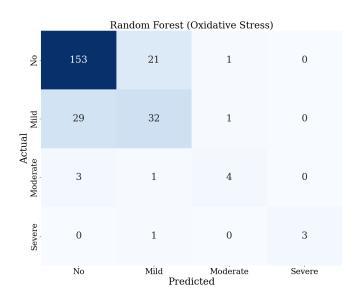


Figure 14: Confusion matrix of RF with oxidative stress biomarkers as the main features.

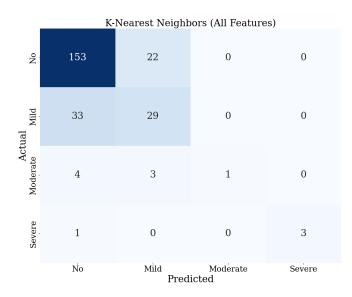


Figure 15: Confusion matrix of KNN with all features.

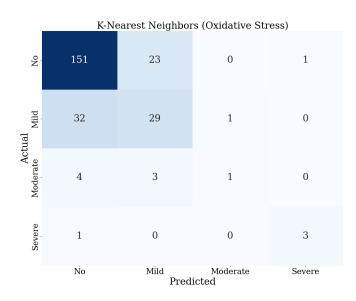


Figure 16: Confusion matrix of KNN with oxidative stress biomarkers as the main features.

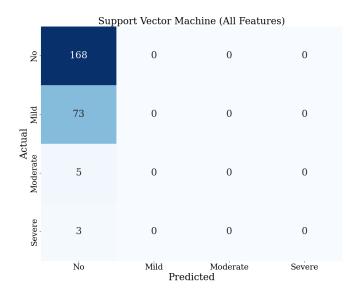


Figure 17: Confusion matrix of SVM with all features.

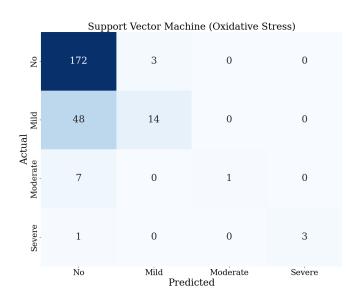


Figure 18: Confusion matrix of SVM with oxidative stress biomarkers as the main features.

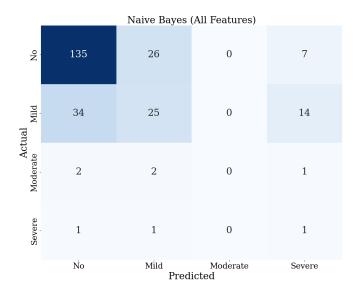


Figure 19: Confusion matrix of NB with all features.

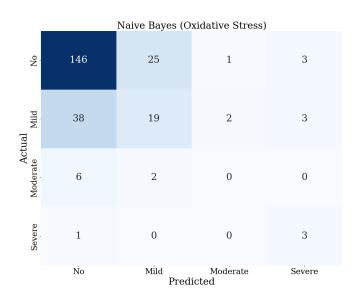


Figure 20: Confusion matrix of NB with oxidative stress biomarkers as the main features.

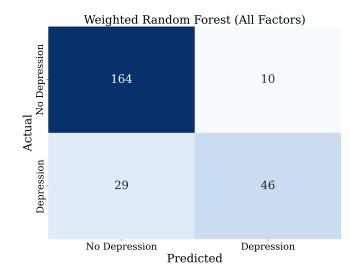


Figure 21: Confusion matrix of weighted RF with all features.

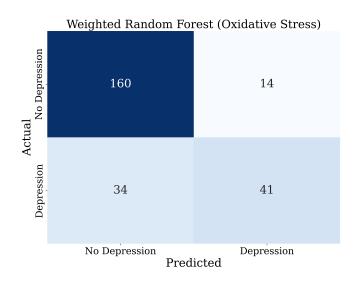


Figure 22: Confusion matrix of weighted RF with oxidative stress biomarkers as the main features.

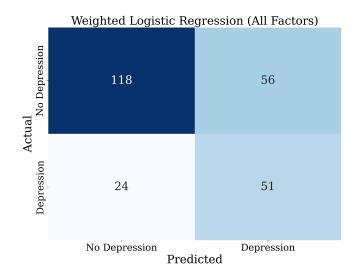


Figure 23: Confusion matrix of weighted LR with all features.

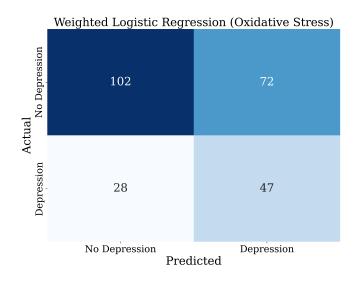


Figure 24: Confusion matrix of weighted LR with oxidative stress biomarkers as the main features.

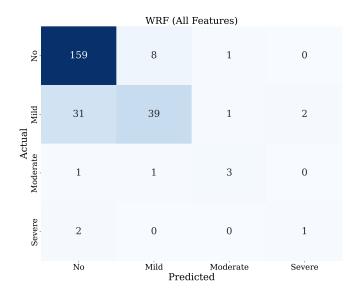


Figure 25: Confusion matrix of weighted RF with all features.

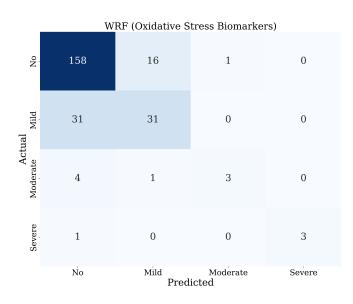


Figure 26: Confusion matrix of weighted RF with oxidative stress biomarkers as the main features.

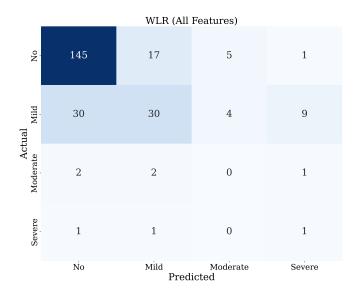


Figure 27: Confusion matrix of weighted LR with all features.

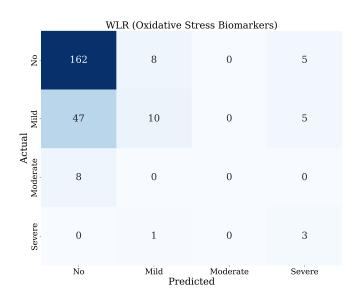


Figure 28: Confusion matrix of weighted LR with oxidative stress biomarkers as the main features.