

## ALGORITHM1: PSEUDO-CODE OF THE MACHINE LEARNING WORKFLOW FOR PREDICTING FOLLOW-UP PERIOD CATEGORY

```

Input: Diabetes dataset  $D$ 
Output: Trained classification models and performance metrics

1  Preprocess  $D$ :
2      Handle missing values
3      Apply one-hot encoding
4      Normalize features using RobustScaler
5  Define Feature Selection with SMOTE:
6       $FS\_set = \{$ Filter method (Mutual Information (MI)),
           Wrapper method (Sequential Feature Selection (SFS), Genetic Algorithm (GA)),
           Embedded method (Random Forest feature importance (RFI), Elastic Net (EN))
            $\}$ 
7  for each feature selection method  $FS\_method$  in  $FS\_set$  do
8      Apply  $FS\_method$  on  $D \rightarrow$  obtain selected features  $FS$ 
9      Initialize  $K$ -fold cross-validation ( $K = 5$ )
10     for each fold  $k = 1$  to  $K$  do
11         Split  $D$  into  $D\_train$  and  $D\_test$ 
12         Apply SMOTE to  $D\_train$ 
13         Extract  $X\_train, y\_train$  using  $FS$ 
14         Extract  $X\_test, y\_test$  using  $FS$ 
15         for each model  $M$  in  $\{SVM, KNN, RF, ETC, AB, ANN\}$  do
16             Train  $M$  on  $X\_train, y\_train$ 
17             Predict  $y\_pred = M(X\_test)$ 
18             Evaluate performance (Accuracy, F1, AUC, etc.)
19             Store results with tag ( $FS\_method, M$ )
           end for
10         end for
7     end for
20 Summarize and compare performance across  $FS\_method \times$  model combinations
21 end

```

Figure S1 Pseudo-code of the machine learning workflow for predicting follow-up period category.

**ALGORITHM2: PSEUDO-CODE OF THE FEATURE SELECTION FRAMEWORK**

```
Input: Dataset  $D$  with feature set  $F$ 
Output: Reduced feature subsets from each selection method

1 Define Feature Selection:
2    $FS\_set = \{Filter: \{Mutual\ Information\ (MI)\},$ 
       $Wrapper: \{Sequential\ Feature\ Selection\ (SFS),\ Genetic\ Algorithm\ (GA)\},$ 
       $Embedded: \{Elastic\ Net\ (EN),\ Random\ Forest\ Feature\ Importance\ (RFFI)\}$ 
       $\}$ 
3 Initialize result dictionary  $F\_selected = \emptyset$ 
4 for each feature selection category  $C$  in  $FS\_set$  do
5   for each method  $M$  in  $FS\_set[C]$  do
6     if  $M = MI$  then
7       Compute MI score for each feature in  $F$ 
8       Select top- $k$  features  $\rightarrow F\_selected[M]$ 
9     else if  $M = SFS$  or  $GA$  then
10      Initialize Decision Tree Classifier
11      Apply wrapper search (forward selection or genetic strategy)
12      Select optimal subset  $\rightarrow F\_selected[M]$ 
13     else if  $M = EN$  then
14      Train Elastic Net model on  $D$ 
15      Rank features by coefficient magnitude
16      Select top- $k$  features  $\rightarrow F\_selected[M]$ 
17     else if  $M = RFFI$  then
18      Train Random Forest Classifier
19      Extract feature importance scores
20      Select top- $k$  features  $\rightarrow F\_selected[M]$ 
21     end if
22   end for
23 end for
Output all reduced feature subsets:  $\{F\_selected[MI], F\_selected[SFS], F\_selected[GA], F\_selected[EN],$ 
 $F\_selected[RFFI]\}$ 
end
```

**Figure S2** Pseudocode of the feature selection framework

**Table S1** Parameter settings of SFS

Parameter	Description	Value
k_features	Number of features to select	10
forward	Direction of feature selection	True
scoring	Evaluation metric	Accuracy
cv	Cross-validation folds	5

**Table S2** Parameter settings of GA

Parameter	Description	Value
k_features	Number of features to select	10
population size	Number of individuals in population	50
ngen	Number of generations	40
cxpb	Crossover probability	0.5
mutpb	Mutation probability	0.2
tournsize	Tournament size for selection	3

**Table S3** Parameter settings of EN

Parameter	Description	Value
alpha	Regularization strength	[0.01, 0.1, 1, 10, 100]
l1_ratio	L1 to L2 penalty ratio	[0.1, 0.5, 0.7, 0.9, 1.0]
max_iter	Maximum iterations	10000

**Table S4** Parameter settings of RF feature importance

Parameter	Description	Value
criterion	The function to measure the quality of a split.	'gini'
max_depth	The maximum depth of the tree.	None
min_samples_split	The minimum number of samples required to split an internal node.	2
min_samples_leaf	The minimum number of samples required to be at a leaf node.	1

**Table S5** Hyperparameter settings for classification algorithm

Classification Algorithm	Hyperparameter setting
SVM	C=1.0, kernel='rbf', degree=3, gamma='scale', coef0=0.0
RF	n_estimators=100, criterion='gini', max_depth=None, min_samples_split=2, min_samples_leaf=1, max_features='sqrt'
KNN	n_neighbors=5, weights='uniform', algorithm='auto', leaf_size=30, p=2, metric='minkowski'
ETC	n_estimators=100, criterion='gini', max_depth=None, min_samples_split=2, min_samples_leaf=1, max_features='sqrt'
AB	n_estimators=50, learning_rate=1.0, algorithm='deprecated'
ANN	hidden_layer_sizes=(100,), activation='relu', solver='adam', alpha=0.0001, batch_size='auto', learning_rate='constant', learning_rate_init=0.001, max_iter=200,