¹ AIFS: A novel perspective, Artificial

- ² Intelligence infused wrapper based
- ³ Feature Selection Algorithm on High
- ⁴ Dimensional data analysis
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12 Abstract

Background: Feature selection is important in high dimensional data analysis. The wrapper approach is one of the ways to perform feature selection, but it is computationally intensive as it builds and evaluates models of multiple subsets of features. The existing wrapper approaches primarily focus on shortening the path to find an optimal feature set. However, these approaches underutilize the capability of feature subset models, which impacts feature selection and its predictive performance.

18 Method and Results: This study proposes a novel Artificial Intelligence infused wrapper based 19 Feature Selection (AIFS), a new feature selection method that integrates artificial intelligence with 20 wrapper based feature selection. The approach creates a Performance Prediction Model (PPM) using 21 artificial intelligence (AI) which predicts the performance of any feature set and allows wrapper 22 based methods to predict and evaluate the feature subset model performance without building 23 actual model. The algorithm can make wrapper based method more relevant for high-dimensional 24 data and is flexible to be applicable in any wrapper based method. We evaluate the performance of 25 this algorithm using simulated studies and real research studies. AIFS shows better or at par feature 26 selection and model prediction performance than standard penalized feature selection algorithms 27 like LASSO and sparse partial least squares.

Conclusion: AIFS approach provides an alternative method to the existing approaches for feature
 selection. The current study focuses on AIFS application in continuous cross-sectional data.
 However, it could be applied to other datasets like longitudinal, categorical and time-to-event
 biological data.

32 Keywords

33 High dimensional data, wrapper feature selection, artificial intelligence, AIFS, machine learning,

34 interaction terms

35 Background

Large feature space (p) is an important aspect of high dimensional data owing to the risk of model overfitting and poor model generalizability [1] and increased computational complexity [2, 3]. Feature selection is a solution which reduces the input feature space to smaller feature space (q) in a given dataset of sample size (n), which provides a parsimonious best fit model for the outcome, y.

$$y = f(q) | q \in (p) \#(1)$$

$$\min \varphi(y, f(q))$$

40 where, *f* represents the model function, and φ represents the error function. The approaches 41 adopted for feature selection can be categorized into two groups. The first and simpler approach 42 uses expert opinion for feature selection where features are selected using domain knowledge [4, 5] 43 and allows feature selection before evaluating the data. This approach has limitation or no 44 applicability if a feature has no or little availability of domain information, high dimensional feature 45 space and/or presence of interactions among the features [6].

46 The second and prominent approach uses the sampled data to perform the feature selection which 47 is broadly classified into filter, embedded and wrapper methods [7–9]. These methods could be used 48 in supervised, semi-supervised or unsupervised learning frameworks [9–11]. Filter methods rely on 49 the internal data structure of the features for selecting features. Commonly, information gain based 50 techniques are used for univariate filtering of features [9, 12] and correlation based techniques are 51 used for multivariate filtering of features [13]. They are computationally efficient, but interactions 52 between the features may hinder the model performance. Embedded methods incorporate feature 53 selection within the model building step by adding a penalization step in the model building process. They are efficient and have the ability to handle interactions between the features. LASSO based 54 55 techniques [14-16] are commonly used for linear combination models, while tree-based algorithm 56 [17] are used in non-linear combination models. Wrapper methods use an iterative approach where 57 a model is built using a subset of features in which the performance is evaluated [18, 19]. The 58 process is repeated until the best performance is obtained. It provides better performance than 59 other methods, but it has a higher computational cost.

60 Most techniques have focused on reducing the computational cost of wrapper based methods by 61 designing algorithms that reduce the optimization route to the target feature set q, i.e., using the 62 minimum number of iterations to get q. The studies achieve this objective by focusing on the 63 sampling of feature subset. Feature subset sampling step is commonly performed using either 64 random sampling, sequential sampling or evolutionary sampling [20-23]. The random sampling 65 approach arbitrarily generates the feature subset [20]. The sequential sampling approach adds or 66 removes a feature sequentially from a feature set like forward sampling and backward sampling [18, 67 21]. The evolutionary sampling approach selects the feature subset based on the performance of 68 features in the previous subset like genetic algorithm [22] and swarm optimization [23]. The number 69 of iterations is an important bottleneck in improving the computation efficiency of the wrapper 70 methods.

The wrapper methods assume that feature subset with target features should provide better performance than other feature subsets. Thus, the wrapper methods build models to estimate the performance for evaluation. The need to build a model for every single feature subset obtained in the sampling step creates another critical bottleneck in reducing computational complexity. Our research suggests that model building may not be the only approach to obtain performance value.

Currently, the existing wrapper methods partially or entirely discard the unselected models of feature subset in selecting the next population of feature subsets. Individually, each model may only be useful in providing performance information, but in combination, these models could help in identifying hidden relationships that could help in predicting the performance of unknown feature subset models. This may eliminate the need for building models for every single feature subset obtained in the sampling step. Accordingly, this study focuses on reducing the number of models

that need to be built for a given number of feature subsets obtained in the sampling step of wrapperbased feature selection.

84 In this study, we propose a novel Artificial Intelligence infused wrapper based Feature Selection 85 (AIFS) algorithm. This algorithm can predict the performance of a feature subset using an existing 86 artificial intelligence (AI) model rather than estimates the performance of a feature subset by 87 building an actual AI model (like LASSO, Random Forest). AIFS is unique in many ways. Firstly, it is 88 unique in its perspective as, unlike classical wrapper approaches of building models for every feature 89 subset provided by feature subset sampling step, it builds models for only a fraction of the feature 90 subset. Secondly, it provides a unique application of AI models, that are used to replace the AI 91 model-based performance estimation step with AI model-based performance prediction step, which 92 may reduce the computation time. Thirdly, AIFS is versatile, which allows its integration with existing 93 statistical and machine learning techniques.

This paper provides the "Conceptual Framework" section to explain the basic framework of AIFS. The "Methodology" section explains the AIFS algorithm used in this paper. The algorithm performance is evaluated and compared against the existing feature selection methodologies for simulations and real studies in the "Simulation Studies" and "Real Studies" sections. Finally, we summarize and provide future directions for research in the "Conclusion and Discussion" section.

99 Results

100 The performance of AIFS is evaluated and compared with standard methods like LASSO, adaptive 101 LASSO, group LASSO, sparse partial least squares, elastic net and adaptive elastic net for both the 102 simulated datasets and real data studies.

103 Simulation Studies

104 We perform simulation studies to evaluate the proposed method and compare its performance with 105 other feature selection methods. The study uses multivariate normal distributions to generate high-

106 dimensional datasets for marginal and interaction models. The regression model, $y = \beta_0 + \sum_{i=1}^{p} \beta_i x_i + \epsilon$ and $y = \beta_0 + \sum_{i=1}^{p} \beta_i x_i + \frac{1}{2} \sum_{i\neq j,i=1,j=1}^{i=p,j=p} \beta_{ij} x_{ij} + \epsilon$ provides the outcome variable of 108 the simulated data for marginal and interaction models, respectively. $\epsilon \sim N(0, \sigma^2)$, $x_i \sim N(0, 1)$ and 109 $\{x_{ij}\}$ represents the pairwise interactions between features $\{(x_1, x_2), (x_1, x_3), \dots, (x_{p-1}, x_p)\}$. In 110 the current study, only two-way interactions are considered for demonstration purposes, but it 111 could be easily extended to higher-order interactions. Correlation is added between the first 15 112 features out of p marginal features using the covariance matrix as given below.

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113 Multiple scenarios are created with the different number of noise features (Table 1). Non-zero β 114 value is assigned only to the true features. The AIFS approach is implemented both with and without 115 a performance-based filter step. The final predictive model from selected features is prepared using 116 either RIDGE regression (AIFS-LR) or non-penalized linear regression (AIFS-LLr). When no 117 performance-based filter step is performed, model obtained from embedded feature selection stage 118 is used as the final predictive model and is referred to as AIFS-L technique.

119 Computation Time estimation

120 We estimate computation time of the AIFS algorithm under different scenarios on a system with 121 processor Intel® Core (TM) i7-8750H CPU@2.20GHz with 16 GB RAM on a Windows 10 64-bit 122 operating system. The computation time is compared with the standard wrapper based approach 123 that did not have the Performance Prediction Model (PPM). Since, standard wrapper (StW) does not 124 have performance-based feature selection step, we compare it with AIFS-L method. Further, we add 125 embedded feature selection step in StW. Thus, any performance difference is only associated with 126 PPM model. Genetic algorithm is used to generate samples in feature subset sampling step with 127 maximum number of iterations fixed to 200. Multiple scenarios are created for the comparative

analysis of two algorithms (Table 1). The training datasets vary from 50-100 samples, while the test datasets contain 500 samples. In each scenario, training samples and test samples are independent samples that came from same distribution. Along with computation time, we evaluated both methods on their ability to select the target features and predictive performance of selected features. F1 score is used to determine the accuracy of selecting target features. Root Mean Square Error (RMSE) from the test data is used to determine the predictive performance of the model obtained from the embedded feature selection step. All the analysis is conducted using R 4.0.3 [24].

135 In both the marginal and interaction models (Table 2), AIFS consumed more time as compared to 136 standard wrapper approach. This is counter intuitive, but this behavior is possible due to the PPM 137 model upgradation step in AIFS. During each upgrade, sample size used for training PPM model 138 increases. The current approach uses random forest to update PPM model and uses LASSO to build 139 the base model. LASSO needs to build the model on a sample size of 50 or 100 but random forest 140 needs to build a PPM model using at least 225 samples (Model 1_I) with sample size increasing 141 during the execution of genetic algorithm.

However, AIFS has a better or at par ability to discriminate between the target and noise features, especially for interaction models as compared to standard wrapper method. Similarly, predictive performance of the features shortlisted from AIFS is better or at par with standard wrapper method, especially for high dimensional data and interaction models. AIFS performance suggests that this methodology framework can be used as an alternative to the standard wrapper framework.

147 AIFS comparison with standard methods

AIFS performance is compared with existing standard penalized regression methods namely LASSO, adaptive LASSO (ALASSO), group LASSO (GLASSO), elastic net (Enet), adaptive elastic net (AEnet) and sparse partial least squares (SPLS) in ten different trials. GLASSO is used only for interaction models. All the analysis is conducted using R 4.0.3 [24]. The standard methods are run using the inbuilt packages in statistical language R. *glmnet* package [25] is used for most methods except GLASSO and SPLS for which *glinternet* [26] and *spls* [27] packages were used. In the case of adaptive models, adaptive weights are obtained from ridge regression [28]. In the case of interaction models, all possible two-way interaction terms were created and entered the model. AIFS is implemented using the algorithm programmed in R.

The AIFS and the standard methods are evaluated on target feature selection and prediction performance. We evaluate the method's ability to discriminate between true and noise features by measuring the selection of true features and rejection of noise features. We use RMSE from the test data as the predictive performance metric.

161 Table 3 shows the feature selection performance of different methods for marginal models. All 162 methods have selected the targeted ten features which means that they can identify the target 163 features in the marginal dataset. However, in most cases, the number of selected features is much 164 higher, indicating that methods also select noise features. Compared to standard methods, the AIFS 165 method selected a similar or lesser number of noise features which suggests that it has better 166 discrimination ability between noise and target features than standard methods. Further, results 167 from Figure 1 indicates better discrimination ability of the AIFS method than the standard methods. 168 It is shown that frequency of selecting a noise feature is consistently lesser than the target features 169 in all methods, but the maximum separation is found only for AIFS method. In addition, the area 170 under curve (AUC) of the features was higher for AIFS method as compared to standard methods. 171 Thus, in the case of marginal datasets, while all methods can identify the target features, AIFS 172 outperforms all other methods with a lesser selection of noise features.

The results from the interaction models reiterate the results of the marginal scenario that the feature selection performance of AIFS is better or at par with the standard methods. Table 4 shows that like marginal models the number of features selected by all methods is more than the number of target features in most cases. This suggests that noise features are selected by all methods, but the number of noise features selected differs with methods. AIFS method selects a similar or lesser number of noise features compared to the standard methods, and results from Figure 2 suggest that AIFS may be selecting a lesser number of noise features compared to other methods. The results show that in low dimensional space, all methods can discriminate between the target and noise features by selecting the target features at a higher frequency as compared to noise features. However, in very high dimensions, only AIFS and GLASSO can perform. AUC performance of different methods also shows better or at par performance of AIFS as it can predict the target and noise features with greater or similar accuracy than other methods.

In AIFS, we used existing classic statistical techniques. The use of statistical techniques could have an important influence on the wrapper method performance [29]. However, a performance comparison between LASSO technique used in AIFS and as a standalone feature selection method clearly showed that AIFS could improve the LASSO performance. The AIFS performance suggests that the proposed methodology could enhance the feature selection performance of the existing statistical techniques by reducing the feature space and increasing the target feature percentage.

191 Table 5 shows the prediction performance of different methods. RMSE performance of the tested 192 methods suggests that AIFS method performs consistently better or at par with the existing 193 methods. In low dimensionality data (2 M, 4 M and 1 I), it is expected that all methods should give 194 similar performance as standard methods are primarily developed for handling low dimensionality 195 data, and results support it. AIFS method can provide better performance even in high dimensional 196 settings (1_M and 3_M) and in the presence of interaction terms (2_I). However, at very high 197 dimensional data (3 I), all methods perform poorly. These findings suggest that the AIFS may 198 provide better or at par prediction performance than existing methods. Overall, the proposed 199 method could expand the capability of existing techniques like non-penalized regression to operate 200 in high-dimensional settings. However, computational intensiveness will be a significant limitation 201 for the proposed methodology compared to standard methods. In summary, when we compare the 202 performance of FS methods across different data dimensionality, performance of all methods deteriorates with an increase in data dimensionality, but performance of most standard methods
 decreases more drastically than AIFS.

205 Real Studies: Population Health Data

206 Four real studies are analyzed to evaluate the performance of AIFS and existing methods. 207 Community Health Status Indicators (CHSI) study focuses on non-communicable diseases from US 208 county with data (n=3141) containing 578 features [30] (Study I). National Social Life, Health and 209 Aging Project (NSHAP) datasets focusing on the health and well-being of aged Americans contains 210 multiple datasets. We chose two datasets (Study II and Study III) containing data for 4377 residents 211 on 1470 features [31] and 3005 residents on 820 features [32]. Study IV is the Study of Women's 212 Health Across the Nation (SWAN), 2006-2008 dataset focusing on 887 physical, biological, 213 psychological and social features in middle-aged women in the USA (n = 2245) [33].

The raw data of the real studies are processed for ease of analysis to obtain final cleaned datasets (Table 6). Features and samples are filtered to remove highly correlated features, non-continuous features, and missing values. Then, each dataset is randomly split into training and testing datasets. As the sample size is large, only 20% of data is used for training while remaining 80% of data is used for testing to create a high dimensional data setting. We compare the performance of different methods for marginal models and interaction models using mean RMSE of the test data in ten trials.

Table 7 summarizes the feature selection results. It is shown that standard methods are selecting a lesser number of features as compared to AIFS methods. However, the results from the previous simulated data studies suggest that standard methods may struggle to discriminate between target and noise features (Figure 1 and Figure 2). Further, the predictive performance results of AIFS method is better than the standard methods for both marginal as well as interaction models (Table 8). The better performance of the proposed method suggests that it may be more reliable than standard methods in identifying the target features.

The results show that in Study III, marginal models performed better than their interaction models for all methods. Better performance of the marginal model compared to the interaction model suggests that AIFS cannot completely reject noise features and is sensitive to an increase in feature space. However, AIFS is still more robust than standard methods and can perform in different dimensions and datasets.

232 Real Studies: Genomic Data

233 AIFS-L method is compared with StW method in the genomic datasets to determine the biological 234 relevance of the solutions obtained from AIFS method. In many cancer studies, it is found that 235 smoking can be detrimental to the cancer patient health [34, 35]. Further, an association between 236 gene expression levels and cancer patient smoking habit has been reported [36]. Thus, it would be 237 relevant to identify the genes in cancer patients which are associated with smoking-related traits. In 238 this study, The Cancer Genomic Atlas (TCGA) program is used to get the data from nine cancer 239 projects (Table 9) which maintained records related to amount smoked and gene expression profile 240 of patients [37]. The sample size n for these projects range from 89 to 592 samples with feature 241 space p of 56602 genes. The gene expression profile is used as the input feature space and number 242 of cigarettes smoked per day (CPD) is used as the outcome.

243 Preliminary processing of all datasets is performed to reduce the input feature space and remove 244 samples with missing values. The input feature space is reduced from 56602 to 50 features through 245 multi-stage processing (Table 9). Step one involved removing the features which are not 246 differentially expressed in cancer patients as compared to normal patients using TCGAbiolinks package [38]. Step two involved supervised dimensionality reduction of the differentially expressed 247 248 genes using partial least squares technique and select top 100 features with highest absolute 249 weights in first latent feature. Step three involved removing correlations among the features. Thus, 250 among any pair of features with more than 0.8 absolute correlation, one feature is randomly 251 selected. Step four involves selecting the top 50 features among the non-correlated features based on their absolute weight in the first latent feature obtained in step two. No interaction effects are
 considered for this analysis.

254 The performance of AIFS and StW in all datasets is compared on three metrics namely predictive 255 performance, computation time and number of genes selected. The results are based on 10-fold 256 cross-validation (Table 10). It observed that in all the datasets the predictive performance of AIFS 257 based features is better or at par with StW based features. Further, it is observed that a smaller set 258 of features are selected by AIFS as compared to StW which suggests AIFS could provide a more 259 parsimonious set of features as compared to StW without compromising on the predictive 260 performance of the features. In terms of computation time, the results are similar to those observed 261 in simulation studies with StW taking less time than AIFS in most cases.

262 In order to assess the biological relevance of the genes selected by each method, selected genes of 263 each dataset are pooled together to create final list of genes selected by each method. The results 264 show that some genes are selected at a very high frequency in dataset during 10-fold feature 265 selection process. Genes need to fulfill one of the two criteria of either having highest selection 266 frequency or selection frequency of more than 80%. Accordingly, across nine datasets, AIFS provided 267 13 genes while StW provided 40 genes. 11 genes (VCX3A, WNT3A, CALHM5, ZMYND10, FOXE1, PLAT, 268 BAAT, WFDC5, CGB5, FADD, APOE) are found to be common across the two methods. Among the 13 269 genes from AIFS method, seven genes (WNT3A [39], TMEM45A [40], BAAT [40], WFDC5 [41], HS3ST5 270 [42], CGB5 and APOE [43]) have been reported in literature to exert influence on tobacco or 271 smoking-related traits. Further, AIFS identified six new genes (VCX3A, CALHM5, ZMYND10, FOXE1, 272 PLAT, FADD) which could be related to smoking in cancer patients, thus providing an opportunity for 273 identifying previously unknown biological functions.

274 Discussion

Building models for each sample feature set obtained during the feature sampling stage of wrapper methods consume computational resources and may not always provide the best results. AIFS allows skipping the model building for many sample feature sets by training an AI model, i.e., the PPM model, which could predict the performance of sample feature sets. AIFS feature selection performance and predictive performance are better or at par than both the standard wrapper approach and penalized standard methods, namely LASSO, adaptive LASSO, group LASSO, Sparse PLS, Elastic net and adaptive elastic net.

282 The proposed method has certain limitations. The current study primarily focuses on testing the 283 concept; thus, the study performed testing on limited datatypes. Future research could focus on 284 evaluating the robustness of the approach using different types of data such as temporal data and 285 categorical data, and outcomes such as binary outcomes and time to event outcomes. Other than 286 data types, the focus could also be directed towards the algorithm used. Currently, the study uses a 287 linear combination function for building actual models, but future studies could also explore the 288 non-linear combination function for model building. Further, the current study reduced the need to 289 build actual models in the wrapper approach but could not eliminate it. Therefore, future research 290 could use other PPM building techniques like an artificial neural network and support vector 291 machines to eliminate the need for actual models.

292 Conclusion

293 In the paper, we propose AIFS, an innovative approach to perform wrapper based feature selection. 294 The method is flexible enough to work with both marginal and interaction terms. The approach 295 could be easily embedded with any of the wrapper techniques as it does not alter existing methods, 296 which allows users to integrate the method in their existing wrapper pipelines. This approach could 297 enhance the performance of existing wrapper techniques available in the literature for high dimensional datasets by accelerating the algorithm. AIFS can identify both the marginal features and
interaction terms without using interaction terms in PPM, which could be critical in reducing the
feature space an algorithm has to process.

The benefits of AIFS comes from using artificial intelligence to learn the dataset performance behavior and build the PPM, which replaces the actual model building process. The studies involving marginal effects with and without interaction effects in simulated data showed that AIFS could outperform existing methods in feature selection and prediction performance. Similar performance in real datasets also demonstrates the practical relevance of AIFS.

306 Conceptual Framework

307 In a wrapper approach, given a dataset D of sample size n with p feature space and outcome y, a 308 subset feature set q is created from p. In the standard wrapper approach (Figure 3a), a model is built 309 for the subset of D containing q features and performance is estimated. This performance is used to 310 select the next subset of p. This dependence of a standard wrapper approach upon model building 311 step for each subset of feature to estimate its performance is targeted in our AIFS algorithm.

312 The conceptual framework used to design AIFS algorithm (Figure 3b) aims at reducing (or removing) 313 the dependence of the wrapper algorithm on model building step for obtaining performance value of q. AIFS algorithm creates a random set $q_{AI} = \left\{ q_{AI_j} \right\} | q_{AI_j} \in \left\{ \{1\}, \dots, \{1, \dots, p\} \right\}, j \in \{1, \dots, k\}$ of 314 315 k feature samples, where each feature sample is a subset of p. The algorithm builds a model for q_{AI} 316 samples to estimate their performance $C = \{C_i\}$. The algorithm creates a Performance Prediction 317 Model (PPM) with q_{AI} as the input and C as the outcome using a machine learning model to enable 318 performance prediction of any subset of p. Finally, the algorithm executes the standard wrapper 319 approach, but uses PPM as a surrogate to the actual model building step that predicts rather than 320 estimates the actual performance of q.

321 Methodology

This section explains the design of AIFS algorithm based on the conceptual framework. The algorithm can be divided into four steps: performance prediction model, wrapper based coarse feature selection, embedded-feature selection and performance-based feature selection (Figure 4).

325 Performance Prediction Model (PPM)

The algorithm generates k random sample datasets containing q_{AI_j} features, and sample size n from D. A set of models $M = \{m_j\}$ are created from k sample datasets for an outcome, y using any modeling technique.

$$m_j: y_j = f\left(q_{AI_j}\right) \mid j \in \{1, \dots, k\} \# (2)$$

329 A performance set $C = \{C_j\}$ contains the performance of M models. The algorithm creates a 330 performance dataset D_{perf} , a matrix of features used in each model of $M(q_f)$ and their 331 performance, C.

$$D_{perf} = |q_{f_{ij}} c_j||q_{f_{ij}} = \begin{cases} 0, & q_{AI_{ij}} \notin \{m_j\}, i \in \{1, p\}, j \in \{1, \dots, k\} \\ 1, & q_{AI_{ij}} \in \{m_j\}, i \in \{1, p\}, j \in \{1, \dots, k\} \end{cases} \#(3)$$

As shown in equation 3, feature matrix (q_f) is a binary matrix that consists of p columns and k rows. The matrix takes the value of 0 for i^{th} column and j^{th} row, if i^{th} feature is not used in m_j model, else i^{th} column and j^{th} row takes the value of 1. PPM is constructed from D_{perf} to provide a predictive model for the outcome, C using any machine learning technique.

$$PPM: C = f(q_f) \# (4)$$

In this study, we have used LASSO to prepare m_j models and random forest to build the PPM. During the preliminary analysis (Additional File 1), it is found that predicted performance and actual performance is strongly and positively correlated, but predicted performance may not match the actual performance, as a result subset corresponding to best predicted performance may not be thebest subset.

341 Wrapper based coarse feature selection

342 The standard wrapper approach as shown in Figure 3a is an iterative process where a subset of 343 feature is evaluated, and performance of the feature subset is used to select the next subset of 344 features. In our work, we used genetic algorithm to search through the feature space iteratively as it 345 is used in wide range of datasets [44–46]. In the proposed algorithm, we use PPM for all iterations to 346 predict the performance C_{pred} of a feature set q. Since, we found that best C_{pred} may correspond to 347 one of the high performing feature sets but not the best feature set, we validate C_{pred} values by 348 building a model using q features to estimate the performance C_{true} (Figure 4). The algorithm uses 349 user-defined criteria val_{crit} to select sample feature sets for validation of C_{pred} values.

In this study, the top quartile of C is used as the val_{crit} criterion, thus q with C_{pred} in top quartile of C are selected for model building. D_{perf} is updated with feature set q whose C_{true} value is available and consequently, is used to update PPM. The iteration stops when we get q_{wrap} features, which provide the best performance.

354 Embedded feature selection

The q_{wrap} features obtained from the wrapper step are processed to obtain the final features because the prediction model does not explicitly provide the non-linear combinations of q_{wrap} features. Thus, an embedded feature selection model is used on q_{wrap} features for an outcome, y which allows the additional features χ like interactions terms to be incorporated. LASSO framework is used as the embedded model in the proposed algorithm.

360 Performance-based feature selection

The features selected from the embedded model q_{embed} undergo the last stage of processing to provide final features q. This step selects features based on their contribution to the model performance. l models m_{perf_l} : $y_j = f(q_{embed} - l) | l \in \{1, ..., q_{embed}\}$ are prepared with each model containing $q_{embed} - 1$ features. l feature importance is determined from the m_{perf_l} performance.

To obtain *l* feature robust importance, we create multiple models using bootstrapping of samples, and their performance \hat{c}^{j} is pooled to get overall model performance $\hat{c}_{pool_{j}}$. In this study, we use RIDGE regression for model building as we are focusing on high dimensional data and non-penalized linear regression could only work for cases with $q_{embed} < n$. Goodness of fit (R^{2}) of out of the bag (OOB) samples is used as the performance metric. Finally, the performance metric is pooled to provide a coefficient of variation of R^{2} as the overall model performance for *l* feature.

A performance threshold c_{cutoff} needs to be defined to select the features. Rather than using an arbitrary threshold, our algorithm uses a dynamic cutoff. The algorithm tries different performance thresholds and selects the threshold which provides the best performance c_{best} for the smallest feature space q_{best} . In the current study, we use genetic algorithm to search through the performance threshold space. Two different techniques, namely non-penalized regression and adaptive RIDGE regression are used for the model building. Pseudo Algorithm summarizes the complete AIFS algorithm.

Pseudo Algorithm: AIFS

Input: Feature data X (p × n) Target feature Y (1 × n) Number of feature samples k PPM performance prediction validation criteria val_{crit} Number of bootstrap replicates B Performance dataset $D_{perf} = \{empty\}$ Wrapper based coarse selected features list $q_{wrap} = \{empty\}$ Embedded method based selected features list $q_{embed} = \{empty\}$ Output: Final Feature set q_{best} **Begin:** # Step I: Performance Prediction Model for i=1 to k do Generate q_{AI}^i random features from p Generate samples $(X^i, Y^i \in R^{n \times (q_{AI}^i + 1)})$

```
Build embedded model (like LASSO) from (X^i, Y^i)
   Compute performance estimate C^i of the model
  Add (q_{AI}^i, C^i) to D_{perf}
end for
Build a supervised machine learning model, PPM from D_{perf}
# Step II: Wrapper based Coarse Feature Selection
Initialize a random sample feature set q
while C^q < C^{best} do
     Predict q performance using PPM
      if q fulfils val<sub>crit</sub>
            Build embedded model (like LASSO) from (X^q, Y^q \in \mathbb{R}^{n \times (q+1)})
            Compute performance estimate C^{q} of the model
             if C^q = C^{best}
                  a^{wrap} = a
                  end while
             else
                   Add (q, C^q) to D_{perf}
                   Update PPM from D_{perf}
# Step III: Embedded Feature Selection
Compute embedded model (like LASSO) estimate \widehat{w}_{efs} from (X, Y \in \mathbb{R}^{n \times (q_{wrap}+1)})
for j=1 to q_{wrap}
   if \widehat{w}_{efs}^j \neq 0
      Add j to q_{embed} feature list
end for
Add missing marginal features for selected interaction terms in q_{embed} to get final feature selection
# Step IV: Performance-based feature selection
for i=1 to q_{embed}
   Select all q_{embed} features q except i feature and its interaction terms
  for j=1 to B
          Compute statistical model (like RIDGE) performance \hat{c}^{j} from (X, Y \in \mathbb{R}^{n \times (q+1)})
   end for
   Compute pooled performance estimate \hat{c}_{pool_i}
   Rank q_{embed} such that feature with highest \hat{c}_{pool_i} is considered best feature
end for
Initialize random performance cut off value c_{cutoff}
while c_{model} < c_{best} do
     Select q features such that \hat{c}_{pool_q} \ge c_{cutoff}
     Compute statistical model (like RIDGE and linear regression) performance c_{model} from
(X, Y \in \mathbb{R}^{n \times (q+1)})
end while
End
```

380 List of abbreviations

- 381 AEnet: Adaptive Elastic Net
- 382 AI: Artificial Intelligence
- 383 AIFS: Artificial Intelligence infused wrapper based Feature Selection
- 384 ALASSO: Adaptive LASSO
- 385 AUC: Area Under Curve
- 386 CHSI: Community Health Status Indicators
- 387 Enet: Elastic Net
- 388 GLASSO: Group LASSO
- 389 NSHAP: National Social Life, Health and Aging Project
- 390 OOB: Out Of the Bag
- 391 PPM: Performance Prediction Model
- 392 RMSE: Root Mean Square Error
- 393 SPLS: Sparse Partial Least Squares
- 394 StW: Standard Wrapper
- 395 SWAN: Study of Women's Health Across the Nation

396 **Declarations**

- 397 Ethics approval and consent to participate
- 398 Not Applicable

- 399 Consent for publication
- 400 Not Applicable
- 401 Availability of data and materials
- 402 All the datasets and code are in the github link: https://github.com/rahijaingithub/AIFS.
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- 404 The authors declare that they have no competing interests
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- 526
- 527

| | | | | Sample | Size (n) | - |
|--------------|----------|---|-----|--------|----------|------|
| Models | Scenario | $oldsymbol{eta}$ (Non-Zero coefficients) | р | Train | Test | σ |
| | 1_M | | 50 | 50 | 500 | 0.25 |
| Marginal | 2_M | $\{\beta_i i = \{1,, 10\}\} =$ | 50 | 100 | 500 | 0.25 |
| Waigina | 3_M | $\{0.5, -0.5, 0.5, -0.5,, 0.5\}$ | 100 | 75 | 500 | 0.25 |
| | 4_M | | 100 | 100 | 500 | 0.25 |
| | 1_ | | 15 | 100 | 500 | 0.25 |
| Interactions | 2_ | $\{\beta_i, \beta_{ij} \mid i = \{1,, 10\}, j = i + 1, j < 11\} = \{0.5, -0.5, 0.5, -0.5,, 0.5\}$ | 25 | 100 | 500 | 0.25 |
| | 3_ | [0.0, 0.0,0.0, 0.0,, 0.0] | 50 | 100 | 500 | 0.25 |

Table 1: Description of the simulation data

| 534 | | | | | performa | nce | | | |
|-----|-------|-----|-----|----------------------------|----------|-------|-------------------------------|----------------------------------|--------|
| | | | | | | Perfo | rmance | | |
| | Model | р | n | Computation Time (minutes) | | • | eature Selection F1 Score) | Predictive Performance (RMSE) | |
| | | | | StW | AIFS-L | StW | AIFS-L | StW | AIFS-L |
| | 1_M | 50 | 50 | 7.57 | 24.85 | 0.48 | 0.47 | 0.55 | 0.43 |
| | 2_M | 50 | 100 | 10.68 | 23.93 | 0.71 | 0.63 | 0.29 | 0.29 |
| | 3_M | 100 | 75 | 11.30 | 18.22 | 0.29 | 0.33 | 0.64 | 0.48 |
| | 4_M | 100 | 100 | 31.52 | 33.07 | 0.42 | 0.43 | 0.36 | 0.36 |
| | 1_ | 15 | 50 | 0.97 | 2.18 | 0.41 | 0.73 | 1.20 | 0.38 |
| | 2_ | 25 | 50 | 2.72 | 7.23 | 0.26 | 0.39 | 1.32 | 0.49 |

533 Table 2: Wrapper methods comparison of computation time, target feature selection and predictive

535

537 Table 3: Feature selection performance of different approaches in simulated scenarios for marginal

538

models

| | | | | Ex | isting Mod | | AIFS | | | |
|----------|--|-----------------|---------|---------|------------|---------|---------|---------|----------|---------|
| Scenario | Performance (Number of Features Selected) | Target Features | ALASSO | LASSO | SPLS | Enet | AEnet | AIFS-L | AIFS-LLr | AIFS-LR |
| Š | 5 E R X | Ë | | | | Mean (| | | | |
| 1 1.4 | Margina | 10 | 24 | 25 | 23 | 27 | 26 | 29 | 15 | 12 |
| 1_M | (p=50) | | (18-32) | (18-37) | (14-35) | (18-36) | (21-30) | (24-33) | (11-22) | (10-16) |
| 2 M | Margina | 10 | 16 | 23 | 16 | 25 | 18 | 24 | 16 | 12 |
| 2_M | (p=50) | 10 | (11-35) | (14-40) | (10-39) | (14-41) | (11-35) | (19-31) | (10-31) | (10-16) |
| 2 M | Margina | 10 | 27 | 32 | 25 | 32 | 28 | 44 | 18 | 14 |
| 3_M | (p=100) | 10 | (20-39) | (16-57) | (12-50) | (21-45) | (20-43) | (29-59) | (10-26) | (10-21) |
| 4 14 | Margina | 10 | 28 | 33 | 19 | 32 | 30 | 44 | 19 | 13 |
| 4_M | (p=100) | TO | (14-46) | (14-55) | (11-47) | (17-55) | (15-48) | (34-51) | (10-45) | (10-22) |

539

541 Table 4: Feature selection performance of different approaches in simulated scenarios for interaction

542

models

| | | | Existing Models | | | | | | | AIFS | | | |
|----------|--|-----------------|-----------------|---------------|---------------|-----------------|---------------|---------------|---------------|---------------|---------------|--|--|
| Scenario | Performance (Number of Features Selected) | Target Features | ALASSO | GLASSO | LASSO | SIdS | Enet | AEnet | AIFS-L | AIFS-LLr | AIFS-LR | | |
| S S | 2 C F S | Та | | | | Ме | an (Range) | 1 | | | | | |
| | Marginal (p=15) | 10 | 15 (15-15) | 15 (14-15) | 15 (15-15) | 14 (12-15) | 15 (15-15) | 15 (15-15) | 12 (12-14) | 12 (12-14) | 12 (12-14) | | |
| 1_ | Interaction (χ=105) | 9 | 31 (20-41) | 40 (22-51) | 33 (18-49) | 36 (16-102) | 34 (21-44) | 32 (24-41) | 34 (20-47) | 30 (8-44) | 34 (20-47) | | |
| | Margina (p=25) | 10 | 24 (22-25) | 25 (24-25) | 24 (22-25) | 19 (9-25) | 22 (14-25) | 24 (22-25) | 18 (14-21) | 16 (10-20) | 18 (14-21) | | |
| 2_ | Interaction (χ =300) | 9 | 46 (32-67) | 66 (39-74) | 45 (30-65) | 65 (6-287) | 39 (11-60) | 44 (31-64) | 50 (26-60) | 36 (5-47) | 50 (26-60) | | |
| 2 1 | Marginal (p=50) | 10 | 32 (2-45) | 47 (45-49) | 16 (1-45) | 38 (6-50) | 29 (2-50) | 37 (2-49) | 29 (27-32) | 24 (8-30) | 28 (24-30) | | |
| 3_ | Interaction (χ =1225) | 9 | 36 (1-67) | 76 (72-81) | 16 (0-71) | 417 (1-1057) | 36 (1-116) | 53 (1-104) | 85 (71-99) | 30 (2-52) | 46 (26-88) | | |

543

545 Table 5: Outcome prediction performance of different approaches in simulated scenarios for the test

546

dataset

| | | | Perf | ormance (RMSE) | | | | | | |
|-----------|-------------|-------------|-----------------|------------------|-------------|-----------------------------|------------|--|--|--|
| Math a da | | Marginal N | lodel Scenarios | | Interac | Interaction Model Scenarios | | | | |
| Methods | 1_M | 2_M | 3_M | 4_M | 1_1 | 2_1 | 3_1 | | | |
| | | | Mean (95) | % Confidence Int | erval) | | | | | |
| ALASSO | 0.44 | 0.28 | 0.39 | 0.30 | 0.44 | 0.94 | 1.36 | | | |
| ALASSU | (0.35-0.54) | (0.23-0.33) | (0.32-0.46) | (0.26-0.35) | (0.36-0.52) | (0.74-1.13) | (1.31-1.41 | | | |
| GLASSO | | | | | 0.36 | 0.65 | 1.20 | | | |
| GLASSO | | | | | (0.3-0.43) | (0.51-0.80) | (1.15-1.26 | | | |
| LASSO | 0.45 | 0.29 | 0.40 | 0.31 | 0.40 | 0.94 | 1.36 | | | |
| LASSO | (0.36-0.54) | (0.24-0.34) | (0.33-0.47) | (0.26-0.36) | (0.33-0.47) | (0.76-1.13) | (1.32-1.40 | | | |
| SPLS | 0.45 | 0.26 | 0.43 | 0.27 | 0.52 | 1.33 | 1.47 | | | |
| SPLS | (0.35-0.55) | (0.21-0.31) | (0.28-0.58) | (0.23-0.31) | (0.38-0.66) | (1.21-1.45) | (1.38-1.56 | | | |
| Enet | 0.45 | 0.29 | 0.42 | 0.32 | 0.41 | 1.02 | 1.34 | | | |
| Enet | (0.36-0.53) | (0.24-0.35) | (0.34-0.5) | (0.27-0.36) | (0.34-0.49) | (0.82-1.22) | (1.29-1.38 | | | |
| AEnet | 0.46 | 0.28 | 0.41 | 0.31 | 0.46 | 0.97 | 1.34 | | | |
| AEIIet | (0.35-0.57) | (0.23-0.33) | (0.33-0.48) | (0.26-0.35) | (0.38-0.54) | (0.79-1.15) | (1.30-1.39 | | | |
| | 0.51 | 0.28 | 0.43 | 0.31 | 0.36 | 0.50 | 1.43 | | | |
| AIFS-L | (0.38-0.65) | (0.23-0.32) | (0.34-0.52) | (0.26-0.36) | (0.29-0.43) | (0.40-0.61) | (1.30-1.57 | | | |
| AIFS-LLr | 0.41 | 0.26 | 0.33 | 0.27 | 0.39 | 0.58 | 1.44 | | | |
| AII" J-LU | (0.26-0.56) | (0.21-0.31) | (0.27-0.39) | (0.22-0.32) | (0.31-0.48) | (0.39-0.77) | (1.33-1.55 | | | |
| AIFS-LR | 0.46 | 0.30 | 0.34 | 0.29 | 0.56 | 0.79 | 1.35 | | | |
| AIF S-LK | (0.33-0.58) | (0.26-0.33) | (0.30-0.38) | (0.26-0.33) | (0.48-0.65) | (0.68-0.91) | (1.28-1.41 | | | |

547

Table 6: Summary of the real datasets

| Real | Marginal | Outcome feature | San | Sample size (n) | | | |
|-----------|--------------|------------------------------|-------|-----------------|------|--|--|
| Studies | Features (p) | Outcome leature | Total | Train | Test | | |
| Study I | 44 | Percentage of unhealthy days | 1471 | 294 | 1177 | | |
| Study II | 19 | Height | 1287 | 257 | 1030 | | |
| Study III | 33 | Height | 943 | 189 | 754 | | |
| Study IV | 26 | Body Mass Index | 1406 | 281 | 1125 | | |

| | Ges | | Exis | ting Mode | els | | | AIFS | | | | | |
|--------------|---|-----------------|--------------|----------------|---------|-----------|---------|---------|-----------|--|--|--|--|
| Real Studies | Performance (Number of Features Selected) | ALASSO | GLASSO | LASSO | SPLS | Enet | AEnet | AIFS-L | AIFS -LLr | | | | |
| - | d un | | Mean (Range) | | | | | | | | | | |
| | 2 | Marginal Models | | | | | | | | | | | |
| | Margina | 7 | | 7 | 23 | 13 | 11 | 13 | 10 | | | | |
| | (p=44) | (4-14) | | (3-16) | (3-44) | (4-22) | (4-21) | (7-21) | (5-16) | | | | |
| | Margina | 5 | | 7 | 9 | 8 | 7 | 9 | 6 | | | | |
| | (p=19) | (1-10) | | (1-12) | (1-15) | (1-15) | (1-12) | (4-13) | (3-9) | | | | |
| | Margina | 8 | | 12 | 11 | 13 | 10 | 13 | 9 | | | | |
| | (p=33) | (4-11) | | (6-16) | (4-33) | (5-18) | (4-18) | (10-18) | (4-13) | | | | |
| IV | Margina | 6 | | 7 | 7 | 8 | 7 | 7 | 5 | | | | |
| · · · | (p=26) | (5-7) | | (5-9) | (5-14) | (5-11) | (5-12) | (5-9) | (3-9) | | | | |
| | | | | | | on Models | | | | | | | |
| | Margina | 13 | 42 | 12 | 12 | 22 | 21 | 21 | 20 | | | | |
| | (p=44) | (7-24) | (41-43) | (7-23) | (3-44) | (10-36) | (7-32) | (15-26) | (14-26) | | | | |
| | Interaction | 4 | 170 | 4 | 63 | 13 | 11 | 23 | 17 | | | | |
| | (χ = 946) | (1-11) | (156-183) | (0-11) | (0-591) | (1-46) | (0-23) | (8-47) | (5-35) | | | | |
| | Margina | 10 | 19 | 9 | 11 | 9 | 10 | 12 | 10 | | | | |
| | (p=19) | (2-18) | (19-19) | (1-16) | (1-19) | (1-15) | (1-16) | (9-15) | (1-14) | | | | |
| | Interaction | 6 | 94 | 4 | 24 | 6 | 6 | 15 | 8 | | | | |
| | (χ =171) | (0-19) | (87-108) | (0-8) | (0-117) | (0-21) | (0-14) | (5-37) | (0-13) | | | | |
| | Marginal | 15 | 33 | 15 | 4 | 14 | 16 | 16 | 15 | | | | |
| | (p=33) | (6-26) | (32-33) | (3-23) | (1-10) | (4-23) | (10-23) | (10-21) | (2-21) | | | | |
| | Interaction | 6 | 125 | 5 | 1 | 4 | 5 | 22 | 19 | | | | |
| | (χ =528) | (1-25) | (113-137) | (0-16) | (0-4) | (0-16) | (1-15) | (1-49) | (1-49) | | | | |
| | Marginal | 5 | 7 | 6 | 9 | 7 | 5 | 10 | 10 | | | | |
| IV | (p=26) | (3-6) | (5-9) | (3-9) | (6-12) | (4-10) | (3-6) | (6-13) | (6-13) | | | | |
| | Interaction | 3 | 7 | 4 | 12 | 5 | 3 | 13 | 13 | | | | |
| | (χ =299) | (1-4) | (5-10) | (2-6) | (7-16) | (2-7) | (1-5) | (7-26) | (7-26) | | | | |

Table 7: Number of features selected by different wrapper methods on the real studies

553

552

Table 8: RMSE performance of different methods on the real studies for test data

| | | | nce (RMSE) | |
|------------|-------------|---------------|-------------------|-------------|
| Methods | | - | odel Scenarios | |
| | 1 | <u> </u> | <u> </u> | IV |
| | | • | fidence Interval) | |
| ALASSO | 0.95 | 3.76 | 3.08 | 0.86 |
| | (0.95-0.96) | (3.67-3.84) | (3.01-3.14) | (0.81-0.90) |
| LASSO | 0.96 | 3.75 | 3.10 | 0.84 |
| 21000 | (0.95-0.97) | (3.65-3.85) | (3.03-3.16) | (0.8-0.87) |
| SPLS | 0.97 | 3.61 | 3.35 | 0.77 |
| 51 65 | (0.95-0.99) | (3.54-3.69) | (3.03-3.66) | (0.76-0.79) |
| Enet | 0.95 | 3.79 | 3.15 | 0.85 |
| Linet | (0.94-0.96) | (3.7-3.87) | (3.08-3.23) | (0.81-0.90) |
| AEnet | 0.96 | 3.76 | 3.11 | 0.84 |
| ALIIEL | (0.94-0.97) | (3.67-3.85) | (3.07-3.15) | (0.8-0.87) |
| AUEC 1 | 0.94 | 3.65 | 3.02 | 0.83 |
| AIFS-L | (0.93-0.94) | (3.59-3.71) | (2.98-3.06) | (0.8-0.86) |
| | 0.96 | 3.59 | 2.97 | 0.75 |
| AIFS-LLr | (0.94-0.97) | (3.55-3.64) | (2.91-3.03) | (0.73-0.78) |
| | 0.95 | 3.80 | 3.19 | 1.20 |
| AIFS-LR | (0.94-0.96) | (3.72-3.87) | (3.11-3.28) | (1.17-1.24) |
| | | | odel Scenarios | . , |
| Methods | 1 | 11 | Ш | IV |
| | | Mean (95% Con | fidence Interval) | |
| ALASSO | 0.94 | 3.69 | 3.12 | 0.52 |
| ALASSU | (0.93-0.95) | (3.61-3.76) | (3.02-3.23) | (0.49-0.55) |
| CLASSO | 1.44 | 4.46 | 8.24 | 0.31 |
| GLASSO | (1.2-1.68) | (4.35-4.57) | (5.37-11.11) | (0.28-0.34) |
| 14660 | 0.95 | 3.74 | 3.15 | 0.43 |
| LASSO | (0.94-0.96) | (3.67-3.81) | (3.02-3.27) | (0.39-0.47) |
| 6016 | 1.03 | 3.81 | 4.34 | 0.24 |
| SPLS | (0.91-1.15) | (3.76-3.86) | (3.26-5.42) | (0.22-0.26) |
| - . | 0.94 | 3.78 | 3.24 | 0.44 |
| Enet | (0.93-0.95) | (3.72-3.84) | (3.13-3.34) | (0.4-0.48) |
| | 0.93 | 3.73 | 3.14 | 0.53 |
| AEnet | (0.92-0.94) | (3.65-3.81) | (3.06-3.21) | (0.5-0.56) |
| | 0.94 | 3.58 | 3.07 | 0.29 |
| AIFS-L | (0.92-0.95) | (3.53-3.63) | (2.98-3.17) | (0.26-0.33) |
| | 1.04 | 3.76 | 3.65 | 0.26 |
| AIFS-LLr | (0.99-1.1) | (3.58-3.93) | (3.26-4.04) | (0.21-0.31) |
| | 0.93 | 3.70 | 3.22 | 1.11 |
| AIFS-LR | (0.92-0.94) | (3.64-3.76) | (3.18-3.26) | (0.99-1.24) |

Table 9: Summary of the genomic datasets

| Datasets | Number of cigarettes smoked per day (μ(σ)) | Sample Size (n) | Feature Space (p) |
|-----------|---|--------------------|----------------------|
| TCGA-BLCA | 1.16 (2.34) | 433 | 56602 |
| TCGA-CESC | 0.30 (0.62) | 307 | 56602 |
| TCGA-ESCA | 0.95 (1.21) | 172 | 56602 |
| TCGA-HNSC | 1.41 (1.89) | 544 | 56602 |
| TCGA-KICH | 0.21 (0.67) | 89 | 56602 |
| TCGA-KIRP | 0.42 (1.04) | 320 | 56602 |
| TCGA-LUAD | 1.53 (1.59) | 592 | 56602 |
| TCGA-LUSC | 2.44 (1.88) | 551 | 56602 |
| TCGA-PAAD | 0.46 (0.88) | 181 | 56602 |

561 Table 10: Wrapper methods comparison of predictive performance, number of genes selected and

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computation time

| | | F | Performance | (μ [95% Cl]) | * | | |
|-----------|------------------|-----------------|-----------------------------|--------------|----------------------------|-----------------|--|
| Dataset | Predictive Perfo | ormance (RMSE) | Number of Genes Selected | | Computation Time (minutes) | | |
| | StW | AIFS-L | StW | AIFS-L | StW | AIFS-L | |
| TCGA-BLCA | 0.79[0.31,1.27] | 0.78[0.30,1.26] | 4[0,9] | 1[0,3] | 5.9[3.2,8.6] | 12.2[10.1,14.3] | |
| TCGA-CESC | 1.00[0.84,1.16] | 0.98[0.84,1.13] | 10[7,13] | 5[4,6] | 11[7.7,14.2] | 14.6[9.9,19.3] | |
| TCGA-ESCA | 1.04[0.87,1.20] | 1.00[0.85,1.15] | 11[5,17] | 8[2,14] | 7.2[4.9,9.5] | 27.9[3.6,52.2] | |
| TCGA-HNSC | 0.99[0.82,1.16] | 0.98[0.81,1.15] | 16[12,20] | 6[3,9] | 11.4[8.7,14] | 20.3[9.3,31.2] | |
| TCGA-KICH | 1.03[0.61,1.46] | 0.82[0.39,1.25] | 11[9,13] | 6[4,8] | 50.2[24.7,75.7] | 10.6[7.5,13.7] | |
| TCGA-KIRP | 0.95[0.66,1.24] | 0.95[0.65,1.24] | 19[18,20] | 15[11,19] | 10.4[8.8,12] | 41.1[12.5,69.8] | |
| TCGA-LUAD | 1.02[0.93,1.11] | 1.02[0.94,1.09] | 25[22,28] | 21[16,26] | 11.6[9.1,14.1] | 42.3[11.6,72.9] | |
| TCGA-LUSC | 0.99[0.91,1.08] | 0.99[0.91,1.08] | 2[1,3] | 1[0,2] | 5.7[4.4,7] | 12[8.8,15.2] | |
| TCGA-PAAD | 1.26[0.74,1.79] | 1.24[0.75,1.73] | 22[20,24] | 14[9,19] | 10.8[7.6,14.1] | 29[0.6,57.4] | |

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- Figure 1: Comparison of different methods' feature selection performance in marginal models a) Frequency of selection of target and noise features. b) AUC for predicting the target and noise features.
- 568 Figure 2: Feature selection performance comparison of different methods in interaction models a)
- 569 Frequency of selection of target and noise features. b) AUC for predicting the target and noise
- 570 features.
- 571 Figure 3: Flow chart of A) Standard wrapper approach and B) Proposed wrapper (AIFS) conceptual 572 approach.
- 573 Figure 4: AIFS algorithm graphical flow chart. Dark Background represents main steps and light
- 574 background represents sub-steps.







