Feature Selection for Huge Data via Minipatch Learning

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Abstract

Feature selection often leads to increased model interpretability, faster computation, and improved model performance by discarding irrelevant or redundant features. While feature selection is a well-studied problem with many widely-used techniques, there are typically two key challenges: i) many existing approaches become computationally intractable in huge-data settings with millions of observations and features; and ii) the statistical accuracy of selected features degrades in high-noise, high-correlation settings, thus hindering reliable model interpretation. We tackle these problems by proposing Stable Minipatch Selection (STAMPS) and Adaptive STAMPS (AdaSTAMPS). These are meta-algorithms that build ensembles of selection events of base feature selectors trained on many tiny, (adaptively-chosen) random subsets of both the observations and features of the data, which we call minipatches. Our approaches are general and can be employed with a variety of existing feature selection strategies and machine learning techniques. In addition, we provide theoretical insights on STAMPS and empirically demonstrate that our approaches, especially AdaSTAMPS, dominate competing methods in terms of feature selection accuracy and computational time.

1 Introduction

Feature selection is critical in statistical machine learning for improving interpretability of machine learning models. While feature selection has been used in a wide variety of applications, there are typically two key challenges: i) many existing techniques can quickly become computationally intractable in huge-data settings on the order of millions of observations and features; and ii) the statistical accuracy of selected features degrades in the presence of noise and/or highly correlated features, thus hampering reliable model interpretation.

In this work, our primary goal is to develop a general feature selection framework that can select statistically accurate features in a computationally efficient manner, even with huge data in high-noise, high-correlation settings. We place great emphasis on selecting the correct features, which is a statistical accuracy problem, because doing so would considerably

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enhance explainability of machine learning methods, enable practitioners to reliably interpret models, and provide insights into the underlying data-generating processes (e.g. candidate gene study (Kohannim et al., 2012)). As stressed in Lipton (2018), interpretability of models is critical in many machine learning applications. Furthermore, we seek to design our feature selection framework to be conducive to efficient computation so that it can be applied in huge-data settings with millions of features and observations. Data of such scale is regularly observed in online marketing, genetics, neuroscience, and text mining, among many other areas of research.

Popular feature selection methods can be divided into three categories - filter, wrapper, and embedded methods (Guyon and Elisseeff, 2003). Filter methods select features based solely on observed characteristics of available data independently of a learning algorithm (Jovic et al., 2015). While filter methods are fast computationally and they might work well for prediction by selecting features related to the output, filter methods usually perform poorly in terms of statistical accuracy of selecting the right features that are critical for reliable model interpretation because they tend to select sets of correlated, redundant features together.

Unlike filter methods, wrapper methods assess the quality of selected features using performance of a predefined learning algorithm. Therefore, wrappers are much slower computationally than filter methods (Jovic et al., 2015; Pirgazi et al., 2019). Despite generally better performance than filter methods, wrappers tend to perform suboptimally in selecting statistically accurate features because they are inherently greedy methods.

Another line of work focuses on embedded methods, which can be further divided into optimization-based methods and methods that select features based on properties of the learner. The Lasso (Tibshirani, 1996) is arguably one of the most widely-used optimization-based feature selection methods. The Lasso tends to do better on statistical accuracy of selected features and is model selection consistent under theoretical conditions including the Irrepresentable Condition (Zhao and Yu, 2006; Meinshausen and Bühlmann, 2006). Yet, such theoretical conditions requiring low correlation among features will be less likely to ever hold in high-dimensional settings in which features tend to become increasingly correlated. Additionally, choosing the right amount of regularization for accurate feature selection is challenging in practice (Meinshausen and Bühlmann, 2010). Fitting the Lasso with model selection procedures (e.g. cross-validation) can be computationally challenging in huge-data settings.

Last but not least, various tree-based algorithms such as Random Forest (RF) (Breiman, 2001) constitute a great proportion of embedded methods that select features using properties of the learner. In particular, numerous importance scores such as the RF permutation importance are used to rank features based on their contributions to improving predictive performance of the learner. However, previous studies have found many such importance measures can be biased and even become unstable in high-dimensional, high-correlation settings (Strobl et al., 2007; Genuer et al., 2010; Nicodemus and Shugart, 2007), thus hindering reliable model interpretation. Additionally, features selected using properties of the learner are often sensitive to the choice of tuning hyperparameters, and systematic hyperparameter search poses great computational challenges in huge-data settings.

The idea of randomly subsampling observations and/or features for model training has appeared in many parts of machine learning, including various ensemble learning techniques.
Breiman, 1996, 2001; Freund and Schapire, 1997; Louppe and Geurts, 2012; Gomes et al., 2019; LeJeune et al., 2020) and the dropout technique (Srivastava et al., 2014) combined with stochastic gradient methods (Hardt et al., 2016) in training deep neural networks. Additionally, there is another line of work focusing on using various data randomization techniques to select more “stable” features for high-dimensional linear regression (Bach, 2008; Meinshausen and Bühlmann, 2010; Wang et al., 2011; Shah and Samworth, 2013; Yu, 2013; Beinrucker et al., 2016; Staerk et al., 2019). Inspired by these approaches, we propose general and flexible strategies that exploit tiny, (adaptively-chosen) random subsets of both the observations and features of the data for considerably enhancing feature selection accuracy and computational efficiency, hence improving model interpretability.

Contributions We summarize our contributions as follows: We develop general feature selection methods named STAMPS and AdaSTAMPS that leverage random data perturbation as well as adaptive feature sampling schemes to select statistically accurate features with dramatically improved robustness and runtime (see Sec. 2). Our proposed approaches are meta-algorithms that can be employed with a variety of existing feature selection strategies and machine learning techniques in practice. In addition, we theoretically show that STAMPS achieves familywise error rate (FWER) control under certain conditions. We empirically demonstrate the effectiveness of our approaches on multiple synthetic and real data sets in Sec. 3, showing that our frameworks, especially AdaSTAMPS, dominate competing methods in terms of both feature selection accuracy and computational time.

2 Minipatch Feature Selection

2.1 Minipatch Learning

People have used random subsets of observations and features for model training in many parts of machine learning (see Sec. 1). Notably in the ensemble learning literature, some have called these “random patches” (Louppe and Geurts, 2012; Gomes et al., 2019). We call tiny, (adaptively-chosen) random subsets of both observations and features “minipatches” to denote the connection with minibatches as well as patches in image processing, and to also denote that these are typically very tiny subsets. Using tiny minipatches can yield major computational advantages for huge data. Formally, given a pair of response vector \( y \in \mathbb{R}^N \) and data matrix \( X \in \mathbb{R}^{N \times M} \) that consists of \( N \) observations each having \( M \) features, a minipatch can be obtained by simultaneously subsampling \( n \) rows (observations) and \( m \) columns (features) without replacement using some form of randomization, with \( n \ll N \) and \( m \ll M \).

2.2 Stable Minipatch Selection (STAMPS)

Inspired by stability selection (Meinshausen and Bühlmann, 2010) and its extensions (Shah and Samworth, 2013; Beinrucker et al., 2016) that keep “stable” features based on selection frequencies, we propose a general and flexible approach to stable feature selection named
Stable Minipatch Selection (STAMPS). STAMPS is a meta-algorithm that can be employed with a variety of existing feature selection strategies and machine learning techniques.

Our proposed STAMPS method is summarized in Algorithm 1. Here, $y_{I_k}$ denotes the subvector of $y$ containing its elements indexed by $I_k$. Similarly, $X_{I_k,F_k}$ denotes the submatrix of $X$ containing its rows indexed by $I_k$ and its columns indexed by $F_k$. For brevity, $[M]$ denotes the set $\{1, 2, \ldots, M\}$.

Specifically, STAMPS fits arbitrary base feature selectors to many tiny, random minipatches and calculates feature selection frequencies by taking an ensemble of estimated feature supports over these minipatches. We define the selection frequency of the $j$th feature at the $k$th iteration, $\hat{\Pi}^{(k)}_j$, to be the number of times it is sampled and then selected by base feature selectors divided by the number of times it is sampled into minipatches after $k$ iterations.

STAMPS finally outputs a set of stable features $\hat{S}_{\text{stable}}$ whose selection frequencies are above a user-specific threshold $\pi_{\text{thr}} \in (0, 1)$. We discuss the choice of $\pi_{\text{thr}}$ in Sec. 2.5.

\textbf{Algorithm 1: STAMPS}

\begin{itemize}
    \item \textbf{Input:} $y \in \mathbb{R}^N$, $X \in \mathbb{R}^{N \times M}$, $n, m, \pi_{\text{thr}} \in (0, 1)$.
    \item \textbf{Initialize:} $\hat{\Pi}^{(0)}_j = 0$, $\forall j \in [M]$;
    \item \textbf{while stopping criterion not met do}
        \begin{enumerate}
            \item Sample a minipatch: subsample $n$ observations $I_k \subset [N]$ and $m$ features $F_k \subset [M]$ uniformly at random without replacement to get a minipatch $(y_{I_k}, X_{I_k,F_k}) \in \mathbb{R}^n \times \mathbb{R}^{n \times m}$;
            \item Fit a selector to minipatch $(y_{I_k}, X_{I_k,F_k})$ to obtain an estimated feature support $\hat{S}_k \subseteq F_k$;
            \item Ensemble feature supports $\{\hat{S}_i\}_{i=1}^k$ to update selection frequencies $\hat{\Pi}^{(k)}_j$, $\forall j \in [M]$:
                \[\hat{\Pi}^{(k)}_j = \frac{1}{\max(1, \sum_{i=1}^k \mathbb{1}(j \in F_i))} \sum_{i=1}^k \mathbb{1}(j \in F_i, j \in \hat{S}_i);\]
        \end{enumerate}
    \item \textbf{end}
    \item \textbf{Output:} $\hat{S}_{\text{stable}} = \{j \in [M] : \hat{\Pi}^{(K)}_j \geq \pi_{\text{thr}}\}$.
\end{itemize}

To avoid a forever STAMPS situation, one should employ some type of stopping criterion for Algorithm 1. While there are many possible choices, we happen to use a simple one that is effective in our empirical studies - stop the algorithm if the rank ordering of the top $\min(\max(|\mathcal{H}|, \tau_l), \tau_u)$ features in terms of selection frequencies $\{\hat{\Pi}^{(k)}_j\}_{j=1}^M$ remain unchanged for the past 100 iterations, where $\mathcal{H} = \{j : \hat{\Pi}^{(k)}_j \geq 0.5\}$. Here, $\tau_l$ and $\tau_u$ are fixed, user-specific parameters. We suggest setting both to well exceed the approximate number of true signal features, with $\tau_u$ being a multiple of $\tau_l$.

\subsection{2.3 Adaptive Stable Minipatch Selection (AdaSTAMPS)}

We propose to adaptively sample minipatches in ways that further boost computational efficiency as well as feature selection accuracy. Replacing uniform feature sampling in step
of Algorithm 1 with any appropriate adaptive feature sampling scheme, we obtain our
Adaptive STAMPS (AdaSTAMPS) framework for feature selection.

While there are many possible adaptive sampling techniques to choose from, we develop
two types of example algorithms in this work. We describe an exploitation and exploration
scheme inspired by the multi-armed bandit literature (Bouneffouf and Rish, 2019; Slivkins,
2019) here in Algorithm 2. For brevity, we use the shorthand AdaSTAMPS (EE) to denote the
specific instance of AdaSTAMPS that employs Algorithm 2 as its adaptive feature sampling
scheme. In addition, we give another probabilistic adaptive feature sampling scheme in the
Appendix, which we abbreviate as AdaSTAMPS (Prob). We use these two in our empirical
studies just to show examples of what can be done with our flexible AdaSTAMPS approach. We
save the investigation of various other adaptive sampling schemes for future work.

In Algorithm 2, \( k \) is the iteration counter, \( E \) denotes the total number of burn-in epochs,
\( \{\gamma^{(k)}\} \) is an increasing geometric sequence in the range \([0.5, 1]\) for controlling the trade-off
between exploitation and exploration of the input feature space, and \( \pi_{\text{active}} \in (0, 1) \) is a fixed,
user-specific threshold for determining the active feature set. We found setting \( \pi_{\text{active}} = 0.1 \)
works well for most problems.

**Algorithm 2: Adaptive Feature Sampling Scheme: Exploitation and Exploration (EE)**

**Input:** \( k, M, m, E, \{\hat{\Pi}^{(k-1)}_j\}_{j=1}^M, \{\gamma^{(k)}\} = \text{geom_seq}([0.5, 1]), \pi_{\text{active}}. \)

**Initialize:** \( G = \lceil \frac{M}{m} \rceil, J = \{1, \ldots, M\}; \)

**if** \( k \leq E \cdot G \) **then** // Burn-in stage

**if** \( \text{mod}_G(k) = 1 \) **then** // A new epoch

1) Randomly reshuffle feature index set \( J \) and partition into disjoint sets
\( \{J_g\}_{g=0}^{G-1}; \)

1) Set \( F_k = J_{\text{mod}_G(k)}; \)

**else** // Adaptive stage

1) Update active feature set: \( A = \{j \in \{1, \ldots, M\} : \hat{\Pi}^{(k-1)}_j \geq \pi_{\text{active}}\}; \)

2) Exploitation step: Sample \( \min(m, \gamma^{(k)}|A|) \) features \( F_{k,1} \subseteq A \) uniformly at random;

3) Exploration step: Sample \( (m - \min(m, \gamma^{(k)}|A|)) \) features \( F_{k,2} \subseteq \{1, \ldots, M\} \setminus A \)
uniformly at random;

4) Set \( F_k = F_{k,1} \cup F_{k,2}; \)

**end**

**Output:** \( F_k. \)

During the initial burn-in stage of Algorithm 2, AdaSTAMPS (EE) explores the entire
input feature space by ensuring each feature is sampled into minipatches for exactly \( E \) times,
one for each epoch. The selection frequency of each feature is continuously updated as feature
selection is repeatedly conducted on each random minipatch. After \( E \) epochs, Algorithm 2
transitions to the adaptive stage during which it exploits a subset of features that are likely
to be true signal features while adaptively exploring the rest of the feature space. Specifically, features with selection frequencies above $\pi_{active}$ are kept in the active feature set $\mathcal{A}$, which is likely a superset of most of the true signal features. As $k$ increases and feature selection frequencies keep polarizing, $\mathcal{A}$ continues to shrink and hone in on the set of true signal features. To exploit the relatively stable features in $\mathcal{A}$, Algorithm 2 samples an increasing proportion of $\mathcal{A}$ into each new minipatch by gradually increasing $\gamma^{(k)}$. Meanwhile, Algorithm 2 continues to explore the rest of the feature space by drawing from $\{1, \ldots, M\} \setminus \mathcal{A}$ into the minipatch, because a tiny fraction of true signal features might still be outside of $\mathcal{A}$ due to noise and correlation among features. Intuitively, the ever-changing active feature set $\mathcal{A}$ and $\{\gamma^{(k)}\}$ work together to adaptively strike a balance between exploitation and exploration of the input feature space, thus further enhancing computational efficiency and feature selection accuracy (e.g. Figure 1).

2.4 Theoretical Properties

Suppose we observe training data $(y, X) \in \mathbb{R}^N \times \mathbb{R}^{N \times M}$ that have $N$ observations and $M$ features. Let $S \subset \{1, \ldots, M\}$ denote the true feature support set, which is the set of indices corresponding to the underlying true signal features. And let $S^c = \{1, \ldots, M\} \setminus S$ be the complement set that contains indices corresponding to the noise features. Both $S$ and $S^c$ are typically unknown. One important goal of feature selection is to infer $S$ from the observed data.

Following the notations in Algorithm 1 an arbitrary minipatch can be represented by the pair of row index set and feature index set $(I_i, F_i)$. The estimated feature support on this arbitrary minipatch by an arbitrary base selector is denoted as $\hat{S}_i \subseteq F_i$. Here, we provide some insights into the theoretical properties of the STAMPS method. The proof is found in the Appendix.

**Assumption (A1)** Each feature $j \in \{1, \ldots, M\}$ is sampled at least once when STAMPS stops after $K$ iterations.

**Assumption (A2)** For a given noise feature $j \in S^c$, $\mathbb{P}(j \in \hat{S}_i)$ is constant across the collection of minipatches $\{(I_i, F_i)\}_{i \in \mathcal{D}}$ that contain this feature, where $\mathcal{D} = \{i \in \{1, 2, \ldots, K\} : j \in F_i\}$.

**Assumption (A3)** $\mathbb{P}\left[j \in \hat{S}_i\right] \leq \alpha \pi_{thr} / |S^c|$ for any noise feature $j \in S^c$ on an arbitrary minipatch $(I_i, F_i)$ that contains this feature.

**Theorem 1** Assume that Assumptions (A1-3) hold. Then STAMPS controls the familywise error rate (FWER) at level $\alpha$:

$$\mathbb{P}[|\hat{S}_{\text{stable}} \cap S^c| \geq 1] \leq \alpha$$

A simple sampling strategy that satisfies Assumption (A1) is to sample non-overlapping blocks of features sequentially such that all features are sampled exactly once before proceeding with uniform random sampling. Assumption (A2) intuitively means that for a given noise feature, the base selector’s decision on whether to select this particular feature does not change across minipatches that contain this feature. In the stability selection literature, Meinshausen and Bühlmann (2010) made use of a similar condition named the exchangeability assumptions.
for false discovery rate (FDR) control. A simple data scenario that fulfills Assumption (A2) is when noise features are independent amongst themselves and are also independent of other true signal features as well as the response. In Assumption (A3), the probability $P$ is taken with respect to the randomness of minipatches and the randomness of the base selector if it is a randomized algorithm. Intuitively, Assumption (A3) requires the base selector to have a low probability to mistakenly select noise features on random minipatches. Putting everything together, under Assumptions (A1-3), Theorem 1 provides guarantee that the probability of STAMPS mistakenly selecting at least one noise feature is at most $\alpha$.

While some of these assumptions may seem stringent, this result is very generic because we are not making any distributional assumptions about the underlying data or considering any specific learner or feature selection algorithm. In future work, we are going to investigate whether these assumptions can be relaxed for particular combinations of learners and selectors. We would also like to consider possible extensions to other forms of error control such as the FDR.

### 2.5 Practical Considerations

![Performance of our approaches on a challenging feature selection task in the regression setting with highly correlated features (as described in the Appendix). (A) Feature selection frequencies by STAMPS versus number of iterations; (B) Feature selection frequencies by AdaSTAMPS (EE) versus number of iterations with the green dotted line denoting the end of the burn-in stage; (C) Feature selection frequencies by AdaSTAMPS (Prob) versus number of iterations. This reveals that well-designed adaptive feature sampling schemes help AdaSTAMPS correctly select all true signal features (red) while discarding all noise features (gray) within much fewer iterations, thus further boosting both feature selection accuracy and computational efficiency in practice.

Our STAMPS and AdaSTAMPS are general meta-algorithms and they have three tuning hyperparameters: minipatch size ($n$ and $m$) and threshold $\pi_{th}$. In general, our methods are quite robust to the choice of these hyperparameters. In our empirical studies in Sec. 3, we use a default threshold $\pi_{th} = 0.5$, which generally works well in practice. We also propose a data-driven way to choose $\pi_{th}$ as detailed in the Appendix. For choosing the minipatch size, we have empirical studies in the Appendix investigating how feature selection accuracy
and runtime vary for various $n$ and $m$ values. In general, we found taking $m$ to well exceed the number of true features (e.g. 3-10 times the number of true features) and then picking $n$ relative to $m$ so that it well exceeds the sample complexity of the feature selector used in step 2 of Algorithm 1 works well in practice. Our empirical results also reveal that our meta-algorithms have stable performance for any sensible range of $n$ and $m$.

We also empirically show that using adaptive feature sampling not only increases computational efficiency but also seems to further improve feature selection accuracy in practice. In Figure 1, we display selection frequencies of all features versus the number of iterations for STAMPS, AdaSTAMPS (EE), and AdaSTAMPS (Prob). We see that both AdaSTAMPS (EE) and AdaSTAMPS (Prob) correctly identify all true signal features while discarding all noise features within 1000 iterations. On the other hand, STAMPS without adaptive feature sampling converges much slower and is unable to clearly separate selection frequencies of signal features from those of noise features. Details of this empirical study are given in the Appendix.

3 Empirical Studies

In this section, we evaluate our proposed STAMPS and AdaSTAMPS approaches on both synthetic and real-world data sets. Because many people study the problem of feature selection in the linear regression setting and it is also easier to simulate synthetic data from linear models for comparisons with existing methods, we first focus on feature selection in the high-dimensional linear regression setting in Sec. 3.1. However, our proposed feature selection frameworks are not limited to either linear or regression settings. We showcase the superior performance of our AdaSTAMPS method on real-world classification problems in Sec. 3.2.

3.1 Feature Selection in High-Dimensional Linear Regression

We compare the feature selection accuracy as well as computational time of our proposed methods with several widely-used competitors including the Lasso (Tibshirani, 1996), Complementary Pairs Stability Selection (CPSS) (Shah and Samworth, 2013), Randomized Lasso (Meinshausen and Bühlmann, 2010), univariate linear regression filter with False Discovery Rate (FDR) control via the Benjamini-Hochberg procedure, recursive feature elimination (RFE) (Guyon et al., 2002), and thresholded Ordinary Least Squares (OLS) (Giurcanu, 2016). Our focus is on high-dimensional cases with $N \ll M$ because these settings are arguably the most difficult for feature selection. In this section, we consider three high-dimensional scenarios with the following data matrices $X \in \mathbb{R}^{N \times M}$:

- **Scenario 1 (S1)** Autoregressive Toeplitz design: the $M = 10000$-dimensional feature vector follows a $\mathcal{N}(0, \Sigma)$ distribution, where $\Sigma_{ij} = \rho^{|i-j|}$ with $\rho = 0.95$; sample size $N = 5000$.

- **Scenario 2 (S2)** Subset of The Cancer Genome Atlas (TCGA) data: this data set ($N = 2834$, $M = 10000$) contains 450K methylation status of the top 10000 most variable CpGs for 2834 patients. This scenario is even more challenging than S1 due to

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strong and complex correlation structure among CpGs. This data set was obtained via TCGA2STAT (Wan et al., 2015).

• Scenario 3 (S3) Fullsize TCGA data: this data set \((N = 2834, M = 335897)\) contains 450K methylation status of 335897 CpGs with complete records for 2834 patients.

Following standard simulation procedures (Meinshausen and Bühlmann, 2010), we create the true feature support set \(S\) by randomly sampling \(S \subset \{1, \ldots, M\}\) with cardinality \(|S| = 20\). The \(M\)-dimensional sparse coefficient vector \(\beta\) is generated by setting \(\beta_j = 0\) for all \(j \notin S\) and \(\beta_j\) is chosen independently and uniformly in \([-3/b, -2/b] \cup [2/b, 3/b]\) for all \(j \in S\). Finally, the response vector \(y \in \mathbb{R}^N\) can be generated by \(y = X\beta + \epsilon\) where the noise vector \((\epsilon_1, \ldots, \epsilon_N)\) is IID \(\mathcal{N}(0, 1)\). For S1 and S2, we vary the positive real number \(b\) to attain various signal-to-noise ratio (SNR) levels. For S3, SNR is fixed at 5.

For each method, commonly-used data-driven model selection procedures are applied to choose tuning hyperparameters and produce a data-driven estimate of the true feature support. In addition, oracle model selection that assumes knowledge of cardinality of the true support \(|S|\) is employed with each method, resulting in an oracle estimate of the true feature support. Detailed descriptions of these data-driven and oracle model selection procedures are given in the Appendix. We evaluate the feature selection accuracy of various methods using the F1 Score, which takes values between 0 and 1 with 1 indicating perfect match between the final estimated feature support and the true feature support \(S\). For every method, we compute F1 Scores for both data-driven and oracle estimates.

We use Scikit-learn (Pedregosa et al., 2011) for all Lasso-based methods, univariate filter, and RFE with Ridge estimator. Due to lack of existing software, we faithfully implemented the thresholded OLS as in Giurcanu (2016). For simplicity, our methods (i.e. STAMPS, AdaSTAMPS (EE), and AdaSTAMPS (Prob)) are employed with the same thresholded OLS as base selector in step 2 of Algorithm 1 using default settings as detailed in the Appendix. All comparisons were run on a VM with 10 Intel (Cascade Lake) vCPUs and 220 GB of memory.

The feature selection accuracy and computational time of various methods from scenarios S1 and S2 are shown in Figure 2. We see that our AdaSTAMPS consistently dominates all competing methods in terms of feature selection accuracy in both scenarios. While some competitors such as RFE and the Lasso achieve comparable accuracy to AdaSTAMPS with oracle model selection at some SNR levels (Figure 2B, 2E), their performance degrades drastically when their tuning hyperparameters are chosen via widely-used data-driven ways (Figure 2A, 2D). On the other hand, performance of AdaSTAMPS doesn’t deteriorate as much across data-driven and oracle model selection. We also point out that our AdaSTAMPS can boost the accuracy of any existing feature selection strategy - AdaSTAMPS with thresholded OLS as base selector outperforms thresholded OLS itself by a large margin, especially in low SNR settings and in the more challenging scenario S2. Furthermore, AdaSTAMPS is computationally the fastest of any method with comparable accuracy, with runtime on the order of seconds to minutes as opposed to hours for many competing methods.

The results for the most challenging scenario S3 are summarized in Table 1. High-dimensional data of such scale is regularly observed in areas such as genetics and neuroimaging. We see that our AdaSTAMPS (EE) perfectly recovers the true feature support across both
Figure 2: Feature selection accuracy (F1 Score) and runtime from simulation Scenario 1 (S1) and Scenario 2 (S2). (A, D) F1 Score for data-driven feature support estimates; (B, E) F1 Score for oracle feature support estimates; (C, F) Computational time in seconds. The fastest runtime is reported for every method with parallelism enabled for methods (LassoCV 1SE, Lasso eBIC, CPSS, Rand. Lasso, RFE) wherever possible. Our AdaSTAMPS dominates all competitors in terms of feature selection accuracy in both scenarios. Moreover, the runtime of our AdaSTAMPS is the best of any method with comparable feature selection accuracy.

data-driven and oracle model selection. In addition, STAMPS and AdaSTAMPS (Prob) also perform favorably. On the other hand, all competing methods attain rather low feature selection accuracy even with oracle selection. Clearly, AdaSTAMPS (EE) is more capable of accurately identifying the true signal features even in high-correlation, high-dimensional settings, thus leading to more reliable model interpretation. In the meantime, AdaSTAMPS (EE) takes less than 20 minutes to run, rather than the 14 hours required to solve Randomized Lasso that achieves the next best data-driven F1 Score among competing methods.

Overall, we empirically demonstrate the effectiveness of our proposed approaches in selecting statistically accurate features with much improved robustness and runtime, even in challenging high-noise, high-correlation scenarios. Even though STAMPS does not perform as well as AdaSTAMPS, STAMPS still performs better than a lot of the competing methods, especially in terms of data-driven F1 Score (e.g. Figure 2A, 2D). In the meantime, our AdaSTAMPS (both (EE) and (Prob)) consistently outperform competing methods in terms of feature selection accuracy while being computationally the fastest of any method with
Table 1: Results of Various Methods for Simulation Scenario 3 (S3). For each method, the number of selected features determined by data-driven procedures, the data-driven F1 Score, the oracle F1 Score, and computational time in minutes are reported. The method with the best accuracy is bold-faced. Note that the fastest runtime is reported for every method with parallelism enabled for methods (*) wherever possible. Our AdaSTAMPS dominates all competing methods in terms of feature selection accuracy.

| Method           | # Selected Features | F1 (Data-Driven) | F1 (Oracle) | Time (Minute) |
|------------------|---------------------|------------------|-------------|---------------|
| AdaSTAMPS (EE)   | 20                  | 1.000            | 1.000       | 19.87         |
| AdaSTAMPS (Prob) | 47                  | 0.597            | 0.750       | 52.10         |
| STAMPS           | 237                 | 0.117            | 0.750       | 74.70         |
| LassoCV 1SE      | 194                 | 0.112            | 0.600       | 176.60*       |
| Lasso eBIC       | 46                  | 0.091            | 0.100       | 61.59*        |
| CPSS             | 81                  | 0.238            | 0.500       | 716.94*       |
| Rand. Lasso      | 9                   | 0.276            | 0.450       | 838.53*       |
| Uni. Filter      | 277261              | 0.000            | 0.050       | 0.05          |
| RFE              | 37035               | 0.001            | 0.350       | 97.82*        |
| Thres. OLS       | 4                   | 0.083            | 0.100       | 1.01          |

comparable accuracy. This suggests that our flexible AdaSTAMPS approach could be a double-win both in terms of statistical accuracy and computation in practice.

3.2 Real-World Classification Data

We now study the performance of our method when employed as a dimensionality-reduction preprocessing step for downstream classification tasks on several real-world data sets - CIFAR10 [Krizhevsky 2009], Gisette [Guyon et al. 2004], and Fashion MNIST [Xiao et al. 2017]. For CIFAR10, we consider two classes (truck, automobile) with \( N = 12000 \) images each having \( M = 3072 \) features. Gisette is a binary classification data set (\( N = 7000, M = 5000 \)). For Fashion MNIST, we consider three classes (T-shirt, coat, shirt) with \( N = 18000 \) and \( M = 784 \). We split each data set into 70% training and 30% test data via stratified sampling.

For every training data set, our AdaSTAMPS (EE) fits Random Forest (RF) classifiers as base selectors to many random minipatches and keeps 10 features with the highest Gini importance from every minipatch. Features are then ranked based on their selection frequencies from AdaSTAMPS (EE), which are computed by taking an ensemble of selection results over these minipatches. For comparisons, we also fit a RF classifier to the full training data and obtain ranks for features based on the RF Gini importance. Moreover, we obtain another set of ranks for features using a RFE with RF as base estimator on the full training data. As the true feature support is unknown in these real-world data, performance of each feature selection method is measured by training a new downstream RF classifier using only its top ranked \( J \) features and then computing classification accuracy on the test data. In all cases, we use the default RF classifier from Scikit-learn. Figure 3 shows the average test accuracy of the downstream RF classifiers over 5 random training/test splits versus number of selected features that are used in training these downstream classifiers.

From Figure 3, we see that AdaSTAMPS (EE) mostly outperforms the competitors
Figure 3: Results for the real-world classification data. The top features selected by each feature selection method are used to train new RF classifiers separately for downstream classification tasks. (A, B, C) Test accuracy of these downstream RF classifiers (averaged over 5 random training/test splits) versus number of selected top features that are used in training the downstream classifiers. Overall, the features selected by our AdaSTAMPS (EE) lead to better classification accuracy on test sets, especially as the number of selected features increases.

because the features selected by AdaSTAMPS (EE) are generally more informative and eventually lead to better test accuracy in all three data sets. Similar to observations from Sec. 3.1, our AdaSTAMPS framework can boost the feature selection performance of any existing selection strategy: AdaSTAMPS (EE) with RF as inner feature selector outperforms RF itself, sometimes by a large margin as in Figure 3B.

4 Conclusion

We have developed general meta-algorithms, STAMPS and AdaSTAMPS, that practitioners can employ with a wide variety of existing learning algorithms and feature selection strategies to identify statistically accurate features in a computationally efficient manner, thus enhancing interpretability of machine learning models. In future work, we look to theoretically analyze the more sophisticated AdaSTAMPS framework and also investigate distributed, memory-efficient implementation of our methods.

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A Probabilistic Adaptive Feature Sampling Scheme

In Algorithm 3, we give our probabilistic adaptive feature sampling scheme. Here, $k$ is the iteration counter, $E$ denotes the total number of burn-in epochs. We use the shorthand AdaSTAMPS (Prob) to denote the specific instance of AdaSTAMPS that employs Algorithm 3 as its adaptive feature sampling scheme.

Similar to our exploitation and exploration scheme (Algorithm 2 in the main paper), AdaSTAMPS (Prob) first explores the entire input feature space during the burn-in stage. After $E$ burn-in epochs, Algorithm 3 transitions to the adaptive stage during which it adaptively updates the sampling probabilities for all features. Specifically, features with higher selection frequencies, which are likely to be the true signal features, have relatively higher probabilities of being sampled into subsequent minipatches. However, since this is a probabilistic sampling scheme, Algorithm 3 still adaptively explores the remaining feature space because many features with relatively lower selection frequencies still have non-zero probabilities of being sampled into minipatches.

**Algorithm 3: Adaptive Feature Sampling Scheme: Probabilistic (Prob)**

```
Input: $k, M, m, E, \{\hat{\Pi}_j^{(k-1)}\}_{j=1}^M$.
Initialize: $G = \lceil \frac{M}{m} \rceil$, $J = \{1, \ldots, M\}$;
if $k \leq E \cdot G$ then // Burn-in stage
    if $\text{mod}_G(k) = 1$ then // Start of a burn-in epoch
        1) Randomly reshuffle feature index set $J$ and partition into disjoint sets
        $\{J_g\}_{g=0}^{G-1}$;
        1) Set $F_k = J_{\text{mod}_G(k)}$;
    else // Adaptive stage
        1) Update feature sampling probabilities using information from previous iterations:
        $$\hat{p}_j^{(k)} = \frac{\hat{\Pi}_j^{(k-1)}}{\sum_{j=1}^M \hat{\Pi}_j^{(k-1)}} \quad \text{for} \quad j \in \{1, \ldots, M\}$$
        2) Sample $m$ features $F_k \subset \{1, \ldots, M\}$ without replacement according to the
        updated feature sampling probabilities $\{\hat{p}_j^{(k)}\}_{j=1}^M$;
end
Output: $F_k$.
```
B Proof of Theorem 1

In this section, we present the detailed proof of Theorem 1. We restate our theoretical statements from the main paper here for convenience.

**Assumption (A1)** Each feature \( j \in \{1, \ldots, M\} \) is sampled at least once when STAMPS stops after \( K \) iterations.

**Assumption (A2)** For a given noise feature \( j \in S^c \), \( \mathbb{1}(j \in \hat{S}_i) \) is constant across the collection of minipatches \( \{(I_i, F_i)\}_{i \in D} \) that contain this feature, where \( D = \{i \in \{1, 2, \ldots, K\} : j \in F_i\} \).

**Assumption (A3)** \( \mathbb{P}[\hat{\Pi}^{(K)}_j \geq \pi_{\text{thr}}] \leq \alpha/|S^c| \) for any noise feature \( j \in S^c \) on an arbitrary minipatch \( (I_i, F_i) \) that contains this feature.

**Theorem 1** Assume that Assumptions (A1-3) hold. Then STAMPS controls the family-wise error rate (FWER) at level \( \alpha \):

\[
\mathbb{P}[|\hat{S}_{\text{stable}} \cap S^c| \geq 1] \leq \alpha
\]

**Proof:** By Markov inequality, we have

\[
\mathbb{P}[|\hat{S}_{\text{stable}} \cap S^c| \geq 1] \leq \frac{\mathbb{E}[|\hat{S}_{\text{stable}} \cap S^c|]}{1} = \mathbb{E}\left[ \sum_{j \in S^c} \mathbb{1}(j \in \hat{S}_{\text{stable}}) \right] = \mathbb{E}\left[ \sum_{j \in S^c} \mathbb{1}(\hat{\Pi}^{(K)}_j \geq \pi_{\text{thr}}) \right] = \sum_{j \in S^c} \mathbb{P}[\hat{\Pi}^{(K)}_j \geq \pi_{\text{thr}}] \tag{1}
\]

By definition, \( \hat{\Pi}^{(K)}_j \geq 0 \) for all \( j = 1, 2, \ldots, M \) and \( \pi_{\text{thr}} \in (0, 1) \). Using Markov inequality, for any noise feature \( j \in S^c \), we have

\[
\mathbb{P}[\hat{\Pi}^{(K)}_j \geq \pi_{\text{thr}}] \leq \frac{\mathbb{E}[\hat{\Pi}^{(K)}_j]}{\pi_{\text{thr}}} \tag{2}
\]

Now we need to evaluate \( \mathbb{E}[\hat{\Pi}^{(K)}_j] \). When STAMPS stops after \( K \) iterations, the selection frequency for any noise feature \( j \in S^c \) is

\[
\hat{\Pi}^{(K)}_j = \frac{1}{\max(1, \sum_{i=1}^K \mathbb{1}(j \in F_i))} \sum_{i=1}^K \mathbb{1}(j \in F_i, j \in \hat{S}_i)
\]

\[
= \frac{1}{\sum_{i=1}^K \mathbb{1}(j \in F_i)} \sum_{i=1}^K \mathbb{1}(j \in F_i, j \in \hat{S}_i) \quad \text{(by Assumption (A1))}
\]

\[
= \frac{\sum_{i=1}^K \mathbb{1}(j \in F_i, j \in \hat{S}_i)}{\sum_{i=1}^K \mathbb{1}(j \in F_i)}
\]
Then for any given noise feature \( j \in S^c \), we have

\[
E[\hat{\Pi}^{(K)}_j] = E\left[ \frac{\sum_{i=1}^{K} 1(j \in F_i, j \in \hat{S}_i)}{\sum_{i=1}^{K} 1(j \in F_i)} \right] \\
= E\left[ \frac{\sum_{i:j \in F_i} 1(j \in \hat{S}_i)}{\sum_{i:j \in F_i} 1} \right] \\
= E\left[ \frac{1(j \in \hat{S}_i)}{\sum_{i:j \in F_i} 1} \right] \\
= E[1(j \in \hat{S}_i)] \\
= \mathbb{P}[j \in \hat{S}_i] 
\]

(3)

Here, \( i \) indexes an arbitrary minipatch \((I_i, F_i)\) that contains this given \( j^{th} \) feature. Then the upper bound in (2) becomes

\[
\mathbb{P}[\hat{\Pi}^{(K)}_j \geq \pi_{\text{thr}}] \leq \mathbb{P}[j \in \hat{S}_i] 
\]

(4)

and combining with (1), we have

\[
\mathbb{P}[|\hat{S}^{\text{stable}} \cap S^c| \geq 1] \leq \sum_{j \in S^c} \mathbb{P}[\hat{\Pi}^{(K)}_j \geq \pi_{\text{thr}}] \\
\leq \sum_{j \in S^c} \frac{\mathbb{P}[j \in \hat{S}_i]}{\pi_{\text{thr}}} \\
\leq \sum_{j \in S^c} \frac{\alpha \pi_{\text{thr}}}{|S^c|\pi_{\text{thr}}} \quad \text{(by Assumption (A3))} \\
= \alpha 
\]

(5)
C More on Practical Considerations

C.1 Data-Driven Rule to Choose $\pi_{thr}$

While there are many possible data-driven approaches to choose the user-specific threshold $\pi_{thr} \in (0, 1)$, we propose a very simple one in Algorithm 4 that we found to be effective in practice. Here, $\{\hat{\Pi}^{(K)}_j\}_{j=1}^{M}$ is the set of feature selection frequencies from the last iteration of STAMPS or AdaSTAMPS, and SD denotes the sample standard deviation. Intuitively, Algorithm 4 tries to find all gaps (local minima of density) among $\{\hat{\Pi}^{(K)}_j\}_{j=1}^{M}$ in order to choose a threshold that separates features with relatively high selection frequencies from those with low selection frequencies. Algorithm 4 does so by first fitting a Gaussian kernel density estimator (KDE) to the selection frequencies and then setting $\pi_{thr}$ to the smallest local minima, if any.

**Algorithm 4: KDE-Based Data-Driven Rule to Choose $\pi_{thr}$**

**Input:** $\{\hat{\Pi}^{(K)}_j\}_{j=1}^{M}$.

1) Fit a Gaussian KDE to $\{\hat{\Pi}^{(K)}_j\}_{j=1}^{M}$:

$$
\hat{f}_h(x) = \frac{1}{M} \sum_{j=1}^{M} \exp \left( - \frac{(x - \hat{\Pi}^{(K)}_j)^2}{2h^2} \right), \forall x \in [0, 1]
$$

where the bandwidth of the Gaussian kernel $h = \text{SD}(\{\hat{\Pi}^{(K)}_1, \hat{\Pi}^{(K)}_2, \ldots, \hat{\Pi}^{(K)}_M\})$;

2) Use method of inflection to find all local minima:

$$
\mathcal{D} = \{x \in [0, 1] \text{ s.t. } \text{sign}(\hat{f}_h(x) - \hat{f}_h(x - \epsilon)) = -1, \text{sign}(\hat{f}_h(x + \epsilon) - \hat{f}_h(x)) = 1 \}
$$

for any $\epsilon > 0$;

if $|\mathcal{D}| \geq 1$ then // Find at least one local minima

Set $\pi_{thr} = \min(\mathcal{D})$;

else // No local minima is found

Set $\pi_{thr} = 0.5$;

end

**Output:** $\pi_{thr}$.

C.2 Additional Empirical Studies to Investigate Effects of Minipatch Size

Here, we empirically investigate how feature selection accuracy and runtime of our meta-algorithm vary for various minipatch sizes (i.e. $n$ and $m$ values). We take a subset of the TCGA data that contains 450K methylation status of the top $M = 2000$ most variable CpGs for $N = 2000$ patients as the data matrix $X \in \mathbb{R}^{N \times M}$. Following similar procedures to Sec. 3.1 of the main paper, we create the true feature support $S$ by randomly sampling $S \subset \{1, \ldots, M\}$ with $|S| = 10$. The $M$-dimensional sparse coefficient vector $\beta$ is generated by setting $\beta_j = 0$ for all $j \notin S$ and $\beta_j = 1$ for all $j \in S$. The true feature importance $\pi^*$ is then calculated as $\pi^* = \sum_{j=1}^{M} \beta_j^2$.
for all $j \notin S$ and $\beta_j$ is chosen independently and uniformly in $[-3/b, -2/b] \cup [2/b, 3/b]$ for all $j \in S$. Finally, the response vector $y \in \mathbb{R}^N$ can be generated by $y = X\beta + \epsilon$ where the noise vector $(\epsilon_1, \ldots, \epsilon_N)$ is IID $\mathcal{N}(0, 1)$. The positive real number $b$ is chosen such that the SNR is fixed at 5. This synthetic data set is also used for Figure 1 in the main paper.

To study how performance varies with minipatch size ($n$ and $m$), we run AdaSTAMPS (EE) using thresholded OLS as the inner base selector for various $n$ and $m$ values. Specifically, we take a sequence of $m$ values that are integer multiples of $|S|$ and then pick $n$ relative to $m$ such that $m/n \in \{0.1, 0.2, \ldots, 0.8\}$. Feature selection accuracy in terms of F1 Scores and computational time are reported for various $n$ and $m$ values in Figure C. We see that our method is quite robust to the choice of minipatch size for it has stable feature selection performance for a sensible range of $n$ and $m$. In general, we found taking $m$ to well exceed the number of true features (e.g. 3-10 times the number of true features $|S|$) and then picking $n$ relative to $m$ so that it well exceeds the sample complexity of the base feature selector used in the meta-algorithm works well in practice.

Figure C: We show how feature selection accuracy and computational time of AdaSTAMPS (EE) vary with minipatch sizes in terms of $m/|S|$ and $m/n$, where $|S|$ is the cardinality of the true feature support. F1 Scores and computational time are averaged over 5 replicates. (A) Feature selection accuracy in terms of F1 Score with data-driven $\pi_{\text{thr}}$; (B) Feature selection accuracy in terms of F1 Score with oracle model selection; and (C) Computational time on $\log_{10}$ scale. Note that gray cells indicate NA’s that result from poor selection performance. We see that our method has stable performance for a sensible range of $n$ and $m$.

D Data-Driven and Oracle Model Selection Procedures for Various Methods in Empirical Studies

In this section, we discuss details of data-driven and oracle model selection procedures that we use for our empirical studies in Sec. 3.1 of the main paper.

For the Lasso, data-driven model selection approaches such as the 10-fold cross-validation with 1 standard error (SE) rule (Hastie et al., 2009) and the extended BIC (eBIC) rule (Chen...
and Chen (2008) are employed to determine the optimal regularization parameter, resulting in $\lambda_{\text{ISE}}$ and $\lambda_{\text{eBIC}}$, respectively. For univariate filter, the data-driven rule keeps all features that are significantly associated with the response with FDR controlled at the default level of 0.05. Additionally, 10-fold cross-validation is carried out for RFE to choose the best tuning hyperparameters and the Bonferroni procedure (Giurcanu, 2016) is used with thresholded OLS to determine the selected features in a data-driven manner. The user-specific threshold $\pi_{\text{thr}}$ is set to 0.5 for STAMPS, AdaSTAMPS (EE), AdaSTAMPS (Prob), CPSS, and Randomized Lasso. Following the default settings in Scikit-learn (Pedregosa et al., 2011), we use a geometric sequence $\Lambda \in (0.001\lambda_{\text{max}}, \lambda_{\text{max}})$ as the set of candidate regularization parameters for all Lasso-based methods, where $\lambda_{\text{max}}$ is the minimum amount of regularization such that all estimated coefficients become zero. $|\Lambda|$ is set to the default 100 for simulation scenarios S1-S2 and to 10 for scenario S3. For AdaSTAMPS (EE), we also use default settings with $E = 10$ burn-in epochs and $\pi_{\text{active}} = 0.1$. For AdaSTAMPS (Prob), $E$ is also set to 10.

Assuming knowledge of the number of true signal features $|S|$, oracle model selection is also applied to every method. Specifically, oracle model selection is applied to STAMPS, AdaSTAMPS (EE), AdaSTAMPS (Prob), CPSS, and Randomized Lasso by keeping the top $|S|$ features with the highest selection frequencies. For the Lasso, oracle selection keeps the top $|S|$ features with the largest magnitude in terms of coefficient estimates at $\lambda_{\text{ISE}}$ and $\lambda_{\text{eBIC}}$, respectively. For univariate filter, the top $|S|$ features with the smallest p-values are kept and the top $|S|$ features with the largest magnitude in terms of OLS coefficient estimates are selected for thresholded OLS. Furthermore, RFE is carried out until exactly $|S|$ features are left for oracle model selection.
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