Learning non-smooth sparse additive models from point queries in high dimensions

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Abstract

We consider the problem of learning a *d*-variate function f defined on the cube $[-1,1]^d \subset \mathbb{R}^d$, where the algorithm is assumed to have black box access to samples of f within this domain. Denote $S_r \subset \binom{[d]}{r}$; $r = 1, \ldots, r_0$ to be sets consisting of unknown *r*-wise interactions amongst the coordinate variables. We then focus on the setting where f has an additive structure, i.e., it can be represented as

$$f = \sum_{\mathbf{j}\in\mathcal{S}_1} \phi_{\mathbf{j}} + \sum_{\mathbf{j}\in\mathcal{S}_2} \phi_{\mathbf{j}} + \dots + \sum_{\mathbf{j}\in\mathcal{S}_{r_0}} \phi_{\mathbf{j}},$$

where each $\phi_{\mathbf{j}}$; $\mathbf{j} \in S_r$ is at most *r*-variate for $1 \leq r \leq r_0$. We derive randomized algorithms that query *f* at carefully constructed set of points, and exactly recover each S_r with high probability. In contrary to the previous work, our analysis does not rely on numerical approximation of derivatives by finite order differences.

Key words: Sparse additive models, sampling, hash functions, sparse recovery

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1 Introduction

Approximating a function from its samples is a fundamental problem with rich theory developed in areas such as numerical analysis and statistics, and which also has numerous practical applications such as in systems biology [15], solving PDEs [7], control systems [45], optimization [34] etc. Concretely, for an unknown *d*-variate function $f : \mathcal{G} \to \mathbb{R}$, one is given information about f in the form of samples $(\mathbf{x}_i, f(\mathbf{x}_i))_{i=1}^n$. Here, the \mathbf{x}_i 's belong to a compact subset $\mathcal{G} \subset \mathbb{R}^d$. The goal is to construct a smooth estimate $\hat{f} : \mathcal{G} \to \mathbb{R}$ such that the error between \hat{f} and f is small. In this paper we focus on the high dimensional setting where d is large. We will consider the scenario where the algorithm has black box access to the function, and can query it at any point within \mathcal{G} . This setting appears for instance in materials science [14], where \mathbf{x} represents some material and $f(\mathbf{x})$ some of its properties of interest (like thermal or electric conductivity). The local-density approximations

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in density functional theory can be used to compute to high accuracy such properties of a given material. The sampling then corresponds to running a costly numerical PDE-solver. Since such simulations are typically expensive to run, one would like to minimize the number of queries made. This setting is different from the regression setting typically considered in statistics wherein the \mathbf{x}_i 's are generated apriori from some unknown distribution over \mathcal{G} .

Curse of dimensionality. It is well known that provided we only make smoothness assumptions on f (such as differentiability or Lipschitz continuity), then the problem is intractable, i.e., has exponential complexity (in the worst case) with respect to the dimension d. For instance if $f \in$ $C^{s}(\mathcal{G})$, then any algorithm needs in the worst case $n = \Omega(\delta^{-d/s})$ samples to uniformly approximate f with error $\delta \in (0,1)$, cf. [27, 40]. Furthermore, the constants behind the Ω -notation may also depend on d. A detailed study of the dependence on d was performed in the field of Information Based Complexity for $f \in C^{\infty}(\mathcal{G})$ in a more recent work [28]. The authors show that even here, n = $\Omega(2^{\lfloor d/2 \rfloor})$ samples are needed in the worst case for uniform approximation within an error $\delta \in (0,1)$ (with no additional dependence on d hidden behind the Ω -notation). This exponential dependence on d is commonly referred to as the *curse of dimensionality*. The above results suggest that in order to get tractable algorithms in the high dimensional regime, one needs to make additional assumptions on f. To this end, a growing line of work over the past decade has focused on the setting where f possesses an intrinsic, albeit unknown, low dimensional structure, with much smaller intrinsic dimension than the ambient dimension d. The motivation is that one could now hope to design algorithms with complexity at most exponential in the intrinsic dimension, but with mild dependence on d.

1.1 Sparse additive models (SPAMs)

A popular class of functions with an intrinsic low dimensional structure are the so-called sparse additive models (SPAMs). These are functions that are decomposable as the sum of a small number of lower dimensional functions. More precisely, for $S_r \subset {\binom{[d]}{r}}; r = 1, \ldots, r_0$, the function $f : \mathcal{G} \to \mathbb{R}$ is of the form

$$f = \sum_{j \in \mathcal{S}_1} \phi_j(x_j) + \sum_{(j_1, j_2) \in \mathcal{S}_2} \phi_{(j_1, j_2)}(x_{j_1}, x_{j_2}) + \dots + \sum_{(j_1, \dots, j_{r_0}) \in \mathcal{S}_{r_0}} \phi_{(j_1, \dots, j_{r_0})}(x_{j_1}, \dots, x_{j_{r_0}})$$
(1.1)

with each $|S_r| \ll {d \choose r}$, and $r_0 \ll d$. We can interpret the tuples in S_r as r^{th} order interactions terms. Let us remark, that usually the terminology Sparse additive models is used for the case $r_0 = 1$, but we prefer to use it here in the general sense of (1.1).

These models appear in optimization under the name *partially separable* models (cf., [16]). They also arise in electronic structure computations in physics (cf., [3]), and problems involving multiagent systems represented as decentralized partially observable Markov decision processes (cf., [12]). There exists a rich line of work that mostly study special cases of the model (1.1). We review them briefly below, leaving a detailed comparison with our results to Section 8.

The case $r_0 = 1$. In this setting, (1.1) reduces to a sparse sum of univariate functions. This model has been studied extensively in the non parametric statistics literature with a range of results on estimation of f (cf., [19, 21, 22, 24, 30, 32]) and also on variable selection, i.e., identifying the support S_1 (cf., [19, 32, 42]). The basic idea behind these approaches is to approximately represent each ϕ_j in a suitable basis of finite size (for eg., splines or wavelets) and then to find the coefficients in the basis expansion by solving a least squares problem with smoothness and sparsity penalty constraints. Koltchinskii et al. [22] and Raskutti et al. [30] proposed a convex program for estimating f in the Reproducing kernel Hilbert space (RKHS) setting, and showed that f lying in a Sobolev space with smoothness parameter $\alpha > 1/2$ can be estimated at the L_2 rate $\frac{k \log d}{n} + kn^{-\frac{2\alpha}{2\alpha+1}}$. This rate was shown to be optimal in [30]. There also exist results for variable selection, i.e., identifying the support S_1 . These results in non parametric statistics are typically asymptotic in the limit of large n, also referred to as sparsistency [19, 32, 42]. Recently, Tyagi et al. [36] derived algorithms that query f, along with non-asymptotic sampling bounds for identifying S_1 . They essentially estimate the (sparse) gradient of f using results from compressed sensing (CS), at few carefully chosen locations in \mathcal{G} .

The case $r_0 = 2$. This setup has received relatively less attention than the aforementioned setting. Radchenko et al. [29] proposed an algorithm VANISH, and showed that it is sparsistent, i.e., recovers S_1, S_2 in the limit of large n. The ACOSSO algorithm [35] can handle this setting, with theoretical guarantees (sparsistency, convergence rates) shown when $r_0 = 1$. Recently, Tyagi et al. [37, 38] derived algorithms that query f, and derived non-asymptotic sampling bounds for recovering S_1, S_2 . Their approach for recovering S_2 was based on estimating the (sparse) Hessian of f using results from CS, at carefully chosen points in \mathcal{G} . The special case where f is *multilinear* has been studied considerably; there exist algorithms that recover S_1, S_2 , along with convergence rates for estimating f in the limit of large n [6, 29, 2]. There also exist non-asymptotic sampling bounds for identifying S_1, S_2 in the noiseless setting (cf., [26, 20]); these works essentially make use of the CS framework.

The general case. Much less is known about the general setup where $r_0 \ge 2$ is possible. Lin et al. [23] were the first to introduce learning SPAMs of the form (1.1), and proposed the COSSO algorithm. Recently, Dalalyan et al. [10] and Yang et al. [44] studied (1.1) in the regression setting and derived non-asymptotic error rates for estimating f. In particular, Dalalyan et al. studied this in the Gaussian white noise model, while Yang et al. considered the Bayesian setup wherein a Gaussian process (GP) prior is placed on f. When f is multilinear, the work of Nazer et al. [26], which is in the CS framework, gives non-asymptotic sampling bounds for recovering S_r , $r = 1, \ldots, r_0$.

1.2 Our contributions and main idea

Before proceeding, we will briefly mention our problem setup to put our results in the context; it is described more formally later on in Section 2. We consider $f: [-1, 1]^d \to \mathbb{R}$ of the form (1.1) and denote by $\mathcal{S}_j^{(1)}$ the variables occurring in \mathcal{S}_j . We assume, that $\mathcal{S}_j^{(1)}$ are disjoint¹ for $1 \leq j \leq r_0$. Each component ϕ is assumed to be Hölder smooth, and is also assumed to be "sufficiently large" at some point within its domain. Our goal is to query f at few locations in $\mathcal{G} = [-1, 1]^d$, and recover the underlying sets of interactions \mathcal{S}_r , for each $r = 1, \ldots, r_0$.

Our results. To our knowledge, we provide the first non-asymptotic sampling bounds for exact identification of S_r , for each $r = 1, \ldots, r_0$, for SPAMs of the form (1.2). In particular, we derive a

¹For $r_0 = 2$, this represents no additional assumption. See discussion after Proposition 2.

randomized algorithm that with high probability recovers each S_r , $r = 1, \ldots, r_0$ with

$$\Omega\left(\sum_{i=3}^{r_0} \left[\underbrace{c_i^i i^{i+2} |\mathcal{S}_i|^2 \log^2 d}_{\text{Identifying } \mathcal{S}_i}\right] + \underbrace{c_2 |\mathcal{S}_2| \log\left(\frac{d^2}{|\mathcal{S}_2|}\right) \log d}_{\text{Identifying } \mathcal{S}_2} + \underbrace{c_1 |\mathcal{S}_1| \log\left(\frac{d}{|\mathcal{S}_1|}\right)}_{\text{Identifying } \mathcal{S}_1}\right)$$
(1.2)

noiseless queries of f within $[-1, 1]^d$. The same bound holds when the queries are corrupted with (a) arbitrary bounded noise provided the noise magnitude is sufficiently small (see Theorem 8), and (b) i.i.d. Gaussian noise, which we handle by resampling each query sufficiently many times, and averaging (see Theorem 9). Here, c_i 's depend on the smoothness parameters of ϕ 's, and in the case of Gaussian noise, also depend on its variance.

We improve on the recent work of Tyagi et al. [37, 38], wherein SPAMs with $r_0 = 2$ were considered, by being able to handle general $r_0 \ge 1$. Moreover, we only require f to be Hölder smooth while the algorithms in [37, 38] necessarily require f to be continuously differentiable. Finally, our bounds improve upon those in [37, 38] when the noise is i.i.d. Gaussian. In this scenario, our bounds are linear in the sparsity $|\mathcal{S}_2| + |\mathcal{S}_1|$ while those in [37, 38] are polynomial in the sparsity.

The sampling scheme that we employ to achieve these bounds is novel, and is specifically tailored to the additive nature of f. We believe this scheme to be of independent interest for other problems involving additive models, such as in optimization of high dimensional functions with partially separable structure.

Main idea. We identify each set S_i in a sequential "top down" manner by first identifying S_{r_0} . Once we find S_{r_0} , the same procedure is repeated on the remaining set of variables (excluding those found in S_{r_0}) to identify S_{r_0-1} , and consequently, each remaining S_i . We essentially perform the following steps for recovering S_{r_0} . Consider some given partition of [d] into r_0 disjoint subsets $\mathcal{A} = (\mathcal{A}_1, \ldots, \mathcal{A}_{r_0})$, a Bernoulli vector $\mathcal{\beta} \in \{-1, 1\}^d$ and some given $\mathbf{x} \in [-1, 1]^d$. We generate 2^{r_0} query points $(\mathbf{x}_i)_{i=1}^{2r_0}$, where each \mathbf{x}_i is constructed using $\mathcal{\beta}, \mathbf{x}$ and \mathcal{A} . Then, for some fixed sequence of signs $s_1, s_2, \ldots, s_{2r_0} \in \{-1, 1\}$ (depending only on r_0), we show for the anchored-ANOVA representation of f (see Section 2 and Lemma 4) that

$$\sum_{i=1}^{2^{r_0}} s_i f(\mathbf{x}_i) = \sum_{(j_1,\dots,j_{r_0})\in\mathcal{A}\cap\mathcal{S}_{r_0}} \beta_{j_1}\dots\beta_{j_{r_0}} \phi_{(j_1,\dots,j_{r_0})}(x_{j_1},\dots,x_{j_{r_0}}).$$
(1.3)

Observe, that (1.3) corresponds to a multilinear measurement of a sparse vector with entries $\phi_{(j_1,\ldots,j_{r_0})}(x_{j_1},\ldots,x_{j_{r_0}})$, indexed by the tuple (j_1,\ldots,j_{r_0}) . Indeed, this vector is $|\mathcal{S}_{r_0}|$ sparse. This suggests that by repeating the above process at sufficiently many random β 's, we can recover an estimate of this $|\mathcal{S}_{r_0}|$ vector by using known results from CS. Thereafter, we repeat the above process for each \mathcal{A} corresponding to a family of perfect hash functions (see Definition 1). The size of this set is importantly at most exponential in r_0 , and only logarithmic in d. The **x**'s are then chosen to be points on a uniform r_0 dimensional grid constructed using \mathcal{A} . This essentially enables us to guarantee that we are able to sample each $\phi_{(j_1,\ldots,j_{r_0})}$ sufficiently fine within its domain, and thus identify (j_1,\ldots,j_{r_0}) by thresholding.

Organization of paper. The rest of the paper is organized as follows. In Section 2, we set up the notation and also define the problem formally. In Section 3, we begin with the case $r_0 = 1$ as warm up, and describe the sampling scheme, along with the algorithm for this setting. Section 4 considers

the bivariate case $r_0 = 2$, while Section 5 consists of the most general setting wherein $r_0 \ge 2$ is possible. Section 6 contains (mostly) known results from compressed sensing for estimating sparse multilinear functions from random samples. Section 7 then puts together the content from the earlier sections, wherein we derive our final theorems. Section 8 consists of a comparison of our results with closely related work, along with some directions for future work.

2 Notation and problem setup

Notation. Scalars will be usually denoted by plain letters (e.g. d), vectors by lowercase boldface letters (e.g., \mathbf{x}), matrices by uppercase boldface letters (e.g., \mathbf{A}) and sets by uppercase calligraphic letters (e.g., \mathcal{S}), with the exception of [n], which denotes the index set $\{1, \ldots, n\}$ for any natural number $n \in \mathbb{N}$. For a (column) vector $\mathbf{x} = (x_1 \dots x_d)^T$ and an r-tuple $\mathbf{j} = (j_1, j_2, \dots, j_r) \in {[d] \choose r}$, we denote $\mathbf{x}_{\mathbf{j}} = (x_{j_1} \dots x_{j_r})^T \in \mathbb{R}^r$ to be the restriction of \mathbf{x} on \mathbf{j} . For any finite set \mathcal{A} , $|\mathcal{A}|$ denotes the cardinality of \mathcal{A} . Moreover, if $\mathcal{A} \subseteq [d]$, then $\Pi_{\mathcal{A}}(\mathbf{x})$ denotes the projection of \mathbf{x} on \mathcal{A} where

$$(\Pi_{\mathcal{A}}(\mathbf{x}))_{i} = \begin{cases} x_{i} ; & i \in \mathcal{A}, \\ 0 ; & i \notin \mathcal{A}, \end{cases} \quad i \in [d].$$

$$(2.1)$$

The ℓ_p norm of a vector $\mathbf{x} \in \mathbb{R}^d$ is defined as $\|\mathbf{x}\|_p := \left(\sum_{i=1}^d |x_i|^p\right)^{1/p}$. A random variable β is called Bernoulli variable if $\beta = +1$ with probability 1/2 and $\beta = -1$ with probability 1/2. A vector $\boldsymbol{\beta} \in \{-1, +1\}^d$ of independent Bernoulli variables is called Bernoulli vector. Similarly, a matrix $\mathbf{B} \in \{-1, +1\}^{n \times d}$ is called Bernoulli matrix, if all its entries are independent Bernoulli variables.

Sparse Additive Models. For an unknown $f : \mathbb{R}^d \to \mathbb{R}$, our aim will be to approximate f uniformly from point queries within a compact domain $\mathcal{G} \subset \mathbb{R}^d$. From now on, we will assume $\mathcal{G} = [-1, 1]^d$. The sets $\mathcal{S}_r \subset {\binom{[d]}{r}}; r = 1, \ldots, r_0$, will represent the interactions amongst the coordinates, with \mathcal{S}_r consisting of *r*-wise interactions. Our interest will be in the setting where each \mathcal{S}_r is sparse, i.e., $|\mathcal{S}_r| \ll d^r$. Given this setting, we assume to have the following structure

$$f = \sum_{\mathbf{j}\in\mathcal{S}_1} \phi_{\mathbf{j}} + \sum_{\mathbf{j}\in\mathcal{S}_2} \phi_{\mathbf{j}} + \dots + \sum_{\mathbf{j}\in\mathcal{S}_{r_0}} \phi_{\mathbf{j}}.$$
 (2.2)

It is important to note here that the components in S_r will be assumed to be truly *r*-variate, in the sense that they cannot be written as the sum of lower dimensional functions. For example, we assume that the components in S_2 cannot be expressed as the sum of univariate functions.

Model Uniqueness and ANOVA-decompositions. We note now that the representation of f in (2.2) is not necessarily unique and some additional assumptions are needed to ensure uniqueness. For instance, one could add constants to each ϕ that sum up to zero, thereby giving the same f. Moreover, if S_2 contains overlapping pairs of variables, then for each such variable – call it p – one could add/subtract functions of the variable x_p to each corresponding ϕ_j such that f remains unaltered. To obtain unique representation of f, we will work with the so-called Anchored ANOVA-decomposition of f. We recall its notation and results in the form needed later and refer to [18] for more details.

The usual notation of an ANOVA-decomposition works with functions indexed by subsets of [d], instead of tuples from [d]. As there is an obvious one-to-one correspondence between r-tuples and subsets of [d] with r elements, we prefer to give the ANOVA-decomposition in its usual form.

Let $\mu_j, j = 1, \ldots, d$ be measures defined on all Borel subsets of [-1, 1] and let $U \subseteq [d]$. We let $d\mu_U(\mathbf{x}_U) = \prod_{i \in U} d\mu_i(x_i)$ be the product measure. We define

$$P_U f(\mathbf{x}_U) = \int_{[-1,1]^{d-|U|}} f(\mathbf{x}) d\mu_{[d]\setminus U}(\mathbf{x}_{[d]\setminus U}).$$

The ANOVA-decomposition of f is then given as

$$f(\mathbf{x}) = f_{\emptyset} + \sum_{i=1}^{d} f_i(x_i) + \sum_{i=1}^{d-1} \sum_{j=i+1}^{d} f_{i,j}(x_i, x_j) + \dots + f_{1,\dots,d}(x_1, \dots, x_d) = \sum_{U \subseteq [d]} f_U(\mathbf{x}_U),$$

where

$$f_U(\mathbf{x}_U) = \sum_{V \subseteq U} (-1)^{|U| - |V|} P_V f(\mathbf{x}_V).$$
(2.3)

In the case of $d\mu_j(x_j) = \delta(x_j)dx_j$, where δ is the Dirac distribution, we obtain the Anchored-ANOVA decomposition

$$f(\mathbf{x}) = \sum_{U \subseteq [d]} f_U(\mathbf{x}_U),$$

where $f_{\emptyset} = f(0)$ and $f_U(\mathbf{x}_U) = 0$ if $x_j = 0$ for some $j \in U$.

The standard theory of ANOVA decompositions is usually based on Hilbert space theory. As we prefer to work with continuous functions, we give the following representation theorem. The proof can be found in the Appendix.

Proposition 1. Let $f \in C([-1,1]^d)$. Then there is a unique collection of $(f_U)_{U \subset [d]}$, such that

- a) $f_U \in C([-1,1]^{|U|}),$
- b) f can be represented as

$$f(\mathbf{x}) = \sum_{U \subseteq [d]} f_U(\mathbf{x}_U), \quad \mathbf{x} \in [-1, 1]^d,$$
(2.4)

where $\mathbf{x}_U \in [-1, 1]^{|U|}$ is the restriction of \mathbf{x} onto indices included in U,

c) $f_U(\mathbf{x}_U) = 0$ if $x_j = 0$ for some $j \in U$.

The Anchored ANOVA-decomposition (2.4) can be used to ensure uniqueness of representation of f of the form (2.2). For the clarity of presentation, we will later distinguish between three settings. The first one is univariate with $r_0 = 1$, the second one with $r_0 = 2$ allows also for bivariate interactions between the variables. Finally, in the multivariate case $r_0 > 2$, arbitrary higher-order interactions can occur. We will present a detailed proposition about the corresponding ANOVA-decomposition in each of the sections separately.

Assumptions. We will specify the assumptions in each of the settings discussed later in more detail. But, in general, we will work with two groups of conditions.

1. Smoothness. We will assume throughout the paper that the components of the ANOVAdecomposition are Hölder smooth with exponent $\alpha \in (0, 1]$ and constant L > 0, i.e.,

$$|\phi(\mathbf{x}) - \phi(\mathbf{y})| \le L \parallel \mathbf{x} - \mathbf{y} \parallel_2^{\alpha}$$

for all admissible \mathbf{x}, \mathbf{y} .

2. Identifiability. Furthermore, our aim is the identification of the possible interactions between the variables. We are therefore not only interested in the approximation of f but also on the identification of the sets $S_1, S_2, \ldots, S_{r_0}$. Naturally, this is only possible if the non-zero functions in the Anchored-ANOVA decomposition are significantly large at some point. We will therefore assume that

$$\|\phi\|_{\infty} = \sup_{\mathbf{x}} |\phi(\mathbf{x})| > D$$

for some D > 0.

Problem parameters and goal. Based on the above setup, we will consider our problem specific parameters to be

- (a) smoothness parameters: $L > 0, \alpha \in (0, 1],$
- (b) identifiability parameters: D,
- (c) intrinsic/extrinsic dimensions: $d, r_0, |\mathcal{S}_1|, \ldots, |\mathcal{S}_{r_0}|$.

These parameters will be assumed to be known by the algorithm. The goal of the algorithm will then be to query f within $[-1,1]^d$, and to identify the sets S_1, \ldots, S_{r_0} exactly. Using standard methods of approximation theory and sampling along canonical subspaces, one may recover also the components in (2.2). We give some more details on this issue in Section 8.

3 The univariate case

As a warm up, we begin with the relatively simple setting where $r_0 = 1$, meaning that f is a sum of only univariate components. It means that f admits the representation

$$f = \mu + \sum_{p \in \mathcal{S}_1} \phi_p(x_p). \tag{3.1}$$

To ensure the uniqueness of this decomposition, we set $\mu = f(0)$ and assume that $\phi_p(0) = 0$ for all $p \in S_1$.

Assumptions. We will make the following assumptions on the model (3.1).

1. Smoothness. The terms in (3.1) are Hölder continuous with parameters $L > 0, \alpha \in (0, 1]$, i.e.,

$$|\phi_p(x) - \phi_p(y)| \le L|x - y|^{\alpha}$$
 for all $p \in \mathcal{S}_1$ and all $x, y \in [-1, 1]$

2. *Identifiability.* For every $p \in S_1$ there is an $x_p^* \in [-1, 1]$, such that $|\phi_p(x_p^*)| > D_1$.

Sampling scheme. Our sampling scheme is motivated by the following simple observation. For any fixed $\mathbf{x} \in [-1, 1]^d$, and some $\boldsymbol{\beta} \in \{-1, +1\}^d$, consider the points $\mathbf{x}^+, \mathbf{x}^- \in [-1, 1]^d$ defined as

$$x_i^+ = \begin{cases} x_i \; ; \quad \beta_i = +1, \\ 0 \; ; \quad \beta_i = -1 \end{cases} \quad \text{and} \quad x_i^- = \begin{cases} 0 \; ; \quad \beta_i = +1, \\ x_i \; ; \quad \beta_i = -1, \end{cases} \quad i \in [d].$$
(3.2)

Upon querying f at $\mathbf{x}^+, \mathbf{x}^-$, we obtain the noisy samples

$$\tilde{f}(\mathbf{x}^+) = f(\mathbf{x}^+) + \eta^+, \quad \tilde{f}(\mathbf{x}^-) = f(\mathbf{x}^-) + \eta^-,$$

where $\eta^+, \eta^- \in \mathbb{R}$ denotes the noise. One can then easily verify that the following identity holds on account of the structure of f

$$\tilde{f}(\mathbf{x}^+) - \tilde{f}(\mathbf{x}^-) = \sum_{i \in \mathcal{S}_1} \beta_i \underbrace{\phi_i(x_i)}_{z_i^*(x_i)} + \eta^+ - \eta^- = \langle \boldsymbol{\beta}, \mathbf{z}^*(\mathbf{x}) \rangle + \eta^+ - \eta^-.$$
(3.3)

Note that $\mathbf{z}^*(\mathbf{x}) = (z_1^*(x_1) \dots z_d^*(x_d))^T$ is $|\mathcal{S}_1|$ sparse, and $\tilde{f}(\mathbf{x}^+) - \tilde{f}(\mathbf{x}^-)$ corresponds to a noisy linear measurement of $\mathbf{z}^*(\mathbf{x})$, with $\boldsymbol{\beta}$. From standard compressive sensing results, we know that a sparse vector can be recovered *stably*, from only a few noisy linear measurements with random vectors, drawn from a suitable distribution. In particular, it is well established that random Bernoulli measurements satisfy this criteria. We discuss this separately later on, for now it suffices to assume that we have at hand an appropriate sparse recovery algorithm: SPARSE-REC.

We thus generate independent Bernoulli vectors $\beta_1, \beta_2, \ldots, \beta_n \in \{-1, +1\}^d$. For each β_i , we create $\mathbf{x}_i^+, \mathbf{x}_i^-$ as described in (3.2) (for some fixed \mathbf{x}), and obtain $\tilde{f}(\mathbf{x}_i^+), \tilde{f}(\mathbf{x}_i^-)$. Then, (3.3) gives us the linear system

$$\underbrace{\begin{pmatrix} \tilde{f}(\mathbf{x}_{1}^{+}) - \tilde{f}(\mathbf{x}_{1}^{-}) \\ \vdots \\ \vdots \\ \tilde{f}(\mathbf{x}_{n}^{+}) - \tilde{f}(\mathbf{x}_{n}^{-}) \end{pmatrix}}_{\mathbf{y}} = \underbrace{\begin{pmatrix} \boldsymbol{\beta}_{1}^{T} \\ \vdots \\ \vdots \\ \boldsymbol{\beta}_{n}^{T} \end{pmatrix}}_{\mathbf{B}} \mathbf{z}^{*}(\mathbf{x}) + \underbrace{\begin{pmatrix} \boldsymbol{\eta}_{1}^{+} - \boldsymbol{\eta}_{1}^{-} \\ \vdots \\ \vdots \\ \boldsymbol{\eta}_{n}^{+} - \boldsymbol{\eta}_{n}^{-} \end{pmatrix}}_{\boldsymbol{\eta}}.$$
(3.4)

SPARSE-REC will take as input \mathbf{y}, \mathbf{B} , and will output an estimate $\widehat{\mathbf{z}^*}(\mathbf{x})$ to $\mathbf{z}^*(\mathbf{x})$. Provided that n is large enough, we will typically have, for some $\epsilon \ge 0$, that $\|\widehat{\mathbf{z}^*}(\mathbf{x}) - \mathbf{z}^*(\mathbf{x})\|_{\infty} \le \epsilon$ holds. In such a case, we will refer to SPARSE-REC as being " ϵ -accurate" at \mathbf{x} .

Given the above, we now describe how to choose $\mathbf{x} \in [-1,1]^d$. To this end, we adopt the approach of [36], where the following grid on the diagonal of $[-1,1]^d$ was considered

$$\chi := \left\{ \mathbf{x} = (x \ x \ \cdots \ x)^T \in \mathbb{R}^d : x \in \left\{ -1, -\frac{m-1}{m}, \dots, \frac{m-1}{m}, 1 \right\} \right\}.$$
 (3.5)

Our aim will be to obtain the estimate $\widehat{\mathbf{z}^*}(\mathbf{x})$ at each $\mathbf{x} \in \chi$. Note that this gives us estimates to $\phi_p(x_p)$ for $p = 1, \ldots, d$, with x_p lying on a uniform one dimensional grid in [-1, 1]. Thus we can see, at least intuitively, that provided ϵ is small enough, and the grid is fine enough (so that we are close to $\phi_p(x_p^*)$ for each $p \in S_1$), we will be able to detect each $p \in S_1$ by thresholding.

Algorithm outline and guarantees The discussion above is outlined formally in the form of Algorithm 1. Lemma 1 below provides formal guarantees for exact recovery of support S_1 .

Lemma 1. Let SPARSE-REC be ϵ -accurate for each $\mathbf{x} \in \chi$ with $\epsilon < D_1/3$, which uses n linear measurements. Then for $m \ge (3L/D_1)^{1/\alpha}$, Algorithm 1 recovers S_1 exactly, i.e., $\widehat{S_1} = S_1$. Moreover, the total number of queries of f is 2(2m+1)n.

Proof. Recall that we denote $\mathbf{z}^*(\mathbf{x}) = (\phi_1(x_1) \dots \phi_d(x_d))^T$, and $z_i^*(x_i) = \phi_i(x_i)$. For any given $p \in S_1$, we know that there exists $x_p^* \in [-1, 1]$ such that $|\phi_p(x_p^*)| > D_1$. Also, on account of the construction of χ , there exists $\mathbf{x} = (x \dots x)^T \in \chi$ such that $|x - x_p^*| \leq 1/m$. Then starting with the fact that SPARSE-REC is ϵ accurate at \mathbf{x} , we obtain

$$\begin{aligned} |\hat{z}_p^*(x)| &\ge |\phi_p(x)| - \epsilon \ge |\phi_p(x_p^*)| - |\phi_p(x_p^*) - \phi_p(x)| - \epsilon \\ &\ge D_1 - \frac{L}{m^{\alpha}} - \epsilon \ge \frac{2D_1}{3} - \epsilon. \end{aligned}$$

Algorithm 1 Algorithm for estimating S_1

1: Input: $d, |\mathcal{S}_1|, m, n, \epsilon$. 2: Initialization: $\widehat{S_1} = \emptyset$. 3: Output: $\widehat{\mathcal{S}_1}$. 4: 5: Construct χ as defined in (3.5) with $|\chi| = 2m + 1$. 6: Generate Bernoulli vectors $\beta_1, \beta_2, \ldots, \beta_n \in \{-1, +1\}^d$. 7: Form $\mathbf{B} \in \mathbb{R}^{n \times d}$ as in (3.4). for $\mathbf{x} \in \chi$ do 8: Generate $\mathbf{x}_i^+, \mathbf{x}_i^- \in [-1, 1]^d$, as in (3.2), using $\mathbf{x}, \boldsymbol{\beta}_i$ for each $i \in [n]$. 9: Using the samples $(\tilde{f}(\mathbf{x}_i^+), \tilde{f}(\mathbf{x}_i^-))_{i=1}^n$, form **y** as in (3.4). 10: Obtain $\widehat{\mathbf{z}^*}(\mathbf{x}) = \text{SPARSE-REC}(\mathbf{y}, \mathbf{B}).$ 11: Update $\widehat{\mathcal{S}}_1 = \widehat{\mathcal{S}}_1 \cup \{ p \in [d] : |(\widehat{\mathbf{z}^*}(\mathbf{x}))_p| > \epsilon \}.$ 12: 13: end for

We used the reverse triangle inequality and the identifiability and smoothness assumptions on ϕ_p . On the other hand, since SPARSE-REC is ϵ accurate at each point in χ , therefore for every $q \notin S_1$ and $(c \ c \ \ldots c) \in \chi$, we know that $|\hat{z}_q^*(c)| \leq \epsilon$. It then follows readily for the stated choice of m, ϵ that $\widehat{S_1}$ contains each variable in S_1 , and none from S_1^c .

4 The bivariate case

Next, we consider the scenario where $r_0 = 2$, i.e., f can be written as a sum of univariate and bivariate functions. We denote by S_2^{var} the set of variables which are part of a 2-tuple in S_2 . Inserting this restriction into Proposition 1, we derive the following uniqueness result (its proof is postponed to the Appendix).

Proposition 2. Let $f \in C([-1, 1]^d)$ be of the form

$$f = \mu + \sum_{p \in \mathcal{S}_1} \phi_p(x_p) + \sum_{\mathbf{j} \in \mathcal{S}_2} \phi_{\mathbf{j}}(x_{\mathbf{j}}) + \sum_{l \in \mathcal{S}_2^{\text{var}}} \phi_l(x_l),$$
(4.1)

where $S_1 \cap S_2^{var} = \emptyset$. Moreover, let

- a) $\mu = f(0),$
- b) $\phi_j(0) = 0$ for all $j \in \mathcal{S}_1 \cup \mathcal{S}_2^{\text{var}}$,
- c) $\phi_{\mathbf{j}}(x_{\mathbf{j}}) = 0$ if $\mathbf{j} = (j_1, j_2) \in S_2$ and $x_{j_1} = 0$ or $x_{j_2} = 0$.

Then the representation (4.1) of f is unique in the sense that each component in (4.1) is uniquely identifiable.

Remark 1. In (4.1), we could have "collapsed" the terms corresponding to variables l in $\sum_{l \in S_2^{\text{var}}} \phi_l(x_l)$ – for l occurring exactly once in S_2 – uniquely into the corresponding component $\phi_j(x_j)$. A similar approach was adopted in [38], and the resulting model was shown to be uniquely identifiable. Yet here, we choose to represent f in the form (4.1) for convenience, and clarity of notation. This also leads to a less cumbersome expression, when we work with general interaction terms later. **Assumptions.** We now make the following assumptions on the model (4.1).

1. Smoothness. We assume each term in (4.1) to be Hölder continuous with parameters $L > 0, \alpha \in (0, 1]$, i.e.,

$$\begin{aligned} |\phi_p(x) - \phi_p(y)| &\leq L|x - y|^{\alpha} \quad \text{for all } p \in \mathcal{S}_1 \cup \mathcal{S}_2^{\text{var}} \text{ and for all } x, y \in [-1, 1], \\ |\phi_{\mathbf{j}}(\mathbf{x}) - \phi_{\mathbf{j}}(\mathbf{y})| &\leq L \parallel \mathbf{x} - \mathbf{y} \parallel_2^{\alpha} \quad \text{for all } \mathbf{j} \in \mathcal{S}_2 \text{ and for all } \mathbf{x}, \mathbf{y} \in [-1, 1]^2. \end{aligned}$$

2. Identifiability of S_1, S_2 . We assume that for each $p \in S_1$, there exists $x_p^* \in [-1, 1]$ so that $|\phi_p(x_p^*)| > D_1$ for some constant $D_1 > 0$. Furthermore, we assume that for each $\mathbf{j} \in S_2$ there exists $\mathbf{x}_{\mathbf{j}}^* \in [-1, 1]^2$ such that $|\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}^*)| > D_2$.

Before describing our sampling scheme, we need some additional notation. For any $\beta \in \{-1, 1\}$, we denote $\bar{\beta} = (-\beta)$. Moreover, $\mathbb{1}_{\beta}$ denotes the indicator variable of β , i.e., $\mathbb{1}_{\beta} = 1$ if $\beta = 1$, and $\mathbb{1}_{\beta} = 0$ if $\beta = -1$. Overall, our scheme proceeds in two stages. We first identify S_2 , and only then S_1 .

Sampling lemma for identifying S_2 . We begin by providing the motivation behind our sampling scheme for identifying S_2 . Consider some fixed mapping $h : [d] \to \{1,2\}$ that partitions [d] into $\mathcal{A}_1 = \{i \in [d] : h(i) = 1\}$ and $\mathcal{A}_2 = \{i \in [d] : h(i) = 2\}$. Then for a given Bernoulli vector $\mathcal{B} \in \{-1,1\}^d$ and $\mathbf{x} = (x_1 \dots x_d)^T \in [-1,1]^d$, consider the points $\mathbf{x}_1, \mathbf{x}_2, \mathbf{x}_3, \mathbf{x}_4 \in [-1,1]^d$ defined as

$$\begin{aligned}
x_{1,i} &= \begin{cases} \mathbbm{1}_{\beta_i} x_i \; ; & i \in \mathcal{A}_1, \\ \mathbbm{1}_{\beta_i} x_i \; ; & i \in \mathcal{A}_2, \end{cases} \quad x_{2,i} = \begin{cases} \mathbbm{1}_{\bar{\beta}_i} x_i \; ; & i \in \mathcal{A}_1, \\ \mathbbm{1}_{\beta_i} x_i \; ; & i \in \mathcal{A}_2, \end{cases} \\
x_{3,i} &= \begin{cases} \mathbbm{1}_{\beta_i} x_i \; ; & i \in \mathcal{A}_1, \\ \mathbbm{1}_{\bar{\beta}_i} x_i \; ; & i \in \mathcal{A}_2, \end{cases} \quad x_{4,i} = \begin{cases} \mathbbm{1}_{\bar{\beta}_i} x_i \; ; & i \in \mathcal{A}_1, \\ \mathbbm{1}_{\bar{\beta}_i} x_i \; ; & i \in \mathcal{A}_2, \end{cases} \quad i \in [d]. \end{aligned} \tag{4.2}
\end{aligned}$$

The following lemma is the key motivation behind our sampling scheme.

Lemma 2. Denote $\mathcal{A} = \left\{ \mathbf{j} \in {\binom{[d]}{2}} : \mathbf{j} \in \{\mathcal{A}_1 \times \mathcal{A}_2\} \cup \{\mathcal{A}_2 \times \mathcal{A}_1\} \right\}$. Then for functions f of the form (4.1), we have that

$$f(\mathbf{x}_{1}) - f(\mathbf{x}_{2}) - f(\mathbf{x}_{3}) + f(\mathbf{x}_{4}) = \sum_{\mathbf{j} \in \mathcal{S}_{2}: j_{1} \in \mathcal{A}_{1}, j_{2} \in \mathcal{A}_{2}} \beta_{j_{1}} \beta_{j_{2}} \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}) + \sum_{\mathbf{j} \in \mathcal{S}_{2}: j_{1} \in \mathcal{A}_{2}, j_{2} \in \mathcal{A}_{1}} \beta_{j_{1}} \beta_{j_{2}} \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}})$$

$$= \sum_{\mathbf{j} \in \mathcal{A} \cap \mathcal{S}_{2}} \beta_{j_{1}} \beta_{j_{2}} \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}).$$
(4.3)

Proof. For any $\mathbf{j} \in S_2$, let us first consider the case where j_1, j_2 lie in different sets. For example, let $j_1 \in \mathcal{A}_1$ and $j_2 \in \mathcal{A}_2$. Then the contribution of $\phi_{\mathbf{j}}$ to the left-hand side of (4.3) turns out to be for all possible values of $\beta_{j_1}, \beta_{j_2} \in \{-1, +1\}$ equal to

$$\begin{split} \phi_{\mathbf{j}}(\mathbf{x}_{1,\mathbf{j}}) &- \phi_{\mathbf{j}}(\mathbf{x}_{2,\mathbf{j}}) - \phi_{\mathbf{j}}(\mathbf{x}_{3,\mathbf{j}}) + \phi_{\mathbf{j}}(\mathbf{x}_{4,\mathbf{j}}) \\ &= \phi_{\mathbf{j}}(\mathbb{1}_{\beta_{j_1}}x_{j_1}, \mathbb{1}_{\beta_{j_2}}x_{j_2}) - \phi_{\mathbf{j}}(\mathbb{1}_{\bar{\beta}_{j_1}}x_{j_1}, \mathbb{1}_{\beta_{j_2}}x_{j_2}) - \phi_{\mathbf{j}}(\mathbb{1}_{\beta_{j_1}}x_{j_1}, \mathbb{1}_{\bar{\beta}_{j_2}}x_{j_2}) + \phi_{\mathbf{j}}(\mathbb{1}_{\bar{\beta}_{j_1}}x_{j_1}, \mathbb{1}_{\bar{\beta}_{j_2}}x_{j_2}) \\ &= \beta_{j_1}\beta_{j_2}(\phi_{\mathbf{j}}(x_{j_1}, x_{j_2}) - \phi_{\mathbf{j}}(x_{j_1}, 0) - \phi_{\mathbf{j}}(0, x_{j_2}) + \phi_{\mathbf{j}}(0, 0)) = \beta_{j_1}\beta_{j_2}\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}). \end{split}$$

In case $j_1 \in \mathcal{A}_2$ and $j_2 \in \mathcal{A}_1$, then the contribution of ϕ_j turns out to be the same as above. Since f is additive over $\mathbf{j} \in \mathcal{S}_2$, thus the total contribution of \mathcal{S}_2 is given by the right-hand side of (4.3).

Now, for all $\mathbf{j} \in S_2$ with j_1, j_2 lying in the same set, the contribution of $\phi_{\mathbf{j}}$ turns out to be zero. Indeed, if $j_1, j_2 \in \mathcal{A}_1$, the contribution of $\phi_{\mathbf{j}}$ is

$$\begin{split} \phi_{\mathbf{j}}(\mathbf{x}_{1,\mathbf{j}}) &- \phi_{\mathbf{j}}(\mathbf{x}_{2,\mathbf{j}}) - \phi_{\mathbf{j}}(\mathbf{x}_{3,\mathbf{j}}) + \phi_{\mathbf{j}}(\mathbf{x}_{4,\mathbf{j}}) \\ &= \phi_{\mathbf{j}}(\mathbbm{1}_{\beta_{j_1}}x_{j_1}, \mathbbm{1}_{\beta_{j_2}}x_{j_2}) - \phi_{\mathbf{j}}(\mathbbm{1}_{\bar{\beta}_{j_1}}x_{j_1}, \mathbbm{1}_{\bar{\beta}_{j_2}}x_{j_2}) - \phi_{\mathbf{j}}(\mathbbm{1}_{\beta_{j_1}}x_{j_1}, \mathbbm{1}_{\beta_{j_2}}x_{j_2}) + \phi_{\mathbf{j}}(\mathbbm{1}_{\bar{\beta}_{j_1}}x_{j_1}, \mathbbm{1}_{\bar{\beta}_{j_2}}x_{j_2}) = 0. \end{split}$$

The same is easily verified if $j_1, j_2 \in \mathcal{A}_2$. Lastly, let us verify that the contribution of ϕ_p for each $p \in \mathcal{S}_1 \cup \mathcal{S}_2^{\text{var}}$ is zero. Indeed, when $p \in \mathcal{A}_1$, we get

$$\phi_p(x_{1,p}) - \phi_p(x_{2,p}) - \phi_p(x_{3,p}) + \phi_p(x_{4,p}) = \phi_p(\mathbb{1}_{\beta_p} x_p) - \phi_p(\mathbb{1}_{\bar{\beta}_p} x_p) - \phi_p(\mathbb{1}_{\beta_p} x_p) + \phi_p(\mathbb{1}_{\bar{\beta}_p} x_p) = 0$$

and the same is true also for $p \in A_2$. This completes the proof.

Denoting $z_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}}) = \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}})$ if $\mathbf{j} \in S_2$ and 0 otherwise, let $\mathbf{z}^*(\mathbf{x}) \in \mathbb{R}^{\binom{[d]}{2}}$ be the corresponding $(|S_2|$ sparse) vector. For $\mathcal{A} \subseteq \binom{[d]}{2}$ we denote $\mathbf{z}^*(\mathbf{x}; \mathcal{A}) \in \mathbb{R}^{\binom{[d]}{2}}$ to be the *projection* of $\mathbf{z}^*(\mathbf{x})$ onto \mathcal{A} . Clearly $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ is at most $|S_2|$ sparse too – it is in fact $|S_2 \cap \mathcal{A}|$ sparse. For a Bernoulli vector $\mathcal{\beta} \in \{-1, +1\}^d$, let $\mathcal{\beta}^{(2)} \in \{-1, +1\}^{\binom{[d]}{2}}$, where $\beta_{\mathbf{j}}^{(2)} = \beta_{j_1}\beta_{j_2}$ for each $\mathbf{j} = (j_1, j_2)$. Hence we see that (4.3) corresponds to a linear measurement of $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ with the Bernoulli vector $\mathcal{\beta}^{(2)}$.

Sampling scheme for identifying S_2 . We first generate independent Bernoulli vectors $\beta_1, \beta_2, \ldots, \beta_n \in \{-1,1\}^d$. Then for some fixed $\mathbf{x} \in [-1,1]^d$ and a mapping $h : [d] \to \{1,2\}$ – the choice of both to be made clear later – we obtain the samples $\tilde{f}(\mathbf{x}_{i,p}) = f(\mathbf{x}_{i,p}) + \eta_{i,p}, i \in [n]$ and $p \in \{1,2,3,4\}$. Here, $\mathbf{x}_{i,1}, \mathbf{x}_{i,2}, \mathbf{x}_{i,3}, \mathbf{x}_{i,4}$ are generated using \mathbf{x}, β_i, h as outlined in (4.2). As a direct implication of Lemma 2, we obtain the linear system

$$\underbrace{\begin{pmatrix} \tilde{f}(\mathbf{x}_{1,1}) - \tilde{f}(\mathbf{x}_{1,2}) - \tilde{f}(\mathbf{x}_{1,3}) + \tilde{f}(\mathbf{x}_{1,4}) \\ \vdots \\ \tilde{f}(\mathbf{x}_{n,1}) - \tilde{f}(\mathbf{x}_{n,2}) - \tilde{f}(\mathbf{x}_{n,3}) + \tilde{f}(\mathbf{x}_{n,4}) \end{pmatrix}}_{\mathbf{y} \in \mathbb{R}^{n}} = \underbrace{\begin{pmatrix} \boldsymbol{\beta}_{1}^{(2)T} \\ \vdots \\ \vdots \\ \boldsymbol{\beta}_{n}^{(2)T} \\ \boldsymbol{\beta}_{n}^{(2)T} \end{pmatrix}}_{\mathbf{B} \in \mathbb{R}^{n \times \binom{d}{2}}} \mathbf{z}^{*}(\mathbf{x}; \mathcal{A}) + \underbrace{\begin{pmatrix} \eta_{1,1} - \eta_{1,2} - \eta_{1,3} + \eta_{1,4} \\ \vdots \\ \eta_{n,1} - \eta_{n,2} - \eta_{n,3} + \eta_{n,4} \end{pmatrix}}_{\boldsymbol{\eta} \in \mathbb{R}^{n}}.$$

$$(4.4)$$

By feeding \mathbf{y} , \mathbf{B} as input to SPARSE-REC, we then obtain the estimate $\widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A})$ to $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$. Assuming SPARSE-REC to be ϵ -accurate at \mathbf{x} , we will have that $\| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \mathbf{z}^*(\mathbf{x}; \mathcal{A}) \|_{\infty} \leq \epsilon$ holds. Let us mention that \mathcal{A} from Lemma 2 is completely determined by h but we avoid denoting this explicitly for clarity of notation.

At this point, it is natural to ask, how one should choose \mathbf{x} and the mapping h. To this end, we borrow the approach of [11], which involves choosing h from a family of hash functions, and creating for each h in the family a uniform grid. To begin with, we introduce the following definition of a family of hash functions.

Definition 1. For some $t \in \mathbb{N}$ and $j = 1, 2, ..., let h_j : [d] \to \{1, 2, ..., t\}$. We call the set $\mathcal{H}_t^d = \{h_1, h_2, ...\}$ a (d, t)-hash family if for any distinct $i_1, i_2, ..., i_t \in [d]$, there exists $h \in \mathcal{H}_t^d$ such that h is an injection when restricted to $i_1, i_2, ..., i_t$.

Hash functions are commonly used in theoretical computer science and are widely used in finding juntas [25]. One can construct \mathcal{H}_t^d of size $O(te^t \log d)$ using a standard probabilistic argument. The reader is for instance referred to Section 5 in [11], where for any constant $C_1 > 1$ the probabilistic

construction yields \mathcal{H}_t^d of size $|\mathcal{H}_t^d| \leq (C_1 + 1)te^t \log d$ with probability at least $1 - d^{-C_1 t}$, in time linear in the output size.

Focusing on the setting t = 2 now, say we have at hand a family \mathcal{H}_2^d of size $O(\log d)$. Then for any $(i, j) \in \binom{[d]}{2}$, there exists $h \in \mathcal{H}_2^d$ so that $h(i) \neq h(j)$. For each $h \in \mathcal{H}_2^d$, let us define $\mathbf{e}_1(h), \mathbf{e}_2(h) \in \mathbb{R}^d$, where

$$(\mathbf{e}_i(h))_q := \begin{cases} 1 ; & h(q) = i, \\ 0 ; & \text{otherwise} \end{cases} \quad \text{for } i = 1, 2 \text{ and } q \in [d].$$

Then we create a two dimensional grid with respect to h

$$\chi(h) := \left\{ \mathbf{x} \in [-1,1]^d : \mathbf{x} = c_1 \mathbf{e}_1(h) + c_2 \mathbf{e}_2(h); c_1, c_2 \in \left\{ -1, -\frac{m-1}{m}, \dots, \frac{m-1}{m}, 1 \right\} \right\}.$$
 (4.5)

Equipped with $\chi(h)$ for each $h \in \mathcal{H}_2^d$, we now possess the following approximation property. For any $\mathbf{j} \in {\binom{[d]}{2}}$ and any $(x_{j_1}^*, x_{j_2}^*) \in [-1, 1]^2$, there exists $h \in \mathcal{H}_2^d$ with $h(j_1) \neq h(j_2)$ and a corresponding $\mathbf{x} \in \chi(h)$ so that $|x_{j_1}^* - x_{j_1}|, |x_{j_2}^* - x_{j_2}| \leq 1/m$. Informally speaking, our idea is the following. Assume that SPARSE-REC is ϵ -accurate for each

Informally speaking, our idea is the following. Assume that SPARSE-REC is ϵ -accurate for each $h \in \mathcal{H}_2^d$, $\mathbf{x} \in \chi(h)$. Also, say m, ϵ are sufficiently large and small respectively. Hence, if we estimate $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ at each $h \in \mathcal{H}_2^d$ and $\mathbf{x} \in \chi(h)$, then for every $\mathbf{j} \in \mathcal{S}_2$, we are guaranteed to have a point \mathbf{x} at which the estimate $|\widehat{z}_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}})|$ is sufficiently large. Moreover, for every $\mathbf{j} \notin \mathcal{S}_2$, we would always (i.e., for each $h \in \mathcal{H}_2^d$ and $\mathbf{x} \in \chi(h)$) have $|\widehat{z}_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}})|$ sufficiently small; more precisely, $|\widehat{z}_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}})| \leq \epsilon$ since $\phi_{\mathbf{j}} \equiv 0$. Consequently, we will be able to identify \mathcal{S}_2 by thresholding, via a suitable threshold.

Sampling scheme for identifying S_1 . Assuming S_2 is identified, the model (4.1) reduces to the univariate case on the reduced set $\mathcal{P} := [d] \setminus S_2^{\text{var}}$ with $S_1 \subset \mathcal{P}$. We can therefore apply Algorithm 1 on \mathcal{P} by setting the coordinates in $\mathcal{P}^c = \widetilde{S_2^{\text{var}}}$ to zero. Indeed, we first construct for some $m \in \mathbb{N}$ the following set

$$\chi = \left\{ (c \ c \ \dots \ c)^T \in \mathbb{R}^{\mathcal{P}} : c \in \left\{ -1, -\frac{m-1}{m}, \dots, \frac{m-1}{m}, 1 \right\} \right\} \subset [-1, 1]^{\mathcal{P}}.$$
 (4.6)

Then, for any given $\boldsymbol{\beta} \in \{-1, 1\}^{\mathcal{P}}$, and $\mathbf{x} \in \chi$, we construct $\mathbf{x}^+, \mathbf{x}^- \in \mathbb{R}^d$ using $\boldsymbol{\beta}, \mathbf{x}$ as follows

$$x_i^+ = \begin{cases} x_i ; & \beta_i = +1 \text{ and } i \in \mathcal{P}, \\ 0 ; & \text{otherwise,} \end{cases} \quad x_i^- = \begin{cases} x_i ; & \beta_i = -1 \text{ and } i \in \mathcal{P}, \\ 0 ; & \text{otherwise,} \end{cases} \quad i \in [d].$$
(4.7)

Note that $x_i^+, x_i^- = 0$ for $i \notin \mathcal{P}$. Then, similarly to (3.3), we have that

$$\tilde{f}(\mathbf{x}^+) - \tilde{f}(\mathbf{x}^-) = \sum_{i \in \mathcal{P}} \beta_i \underbrace{\phi_i(x_i)}_{z_i^*(x_i)} + \eta^+ - \eta^- = \langle \boldsymbol{\beta}, \mathbf{z}_{\mathcal{P}}^*(\mathbf{x}) \rangle + \eta^+ - \eta^-, \qquad (4.8)$$

where $\mathbf{z}_{\mathcal{P}}^*(\mathbf{x}) \in \mathbb{R}^{\mathcal{P}}$ is the restriction of $\mathbf{z}^*(\mathbf{x})$ onto \mathcal{P} , and is $|\mathcal{S}_1|$ sparse. Thereafter, we proceed as in Algorithm 1 by forming a linear system as in (3.4) (where now $\mathbf{B} \in \mathbb{R}^{n \times |\mathcal{P}|}$) at each $\mathbf{x} \in \chi$, and employing an ϵ -accurate SPARSE-REC to estimate $\mathbf{z}_{\mathcal{P}}^*(\mathbf{x})$.

Algorithm outline and guarantees. Our scheme for identifying S_2 is outlined formally as the first part of Algorithm 2. The second part involves the estimation of S_1 . Lemma 3 provides exact recovery guarantees for S_2 and S_1 by Algorithm 2.

Algorithm 2 Algorithm for estimating S_2, S_1

// Estimation of S_2 1: Input: $d, |S_2|, m_2, n_2, \epsilon_2$. 2: Initialization: $\widehat{\mathcal{S}_2} = \emptyset$. 3: Output: $\widehat{\mathcal{S}_2}$. 4: 5: Generate independent Bernoulli vectors $\beta_1, \beta_2, \ldots, \beta_{n_2} \in \{-1, +1\}^d$. 6: Form $\mathbf{B} \in \mathbb{R}^{n_2 \times \binom{d}{2}}$ as in (4.4). 7: Construct a (d, 2) hash family: \mathcal{H}_2^d . 8: for $h \in \mathcal{H}_2^d$ do Construct $\chi(h)$ as defined in (4.5) with $|\chi(h)| = (2m_2 + 1)^2$. 9: for $\mathbf{x} \in \chi(h)$ do 10:Generate $\mathbf{x}_{i,1}, \mathbf{x}_{i,2}, \mathbf{x}_{i,3}, \mathbf{x}_{i,4} \in [-1, 1]^d$, as in (4.2), using $\mathbf{x}, \boldsymbol{\beta}_i$ for each $i \in [n_2]$. 11: Using the samples $(\tilde{f}(\mathbf{x}_{i,1}), \tilde{f}(\mathbf{x}_{i,2}), \tilde{f}(\mathbf{x}_{i,3}), \tilde{f}(\mathbf{x}_{i,4}))_{i=1}^{n_2}$, form **y** as in (4.4). 12:Obtain $\widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) = \text{SPARSE-REC}_2(\mathbf{y}, \mathbf{B}).$ 13:Update $\widehat{\mathcal{S}}_2 = \widehat{\mathcal{S}}_2 \cup \left\{ \mathbf{j} \in \mathcal{A} : |\widehat{z_{\mathbf{j}}^*}(\mathbf{x}_{\mathbf{j}})| > \epsilon_2 \right\}.$ 14: end for 15:16: end for 17:18: Input: d, $|S_1|$, $\widehat{S_2^{\text{var}}}$, m_1 , n_1 , ϵ_1 . 19: Initialization: $\widehat{S}_1 = \emptyset$, $\mathcal{P} = [d] \setminus \widehat{S_2^{\text{var}}}$. // Estimation of S_1 20: Output: $\widehat{\mathcal{S}_1}$. 21: Construct $\chi \subset [-1, 1]^{\mathcal{P}}$ with $|\chi| = 2m_1 + 1$, as in (4.6). 22: Generate independent Bernoulli vectors $\beta_1, \beta_2, \ldots, \beta_{n_1} \in \{-1, +1\}^{\mathcal{P}}$. 23: Form $\mathbf{B} \in \mathbb{R}^{n_1 \times |\mathcal{P}|}$ as in (3.4). 24:for $\mathbf{x} \in \chi$ do Generate $\mathbf{x}_i^+, \mathbf{x}_i^- \in [-1, 1]^d$, as in (4.7), using $\mathbf{x}, \boldsymbol{\beta}_i$ for each $i \in [n_1]$. 25:Using the samples $(\tilde{f}(\mathbf{x}_i^+), \tilde{f}(\mathbf{x}_i^-))_{i=1}^{n_1}$, form **y** as in (3.4). 26:Obtain $\widehat{\mathbf{z}^*}(\mathbf{x}) = \text{SPARSE-REC}_1(\mathbf{y}, \mathbf{B})$ where $\widehat{\mathbf{z}^*}(\mathbf{x}) \in \mathbb{R}^{\mathcal{P}}$. 27:Update $\widehat{\mathcal{S}}_1 = \widehat{\mathcal{S}}_1 \cup \left\{ p \in \mathcal{P} : |(\widehat{\mathbf{z}^*}(\mathbf{x}))_p| > \epsilon_1 \right\}.$ 28:29: end for

Lemma 3. Let \mathcal{H}_2^d be a (d, 2) hash family, and let $SPARSE-REC_2$ be ϵ_2 -accurate for each $h \in \mathcal{H}_2^d$, $\mathbf{x} \in \chi(h)$ with $\epsilon_2 < D_2/3$, which uses n_2 linear measurements. If $m_2 \ge \sqrt{2} \left(\frac{3L}{D_2}\right)^{1/\alpha}$, then Algorithm 2 recovers S_2 exactly, i.e., $\widehat{S_2} = S_2$. Moreover, assuming $\widehat{S_2} = S_2$ holds, and $SPARSE-REC_1$ is ϵ_1 -accurate (using n_1 measurements), then if m_1, n_1, ϵ_1 satisfy the conditions of Lemma 1, we have $\widehat{S_1} = S_1$. Lastly, the total number of queries of f made is $4(2m_2+1)^2n_2|\mathcal{H}_2^d| + 2(2m_1+1)n_1$.

Proof. For any given $\mathbf{j} \in S_2$ there exists $\mathbf{x}_{\mathbf{j}}^* \in [-1, 1]^2$ with $|\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}^*)| \geq D_2$. Moreover, since \mathcal{H}_2^d is a (d, 2) hash family, there exists $h \in \mathcal{H}_2^d$ that is an injection on \mathbf{j} . Consequently, there exists $\mathbf{x} \in \chi(h)$ such that $\|\mathbf{x}_{\mathbf{j}} - \mathbf{x}_{\mathbf{j}}^*\|_2 \leq \frac{\sqrt{2}}{m_2}$. This in turn implies by Hölder continuity of $\phi_{\mathbf{j}}$ that

$$|\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}) - \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}^*)| \le L \frac{2^{\alpha/2}}{m_2^{\alpha}}.$$
(4.9)

Since SPARSE-REC₂ is ϵ_2 -accurate for each $h \in \mathcal{H}_2^d$, $\mathbf{x} \in \chi(h)$, we know that at the aforementioned

 \mathbf{x} , the following holds via reverse triangle inequality

$$|\widehat{z_{\mathbf{j}}^*}(\mathbf{x}_{\mathbf{j}})| \ge |\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}})| - \epsilon_2.$$
(4.10)

Using (4.9), (4.10) and the reverse triangle inequality, we get by the choice of ϵ_2 and m_2

$$|\widehat{z_{\mathbf{j}}^{*}}(\mathbf{x}_{\mathbf{j}})| \ge |\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}^{*})| - L\frac{2^{\alpha/2}}{m_{2}^{\alpha}} - \epsilon_{2} \ge D_{2} - L\frac{2^{\alpha/2}}{m_{2}^{\alpha}} - \epsilon_{2} \ge \frac{2D_{2}}{3} - L\frac{2^{\alpha/2}}{m_{2}^{\alpha}} \ge \frac{D_{2}}{3}.$$

Also, for any $\mathbf{j} \notin S_2$, we have for all $h \in \mathcal{H}_2^d$, $\mathbf{x} \in \chi(h)$ that $|\widehat{z}_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}})| \leq \epsilon_2 < D_2/3$ (since $\phi_{\mathbf{j}} \equiv 0$). Hence, the stated choice of ϵ_2 guarantees identification of each $\mathbf{j} \in S_2$, and none from $\binom{[d]}{2} \setminus S_2$. The proof for recovery of S_1 is identical to Lemma 1, and hence omitted.

Remark 2. On a top level, Algorithm 2 is similar to [38, Algorithms 3,4] in the sense that they all involve solving ℓ_1 minimization problems at base points lying in $\chi(h)$ defined in (4.5) (for identification of S_2), and χ defined in (4.6) (for identification of S_1). The difference however lies in the nature of the sampling schemes. The scheme in [38, Algorithms 3,4] relies on estimating sparse Hessians, gradients of f via their linear measurements, through random samples in the neighborhood of the base point. In contrast, the sampling scheme in Algorithm 2 is not local; for instance during the identification of S_2 , at each base point $\mathbf{x} \in \chi(h)$, the points $\mathbf{x}_{i,1}, \mathbf{x}_{i,2}, \mathbf{x}_{i,3}, \mathbf{x}_{i,4}$ for any given $i \in [n_2]$ can be arbitrarily far from each other. The same is true during the identification of S_1 .

5 The multivariate case

Finally, we treat also the general case where f consists of at most r_0 -variate components, where $r_0 > 2$ is possible. To begin with, let $S_1, S_2, \ldots, S_{r_0}$ be such that $S_r \subset {\binom{[d]}{r}}$ for $r \in [r_0]$. Here S_r represents the r wise interaction terms. We now need some additional notation and assumptions.

- 1. For $r \geq 1$, let $\mathcal{S}_r^{(1)}$ denote the set of variables occurring in \mathcal{S}_r with $\mathcal{S}_1^{(1)} = \mathcal{S}_1$. We assume that $\mathcal{S}_p^{(1)} \cap \mathcal{S}_q^{(1)} = \emptyset$ for all $p \neq q \in [r_0]$.
- 2. For each $1 \le i < r \le r_0$, denote $\mathcal{S}_r^{(i)} = {\mathcal{S}_r^{(1)} \choose i}$ to be the sets of i^{th} order tuples induced by \mathcal{S}_r .

The multivariate analogue of Proposition 2 is provided by the following result.

Proposition 3. Let $1 \le r_0 \le d$ and let $f \in C([-1,1]^d)$ be of the form

$$f(\mathbf{x}) = \mu + \sum_{j \in \bigcup_{r=2}^{r_0} S_r^{(1)} \cup S_1} \phi_j(x_j) + \sum_{\mathbf{j} \in \bigcup_{r=3}^{r_0} S_r^{(2)} \cup S_2} \phi_{\mathbf{j}}(x_{\mathbf{j}}) + \dots + \sum_{\mathbf{j} \in S_{r_0}^{(r_0-1)} \cup S_{r_0-1}} \phi_{\mathbf{j}}(x_{\mathbf{j}}) + \sum_{\mathbf{j} \in S_{r_0}} \phi_{\mathbf{j}}(x_{\mathbf{j}}),$$
(5.1)

where all the functions ϕ_i are not identically zero. Moreover, let

- (a) $\mu = f(0)$.
- (b) For each $1 \le l \le r_0 1$, $\phi_{\mathbf{j}}(x_{\mathbf{j}}) = 0$ if $\mathbf{j} = (j_1, \dots, j_l) \in \bigcup_{r=l+1}^{r_0} \mathcal{S}_r^{(l)} \cup \mathcal{S}_l$, and $x_{j_i} = 0$ for some $i \in [l]$.

(c) $\phi_j(0) = 0$ if $\mathbf{j} = (j_1, \dots, j_{r_0}) \in \mathcal{S}_{r_0}$, and $x_{j_i} = 0$ for some $i \in [r_0]$.

Then the representation (5.1) of f is unique in the sense that each component in (5.1) is uniquely identifiable.

The proof of this result is similar to the proof of Proposition 2, and we leave it to the reader.

Remark 3. Let us note that for the special cases $r_0 \in \{1, 2\}$, the statement of Proposition 3 reduces to that of Proposition 2 for univariate/bivariate SPAMs. If $r_0 = 2$, we observed earlier in Section 4 that the assumption $S_1 \cap S_2^{(1)} = \emptyset$ can be made without loss of generality. However, for $r_0 > 2$, this is an additional assumption. It will allow to structure the recovery algorithm into recursive steps.

Assumptions. We will make the following assumptions on the model (5.1).

1. Smoothness. Each term in (5.1) is Hölder continuous with parameters $L > 0, \alpha \in (0, 1]$, i.e., for each $i \in [r_0]$,

$$|\phi_{\mathbf{j}}(\mathbf{x}) - \phi_{\mathbf{j}}(\mathbf{y})| \le L \parallel \mathbf{x} - \mathbf{y} \parallel_2^{\alpha} \quad \text{for all } \mathbf{j} \in \mathcal{S}_i \cup \bigcup_{l=i+1}^{r_0} \mathcal{S}_l^{(i)} \text{ and for all } \mathbf{x}, \mathbf{y} \in [-1, 1]^i.$$
(5.2)

2. Identifiability of S_i , $i \in [r_0]$. We assume that for each $i \in [r_0]$ there exists a constant $D_i > 0$, such that for every $\mathbf{j} \in S_i$ there exists $\mathbf{x}_{\mathbf{j}}^* \in [-1, 1]^i$ with $|\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}^*)| > D_i$.

We now generalize the sampling scheme given before for bivariate components to the setting of multivariate components. Let us fix some mapping $h : [d] \to [r_0]$ that partitions [d] into $\mathcal{A}_1 = \{i \in [d] : h(i) = 1\}, \mathcal{A}_2 = \{i \in [d] : h(i) = 2\}, \ldots, \mathcal{A}_{r_0} = \{i \in [d] : h(i) = r_0\}$. Let us fix a Bernoulli vector $\boldsymbol{\beta} \in \{-1, 1\}^d$ and $\mathbf{x} = (x_1 \dots x_d)^T \in [-1, 1]^d$. For $z \in [2^{r_0}]$ and $i \in [d]$, we define

$$(\mathbf{x}_z)_i = x_{z,i} = \begin{cases} x_i & \text{if } \beta_i = (-1)^{\text{digit}(z-1,h(i)-1)}, \\ 0 & \text{otherwise}, \end{cases}$$
(5.3)

where $\operatorname{digit}(a, b) \in \{0, 1\}$ is the b^{th} digit of the dyadic decomposition of a for $a, b \in \mathbb{N}_0$. Furthermore, we denote by $\operatorname{digit}(a)$ the sum of digits of $a \in \mathbb{N}_0$, i.e.

$$a = \sum_{i=0}^\infty \operatorname{digit}(a,i) \, \cdot \, 2^i, \qquad \operatorname{digit}(a) = \sum_{i=0}^\infty \operatorname{digit}(a,i).$$

Remark 4. It $r_0 = 1$, it is easily verified, that the points $(\mathbf{x}_z)_{z=1}^2$ in (5.3) coincide with the points $\mathbf{x}^+, \mathbf{x}^-$ defined in (3.2) for univariate SPAMs. Similarly, for $r_0 = 2$, the points $(\mathbf{x}_z)_{z=1}^4$ from (5.3) agree with those defined in (4.2) for bivariate SPAMs. In the same way, the following lemma is a generalization of (3.3) and Lemma 2 for (5.1). Indeed, if $r_0 = 1$, there is only one mapping $h: [d] \to \{1\}$, and so $\mathcal{A} = [d]$.

Lemma 4. Denote $\mathcal{A} = \left\{ \mathbf{j} \in {\binom{[d]}{r_0}} : h \text{ is injective on } \{j_1, \ldots, j_{r_0}\} \right\}$. Then for functions f of the form (5.1), we have that

$$\sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} f(\mathbf{x}_z) = \sum_{\mathbf{j} \in \mathcal{A} \cap \mathcal{S}_{r_0}} \beta_{j_1} \dots \beta_{j_{r_0}} \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}).$$
(5.4)

Proof. We plug (5.1) into the left-hand side of (5.4) and obtain

$$\begin{split} &\sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} f(\mathbf{x}_z) \\ &= \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} \Big[\mu + \sum_{j \in \bigcup_{r=2}^{r_0} S_r^{(1)} \cup S_1} \phi_j(x_{z,j}) + \sum_{\mathbf{j} \in \bigcup_{r=3}^{r_0} S_r^{(2)} \cup S_2} \phi_{\mathbf{j}}((\mathbf{x}_z)_{\mathbf{j}}) + \dots + \sum_{\mathbf{j} \in S_{r_0}} \phi_{\mathbf{j}}((\mathbf{x}_z)_{\mathbf{j}}) \Big] \\ &= \mu \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} + \sum_{j \in \bigcup_{r=2}^{r_0} S_r^{(1)} \cup S_1} \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} \phi_j(x_{z,j}) + \dots + \sum_{\mathbf{j} \in S_{r_0}} \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} \phi_{\mathbf{j}}((\mathbf{x}_z)_{\mathbf{j}}) \\ &= I_0 + I_1 + \dots + I_{r_0}. \end{split}$$

We show first that $I_0 = I_1 = \cdots = I_{r_0-1} = 0$. Indeed,

$$I_0 = \mu \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} = \mu \sum_{z=1}^{2^{r_0-1}} \left((-1)^{\operatorname{digit}(2z-2)} + (-1)^{\operatorname{digit}(2z-1)} \right)$$

and the last expression vanishes as $\operatorname{digit}(2z-1) = \operatorname{digit}(2z-2) + 1$ for every $z \in [2^{r_0-1}]$.

If $j \in \bigcup_{r=2}^{r_0} \mathcal{S}_r^{(1)} \cup \mathcal{S}_1$, we define the set $U_j = \left\{z \in [2^{r_0}] : \beta_j = (-1)^{\operatorname{digit}(z-1,h(j)-1)}\right\}$ and write

$$\begin{split} I_1 &= \sum_{j \in \bigcup_{r=2}^{r_0} \mathcal{S}_r^{(1)} \cup \mathcal{S}_1} \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} \phi_j(x_{z,j}) = \sum_{j \in \bigcup_{r=2}^{r_0} \mathcal{S}_r^{(1)} \cup \mathcal{S}_1} \phi_j(x_j) \sum_{z \in [2^{r_0}] \atop \beta_j = (-1)^{\operatorname{digit}(z-1,h(j)-1)}} (-1)^{\operatorname{digit}(z-1)} \\ &= \sum_{j \in \bigcup_{r=2}^{r_0} \mathcal{S}_r^{(1)} \cup \mathcal{S}_1} \phi_j(x_j) \sum_{z \in U_j} (-1)^{\operatorname{digit}(z-1)}. \end{split}$$

Each of the sums over U_j contains 2^{r_0-1} number of summands, half of which is equal to 1 and the other half to -1. Therefore also $I_1 = 0$. Similarly, if $\mathbf{j} = (j_1, j_2) \in \bigcup_{r=3}^{r_0} \mathcal{S}_r^{(2)} \cup \mathcal{S}_2$, we set

$$U_{\mathbf{j}} = \left\{ z \in [2^{r_0}] : \beta_{j_1} = (-1)^{\operatorname{digit}(z-1,h(j_1)-1)} \text{ and } \beta_{j_2} = (-1)^{\operatorname{digit}(z-1,h(j_2)-1)} \right\}$$

and obtain

$$I_2 = \sum_{\mathbf{j} \in \bigcup_{r=3}^{r_0} \mathcal{S}_r^{(2)} \cup \mathcal{S}_2} \phi_{\mathbf{j}}((\mathbf{x}_z)_{\mathbf{j}}) \sum_{z \in U_{\mathbf{j}}} (-1)^{\mathtt{digit}(z-1)}.$$

Half of the terms in the last sum is again equal to 1, the other half being equal to -1. Hence, $I_2 = 0$. Arguing along the same lines, we conclude $I_0 = I_1 = \cdots = I_{r_0-1} = 0$.

Finally, if $\mathbf{j} = (j_1, \ldots, j_{r_0}) \in \mathcal{S}_{r_0}$, we define

$$U_{\mathbf{j}} = \left\{ z \in [2^{r_0}] : \beta_{j_i} = (-1)^{\texttt{digit}(z-1,h(j_i)-1)} \text{ for all } i \in [r_0] \right\}.$$

If h is an injection on $\{j_1, \ldots, j_{r_0}\}$, we get that $\{h(j_1), \ldots, h(j_{r_0})\} = [r_0]$ and $U_{\mathbf{j}} = \{z^{\mathbf{j}}\}$ is a singleton with

$$\sum_{z \in U_{\mathbf{j}}} (-1)^{\mathtt{digit}(z-1)} = (-1)^{\mathtt{digit}(z^{\mathbf{j}}-1)} = \prod_{i=1}^{r_0} (-1)^{\mathtt{digit}(z^{\mathbf{j}}-1,i-1)} = \prod_{i=1}^{r_0} (-1)^{\mathtt{digit}(z^{\mathbf{j}}-1,h(j_i)-1)} = \prod_{i=1}^{r_0} \beta_{j_i}.$$

If, on the other hand, h is no injection on $\{j_1, \ldots, j_{r_0}\}$, U_j has even number of elements and using the same argument as above we obtain

$$\sum_{z \in U_{\mathbf{i}}} (-1)^{\operatorname{digit}(z-1)} = 0.$$

We conclude that

$$\begin{split} \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} f(\mathbf{x}_z) &= I_0 + I_1 + \dots + I_{r_0} = \sum_{\mathbf{j} \in \mathcal{S}_{r_0}} \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}) \sum_{z \in U_{\mathbf{j}}} (-1)^{\operatorname{digit}(z-1)} \\ &= \sum_{\mathbf{j} \in \mathcal{A} \cap \mathcal{S}_{r_0}} \beta_{j_1} \dots \beta_{j_{r_0}} \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}). \end{split}$$

We denote again $z_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}}) = \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}})$ if $\mathbf{j} \in S_{r_0}$ and 0 otherwise. Similarly, $\mathbf{z}^*(\mathbf{x}) \in \mathbb{R}^{\binom{[d]}{r_0}}$ stands for the corresponding $|S_{r_0}|$ -sparse vector and $\mathbf{z}^*(\mathbf{x}; \mathcal{A}) \in \mathbb{R}^{\binom{[d]}{r_0}}$ for the projection of $\mathbf{z}^*(\mathbf{x})$ onto \mathcal{A} . Again, $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ is $|S_{r_0} \cap \mathcal{A}|$ -sparse. Finally, for a Bernoulli vector $\boldsymbol{\beta} \in \{-1, +1\}^d$, let $\boldsymbol{\beta}^{(r_0)} \in \{-1, +1\}^{\binom{[d]}{r_0}}$ where $\beta_{\mathbf{j}}^{(r_0)} = \beta_{j_1}\beta_{j_2}\dots\beta_{j_{r_0}}$ for each $\mathbf{j} = (j_1, j_2, \dots, j_{r_0})$. Hence, (5.4) corresponds to a linear measurement of $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ with $\boldsymbol{\beta}^{(r_0)}$.

Sampling scheme for identifying S_{r_0} . Similarly to the sampling scheme for identifying S_2 in the bivariate case, we generate independent Bernoulli vectors $\beta_1, \beta_2, \ldots, \beta_n \in \{-1, 1\}^d$. For fixed $\mathbf{x} \in [-1, 1]^d$ and $h : [d] \to [r_0]$, we obtain the samples $\tilde{f}(\mathbf{x}_{i,z}) = f(\mathbf{x}_{i,z}) + \eta_{i,z}$, where $i \in [n]$ and $z \in [2^{r_0}]$. Here, $\mathbf{x}_{i,z}$ are generated using \mathbf{x}_i, β_i, h as outlined in (5.3).

As a direct implication of Lemma 4, we obtain the linear system

$$\underbrace{\begin{pmatrix} \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} \tilde{f}(\mathbf{x}_{1,z}) \\ \vdots \\ \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} \tilde{f}(\mathbf{x}_{n,z}) \end{pmatrix}}_{\mathbf{y} \in \mathbb{R}^n} = \underbrace{\begin{pmatrix} \boldsymbol{\beta}_1^{(r_0)^T} \\ \vdots \\ \vdots \\ \boldsymbol{\beta}_n^{(r_0)^T} \end{pmatrix}}_{\mathbf{B} \in \mathbb{R}^{n \times \binom{d}{r_0}}} \mathbf{z}^*(\mathbf{x}; \mathcal{A}) + \underbrace{\begin{pmatrix} \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} \eta_{1,z} \\ \vdots \\ \sum_{z=1}^{2^{r_0}} (-1)^{\operatorname{digit}(z-1)} \eta_{n,z} \end{pmatrix}}_{\boldsymbol{\eta} \in \mathbb{R}^n}.$$
(5.5)

By feeding \mathbf{y}, \mathbf{B} as input to SPARSE-REC, we will obtain the estimate $\widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A})$ to $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$. Assuming SPARSE-REC to be ϵ -accurate at \mathbf{x} , we will have that $\| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \mathbf{z}^*(\mathbf{x}; \mathcal{A}) \|_{\infty} \leq \epsilon$ holds.

The choice of \mathbf{x}, h is along similar lines as in the previous section. Indeed we first construct a (d, r_0) hash family $\mathcal{H}_{r_0}^d$ so that for any $\mathbf{j} = (j_1, \ldots, j_{r_0}) \in {\binom{[d]}{r_0}}$, there exists $h \in \mathcal{H}_{r_0}^d$ which is injective on [**j**]. For each $h \in \mathcal{H}_{r_0}^d$, let us define $\mathbf{e}_1(h), \mathbf{e}_2(h), \ldots, \mathbf{e}_{r_0}(h) \in \mathbb{R}^d$, where

$$(\mathbf{e}_i(h))_q := \begin{cases} 1 ; & h(q) = i, \\ 0 ; & \text{otherwise} \end{cases} \text{ for } i \in [r_0] \text{ and } q \in [d].$$

We then create the following r_0 dimensional grid with respect to h.

$$\chi(h) := \left\{ \mathbf{x} \in [-1,1]^d : \mathbf{x} = \sum_{i=1}^{r_0} c_i \mathbf{e}_i(h); c_1, c_2, \dots, c_{r_0} \in \left\{ -1, -\frac{m-1}{m}, \dots, \frac{m-1}{m}, 1 \right\} \right\}.$$
 (5.6)

Equipped with $\chi(h)$ for each $h \in \mathcal{H}_{r_0}^d$, we now possess the following approximation property. For any $\mathbf{j} \in {\binom{[d]}{r_0}}$ and any $(x_{j_1}^*, x_{j_2}^*, \dots, x_{j_{r_0}}^*) \in [-1, 1]^{r_0}$, there exists $h \in \mathcal{H}_{r_0}^d$ and a corresponding $\mathbf{x} \in \chi(h)$ so that $|x_{j_1}^* - x_{j_1}|, |x_{j_2}^* - x_{j_2}|, \dots, |x_{j_{r_0}}^* - x_{j_{r_0}}| \leq 1/m$. Here on, our idea for estimating \mathcal{S}_{r_0} is based on the same principle that we followed in the

Here on, our idea for estimating S_{r_0} is based on the same principle that we followed in the preceding section. Assume that SPARSE-REC is ϵ accurate for each $h \in \mathcal{H}_{r_0}^d$, $\mathbf{x} \in \chi(h)$, and that m, ϵ are sufficiently large and small respectively. Hence, if we estimate $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ at each $h \in \mathcal{H}_{r_0}^d$ and $\mathbf{x} \in \chi(h)$, then for every $\mathbf{j} \in S_{r_0}$ we are guaranteed to have a point \mathbf{x} at which the estimate $|\hat{z}_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}})|$ is sufficiently large. Moreover, for every $\mathbf{j} \notin S_{r_0}$, we would always (i.e., for each $h \in \mathcal{H}_{r_0}^d$, $\mathbf{x} \in \chi(h)$) have that $|\hat{z}_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}})|$ is sufficiently small; more precisely, $|\hat{z}_{\mathbf{j}}^*(\mathbf{x}_{\mathbf{j}})| \leq \epsilon$ since $\phi_{\mathbf{j}} \equiv 0$. Consequently, we will be able to identify S_{r_0} by thresholding, via a suitable threshold.

Sampling scheme for identifying S_{r_0-1} . Say we have an estimate for S_{r_0} , lets call it $\widehat{S_{r_0}}$, and assume $\widehat{S_{r_0}}$ was identified correctly, so $\widehat{S_{r_0}} = S_{r_0}$. Then, we now have a SPAM of order $r_0 - 1$ on the reduced set of variables $\mathcal{P} = [d] \setminus \widehat{S_{r_0}^{(1)}}$. Therefore, in order to estimate S_{r_0-1} , we simply repeat the above procedure on the reduced set \mathcal{P} by freezing the variables in $\widehat{S_{r_0}^{(1)}}$ to 0. More precisely, we have the following steps.

- We will construct a $(\mathcal{P}, r_0 1)$ hash family $\mathcal{H}_{r_0-1}^{\mathcal{P}}$, hence each $h \in \mathcal{H}_{r_0-1}^{\mathcal{P}}$ is a mapping $h : \mathcal{P} \to [r_0 1]$.
- For each $h \in \mathcal{H}_{r_0-1}^{\mathcal{P}}$, define $\mathbf{e}_1(h), \mathbf{e}_2(h), \dots, \mathbf{e}_{r_0-1}(h) \in \mathbb{R}^{\mathcal{P}}$, where

$$(\mathbf{e}_i(h))_q := \begin{cases} 1 ; & h(q) = i \text{ and } q \in \mathcal{P}, \\ 0 ; & \text{otherwise,} \end{cases} \quad \text{for } i \in [r_0 - 1] \text{ and } q \in \mathcal{P}, \end{cases}$$

and use $(\mathbf{e}_i(h))_{i=1}^{r_0-1}$ to create a r_0-1 dimensional grid $\chi(h) \subset [-1,1]^{\mathcal{P}}$ in the same manner as in (5.6).

• For $h \in \mathcal{H}_{r_0-1}^{\mathcal{P}}$, a Bernoulli vector $\boldsymbol{\beta} \in \{-1,1\}^{\mathcal{P}}$ and $\mathbf{x} \in [-1,1]^{\mathcal{P}}$, we define $\mathbf{x}_z \in \mathbb{R}^d$ in (5.3) as follows

$$(\mathbf{x}_z)_i = x_{z,i} = \begin{cases} x_i ; & \text{if } \beta_i = (-1)^{\text{digit}(z-1,h(i)-1)} \text{ and } i \in \mathcal{P}, \\ 0 ; & \text{otherwise,} \end{cases} \quad \text{for } i \in [d] \quad \text{and} \quad z \in [2^{r_0-1}].$$

$$(5.7)$$

Hence denoting $\mathcal{A} = \left\{ \mathbf{j} \in \binom{\mathcal{P}}{r_0 - 1} : h \text{ is injective on } \{j_1, \ldots, j_{r_0 - 1}\} \right\}$, since $\widehat{\mathcal{S}_{r_0}} = \mathcal{S}_{r_0}$, we obtain as a result of Lemma 4 that

$$\sum_{z=1}^{2^{r_0-1}} (-1)^{\operatorname{digit}(z-1)} f(\mathbf{x}_z) = \sum_{\mathbf{j} \in \mathcal{A} \cap \mathcal{S}_{r_0-1}} \beta_{j_1} \dots \beta_{j_{r_0-1}} \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}).$$
(5.8)

Consequently, in the linear system in (5.5), we have $\mathbf{B} \in \mathbb{R}^{n \times \binom{|\mathcal{P}|}{r_0 - 1}}$ where the i^{th} row of \mathbf{B} is $\boldsymbol{\beta}^{(r_0 - 1)} \in \{-1, +1\}^{\binom{\mathcal{P}}{r_0 - 1}}$ with $\beta_{\mathbf{j}}^{(r_0 - 1)} = \beta_{j_1}\beta_{j_2}\dots\beta_{j_{r_0 - 1}}$ for each $\mathbf{j} = (j_1, j_2, \dots, j_{r_0 - 1})$. Note that $\mathbf{z}^*(\mathbf{x}; \mathcal{A}) \in \mathbb{R}^{\binom{\mathcal{P}}{r_0 - 1}}$ is the $|\mathcal{S}_{r_0 - 1}|$ sparse vector to be estimated.

• Finally, we will estimate $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ at each $h \in \mathcal{H}_{r_0-1}^{\mathcal{P}}$ and $\mathbf{x} \in \chi(h)$. If SPARSE-REC is ϵ accurate, with ϵ sufficiently small, then by choosing the number of points m to be sufficiently large, we will be able to identify \mathcal{S}_{r_0-1} via thresholding.

By repeating the above steps for all $i = r_0, r_0 - 1, ..., 1$, we arrive at a procedure for estimating the supports $S_i, i \in [r_0]$; this is outlined formally in the form of the Algorithm 3 below. Lemma 5

Algorithm 3 Algorithm for estimating $S_1, S_2, \ldots, S_{r_0}$

1: Input: $d, |S_i|, (m_i, n_i, \epsilon_i)$ for $i = 1, ..., r_0$. 2: Initialization: $\widehat{S}_i = \emptyset$ for $i = 1, \ldots, r_0$. $\mathcal{P} = [d]$. 3: **Output:** \widehat{S}_i for $i = 1, \ldots, r_0$. 4: 5: **for** $i = r_0, r_0 - 1, \dots, 1$ **do** // Estimation of S_i Generate Bernoulli random vectors $\beta_1, \beta_2, \ldots, \beta_{n_i} \in \{-1, 1\}^{\mathcal{P}}$. 6: Form $\mathbf{B} \in \mathbb{R}^{n_i \times |\mathcal{P}|}$ as in (5.5). 7: Construct a (\mathcal{P}, i) hash family $\mathcal{H}_i^{\mathcal{P}}$. 8: for $h \in \mathcal{H}_i^{\mathcal{P}}$ do 9: Construct $\chi(h) \subset [-1,1]^{\mathcal{P}}$ in the same manner as in (5.6) with $|\chi(h)| = (2m_i + 1)^i$. 10: for $\mathbf{x} \in \chi(h)$ do 11: Generate $\mathbf{x}_z \in [-1, 1]^d$, with $z \in [2^i]$ as in (5.7), using $\mathbf{x}, \boldsymbol{\beta}_u$ for each $u \in [n_i]$. 12:Using the samples $(\tilde{f}(\mathbf{x}_z))_{z=1}^{2^i}$, form $\mathbf{y} \in \mathbb{R}^{n_i}$ as in (5.5). 13:Obtain $\widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) = \text{SPARSE-REC}_i(\mathbf{y}, \mathbf{B}).$ 14: Update $\widehat{\mathcal{S}}_i = \widehat{\mathcal{S}}_i \cup \left\{ \mathbf{j} \in \mathcal{A} : |\widehat{z_{\mathbf{j}}^*}(\mathbf{x}_{\mathbf{j}})| > \epsilon_i \right\}.$ 15:end for 16:end for 17:Update $\mathcal{P} = \mathcal{P} \setminus \widehat{\mathcal{S}}_i^{(1)}$. 18:19: end for

below provides sufficient conditions on the sampling parameters in Algorithm 3 for exact recovery of all S_i 's.

Lemma 5. For each $i \in [r_0]$ assume that the following hold:

- 1. $m_i \geq \sqrt{i} \left(\frac{3L}{D_i}\right)^{1/\alpha}$.
- 2. SPARSE-REC_i is ϵ_i accurate with $\epsilon_i < D_i/3$ for all $h \in \mathcal{H}_i^{\mathcal{P}_i}$, $\mathbf{x} \in \chi(h)$, where \mathcal{P}_i denotes the set \mathcal{P} at the beginning of iteration i (so $\mathcal{P}_{r_0} = [d]$). The number of measurements used by SPARSE-REC_i is denoted by n_i .
- 3. $\mathcal{H}_i^{\mathcal{P}_i}$ is a (\mathcal{P}_i, i) hash family.

Then $\widehat{S}_i = S_i$ for all $i = r_0, r_0 - 1, ..., 1$ in Algorithm 3. Moreover, the total number of queries of f made is

$$\sum_{i=1}^{r_0} 2^i (2m_i + 1)^i n_i |\mathcal{H}_i^{\mathcal{P}_i}|.$$

Proof. The proof outline builds on what we have seen in the preceding sections. Say we are at the beginning of iteration $i \in [r_0]$ with $\widehat{\mathcal{S}}_l = \mathcal{S}_l$ holding true for each l > i. Hence, the model has reduced to an order i sparse additive model on the set $\mathcal{P}_i \subset [d]$, with $\mathcal{S}_i^{(1)}, \mathcal{S}_{i-1}^{(1)}, \ldots, \mathcal{S}_1 \subset \mathcal{P}_i$. By identifiability assumption, we know that for any given $\mathbf{j} \in \mathcal{S}_i$, there exists $\mathbf{x}_{\mathbf{j}}^* \in [-1, 1]^i$ such

By identifiability assumption, we know that for any given $\mathbf{j} \in S_i$, there exists $\mathbf{x}_{\mathbf{j}}^* \in [-1,1]^i$ such that $|\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}^*)| \geq D_i$ holds. Moreover, since $\mathcal{H}_i^{\mathcal{P}_i}$ is a (\mathcal{P}_i, i) hash family, there exists a $h \in \mathcal{H}_i^{\mathcal{P}_i}$ that

is an injection on **j**. Consequently, there exists $\mathbf{x} \in \chi(h)$ such that $\|\mathbf{x}_{\mathbf{j}} - \mathbf{x}_{\mathbf{j}}^*\|_2 \leq (\sum_{p=1}^{i} \frac{1}{m_i^2})^{1/2} = \sqrt{i/m_i}$. By Hölder continuity of $\phi_{\mathbf{j}}$, this means

$$|\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}) - \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}^*)| \le L \frac{i^{\alpha/2}}{m_i^{\alpha}}.$$
(5.9)

Since SPARSE-REC_i is ϵ_i accurate for each $h \in \mathcal{H}_i^{\mathcal{P}_i}$, $\mathbf{x} \in \chi(h)$, we know that at the aforementioned \mathbf{x} , the following holds via reverse triangle inequality

$$|\widehat{z_{\mathbf{j}}^*}(\mathbf{x_j})| \ge |\phi_{\mathbf{j}}(\mathbf{x_j})| - \epsilon_i.$$
(5.10)

Using (5.9), (5.10), reverse triangle inequality and the choice of ϵ_i and m_i , we obtain

$$|\widehat{z_{\mathbf{j}}^*}(\mathbf{x_j})| \ge |\phi_{\mathbf{j}}(\mathbf{x_j}^*)| - L\frac{i^{\alpha/2}}{m_i^{\alpha}} - \epsilon_i \ge D_i - L\frac{i^{\alpha/2}}{m_i^{\alpha}} - \epsilon_i \ge \frac{2D_i}{3} - L\frac{i^{\alpha/2}}{m_i^{\alpha}} \ge \frac{D_i}{3}$$

For any $\mathbf{j} \notin S_i$, we have for all $h \in \mathcal{H}_i^{\mathcal{P}_i}$, $\mathbf{x} \in \chi(h)$ that $|\widehat{z_j^*}(\mathbf{x_j})| \leq \epsilon_i < D_i/3$ (since $\phi_j \equiv 0$). Hence clearly, the stated choice of ϵ_i guarantees identification of each $\mathbf{j} \in S_i$, and none from $\binom{\mathcal{P}_i}{i} \setminus S_i$. This means that we will recover S_i exactly. As this is true for each $i \in [r_0]$, it also completes the proof for exact recovery of S_i for each $i \in [r_0]$.

The expression for the total number of queries made follows from a simple calculation where we note that at iteration *i*, and corresponding to each $\mathbf{x} \in \bigcup_{h \in \mathcal{H}_i^{\mathcal{P}_i}} \chi(h)$, we make $2^i n_i$ queries of f. \Box

6 Estimating sparse multilinear functions from few samples

In this section, we provide results from the sparse recovery literature for estimating sparse multilinear forms from random samples. In particular, these results cover arbitrary bounded noise and i.i.d. Gaussian noise models.

6.1 Sparse linear functions

Consider a linear function $g : \mathbb{R}^d \to \mathbb{R}$, where $g(\beta) = \beta^T \mathbf{a}$. Our interest is in recovering the unknown coefficient vector \mathbf{a} from n noisy samples $y_i = g(\beta_i) + \eta_i$, $i \in [n]$, where η_i refers to the noise in the i^{th} sample. Arranging the samples together, we arrive at the linear system $\mathbf{y} = \mathbf{B}\mathbf{a} + \boldsymbol{\eta}$, where

$$\underbrace{\begin{pmatrix} y_1 \\ \vdots \\ y_n \end{pmatrix}}_{\mathbf{y} \in \mathbb{R}^n} = \underbrace{\begin{pmatrix} \beta_1^T \\ \vdots \\ \beta_n^T \end{pmatrix}}_{\mathbf{B} \in \mathbb{R}^{n \times d}} \mathbf{a} + \underbrace{\begin{pmatrix} \eta_1 \\ \vdots \\ \vdots \\ \eta_n \end{pmatrix}}_{\boldsymbol{\eta}}.$$
(6.1)

Denoting by $S := \{j \in [d] : a_j \neq 0\}$ the support of **a**, our interest is in the setting where **a** is sparse, i.e., $|S| = k \ll d$, and consequently to estimate **a** from a small number of samples *n*. To begin with, we will require **B** in (6.1) to satisfy the so called ℓ_2/ℓ_2 RIP, defined below.

Definition 2. A matrix $\mathbf{A} \in \mathbb{R}^{n \times d}$ is said to satisfy the ℓ_2/ℓ_2 Restricted Isometry Property (RIP) of order k with constant $\delta_k \in (0, 1)$ if

$$(1 - \delta_k) \parallel \mathbf{x} \parallel_2^2 \le \frac{1}{n} \parallel \mathbf{A} \mathbf{x} \parallel_2^2 \le (1 + \delta_k) \parallel \mathbf{x} \parallel_2^2$$

holds for all k-sparse \mathbf{x} .

Bounded noise model. Let us consider the scenario where the noise is bounded in the ℓ_2 norm, i.e., $\|\boldsymbol{\eta}\|_2 \leq \nu$. We will recover an estimate $\hat{\mathbf{a}}$ to \mathbf{a} as a solution of the following quadratically constrained ℓ_1 minimization program [4]

(P1)
$$\min_{\mathbf{z}\in\mathbb{R}^d} \|\mathbf{z}\|_1$$
 s.t $\|\mathbf{y} - \mathbf{B}\mathbf{z}\|_2 \le \nu.$ (6.2)

The following result provides a bound on the estimation error $\| \hat{\mathbf{a}} - \mathbf{a} \|_2$ for (P1).

Theorem 1. Consider the sampling model in (6.1), where $\mathbf{B} \in \{-1, +1\}^{n \times d}$ is a Bernoulli matrix. Then the following hold.

1. ([1]) For any constant $\delta \in (0,1)$, there exist constants $c_1, c_2 > 0$ depending on δ such that if

$$n \ge c_1 k \log(d/k),$$

then with probability at least $1 - 2\exp(-c_2n)$, the matrix **B** satisfies ℓ_2/ℓ_2 RIP of order k, with $\delta_k \leq \delta$.

2. ([4, Theorem 1.2]) Let **B** satisfy the ℓ_2/ℓ_2 RIP with $\delta_{2k} < \sqrt{2} - 1$. Then there exist constants $C_1, C_2 > 0$ such that, simultaneously for all vectors $\mathbf{a} \in \mathbb{R}^d$, any solution $\hat{\mathbf{a}}$ to (P1) satisfies

$$\| \widehat{\mathbf{a}} - \mathbf{a} \|_2 \le C_1 \frac{\| \mathbf{a} - \mathbf{a}_k \|_1}{\sqrt{k}} + C_2 \frac{\nu}{\sqrt{n}}$$

Here, \mathbf{a}_k denotes the best k-term approximation of \mathbf{a} .

Gaussian noise model. We now consider the scenario where the noise samples are i.i.d. Gaussian with variance σ^2 , i.e., $\eta_i \sim \mathcal{N}(0, \sigma^2)$ i.i.d. for all $i \in [n]$. Using standard concentration inequalities for sub-exponential random variables (see Proposition 6(1)), one can show that $\| \boldsymbol{\eta} \|_2 = \Theta(\sigma \sqrt{n})$ with high probability. This leads to the following straightforward corollary of Theorem 1.

Corollary 1. Consider the sampling model in (6.1) for some given vector $\mathbf{a} \in \mathbb{R}^n$, and let $\eta_i \sim \mathcal{N}(0, \sigma^2)$ i.i.d. for all $i \in [n]$. Say **B** satisfies ℓ_2/ℓ_2 RIP with $\delta_{2k} < \sqrt{2} - 1$. For some $\varepsilon \in (0, 1)$, let $\hat{\mathbf{a}}$ be a solution to (P1) with $\nu = (1 + \varepsilon)\sigma\sqrt{n}$. Then there exists a constant $c_3 > 0$ so that any solution $\hat{\mathbf{a}}$ to (P1) satisfies

$$\| \widehat{\mathbf{a}} - \mathbf{a} \|_{2} \leq C_{1} \frac{\| \mathbf{a} - \mathbf{a}_{k} \|_{1}}{\sqrt{k}} + C_{2}(1 + \varepsilon)\sigma$$

with probability at least $1 - 2\exp(-c_3\varepsilon^2 n)$. Here C_1, C_2 are the constants from Theorem 1.

Proof. Use Proposition 6(1) with Theorem 1.

6.2 Sparse bilinear functions

Let $\mathbf{a} \in \mathbb{R}^{\binom{d}{2}}$ be a vector of length $\binom{d}{2}$ with entries indexed by $\binom{[d]}{2}$ (sorted in lexicographic order). The entry of \mathbf{a} at index $\mathbf{j} = (j_1, j_2) \in \binom{[d]}{2}$ will be denoted by $a_{\mathbf{j}}$. We now consider the setting where $g : \mathbb{R}^d \to \mathbb{R}$ is a second order multilinear function, i.e., $g(\boldsymbol{\beta}) = \langle \boldsymbol{\beta}^{(2)}, \mathbf{a} \rangle$ with $\boldsymbol{\beta}^{(2)}, \mathbf{a} \in \mathbb{R}^{\binom{d}{2}}$, and

 $\beta_{(j_1,j_2)}^{(2)} = \beta_{j_1}\beta_{j_2}$. As before, our goal is to recover **a** from *n* noisy samples $y_i = g(\beta_i) + \eta_i$, $i \in [n]$ resulting in the linear system

$$\begin{pmatrix}
y_1 \\
\vdots \\
y_n
\end{pmatrix}_{\mathbf{y}\in\mathbb{R}^n} = \underbrace{\begin{pmatrix}
\beta_1^{(2)^T} \\
\vdots \\
\beta_n^{(2)^T}
\end{pmatrix}}_{\mathbf{B}\in\mathbb{R}^{n\times\binom{d}{2}}} \mathbf{a} + \underbrace{\begin{pmatrix}
\eta_1 \\
\vdots \\
\vdots \\
\eta_n
\end{pmatrix}}_{\boldsymbol{\eta}}.$$
(6.3)

Observe that

$$\langle \boldsymbol{\beta}^{(2)}, \mathbf{a} \rangle = \boldsymbol{\beta}^T \mathbf{A} \boldsymbol{\beta} = \langle \boldsymbol{\beta} \boldsymbol{\beta}^T, \mathbf{A} \rangle,$$
 (6.4)

where $\mathbf{A} \in \mathbb{R}^{d \times d}$ is a symmetric matrix with zero on the diagonal and $A_{i,j} = a_{(i,j)}/2$ if $i \neq j$. This simple observation allows us to rewrite (6.3) as

$$y_i = \langle \boldsymbol{\beta}_i \boldsymbol{\beta}_i^T, \mathbf{A} \rangle + \eta_i, \quad i \in [n].$$
 (6.5)

Chen et al. [5] recently showed that a sparse symmetric matrix – not necessarily being zero on the diagonal – can be recovered from its measurements of the form (6.5), provided that β_j is sampled in an i.i.d. manner from a distribution satisfying

$$\mathbb{E}[\beta_i] = 0, \quad \mathbb{E}[\beta_i^2] = 1 \quad \text{and} \quad \mathbb{E}[\beta_i^4] > 1.$$
(6.6)

Their recovery program² is essentially constrained ℓ_1 minimization ([5, Eq.(4)]), the guarantees for which rely on the ℓ_2/ℓ_1 RIP for sparse symmetric matrices ([5, Def. 2]).

Since our matrix \mathbf{A} in (6.5) is sparse and symmetric, it is natural for us to use their scheme. We do this, albeit with some technical changes:

- We will see that if **A** is known to be zero on the diagonal (which is the case here), the fourth order moment condition in (6.6) is not needed. Hence one could, for instance, sample β from the symmetric Bernoulli distribution.
- Instead of optimizing over the set of symmetric matrices with zeros on the diagonal, we will perform ℓ_1 minimization over the upper triangular entries of **A**, represented by **a**. These approaches are equivalent, but the latter has the computational advantage of having fewer constraints.

The analysis is based on the notion of ℓ_2/ℓ_1 RIP of the matrix **B** in (6.3), which is defined as follows.

Definition 3. A matrix $\mathbf{B} \in \mathbb{R}^{n \times N}$ is said to satisfy the ℓ_2/ℓ_1 Restricted Isometry Property (RIP) of order k with constants $\gamma_k^{\text{lb}} \in (0, 1)$ and $\gamma_k^{\text{ub}} > 0$ if

$$(1 - \gamma_k^{\text{lb}}) \parallel \mathbf{x} \parallel_2 \le \frac{1}{n} \parallel \mathbf{B} \mathbf{x} \parallel_1 \le (1 + \gamma_k^{\text{ub}}) \parallel \mathbf{x} \parallel_2$$

holds for all k-sparse $\mathbf{x} \in \mathbb{R}^N$.

The above definition is analogous to the one in [5, Def. 2] for sparse symmetric matrices.

²Note that one can remove the positive semi-definite constraint in their program and replace it with a symmetry enforcing constraint, the result remains unchanged (cf. remark in [5, Section E])

Bounded noise model. Let us first consider the setting where the noise is bounded in the ℓ_1 norm, i.e., $\|\boldsymbol{\eta}\|_1 \leq \nu$. We recover the estimate $\hat{\mathbf{a}}$ as a solution to the following program

(P2)
$$\min_{\mathbf{z}\in\mathbb{R}^{\binom{d}{2}}} \| \mathbf{z} \|_{1} \quad \text{s.t} \quad \| \mathbf{y} - \mathbf{B}\mathbf{z} \|_{1} \le \nu.$$
(6.7)

The next result shows that **B** satisfies ℓ_2/ℓ_1 RIP with high probability if the rows of **B** are formed by independent Bernoulli vectors. Consequently, the above program stably recovers **a**.

Theorem 2. Consider the sampling model in (6.3), where the rows of **B** are formed by independent Bernoulli vectors. Then the following hold.

1. There exist absolute constants $c_1, c_2, c_3 > 0$, such that the following is true. Let $\mathbf{a} \in \mathbb{R}^{\binom{d}{2}}$. Then

$$c_1 \parallel \mathbf{a} \parallel_2 \le \frac{1}{n} \parallel \mathbf{B}\mathbf{a} \parallel_1 \le c_2 \parallel \mathbf{a} \parallel_2$$
 (6.8)

holds with probability at least $1 - \exp(-c_3 n)$.

2. With constants c_1, c_2 from (6.8), there exist constants $c'_1, c'_2, c'_3 > 0$ such that if $n > c'_3 k \log(d^2/k)$, then **B** satisfies ℓ_2/ℓ_1 RIP of order k with probability at least $1 - c'_1 \exp(-c'_2 n)$ with constants γ_k^{lb} and γ_k^{ub} , which fulfill

$$1 - \gamma_k^{\text{lb}} \ge \frac{c_1}{2} \quad and \quad 1 + \gamma_k^{\text{ub}} \le 2c_2.$$
 (6.9)

3. If there exists a number K > 2k such that **B** satisfies

$$\frac{1-\gamma^{\rm lb}_{k+K}}{\sqrt{2}} - (1+\gamma^{\rm ub}_K)\sqrt{\frac{k}{K}} \ge \beta > 0$$

for some $\beta > 0$, then the solution $\hat{\mathbf{a}}$ to (P2) satisfies

$$\| \widehat{\mathbf{a}} - \mathbf{a} \|_{2} \leq \left(\frac{\tilde{C}_{1}}{\beta} + \tilde{C}_{3} \right) \frac{\| \mathbf{a} - \mathbf{a}_{k} \|_{1}}{\sqrt{K}} + \frac{\tilde{C}_{2}}{\beta} \frac{\nu}{n},$$

where \mathbf{a}_k denotes the best k-term approximation of \mathbf{a} and $\tilde{C}_1, \tilde{C}_2, \tilde{C}_3$ are universal positive constants.

4. There exist absolute constants $\tilde{c}_3, \tilde{c}_4, C_3, C_4 > 0$ such that if $n \geq \tilde{c}_3 k \log(d^2/k)$, then the solution $\hat{\mathbf{a}}$ to (P2) satisfies

$$\| \widehat{\mathbf{a}} - \mathbf{a} \|_2 \le C_3 \frac{\| \mathbf{a} - \mathbf{a}_k \|_1}{\sqrt{k}} + C_4 \frac{\nu}{n},$$

simultaneously for all $\mathbf{a} \in \mathbb{R}^{\binom{d}{2}}$ with probability at least $1 - \exp(-\tilde{c}_4 n)$.

Gaussian noise model. We now consider the scenario where the noise samples are i.i.d. Gaussian with variance σ^2 , i.e. $\eta_i \sim \mathcal{N}(0, \sigma^2)$ i.i.d. for all $i \in [n]$. Using standard concentration inequalities for sub-Gaussian random variables (see Proposition 6(2)), one can show that $\| \boldsymbol{\eta} \|_1 = \Theta(\sigma n)$ with high probability. This leads to the following straightforward corollary of Theorem 2.

Corollary 2. For constants $\tilde{c}_3, \tilde{c}_4, C_3, C_4 > 0$ defined in Theorem 2, the following is true. Consider the sampling model in (6.3) for a given $\mathbf{a} \in \mathbb{R}^{\binom{d}{2}}$, where the rows of **B** are formed by independent Bernoulli vectors, and $\eta_i \sim \mathcal{N}(0, \sigma^2)$ i.i.d. for all $i \in [n]$ with $n \geq \tilde{c}_3 k \log(d^2/k)$. For some $\varepsilon \in (0, 1)$, let $\hat{\mathbf{a}}$ be a solution to (P2) with $\nu = (1 + \varepsilon)\sigma n$. Then

$$\| \widehat{\mathbf{a}} - \mathbf{a} \|_{2} \leq C_{3} \frac{\| \mathbf{a} - \mathbf{a}_{k} \|_{1}}{\sqrt{k}} + C_{4} \sqrt{\frac{2}{\pi}} (1 + \varepsilon) \sigma$$

with probability at least $1 - \exp(-\tilde{c}_4 n) - e \cdot \exp(-\tilde{c}_5 \varepsilon^2 n)$, for some constant $\tilde{c}_5 > 0$. Here, \mathbf{a}_k denotes the best k-term approximation of \mathbf{a} .

Proof. Use Proposition 6(2) with Theorem 2.

6.3 Sparse multilinear functions

For $p, d \in \mathbb{N}$ with $p \leq d$, let $\mathbf{a} \in \mathbb{R}^{\binom{d}{p}}$ be a vector with entries indexed by $\binom{[d]}{p}$ (sorted in lexicographic order). We denote again the entry of \mathbf{a} at index $\mathbf{j} = (j_1, \ldots, j_p) \in \binom{[d]}{p}$ by $a_{\mathbf{j}}$ and consider a multilinear function $g : \mathbb{R}^d \to \mathbb{R}$ in d variables $\boldsymbol{\beta} = (\beta_1 \ldots \beta_d)^T$ such that

$$g(\boldsymbol{\beta}) = \sum_{\mathbf{j} = (j_1, \dots, j_p) \in \binom{[d]}{p}} \beta_{j_1} \beta_{j_2} \cdots \beta_{j_p} a_{\mathbf{j}}.$$
(6.10)

We will refer to (6.10) as a multilinear function of order p. For clarity of notation, we will write (6.10) as $g(\boldsymbol{\beta}) = \langle \boldsymbol{\beta}^{(p)}, \mathbf{a} \rangle$, where $\boldsymbol{\beta}^{(p)} \in \mathbb{R}^{\binom{d}{p}}$, and the entry of $\boldsymbol{\beta}^{(p)}$ at index $\mathbf{j} = (j_1, \ldots, j_p) \in \binom{[d]}{p}$ being $\beta_{j_1}\beta_{j_2}\cdots\beta_{j_p}$.

We are again interested in recovering the unknown coefficient vector **a** from *n* noisy samples $y_i = g(\beta_i) + \eta_i, i \in [n]$, where η_i refers to the noise in the *i*th sample. Arranging the samples together, we arrive at the linear system $\mathbf{y} = \mathbf{B}\mathbf{a} + \boldsymbol{\eta}$, where

$$\underbrace{\begin{pmatrix} y_1 \\ \vdots \\ y_n \end{pmatrix}}_{\mathbf{y} \in \mathbb{R}^n} = \underbrace{\begin{pmatrix} \boldsymbol{\beta}_1^{(p)^T} \\ \vdots \\ \vdots \\ \boldsymbol{\beta}_n^{(p)^T} \end{pmatrix}}_{\mathbf{B} \in \mathbb{R}^{n \times \binom{d}{p}}} \mathbf{a} + \underbrace{\begin{pmatrix} \eta_1 \\ \vdots \\ \vdots \\ \eta_n \end{pmatrix}}_{\boldsymbol{\eta}}.$$
(6.11)

We denote by $S = \left\{ (j_1, j_2, \dots, j_p) \in {\binom{[d]}{p}} : a_{(j_1, j_2, \dots, j_p)} \neq 0 \right\}$ the support of **a** and we are especially interested in the setting where **a** is sparse, i.e., $|S| = k \ll {\binom{d}{p}}$. Our aim is to estimate **a** with a small number of samples n.

Bounded noise model. Let us consider the scenario where the noise is bounded in the ℓ_2 norm, i.e., $\|\boldsymbol{\eta}\|_2 \leq \nu$. We will recover an estimate $\hat{\mathbf{a}}$ to \mathbf{a} as a solution of (P1). The following result provides a bound on the estimation error $\|\hat{\mathbf{a}} - \mathbf{a}\|_2$.

Theorem 3. ([26, Theorem 4; Lemmas 4, 5]) Let $D = \binom{d}{p}$ and let **B** be defined as in (6.11) with rows formed by independent Bernoulli vectors. Then, for any $\delta \in (0,1)$, there exist constants $c_6, c_7 > 0$ depending on δ such that the matrix **B** satisfies ℓ_2/ℓ_2 RIP of order k, with $\delta_k \leq \delta$,

- (a) with probability at least $1 \exp(-c_7 n/k^2)$ if $n \ge c_6 k^2 \log D$
- (b) and with probability at least $1 \exp(-c_7 \min\{n/3^{2p}, n/k\})$ if

 $n \ge c_6 \max \left\{ 3^{2p} k \log(D/k), k^2 \log(D/k) \right\}.$

The bounds on n in the theorem are obtained via the application of two very different methodologies. The bound in part (a) is a consequence of bounding the eigenvalues of the $k \times k$ Gram matrices $\frac{1}{n} \mathbf{B}_{\mathcal{S}}^T \mathbf{B}_{\mathcal{S}}$ for all \mathcal{S} (here $\mathbf{B}_{\mathcal{S}}$ is the submatrix of \mathbf{B} with column indices in \mathcal{S}) using Gershgorin's disk theorem, along with standard concentration inequalities [26, Theorem 4]. The bound in part (b) involves the usage of tail estimates for Rademacher chaos variables and follows from [26, Lemmas 4, 5].

Remark 5. The analysis in [26, Section A] derives RIP bounds in terms of the so-called combinatorial dimension. However in our opinion, this analysis has several inaccuracies because of which we are not sure if the corresponding bounds are correct. Hence we do not state those bounds here.

Gaussian noise model. In the scenario where the noise in the samples is i.i.d. Gaussian, we arrive at a statement similar to Corollary 1, hence we do not discuss this further.

7 Putting it together: final theoretical guarantees

We are now in a position to combine our efforts from the preceding sections and to state the final results for recovery of the support sets. As before, we state this separately for the univariate, bivariate, and general multivariate cases.

7.1 Univariate case

We begin with the univariate case considered in Section 3. Recall Algorithm 1 for recovering the support S_1 . The ensuing Lemma 1 gave sufficient conditions for exact recovery provided SPARSE-REC is ϵ -accurate at each $\mathbf{x} \in \chi$, for small enough ϵ . Instantiating SPARSE-REC with (P1) in (6.2) gives the final results below. Let us start with the bounded noise model.

Bounded noise model. In this noise model, querying f at \mathbf{x} returns $f(\mathbf{x}) + \eta$, where $|\eta| \leq \Delta$. For the linear system (3.4), this means that $|\eta_i^+|, |\eta_i^-| \leq \Delta$, $i \in [n]$, and hence $\|\boldsymbol{\eta}\|_{\infty} \leq 2\Delta$. The following theorem shows that if Δ is sufficiently small, then Algorithm 1 recovers S_1 exactly provided the parameters m, n, ϵ are chosen in a suitable way.

Theorem 4. For the bounded noise model with the noise uniformly bounded by \triangle , consider Algorithm 1 with SPARSE-REC instantiated with (P1) in (6.2). If $\nu = 2 \triangle \sqrt{n}$ in (P1),

$$\Delta < \frac{D_1}{6C_2}, \quad m \ge (3L/D_1)^{1/\alpha}, \quad and \quad n \ge \tilde{c}_1|\mathcal{S}_1|\log(d/|\mathcal{S}_1|)$$

are satisfied, it follows for the choice $\epsilon = 2C_2 \triangle$ that $\widehat{S_1} = S_1$, with probability at least $1 - 2\exp(-\tilde{c}_2 n)$. The total number of queries made is $2(2m+1)n = \Omega\left(|S_1|\log\left(\frac{d}{|S_1|}\right)\right)$.

Proof. The proof follows by combining Lemma 1 with Theorem 1. Since $|\eta_i^+|, |\eta_i^-| \leq \Delta$ in (3.4) for $i \in [n]$, we obtain $\|\boldsymbol{\eta}\|_2 \leq \sqrt{n} \|\boldsymbol{\eta}\|_{\infty} \leq 2\Delta\sqrt{n}$. Therefore we set SPARSE-REC = (P1) with $\nu = 2\Delta\sqrt{n}$.

As a consequence of Theorem 1, there exist constants $\tilde{c}_1, \tilde{c}_2 > 0$ (depending only on $c_1, c_2 > 0$ defined therein) so that, for $n \geq \tilde{c}_1 |\mathcal{S}_1| \log(d/|\mathcal{S}_1|)$, **B** satisfies ℓ_2/ℓ_2 RIP with $\delta_{2|\mathcal{S}_1|} < \sqrt{2} - 1$ with probability at least $1 - 2 \exp(-\tilde{c}_2 n)$. Conditioning on this event, it follows from Theorem 1 that

$$\| \widehat{\mathbf{z}^*}(\mathbf{x}) - \mathbf{z}^*(\mathbf{x}) \|_{\infty} \le \| \widehat{\mathbf{z}^*}(\mathbf{x}) - \mathbf{z}^*(\mathbf{x}) \|_2 \le C_2(2\Delta) \quad \text{for all} \quad \mathbf{x} \in \chi_2$$

Hence, we see that with probability at least $1 - 2 \exp(-\tilde{c}_2 n)$, SPARSE-REC is $\epsilon = 2C_2 \triangle$ accurate at each $\mathbf{x} \in \chi$. Now invoking Lemma 1, it follows that if

$$2C_2 \triangle < \frac{D_1}{3} \Leftrightarrow \triangle < \frac{D_1}{6C_2}$$

holds, then the stated choice of ϵ and m ensures exact recovery. This completes the proof.

Gaussian noise model. We now move to the Gaussian noise model, wherein querying f at \mathbf{x} returns $f(\mathbf{x}) + \eta$; $\eta \sim \mathcal{N}(0, \sigma^2)$. Moreover, the noise samples are independent across the queries. For the linear system (3.4), this means that $\eta_i \sim \mathcal{N}(0, 2\sigma^2), i \in [n]$ are i.i.d. random variables. The following theorem essentially shows that if the noise variance σ^2 is sufficiently small, then Algorithm 1 recovers S_1 exactly provided the parameters m, n, ϵ are chosen properly. The reduction in the variance is handled via re-sampling each query N times and averaging the values. Essentially, this leads to the same sampling model with i.i.d. $\eta_i \sim \mathcal{N}(0, 2\sigma^2/N), i \in [n]$.

Theorem 5. For the Gaussian noise model with i.i.d. noise samples with variance σ^2 and $N \in \mathbb{N}$, we resample each query N times, and average the values. Consider Algorithm 1 with SPARSE-REC instantiated with (P1) in (6.2), wherein $\nu = \sqrt{2}(1 + \varepsilon)\sigma\sqrt{n/N}$ for some $\varepsilon \in (0, 1)$, and with

$$N \ge \left\lfloor \frac{18C_2^2(1+\varepsilon)^2 \sigma^2}{D_1^2} \right\rfloor + 1, \ m \ge \left(\frac{3L}{D_1}\right)^{1/\alpha}, \ n \ge \max\left\{\tilde{c}_1 |\mathcal{S}_1| \log\left(\frac{d}{|\mathcal{S}_1|}\right), \frac{2\log(2m+1)}{c_3\varepsilon^2}\right\} (7.1)$$

being satisfied. Then $\widehat{S_1} = S_1$ with probability at least $1-2\exp(-\tilde{c}_2n)-2\exp(-\frac{c_3\varepsilon^2n}{2})$. The constants $\tilde{c}_1, \tilde{c}_2 > 0$ are as in Theorem 4; $C_2, c_3 > 0$ come from Theorem 1 and Corollary 1, respectively. The total number of queries made is $2(2m+1)nN = \Omega\left(|S_1|\log\left(\frac{d}{|S_1|}\right)\right)$.

Proof. First note that in (3.4), as a consequence of resampling N times and averaging, we have $\eta_i^+, \eta_i^- \sim \mathcal{N}(0, \sigma^2/N)$, and $\eta_i = \eta_i^+ - \eta_i^- \sim \mathcal{N}(0, 2\sigma^2/N)$, $i \in [n]$. Hence we set SPARSE-REC = (P1) with $\nu = \sqrt{2}(1 + \varepsilon)\sigma\sqrt{n/N}$.

From Theorem 1, there exist constants $\tilde{c}_1, \tilde{c}_2 > 0$ (depending only on $c_1, c_2 > 0$ defined therein) so that for the choice $n \geq \tilde{c}_1 |\mathcal{S}_1| \log(d/|\mathcal{S}_1|)$, **B** satisfies ℓ_2/ℓ_2 RIP with $\delta_{2|\mathcal{S}_1|} < \sqrt{2} - 1$ with probability at least $1 - 2 \exp(-\tilde{c}_2 n)$. Conditioning on this event, and invoking Corollary 1 for the stated choice of ν , we have for any given $\mathbf{x} \in \chi$ that with probability at least $1 - 2 \exp(-c_3 \varepsilon^2 n)$, the following holds

$$\| \widehat{\mathbf{z}^*}(\mathbf{x}) - \mathbf{z}^*(\mathbf{x}) \|_{\infty} \leq \| \widehat{\mathbf{z}^*}(\mathbf{x}) - \mathbf{z}^*(\mathbf{x}) \|_2 \leq \sqrt{2}C_2(1+\varepsilon)\sigma/\sqrt{N}.$$
(7.2)

By the union bound over the 2m + 1 elements of χ , it follows that (7.2) holds for all $\mathbf{x} \in \chi$ with probability at least

$$1 - 2|\chi| \exp(-c_3 \varepsilon^2 n) = 1 - 2 \exp(\log(2m + 1) - c_3 \varepsilon^2 n) \ge 1 - 2 \exp\left(-\frac{c_3 \varepsilon^2 n}{2}\right)$$

if $n \geq \frac{2\log(2m+1)}{c_3\varepsilon^2}$ holds. By (7.1), this condition is indeed satisfied and furthermore, we see that SPARSE-REC is $\epsilon = \sqrt{2}C_2(1+\varepsilon)\sigma/\sqrt{N}$ accurate for all $\mathbf{x} \in \chi$ with probability at least $1-2\exp(-\tilde{c}_2n)-2\exp(-\frac{c_3\varepsilon^2n}{2})$. Now invoking Lemma 1, and using (7.2), it follows that if

$$\frac{\sqrt{2}C_2(1+\varepsilon)\sigma}{\sqrt{N}} < \frac{D_1}{3} \tag{7.3}$$

holds, then the stated choice of ϵ and m ensures exact recovery. Finally, (7.1) ensures that (7.3) holds and this completes the proof.

7.2 Bivariate case

In the bivariate case from Section 4, we use Algorithm 2 for recovering the supports S_2, S_1 and with the Lemma 3 giving sufficient conditions for their exact recovery. Instantiating SPARSE-REC₂ with (P2) in (6.7), and SPARSE-REC₁ with (P1) in (6.2) gives then the results below.

Bounded noise model. Following theorem shows that if \triangle is sufficiently small in the bounded noise model described before, then Algorithm 2 recovers S_2 and S_1 exactly provided the parameters $m_i, n_i, \epsilon_i, i = 1, 2$ are chosen in a suitable way.

Theorem 6. For the bounded noise model with the noise uniformly bounded by \triangle , consider Algorithm 2 with SPARSE-REC₂ instantiated with (P2) with $\nu_2 = 4 \triangle n_2$ in (6.7), and SPARSE-REC₁ realized by (P1) with $\nu_1 = 2 \triangle \sqrt{n_1}$ in (6.2), respectively. Let \mathcal{H}_2^d be a (d, 2) hash family. If

$$\Delta < \min\left\{\frac{D_2}{12C_4}, \frac{D_1}{6C_2}\right\}, \ m_2 \ge \sqrt{2}\left(\frac{3L}{D_2}\right)^{1/\alpha}, \ n_2 \ge \tilde{c}_3|\mathcal{S}_2|\log(d^2/|\mathcal{S}_2|), \\ m_1 \ge (3L/D_1)^{1/\alpha} \quad and \quad n_1 \ge \tilde{c}_1|\mathcal{S}_1|\log\left(\frac{d-|\widehat{\mathcal{S}_2^{\text{var}}|}}{|\mathcal{S}_1|}\right)$$

are satisfied, then $\widehat{S}_2 = S_2$ and $\widehat{S}_1 = S_1$ with probability at least $1 - \exp(-\tilde{c}_4 n_2) - 2\exp(-\tilde{c}_2 n_1)$. Here, the constants $C_4, \tilde{c}_3, \tilde{c}_4 > 0$ are from Theorem 2, while $\tilde{c}_1, \tilde{c}_2, C_2 > 0$ are as defined in Theorem 4. The total number of queries made is

$$4(2m_2+1)^2 n_2 |\mathcal{H}_2^d| + 2(2m_1+1)n_1 = \Omega\left(|\mathcal{S}_2|\log\left(\frac{d^2}{|\mathcal{S}_2|}\right)|\mathcal{H}_2^d| + |\mathcal{S}_1|\log\left(\frac{d-|\widehat{\mathcal{S}_2^{\text{var}}|}}{|\mathcal{S}_1|}\right)\right).$$

Proof. We focus on the proof of the exact recovery of S_2 . Once S_2 is recovered exactly, the model reduces to a univariate SPAM on the variable set $\mathcal{P} = [d] \setminus S_2^{\text{var}}$, with $S_1 \subset \mathcal{P}$. Thereafter, the proof of the exact recovery of S_1 is identical to the proof of Theorem 4.

Since $|\eta_{i,1}|, |\eta_{i,2}|, |\eta_{i,3}|, |\eta_{i,4}| \leq \Delta$ in (4.4) for $i \in [n_2]$, we obtain $\|\boldsymbol{\eta}\|_1 \leq n_2 \|\boldsymbol{\eta}\|_{\infty} \leq 4n_2 \Delta$. Therefore we set SPARSE-REC₂ = (P2) with $\nu = 4\Delta n_2$.

Now, as a consequence of Theorem 2, there exist constants $\tilde{c}_3, \tilde{c}_4, C_4 > 0$ such that if $n_2 \geq \tilde{c}_3|\mathcal{S}_2|\log(d^2/|\mathcal{S}_2|)$, then with probability at least $1 - \exp(-\tilde{c}_4 n_2)$,

$$\| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \mathbf{z}^*(\mathbf{x}; \mathcal{A}) \|_{\infty} \le \| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \mathbf{z}^*(\mathbf{x}; \mathcal{A}) \|_2 \le 4C_4 \Delta \quad \text{for all } \mathbf{x} \in \bigcup_{h \in \mathcal{H}_2^d} \chi(h).$$

This holds since $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ is always at most $|\mathcal{S}_2|$ sparse. Thus, with probability at least $1 - \exp(-\tilde{c}_4 n_2)$, SPARSE-REC₂ is $\epsilon_2 = 4C_4 \Delta$ accurate for each $h \in \mathcal{H}_2^d$, $\mathbf{x} \in \chi(h)$. Now invoking Lemma 3 reveals that if

$$4C_4 \triangle < \frac{D_2}{3} \Leftrightarrow \triangle < \frac{D_2}{12C_4}$$

holds, then the stated choice of ϵ_2 and m_2 ensures $\widehat{S_2} = S_2$. Lastly, the bound on the total number of queries follows from Lemma 3 by plugging in the stated bounds on m_i, n_i ; i = 1, 2.

Gaussian noise model. Next, we consider the Gaussian noise model with the noise samples i.i.d. Gaussian (~ $\mathcal{N}(0, \sigma^2)$) across queries. Similarly to Theorem 5, we show that if the noise variance σ^2 is sufficiently small, then Algorithm 2 recovers S_2 and S_1 for a careful choice of m_i, n_i , i = 1, 2. We again reduce the variance via re-sampling each query either N_2 times (during the estimation of S_2) or N_1 times (during the estimation of S_1), and averaging the values.

Theorem 7. For the Gaussian noise model with i.i.d noise samples ~ $\mathcal{N}(0, \sigma^2)$, say we resample each query N_2 times (during estimation of S_2) or N_1 times (during estimation of S_1), and average the values. Consider Algorithm 2, where SPARSE-REC₂ and SPARSE-REC₁ are instantiated with (P2) with $\nu_2 = 2\sigma(1+\varepsilon)n_2/\sqrt{N_2}$ in (6.7), and (P1) with $\nu_1 = \sqrt{2}(1+\varepsilon)\sigma\sqrt{n_1/N_1}$ in (6.2), respectively, for some $\varepsilon \in (0, 1)$. Let \mathcal{H}_2^d be a (d, 2) hash family. If

$$N_{2} \geq \left\lfloor \frac{72C_{4}^{2}(1+\varepsilon)^{2}\sigma^{2}}{\pi D_{2}^{2}} \right\rfloor + 1, \ m_{2} \geq \sqrt{2} \left(\frac{3L}{D_{2}}\right)^{1/\alpha}, \ n_{2} \geq \max\left\{\tilde{c}_{3}|\mathcal{S}_{2}|\log\left(\frac{d^{2}}{|\mathcal{S}_{2}|}\right), \frac{2\log[(2m_{2}+1)^{2}e|\mathcal{H}_{2}^{d}|]}{\tilde{c}_{5}\varepsilon^{2}}\right\}$$
$$N_{1} \geq \left\lfloor \frac{18C_{2}^{2}(1+\varepsilon)^{2}\sigma^{2}}{D_{1}^{2}} \right\rfloor + 1, \ m_{1} \geq \left(\frac{3L}{D_{1}}\right)^{1/\alpha}, \ n_{1} \geq \max\left\{\tilde{c}_{1}|\mathcal{S}_{1}|\log\left(\frac{d}{|\mathcal{S}_{1}|}\right), \frac{2\log(2m_{1}+1)}{c_{3}\varepsilon^{2}}\right\}$$

hold, then $\widehat{\mathcal{S}_2} = \mathcal{S}_2$ and $\widehat{\mathcal{S}_1} = \mathcal{S}_1$ with probability at least

$$1 - \exp(-\tilde{c}_4 n_2) - \exp\left(-\frac{\tilde{c}_5 n_2 \varepsilon^2}{2}\right) - 2\exp(-\tilde{c}_2 n_1) - 2\exp\left(-\frac{c_3 \varepsilon^2 n_1}{2}\right).$$

The total number of queries made is

$$4(2m_2+1)^2 n_2 N_2 |\mathcal{H}_2^d| + 2(2m_1+1)n_1 N_1 = \Omega\left(|\mathcal{S}_2|\log(d^2/|\mathcal{S}_2|)|\mathcal{H}_2^d| + |\mathcal{S}_1|\log(d/|\mathcal{S}_1|)\right).$$

Proof. As in the bounded noise model, we only prove the exact recovery of S_2 . First, we note that in (4.4), as a consequence of resampling each query N_2 times and averaging, $\eta_{i,1}, \eta_{i,2}, \eta_{i,3}, \eta_{i,4} \sim \mathcal{N}(0, \frac{\sigma^2}{N_2}), i \in [n_2]$ are independent. Therefore $\eta_i \sim \mathcal{N}(0, \frac{4\sigma^2}{N_2}), i \in [n_2]$ are also independent and we set SPARSE-REC₂ = (P2) with $\nu_2 = 2\sigma(1 + \varepsilon)n_2/\sqrt{N_2}$.

From Corollary 2, we know that there exist constants $\tilde{c}_3, \tilde{c}_4, \tilde{c}_5, C_4 > 0$ such that if $n_2 \geq \tilde{c}_3|\mathcal{S}_2|\log(d^2/|\mathcal{S}_2|)$, then for any given $h \in \mathcal{H}_2^d$, $\mathbf{x} \in \chi(h)$, we have for the stated choice of ν_2 that

$$\| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \mathbf{z}^*(\mathbf{x}; \mathcal{A}) \|_{\infty} \le \| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \mathbf{z}^*(\mathbf{x}; \mathcal{A}) \|_1 \le C_4 \sqrt{\frac{2}{\pi}} (1+\varepsilon) \cdot \frac{2\sigma}{\sqrt{N_2}} =: \epsilon_2, \qquad (7.4)$$

with probability at least $1 - \exp(-\tilde{c}_4 n_2) - e \cdot \exp(-\tilde{c}_5 \varepsilon^2 n_2)$. This is true since $\mathbf{z}^*(\mathbf{x}; \mathcal{A})$ is always at most $|\mathcal{S}_2|$ sparse. Therefore, by taking the union bound, (7.4) holds uniformly for all $h \in \mathcal{H}_2^d$, $\mathbf{x} \in \chi(h)$, with probability at least

$$1 - \exp(-\tilde{c}_4 n_2) - (2m_2 + 1)^2 |\mathcal{H}_2^d| e \cdot \exp\left(-\tilde{c}_5 \varepsilon^2 n_2\right)$$

=1 - exp(- $\tilde{c}_4 n_2$) - exp $\left(\log[(2m_2 + 1)^2 |\mathcal{H}_2^d| e] - \tilde{c}_5 \varepsilon^2 n_2\right)$
 $\geq 1 - \exp(-\tilde{c}_4 n_2) - \exp\left(-\frac{\tilde{c}_5 n_2 \varepsilon^2}{2}\right)$ (7.5)

if $n_2 \geq \frac{2 \log[(2m_2+1)^2 e|\mathcal{H}_2^d]|}{\tilde{c}_5 \varepsilon^2}$ holds. We conclude that SPARSE-REC₂ is ϵ_2 -accurate for each $h \in \mathcal{H}_2^d$, $\mathbf{x} \in \chi(h)$ with probability at least (7.5).

By the stated choice of N_2 , we have $\epsilon_2 < \frac{D_2}{3}$ and using Lemma 3 and the condition on m_2 , we obtain $\widehat{S}_2 = S_2$. Lastly, the bound on the total number of queries follows from the expression in Lemma 3 (taking the resampling into account) by plugging in the stated bounds on m_i, n_i, N_i ; i = 1, 2.

7.3 Multivariate case

Finally, we consider the most general multivariate setting of Section 5. Based on Lemma 5, we analyze Algorithm 3 recovering the supports S_i ; $i \in [r_0]$. The recovery routines SPARSE-REC_i are realized by (P1) in (6.2) for i = 1 and for each $3 \le i \le r_0$ and SPARSE-REC₂ is instantiated with (P2) in (6.7). This leads to the results below.

Bounded noise model. This is the same noise model as described in the preceding subsections. The following theorem shows that if \triangle is sufficiently small, then Algorithm 3 recovers S_i exactly for all $i \in [r_0]$ provided the parameters $(m_i, n_i)_{i=1}^{r_0}$ are well chosen.

Theorem 8. For the bounded noise model with noise uniformly bounded by \triangle , consider Algorithm 3 with

- (a) SPARSE-REC_i instantiated with (P1) with $\nu_i = 2^i \triangle \sqrt{n_i}$ in (6.2) for $i \in \{3, \ldots, r_0\}$,
- (b) SPARSE-REC₂ instantiated with (P2) with $\nu_2 = 4 \triangle n_2$ in (6.7),
- (c) SPARSE-REC₁ instantiated with (P1) with $\nu_1 = 2 \triangle \sqrt{n_1}$ in (6.2), respectively.

Let $\mathcal{H}_i^{\mathcal{P}_i}$ be a (\mathcal{P}_i, i) hash family for each $2 \leq i \leq r_0$ and let \mathcal{P}_i denote the set $\mathcal{P} \subseteq [d]$ at the beginning of i^{th} iteration (with $\mathcal{P}_{r_0} = [d]$). If

$$\Delta < \min\left\{\min_{i \in \{3, \dots, r_0\}} \frac{D_i}{2^i 3 C_2}, \frac{D_2}{12 C_4}, \frac{D_1}{6 C_2}\right\}, \quad m_i \ge \left(\frac{3L(\sqrt{i})^{\alpha}}{D_i}\right)^{1/\alpha}; \quad i \in [r_0],$$

$$n_i \ge \tilde{c}_6 |\mathcal{S}_i|^2 \log \binom{|\mathcal{P}_i|}{i}; \quad i \in \{3, \dots, r_0\},$$

$$n_2 \ge \tilde{c}_3 |\mathcal{S}_2| \log(|\mathcal{P}_2|^2/|\mathcal{S}_2|) \quad and \quad n_1 \ge \tilde{c}_1 |\mathcal{S}_1| \log(|\mathcal{P}_1|/|\mathcal{S}_1|)$$

are satisfied, then $\widehat{S}_i = S_i$ for all $i \in [r_0]$ with probability at least

$$1 - \sum_{i=3}^{r_0} \exp\left(-\frac{\tilde{c}_7 n_i}{|\mathcal{S}_i|^2}\right) - \exp(-\tilde{c}_4 n_2) - 2\exp(-\tilde{c}_2 n_1).$$

Here, the constants C_4 , \tilde{c}_3 , $\tilde{c}_4 > 0$ are from Theorem 2, while \tilde{c}_1 , \tilde{c}_2 , $C_2 > 0$ are as defined in Theorem 4. The constants \tilde{c}_6 , $\tilde{c}_7 > 0$ depend only on the constants c_6 , $c_7 > 0$ defined in Theorem 3. The total number of queries made is

$$\Omega\left(\sum_{i=3}^{r_0} \left[c_i^i i^{i+1} |\mathcal{S}_i|^2 \log(|\mathcal{P}_i|) |\mathcal{H}_i^{\mathcal{P}_i}|\right] + |\mathcal{S}_2| \log\left(\frac{|\mathcal{P}_2|^2}{|\mathcal{S}_2|}\right) |\mathcal{H}_2^d| + |\mathcal{S}_1| \log\left(\frac{|\mathcal{P}_1|}{|\mathcal{S}_1|}\right)\right)$$

where each $c_i > 1$ depends on D_i, L, α .

Proof. Say we are at the beginning of i^{th} iteration with $3 \leq i \leq r_0$ and $\widehat{\mathcal{S}}_l = \mathcal{S}_l$ holds true for each l > i. Hence, the model has reduced to an order i sparse additive model on the set $\mathcal{P}_i \subseteq [d]$, with $\mathcal{S}_i^{(1)}, \mathcal{S}_{i-1}^{(1)}, \ldots, \mathcal{S}_1 \subset \mathcal{P}_i$.

From (5.5), we see for the noise vector $\boldsymbol{\eta} \in \mathbb{R}^{n_i}$ that

$$\eta_s = \sum_{z=1}^{2^i} (-1)^{\text{digit}(z-1)} \eta_{s,z} \quad \text{for all } s \in [n_i].$$
(7.6)

Since $|\eta_{s,z}| \leq \Delta$, this implies $\|\boldsymbol{\eta}\|_{\infty} \leq 2^i \Delta$ and thus $\|\boldsymbol{\eta}\|_2 \leq 2^i \Delta \sqrt{n_i}$. This bound holds uniformly for each **x** at which the linear system is formed. So we now instantiate SPARSE-REC_i with (P1) in (6.2), with $\nu_i = 2^i \Delta \sqrt{n_i}$.

As a consequence of part 1 of Theorem 3, there exists constants $\tilde{c}_6, \tilde{c}_7 > 0$ depending on $c_6, c_7 > 0$ (as defined in Theorem 3) so that if $n_i \geq \tilde{c}_6 |\mathcal{S}_i|^2 \log \binom{|\mathcal{P}_i|}{i}$, then with probability at least $1 - \exp\left(-\frac{\tilde{c}_7 n_i}{|\mathcal{S}_i|^2}\right)$, the matrix $\mathbf{B} \in \mathbb{R}^{n_i \times \binom{|\mathcal{P}_i|}{i}}$ satisfies ℓ_2/ℓ_2 RIP with $\delta_{2|\mathcal{S}_i|} < \sqrt{2} - 1$. Conditioning on this event, it follows from Theorem 3 that

$$\| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \underbrace{\mathbf{z}^*(\mathbf{x}; \mathcal{A})}_{|\mathcal{S}_i| \text{ sparse}} \|_{\infty} \leq \| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \mathbf{z}^*(\mathbf{x}; \mathcal{A}) \|_2 \leq 2^i \triangle C_2 =: \epsilon_i \text{ for all } \mathbf{x} \in \bigcup_{h \in \mathcal{H}_i^{\mathcal{P}_i}} \chi(h).$$

Thus, with probability at least $1 - \exp(-\frac{\tilde{c}_7 n_i}{|S_i|^2})$, SPARSE-REC_i is ϵ_i -accurate for each $h \in \mathcal{H}_i^{\mathcal{P}_i}$, $\mathbf{x} \in \chi(h)$. The assumption on \triangle ensures that $\epsilon_i < D_i/3$ and by Lemma 5 it follows that the stated choice of (m_i, ϵ_i) ensures exact recovery of S_i .

Hence if

$$\triangle < \min_{i \in \{3,\dots,r_0\}} \frac{D_i}{2^i 3 C_2}$$

is satisfied and m_i, n_i, ϵ_i satisfy their stated bounds (for $3 \le i \le r_0$), it follows by the union bound that $\widehat{S}_i = S_i$ holds for all $3 \le i \le r_0$ with probability at least

$$1 - \sum_{i=3}^{r_0} \exp\left(-\frac{\tilde{c}_7 n_i}{|\mathcal{S}_i|^2}\right).$$

Once S_i are identified exactly for all $3 \leq i \leq r_0$, we are left with a bivariate SPAM on the set \mathcal{P}_2 , with $S_1, S_2^{(1)} \subset \mathcal{P}_2$. Therefore by invoking Theorem 6, we see that if furthermore $\Delta < \min\left\{\frac{D_2}{12C_4}, \frac{D_1}{6C_2}\right\}$ holds, and m_1, n_1, m_2, n_2 satisfy their respective stated conditions, then the stated instantiations of SPARSE-REC₁, SPARSE-REC₂ (along with the stated choices of ν_1, ν_2) ensures $\widehat{S}_1 = S_1$ and $\widehat{S}_2 = S_2$ with probability at least $1 - \exp(-\tilde{c}_4 n_2) - 2\exp(-\tilde{c}_2 n_1)$. This completes the proof for exact recovery of S_i 's for all $i \in [r_0]$.

Finally, the stated sample complexity bound for the total number of queries made by Algorithm 3 follows in a straightforward manner by plugging in

$$m_i = \Omega\left(iL^{1/\alpha}D_i^{-1/\alpha}\right), \quad i \in [r_0]; \quad n_i = \Omega(i|\mathcal{S}_i|^2 \log |\mathcal{P}_i|), \quad 3 \le i \le r_0$$

along with the complexity bounds for n_1, n_2 into the expression for total number of samples from Lemma 5. This completes the proof.

Gaussian noise model. In the Gaussian noise model with noise samples i.i.d. Gaussian (~ $\mathcal{N}(0, \sigma^2)$) across queries, we again reduce the variance via re-sampling each query N_i times (during the estimation of S_i) for every $i \in [r_0]$ and averaging the values. We show that if the noise variance σ^2 is sufficiently small, then Algorithm 3 recovers S_i exactly for each $i \in [r_0]$, provided the parameters $m_i, n_i; i \in [r_0]$, are well chosen.

Theorem 9. For the Gaussian noise model with i.i.d. noise samples $\sim \mathcal{N}(0, \sigma^2)$, consider Algorithm 3 wherein we resample each query N_i times during estimation of S_i and average the values. Let

- (a) SPARSE-REC_i = (P1) with $\nu_i = 2^{i/2}(1+\varepsilon)\sigma\sqrt{n_i/N_i}$ in (6.2) for $3 \le i \le r_0$,
- (b) SPARSE-REC₂ = (P2) with $\nu_2 = 2(1 + \varepsilon)\sigma n_2/\sqrt{N_2}$ in (6.7) and
- (c) SPARSE-REC₁ = (P1) with $\nu_1 = \sqrt{2}(1+\varepsilon)\sigma\sqrt{n_1/N_1}$ in (6.2), respectively,

for some $\varepsilon \in (0,1)$. Let $\mathcal{H}_i^{\mathcal{P}_i}$ be a (\mathcal{P}_i, i) hash family for each $2 \leq i \leq r_0$, and denote \mathcal{P}_i to be the set $\mathcal{P} \subseteq [d]$ at the beginning of i^{th} iteration with $\mathcal{P}_{r_0} = [d]$. If

$$\begin{split} N_{i} &\geq \left\lfloor \frac{9C_{2}^{2}(1+\varepsilon)^{2}2^{i}\sigma^{2}}{D_{i}^{2}} \right\rfloor + 1, \quad n_{i} \geq \max\left\{ \tilde{c}_{6}|\mathcal{S}_{i}|^{2}\log\binom{|\mathcal{P}_{i}|}{i}, \frac{2\log[(2m_{i}+1)^{i}|\mathcal{H}_{i}^{\mathcal{P}_{i}}|]}{c_{3}\varepsilon^{2}} \right\}; \quad 3 \leq i \leq r_{0} \\ m_{i} &\geq \left(\frac{3L(\sqrt{i})^{\alpha}}{D_{i}}\right)^{1/\alpha}; \quad i \in [r_{0}], \\ N_{2} &\geq \left\lfloor \frac{72C_{4}^{2}(1+\varepsilon)^{2}\sigma^{2}}{\pi D_{2}^{2}} \right\rfloor + 1, \quad n_{2} \geq \max\left\{ \tilde{c}_{3}|\mathcal{S}_{2}|\log\left(\frac{|\mathcal{P}_{2}|^{2}}{|\mathcal{S}_{2}|}\right), \frac{2\log[(2m_{2}+1)^{2}e|\mathcal{H}_{2}^{\mathcal{P}_{2}}|]}{\tilde{c}_{5}\varepsilon^{2}} \right\}, \\ N_{1} &\geq \left\lfloor \frac{18C_{2}^{2}(1+\varepsilon)^{2}\sigma^{2}}{D_{1}^{2}} \right\rfloor + 1, \quad n_{1} \geq \max\left\{ \tilde{c}_{1}|\mathcal{S}_{1}|\log\left(\frac{|\mathcal{P}_{1}|}{|\mathcal{S}_{1}|}\right), \frac{2\log(2m_{1}+1)}{c_{3}\varepsilon^{2}} \right\} \end{split}$$

hold, then $\widehat{\mathcal{S}}_i = \mathcal{S}_i$ for all $i \in [r_0]$ with probability at least

$$1 - \sum_{i=3}^{r_0} \left[\exp\left(-\frac{\tilde{c}_7 n_i}{|\mathcal{S}_i|^2}\right) + 2\exp\left(-\frac{c_3\varepsilon^2 n_i}{2}\right) \right] - \exp(-\tilde{c}_4 n_2) - \exp\left(-\frac{\tilde{c}_5 n_2\varepsilon^2}{2}\right) - 2\exp(-\tilde{c}_2 n_1) - 2\exp\left(-\frac{c_3\varepsilon^2 n_1}{2}\right).$$
(7.7)

The constants $C_4, \tilde{c}_3, \tilde{c}_4, \tilde{c}_1, \tilde{c}_2, C_2, \tilde{c}_6, \tilde{c}_7 > 0$ are as explained in Theorem 8, while $\tilde{c}_5, c_3 > 0$ come from Corollaries 2, 1 respectively. The total number of queries made is

$$\sum_{i=1}^{r_0} N_i 2^i (2m_i + 1)^i n_i |\mathcal{H}_i^{\mathcal{P}_i}|$$

= $\Omega \left(\sum_{i=3}^{r_0} \left[\bar{c}_i^i i^{i+1} |\mathcal{S}_i|^2 \log(|\mathcal{P}_i|) |\mathcal{H}_i^{\mathcal{P}_i}| \right] + |\mathcal{S}_2| \log\left(\frac{|\mathcal{P}_2|^2}{|\mathcal{S}_2|}\right) |\mathcal{H}_2^{\mathcal{P}_2}| + |\mathcal{S}_1| \log\left(\frac{|\mathcal{P}_1|}{|\mathcal{S}_1|}\right) \right)$

where each $\bar{c}_i > 1$ depends on D_i, L, α .

Proof. Again, in the beginning of i^{th} iteration $3 \leq i \leq r_0$ with $\widehat{\mathcal{S}}_l = \mathcal{S}_l$ for each l > i, the model has reduced to an order i sparse additive model on the set $\mathcal{P}_i \subseteq [d]$, with $\mathcal{S}_i^{(1)}, \mathcal{S}_{i-1}^{(1)}, \ldots, \mathcal{S}_1 \subset \mathcal{P}_i$.

The noise vector is again given by (7.6) As a consequence of resampling each query point N_i times and averaging, we get $\eta_{s,z} \sim \mathcal{N}(0, \frac{\sigma^2}{N_i})$ i.i.d. for all s, z and, therefore, $\eta_s \sim \mathcal{N}(0, \frac{2^i \sigma^2}{N_i})$ i.i.d. for each s. From part 1 of Theorem 3, we know that if $n_i \geq \tilde{c}_6 |\mathcal{S}_i|^2 \log \binom{|\mathcal{P}_i|}{i}$, then with probability at least $1 - \exp\left(-\frac{\tilde{c}_7 n_i}{|\mathcal{S}_i|^2}\right)$ (with \tilde{c}_6, \tilde{c}_7 depending only on c_6, c_7), the matrix $\mathbf{B} \in \mathbb{R}^{n_i \times \binom{|\mathcal{P}_i|}{i}}$ satisfies ℓ_2/ℓ_2 RIP with $\delta_{2|\mathcal{S}_i|} < \sqrt{2} - 1$. Let us condition on this event. Then by setting SPARSE-REC_i = (P1) with $\nu_i = (1 + \varepsilon) \left(\frac{2^{i/2}\sigma\sqrt{n_i}}{\sqrt{N_i}}\right)$, and invoking Corollary 1, it follows for any given $h \in \mathcal{H}_i^{\mathcal{P}_i}$, $\mathbf{x} \in \chi(h)$ that

$$\| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \underbrace{\mathbf{z}^*(\mathbf{x}; \mathcal{A})}_{|\mathcal{S}_i| \text{ sparse}} \|_{\infty} \le \| \widehat{\mathbf{z}^*}(\mathbf{x}; \mathcal{A}) - \mathbf{z}^*(\mathbf{x}; \mathcal{A}) \|_2 \le 2^{i/2} C_2 (1+\varepsilon) \sigma \sqrt{n_i/N_i} =: \epsilon_i$$
(7.8)

with probability at least $1 - 2 \exp(-c_3 \varepsilon^2 n_i)$. By the union bound, it follows that (7.8) holds for all $h \in \mathcal{H}_i^{\mathcal{P}_i}$, $\mathbf{x} \in \chi(h)$, with probability at least

$$1 - 2(2m_i + 1)^i |\mathcal{H}_i^{\mathcal{P}_i}| \exp(-c_3 \varepsilon^2 n_i) = 1 - 2 \exp[\log[(2m_i + 1)^i |\mathcal{H}_i^{\mathcal{P}_i}|] - c_3 \varepsilon^2 n_i]$$

$$\geq 1 - 2 \exp\left(-\frac{c_3 \varepsilon^2 n_i}{2}\right)$$

if $n_i \geq \frac{2\log[(2m_i+1)^i|\mathcal{H}_i^{\mathcal{P}_i}|]}{c_3\varepsilon^2}$ holds. This gives us the stated condition on n_i for $3 \leq i \leq r_0$. Hence for the aforementioned choice of n_i , SPARSE-REC_i is ϵ_i -accurate for each $h \in \mathcal{H}_i^{\mathcal{P}_i}$, $\mathbf{x} \in \chi(h)$, with probability at least $1-2\exp\left(-\frac{c_3\varepsilon^2n_i}{2}\right) - \exp\left(-\frac{\tilde{c}_7n_i}{|\mathcal{S}_i|^2}\right)$. By the condition on N_i , we obtain $\epsilon_i < D_i/3$ and from Lemma 5, it follows that for the stated choice of m_i and ϵ_i , we have $\widehat{\mathcal{S}}_i = \mathcal{S}_i$. Thus we conclude by a union bound that $\widehat{\mathcal{S}}_i = \mathcal{S}_i$ holds for all $3 \leq i \leq r_0$, for the stated choice of $m_i, n_i, \epsilon_i, N_i$.

Finally, say $\widehat{\mathcal{S}}_i = \mathcal{S}_i$ holds for all $3 \leq i \leq r_0$. Then we are left with a bivariate SPAM on the set \mathcal{P}_2 , with $\mathcal{S}_1, \mathcal{S}_2^{(1)} \subset \mathcal{P}_2$. Thereafter, we only need to invoke Theorem 7, which guarantees $\widehat{\mathcal{S}}_2 = \mathcal{S}_2$ and $\widehat{\mathcal{S}}_1 = \mathcal{S}_1$ for the stated choices of SPARSE-REC_i, $\nu_i, m_i, n_i, \epsilon_i, N_i$; i = 1, 2. This completes the proof for the exact recovery of \mathcal{S}_i 's.

Since each query is resampled N_i times, with $N_i = \Omega(2^i)$, we obtain the stated sample complexity bound by proceeding as in the proof of Theorem 8.

8 Discussion

We now discuss how the components ϕ can be identified, and also compare our results with closely related work. Finally, we discuss an alternative approach described in [38] in more detail.

Identification of $\phi_{\mathbf{j}}$'s. Once the sets $\mathcal{S}_1, \mathcal{S}_2, \ldots, \mathcal{S}_{r_0}$ are known, then one can identify each component in the representation (5.1) by querying f along the corresponding canonical subspaces. Indeed, for a given $1 \leq p \leq r_0$, and $1 \leq r \leq p$, let us see how we can identify the *r*-variate component $\phi_{\mathbf{j}}$ for a given $\mathbf{j} \in \mathcal{S}_p^{(r)}$. Consider any $\mathbf{x} = (x_1 \ldots x_d)^T \in [-1, 1]^d$ which is supported on $\mathcal{S}_p^{(1)}$, i.e., $x_l = 0$ if $l \notin \mathcal{S}_p^{(1)}$. We then have from (5.1) that

$$f(\mathbf{x}) = \mu + \sum_{u \in \mathcal{S}_p^{(1)}} \phi_u(x_u) + \sum_{\mathbf{u} \in \mathcal{S}_p^{(2)}} \phi_{\mathbf{u}}(x_u) + \cdots + \sum_{\mathbf{u} \in \mathcal{S}_p^{(p-1)}} \phi_{\mathbf{u}}(x_u) + \sum_{\mathbf{u} \in \mathcal{S}_p} \phi_{\mathbf{u}}(x_u).$$

In particular, it follows from (2.3) that

$$\sum_{\mathbf{i} \subseteq \mathbf{j}} (-1)^{|\mathbf{j}| - |\mathbf{i}|} f(\Pi_{\mathbf{i}}(\mathbf{x})) = \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}),$$
(8.1)

with $\Pi_{\mathbf{i}}(\mathbf{x})$ denoting the projection of \mathbf{x} on the set of variables \mathbf{i} . Hence by querying f at the $2^{|\mathbf{j}|}$ points $\{\Pi_{\mathbf{i}}(\mathbf{x}) : \mathbf{i} \subseteq \mathbf{j}\}$, we can obtain the sample $\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}})$ via (8.1). Consequently, by choosing \mathbf{x}_{j} from a regular grid on $[-1, 1]^{\mathbf{j}}$, we can estimate $\phi_{\mathbf{j}}$ from its corresponding samples via standard quasi interpolants (in the noiseless case) or tools from non parametric regression (in the noisy case).

SPAMs. To begin with, we note that our work generalizes the recent results of Tyagi et al. [37, 38] in two fundamental ways. Firstly, we provide an algorithm for the general case where $r_0 \ge 2$ is possible, the results in [37, 38] are for the case $r_0 = 2$. Secondly, our results only require f to be Hölder continuous and not continuously differentiable as is the case in [37, 38]. We also mention that our sampling bounds for the case $r_0 = 2$ are linear in the sparsity $|S_1| + |S_2|$ even when the noise samples are i.i.d. Gaussian. However the algorithms in [37, 38] due to the localized nature of the sampling scheme, wherein finite difference operations are used to obtain linear measurements of the sparse gradient and Hessian of f. In the presence of noise, this essentially leads to the noise level getting scaled up by the step size parameter, and thus reducing the variance of noise necessarily leads to a resampling factor which is super linear in sparsity.

Dalalyan et al. [10] recently studied models of the form (1.1) with a small number m of swise interaction terms. They considered the Gaussian white noise model, which while not the same as the usual regression setup, is known to be asymptotically equivalent to the same. They derived non-asymptotic L_2 error rates in expectation for an estimator, with f lying in a Sobolev space, and showed the rates to be minimax optimal. However, they do not consider the problem of identification of the interaction terms. Moreover, as noted in [10], the computational cost of their method typically scales exponentially in m, s, d. Yang et al. [44] also studied models of the form (1.1) in a Bayesian setting, wherein they place a Gaussian prior (GP) on f and can carry out inference via the posterior probability distribution. They derive an estimator and provide error rates in the empirical L_2 norm for Hölder smooth f, but do not address the problem of identifying the interaction terms.

Functions with few active variables. There has been a fair amount of work in the literature on functions which intrinsically depend on a small subset of $k \ll d$ variables [11, 33, 9, 8]. To our knowledge, this model was first considered in [11], and in fact, our idea of using a family of hash functions is essentially motivated from [11] wherein such a family was used to construct the query points. A prototypical result in [11] is an algorithm that identifies the set of active variables with $(L+1)^k |\mathcal{H}_k^d| + k \log d$ queries, with L > 0 being the number of points on a uniform grid along a coordinate. The exponential dependence on k is unavoidable in the worst case, and indeed our bounds are also exponential in r_0 (see (1.2)).

- When $r_0 \ge k$, (1.1) is clearly a generalization of this model.
- In general, the model (1.1) is also a function of few active variables (those part of S_i 's); more precisely, at most $\sum_{i=1}^{r_0} i |S_i|$ variables. However, using a method that is designed generically for learning intrinsically k-variate functions would typically have sample complexity scaling exponentially with $\sum_{i=1}^{r_0} i |S_i|$. This is clearly suboptimal; our bounds in general depend at most polynomially on the size of S_i 's. This dependence is actually linear for the case $r_0 = 2$.

An alternative approach. Next, we discuss an alternative approach for learning the model (5.1) which was mentioned already in [38]. It is based on a truncated expansion in a bounded orthonormal system. For simplicity, we assume that f takes the form

$$f(\mathbf{x}) = \sum_{\mathbf{j} \in \mathcal{S}_{r_0}} \phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}), \quad \mathbf{x} \in [-1, 1]^d.$$
(8.2)

Let $\{\psi_k\}_{k\in\mathbb{Z}}$ be an orthonormal basis in $L_2([-1,1])$ with respect to the normalized Lebesgue measure on [-1,1]. We assume that $\psi_0 \equiv 1$ and we define $\{\psi_i\}_{i \in \mathbb{Z}^d}$ to be the tensor product orthonormal basis in $L_2([-1,1]^d)$, where

$$\psi_{\mathbf{i}}(\mathbf{x}) = \bigotimes_{l=1}^{d} \psi_{i_l}(x_l), \quad \mathbf{x} \in [-1, 1]^d.$$

For the components of (8.2) we obtain (for each $\mathbf{j} \in S_{r_0}$) the decomposition

$$\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}) = \sum_{\mathbf{i} \in \mathbb{Z}^d} a_{\mathbf{j},\mathbf{i}} \psi_{\mathbf{i}}(\mathbf{x}_{\mathbf{j}}) = \sum_{\mathbf{i} \in \mathbb{Z}^d: \text{supp } \mathbf{i} \subseteq \mathbf{j}} a_{\mathbf{j},\mathbf{i}} \psi_{\mathbf{i}}(\mathbf{x}_{\mathbf{j}}).$$
(8.3)

In the last identity, we used that $\int_{-1}^{1} \psi_k(t) dt = 0$ for every $k \in \mathbb{Z} \setminus \{0\}$ and, therefore, $a_{\mathbf{j},\mathbf{i}} =$ $\langle \phi_{\mathbf{i}}, \psi_{\mathbf{i}} \rangle = 0$ if $i_l \neq 0$ for some $l \notin \mathbf{j}$.

Using the smoothness of ϕ_i , we can truncate (8.3) at level $N \in \mathbb{N}$ (which we will determine later) and obtain

$$\phi_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}}) = \sum_{\substack{\mathbf{i} \in \mathbb{Z}^d: \|\mathbf{i}\|_{\infty} \le N \\ \text{supp} \, \mathbf{i} \subseteq \mathbf{j}}} a_{\mathbf{j}, \mathbf{i}} \psi_{\mathbf{i}}(\mathbf{x}_{\mathbf{j}}) + r_{\mathbf{j}}(\mathbf{x}_{j}) \quad \text{for all} \quad \mathbf{j} \in \mathcal{S}_{r_0}.$$
(8.4)

Summing up over $\mathbf{j} \in \mathcal{S}_{r_0}$ we arrive at

$$f(\mathbf{x}) = \sum_{\mathbf{j} \in \mathcal{S}_{r_0}} \left(\sum_{\substack{\mathbf{i} \in \mathbb{Z}^d: \|\mathbf{i}\|_{\infty} \leq N \\ \text{supp } \mathbf{i} \subseteq \mathbf{j}}} a_{\mathbf{j}, \mathbf{i}} \psi_{\mathbf{i}}(\mathbf{x}_{\mathbf{j}}) \right) + r(\mathbf{x}) \quad \text{with} \quad r(\mathbf{x}) = \sum_{\mathbf{j} \in \mathcal{S}_{r_0}} r_{\mathbf{j}}(\mathbf{x}_j).$$
(8.5)

The worst-case error of the uniform approximation of Hölder continuous functions with exponent $\alpha > 0$ (i.e., functions from the unit ball of C^{α}) is bounded from below by the Kolmogorov numbers of the embedding of C^{α} into L_{∞} , cf. [41],

$$||r_{\mathbf{j}}(\mathbf{x}_{\mathbf{j}})||_{\infty} \approx [(2N)^{r_0}]^{-\alpha/r_0} = (2N)^{-\alpha} \text{ and } ||r(\mathbf{x})||_{\infty} \approx |\mathcal{S}_{r_0}|(2N)^{-\alpha}.$$

Thus for any $\epsilon \in (0, 1)$, we typically require $N \gtrsim \left(\frac{|S_{r_0}|}{\epsilon}\right)^{1/\alpha}$ to ensure $||r(\mathbf{x})||_{\infty} \lesssim \epsilon$. Furthermore, the number of degrees of freedom in (8.5) is bounded by

$$D \ge \binom{d}{r_0} (2N)^{r_0}.$$

If the basis functions $\psi_{\mathbf{i}}$ are uniformly bounded (i.e. they form a Bounded Orthonormal System (BOS)), it was shown in [13, Theorem 12.31] or [31, Theorem 4.4], that one can recover $\mathbf{a} = (a_{\mathbf{j},\mathbf{i}})$ in (8.5) from $m \gtrsim |\mathcal{S}_{r_0}|(2N)^{r_0} \log^4(D)$ random samples of f by ℓ_1 -minimization. Plugging in our estimates on D and N, we arrive at $m \gtrsim |\mathcal{S}_{r_0}|^{1+r_0/\alpha} \cdot \log^4(d)$. This bound is always superlinear in $|\mathcal{S}_{r_0}|$ and (if $\alpha \in (0,1]$) with the power of dependence at least $1 + r_0$.

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A Proofs for Section 2

Proof of Proposition 1. For $U \subseteq [d]$, we define

$$(P_U f)(\mathbf{x}_U) = f\left(\sum_{j \in U} x_j \mathbf{e}_j\right)$$
 and $f_U(\mathbf{x}_U) = \sum_{V \subseteq U} (-1)^{|U| - |V|} (P_V f)(\mathbf{x}_V).$

Here, $\{\mathbf{e}_1, \ldots, \mathbf{e}_d\}$ is the canonical basis of \mathbb{R}^d .

By its definition, f_U is a continuous function. Furthermore,

$$\sum_{U \subseteq [d]} f_U(\mathbf{x}_U) = \sum_{U \subseteq [d]} \sum_{V \subseteq U} (-1)^{|U| - |V|} (P_V f)(\mathbf{x}_V) = \sum_{V \subseteq [d]} (P_V f)(\mathbf{x}_V) \sum_{U \supseteq V} (-1)^{|U| - |V|}$$
$$= \sum_{V \subseteq [d]} (P_V f)(\mathbf{x}_V) \sum_{W \subseteq [d] \setminus V} (-1)^{|W|}.$$

The last sum is equal to zero for all $V \subseteq [d]$ except V = [d]. This leads to (2.4).

If $x_j = 0$ for some $j \in U$, we get

$$f_U(\mathbf{x}_U) = \sum_{V \subseteq U} (-1)^{|U| - |V|} (P_V f)(\mathbf{x}_V)$$

=
$$\sum_{V \subseteq U \setminus \{j\}} \left[(-1)^{|U| - |V|} (P_V f)(\mathbf{x}_V) + (-1)^{|U| - |V \cup \{j\}|} (P_{V \cup \{j\}} f)(\mathbf{x}_{V \cup \{j\}}) \right] = 0$$

as all the terms in the last sum are equal to zero.

Finally, the uniqueness follows by induction. The statement is obvious for d = 1. Let now d > 1and let us assume that a given function $f \in C([-1, 1]^d)$ allows a decomposition

$$f(\mathbf{x}) = \sum_{U \subseteq [d]} f_U(\mathbf{x}_U), \quad \mathbf{x} \in [-1, 1]^d,$$

which satisfies the properties a)-c) of Proposition 1.

Let W be a proper subset of [d] and put for $\mathbf{x}_W \in [-1, 1]^{|W|}$

$$g_W(\mathbf{x}_W) = f\Big(\sum_{j \in W} x_j \mathbf{e}_j\Big).$$

Then $g_W \in C([-1, 1]^{|W|})$ and using c) of Proposition 1 we obtain

$$g_W(\mathbf{x}_W) = \sum_{U \subseteq [d]} f_U\left(\left(\sum_{j \in W} x_j \mathbf{e}_j\right)_U\right) = \sum_{U \subseteq [d]} f_U\left(\sum_{j \in W} x_j (\mathbf{e}_j)_U\right) = \sum_{U \subseteq W} f_U(\mathbf{x}_U).$$

This decomposition of g_W satisfies a)-c) of Proposition 1 with $|W| \leq d - 1$. By the induction assumption, this decomposition is therefore unique. We conclude, that f_U is uniquely determined for all $U \subset [d]$. Finally, also

$$f_{[d]}(\mathbf{x}) = f(\mathbf{x}) - \sum_{U \subset [d]} f_U(\mathbf{x}_U)$$

is uniquely determined.

Proof of Proposition 2. Let f be given by (4.1) and let us assume that it satisfies all the assumptions of Proposition 2. We show that (4.1) coincides with its Anchored-ANOVA decomposition as described in Proposition 1 and (4.1) is therefore unique.

Let $U \subseteq [d]$ with $|U| \ge 3$. Then

$$\begin{split} f_{U}(\mathbf{x}_{U}) &= \sum_{V \subseteq U} (-1)^{|U| - |V|} (P_{V}f)(\mathbf{x}_{V}) \\ &= \sum_{V \subseteq U} (-1)^{|U| - |V|} \Big\{ \mu + \sum_{j \in \mathcal{S}_{1} \cup \mathcal{S}_{2}^{\text{var}}} \phi_{j}((x_{V})_{j}) + \sum_{\mathbf{j} = (j_{1}, j_{2}) \in \mathcal{S}_{2}} \phi_{\mathbf{j}}((x_{V})_{j_{1}}, (x_{V})_{j_{2}}) \Big\} \\ &= \mu \sum_{V \subseteq U} (-1)^{|U| - |V|} + \sum_{j \in \mathcal{S}_{1} \cup \mathcal{S}_{2}^{\text{var}}} \phi_{j}(x_{j}) \sum_{\substack{V \subseteq U \\ j \in V}} (-1)^{|U| - |V|} + \sum_{j \in \mathcal{S}_{1} \cup \mathcal{S}_{2}^{\text{var}}} \phi_{j}(0) \sum_{\substack{V \subseteq U \\ j \notin V}} (-1)^{|U| - |V|} \\ &+ \sum_{\mathbf{j} = (j_{1}, j_{2}) \in \mathcal{S}_{2}} \sum_{V \subseteq U} (-1)^{|U| - |V|} \phi_{\mathbf{j}}((x_{V})_{j_{1}}, (x_{V})_{j_{2}}). \end{split}$$

It is easy to see, that the first three terms are zero. Finally, the last term can be split into a sum of four terms depending on if $j_1 \in V$ or $j_2 \in V$, i.e. terms of the kind

$$\sum_{\mathbf{j}=(j_1,j_2)\in\mathcal{S}_2} \phi_{\mathbf{j}}(x_{j_1},x_{j_2}) \sum_{\substack{V\subseteq U\\ j_1,j_2\in V}} (-1)^{|U|-|V|},$$

which also vanish. Therefore, $f_U(\mathbf{x}_U) = 0$.

If $U = \{j_1, j_2\}$ with $1 \le j_1 < j_2 \le d$, then $\emptyset, \{j_1\}, \{j_2\}$ and $\{j_1, j_2\}$ are the only subsets of U and we obtain

$$f_{U}(\mathbf{x}_{U}) = \sum_{V \subseteq U} (-1)^{|U| - |V|} (P_{V}f)(\mathbf{x}_{V})$$

$$= (P_{\emptyset}f)(0) - (P_{j_{1}}f)(x_{j_{1}}) - (P_{j_{2}}f)(x_{j_{2}}) + (P_{\{j_{1},j_{2}\}}f)(x_{j_{1}}, x_{j_{2}})$$

$$= f(0) - f(x_{j_{1}}\mathbf{e}_{j_{1}}) - f(x_{j_{2}}\mathbf{e}_{j_{2}}) + f(x_{j_{1}}\mathbf{e}_{j_{1}} + x_{j_{2}}\mathbf{e}_{j_{2}})$$

$$= \sum_{j \in \mathcal{S}_{1} \cup \mathcal{S}_{2}^{\text{var}}} [\phi_{j}(0) - \phi_{j}((x_{j_{1}}\mathbf{e}_{j_{1}})_{j}) - \phi_{j}((x_{j_{2}}\mathbf{e}_{j_{2}})_{j}) + \phi_{j}((x_{j_{1}}\mathbf{e}_{j_{1}} + x_{j_{2}}\mathbf{e}_{j_{2}})_{j})]$$

$$+ \sum_{\mathbf{j} \in \mathcal{S}_{2}} [\phi_{\mathbf{j}}(0) - \phi_{\mathbf{j}}((x_{j_{1}}\mathbf{e}_{j_{1}})_{\mathbf{j}}) - \phi_{\mathbf{j}}((x_{j_{2}}\mathbf{e}_{j_{2}})_{\mathbf{j}}) + \phi_{\mathbf{j}}((x_{j_{1}}\mathbf{e}_{j_{1}} + x_{j_{2}}\mathbf{e}_{j_{2}})_{\mathbf{j}})].$$

The first sum vanishes for all $j \in S_1 \cup S_2^{\text{var}}$, which can be easily observed by considering the options $j \notin \{j_1, j_2\}, j = j_1$, or $j = j_2$. If $(j_1, j_2) \notin S_2$, then also the second sum vanishes. If $(j_1, j_2) \in S_2$, then

$$f_{j_1,j_2}(x_{j_1},x_{j_2}) = \phi_{j_1,j_2}(x_{j_1},x_{j_2}) - \phi_{j_1,j_2}(x_{j_1},0) - \phi_{j_1,j_2}(0,x_{j_2}) + \phi_{j_1,j_2}(0,0) = \phi_{j_1,j_2}(x_{j_1},x_{j_2}).$$

Finally, if $l \in [d]$ then \emptyset and $\{l\}$ are the only subsets of $\{l\}$. If furthermore $l \notin S_1 \cup S_2^{\text{var}}$, we get

$$f_{\{l\}}(x_l) = -f(0) + f(x_l \mathbf{e}_l) = -\left(\mu + \sum_{j \in \mathcal{S}_1 \cup \mathcal{S}_2^{\text{var}}} \phi_j(0) + \sum_{\mathbf{j} \in \mathcal{S}_2} \phi_{\mathbf{j}}(0)\right) + \left(\mu + \sum_{j \in \mathcal{S}_1 \cup \mathcal{S}_2^{\text{var}}} \phi_j(0) + \sum_{\mathbf{j} \in \mathcal{S}_2} \phi_{\mathbf{j}}(0)\right) = 0.$$

Similarly, $f_{\{l\}}(x_l) = \phi_l(x_l)$ if $l \in \mathcal{S}_1 \cup \mathcal{S}_2^{\text{var}}$.

B Some standard concentration results

First, we recall that the sub-Gaussian norm of a random variable X is defined as

$$\|X\|_{\psi_2} = \sup_{p \ge 1} p^{-1/2} (\mathbb{E}|X|^p)^{1/p}$$

and X is called sub-Gaussian random variable if $|| X ||_{\psi_2}$ is finite. Similarly, the sub-exponential norm of a random variable is the quantity

$$\| X \|_{\psi_1} = \sup_{p \ge 1} p^{-1} (\mathbb{E} |X|^p)^{1/p}$$

and X is called sub-exponential if $||X||_{\psi_1}$ is finite. Next, we recall the following concentration results for sums of i.i.d. sub-Gaussian and sub-exponential random variables.

Proposition 4. [39, Proposition 5.16] Let X_1, \ldots, X_N be independent centered sub-exponential random variables, and $K = \max_i || X_i ||_{\psi_1}$. Then for every $\mathbf{a} = (a_1, \ldots, a_N) \in \mathbb{R}^N$ and every $t \ge 0$, we have

$$\mathbb{P}\left(\left|\sum_{i} a_{i} X_{i}\right| \geq t\right) \leq 2 \exp\left[-c \min\left\{\frac{t^{2}}{K^{2} \parallel \mathbf{a} \parallel_{2}^{2}}, \frac{t}{K \parallel \mathbf{a} \parallel_{\infty}}\right\}\right],$$

where c > 0 is an absolute constant.

Proposition 5. [39, Proposition 5.10] Let X_1, \ldots, X_N be independent centered sub-Gaussian random variables, and $K = \max_i \parallel X_i \parallel_{\psi_2}$. Then for every $\mathbf{a} = (a_1, \ldots, a_N) \in \mathbb{R}^N$ and every $t \ge 0$, we have

$$\mathbb{P}\left(\left|\sum_{i} a_{i} X_{i}\right| \geq t\right) \leq e \cdot \exp\left[-\frac{ct^{2}}{K^{2} \parallel \mathbf{a} \parallel_{2}^{2}}\right],$$

where c > 0 is an absolute constant.

Let $\boldsymbol{\eta} = (\eta_1 \ \eta_2 \ \dots \ \eta_n)^T$, where $\eta_i \sim \mathcal{N}(0, \sigma^2)$ are i.i.d. for each *i*. The Proposition below is a standard concentration result, stating that $\| \boldsymbol{\eta} \|_2 = \Theta(\sigma \sqrt{n})$ and $\| \boldsymbol{\eta} \|_1 = \Theta(\sigma n)$, with high probability. We provide proofs for completeness.

Proposition 6. Let $\boldsymbol{\eta} = (\eta_1 \ \eta_2 \ \dots \ \eta_n)^T$ where $\eta_i \sim \mathcal{N}(0, \sigma^2)$ i.i.d for each *i*. Then, there exists constants $c_1, c_2 > 0$ so that for any $\epsilon \in (0, 1)$, we have:

1.
$$\mathbb{P}(\parallel \boldsymbol{\eta} \parallel_2 \in [(1-\epsilon)\sigma\sqrt{n}, (1+\epsilon)\sigma\sqrt{n}]) \ge 1 - 2\exp\left(-c_1\epsilon^2n\right), and$$

2.
$$\mathbb{P}\left(\parallel \boldsymbol{\eta} \parallel_{1} \in \left[(1-\epsilon)n\sigma\sqrt{\frac{2}{\pi}}, (1+\epsilon)n\sigma\sqrt{\frac{2}{\pi}}\right]\right) \geq 1-e \cdot \exp\left(-\frac{2c_{2}\epsilon^{2}n}{\pi}\right).$$

Proof. 1. Note that η_i are i.i.d. sub-Gaussian random variables with $\| \eta_i \|_{\psi_2} \leq C_1 \sigma$. Hence η_i^2 are i.i.d. sub-exponential³ with

$$\| \eta_i^2 \|_{\psi_1} \le 2 \| \eta_i \|_{\psi_2}^2 \le C_2 \sigma^2$$

for some constant $C_2 > 0$. This implies that $\eta_i^2 - \mathbb{E}[\eta_i^2]$ are i.i.d. sub-exponential with

$$\parallel \eta_i^2 - \mathbb{E}[\eta_i^2] \parallel_{\psi_1} \leq 2 \parallel \eta_i^2 \parallel_{\psi_1} \leq C_3 \sigma^2$$

for some constant $C_3 > 0$. Using Proposition 4 with $t = n\epsilon\sigma^2$ for $0 < \epsilon < 1$, we obtain for some constant $c_1 > 0$

$$\mathbb{P}\left(\left|\sum_{i=1}^{n} (\eta_i^2 - \mathbb{E}[\eta_i^2])\right| \le n\epsilon\sigma^2\right) \ge 1 - 2\exp\left[-c_1\min\left\{\epsilon^2, \epsilon\right\}n\right].$$

Together with some standard manipulations this completes the proof.

2. Note that $|\eta_i| - \mathbb{E}[|\eta_i|]$ is sub-Gaussian with

$$\| |\eta_i| - \mathbb{E}[|\eta_i|] \|_{\psi_2} \le 2 \| \eta_i \|_{\psi_2} \le C\sigma,$$

for some constant C > 0. Using Proposition 5 with $t = n \epsilon \mathbb{E}[|\eta_1|]$ for $\epsilon > 0$, we hence obtain

$$\mathbb{P}\left(\left|\sum_{i=1}^{n} (|\eta_i| - \mathbb{E}[|\eta_i|])\right| \le n\epsilon \mathbb{E}[|\eta_1|]\right) \ge 1 - e \cdot \exp\left[-\frac{c_2 n\epsilon^2 (\mathbb{E}[|\eta_1|])^2}{\sigma^2}\right]$$

³A random variable X is sub-Gaussian iff X^2 is sub-exponential. Moreover, $||X||_{\psi_2}^2 \leq ||X^2||_{\psi_1} \leq 2 ||X||_{\psi_2}^2$ (cf. [39, Lemma 5.14]).

for some constant $c_2 > 0$. Finally, observing that $\mathbb{E}[|\eta_1|] = \sigma \sqrt{\frac{2}{\pi}}$ (cf., [43]) completes the proof.

C Proof of Theorem 2

The proof of part (1) follows in the same manner as [5, Proposition 1], with minor differences in calculation at certain parts. For completeness, we outline the main steps below.

Invoking the Hanson-Wright inequality for quadratic forms [17], we get for some constant c > 0and all t > 0

$$\mathbb{P}(|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta} - \mathbb{E}[\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}]| > t) \le 2\exp\left[-c\min\left\{\frac{t^{2}}{K^{4} \parallel \mathbf{A} \parallel_{F}^{2}}, \frac{t}{K^{2} \parallel \mathbf{A} \parallel}\right\}\right],$$
(C.1)

where $\parallel \beta_i \parallel_{\psi_2} \leq K$. For β_i Bernoulli symmetric, K = 1 and

$$\mathbb{E}[\boldsymbol{\beta}^T \mathbf{A} \boldsymbol{\beta}] = \mathbb{E} \sum_{i \neq j} \beta_i \beta_j A_{ij} = 0.$$

Using $\| \mathbf{A} \| \leq \| \mathbf{A} \|_{F}$, (C.1) implies for every t > 0

$$\mathbb{P}(|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}| > t) \le 2\exp\left[-c\min\left\{\frac{t^{2}}{\|\mathbf{A}\|_{F}^{2}}, \frac{t}{\|\mathbf{A}\|_{F}}\right\}\right].$$
(C.2)

We will now find upper and lower bounds on $\mathbb{E}[|\boldsymbol{\beta}^T \mathbf{A}\boldsymbol{\beta}|]$. The upper bound is easy since (C.2) implies

$$\mathbb{E}[|\boldsymbol{\beta}^T \mathbf{A}\boldsymbol{\beta}|] = \int_0^\infty \mathbb{P}(|\boldsymbol{\beta}^T \mathbf{A}\boldsymbol{\beta}| > t) dt \le c' \parallel \mathbf{A} \parallel_F.$$
(C.3)

In order to find the lower bound, we have via repeated application of Cauchy Schwartz inequality

$$\left(\mathbb{E}[|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}|^{2}]\right)^{2} \leq \mathbb{E}[|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}|] \cdot \mathbb{E}[|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}|^{3}] \leq \mathbb{E}[|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}|] \cdot \left(\mathbb{E}[|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}|^{2}]\right)^{1/2} \cdot \left(\mathbb{E}[|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}|^{4}]\right)^{1/2}$$

and

$$\mathbb{E}[|\boldsymbol{\beta}^T \mathbf{A} \boldsymbol{\beta}|] \geq \sqrt{\frac{(\mathbb{E}[|\boldsymbol{\beta}^T \mathbf{A} \boldsymbol{\beta}|^2])^3}{\mathbb{E}[|\boldsymbol{\beta}^T \mathbf{A} \boldsymbol{\beta}|^4]}}$$

Since β consists of i.i.d. Bernoulli symmetric random variables, we obtain

$$\mathbb{E}[|\boldsymbol{\beta}^T \mathbf{A} \boldsymbol{\beta}|^2] = 2 \| \mathbf{A} \|_F^2 = \| \mathbf{a} \|_2^2$$

Moreover, an argument similar to (C.3) gives $\mathbb{E}[|\boldsymbol{\beta}^T \mathbf{A} \boldsymbol{\beta}|^4] \leq c'' \parallel \mathbf{A} \parallel_F^4$. Hence,

$$\mathbb{E}[|\boldsymbol{\beta}^{T}\mathbf{A}\boldsymbol{\beta}|] \geq \sqrt{\frac{8 \|\mathbf{A}\|_{F}^{6}}{c'' \|\mathbf{A}\|_{F}^{4}}} = \tilde{c} \|\mathbf{A}\|_{F}.$$
(C.4)

Eqs. (C.3), (C.4) give us upper and lower bounds for $\mathbb{E}[|\boldsymbol{\beta}^T \mathbf{A}\boldsymbol{\beta}|]$. As a last step, we consider the zero mean random variables X_1, \ldots, X_n , where $X_i = |\boldsymbol{\beta}_i^T \mathbf{A}\boldsymbol{\beta}_i| - \mathbb{E}|\boldsymbol{\beta}_i^T \mathbf{A}\boldsymbol{\beta}_i|$. A simple modification of (C.3) shows that they are sub-exponential with $||X_i||_{\psi_1} \leq c ||\mathbf{A}||_F$. We can therefore apply a standard concentration bound (see [39, Proposition 5.16]) to bound the deviation

$$\left|\frac{1}{n} \parallel \mathbf{B}\mathbf{a} \parallel_1 - \frac{1}{n} \mathbb{E}[\parallel \mathbf{B}\mathbf{a} \parallel_1]\right| = \frac{1}{n} \left|\sum_{i=1}^n X_i\right|.$$

A straightforward calculation then yields the statement of part (1) of the Theorem.

Part (2) follows from standard arguments based on ϵ -nets, detailed for instance in [1].

The proof of part (3) copies that of [5, Theorem 3], which again is inspired by [4]. We sketch the main steps below for completeness. Denoting $\hat{\mathbf{a}} = \mathbf{a} + \mathbf{h}$, we have by feasibility of \mathbf{a} that $\frac{1}{n} \| \mathbf{Bh} \|_{1} \leq \frac{2\nu}{n}$. Denoting Ω_{0} to be set of indices corresponding to the k largest entries of \mathbf{a} , we get $\mathbf{a} = \mathbf{a}_{\Omega_{0}} + \mathbf{a}_{\Omega_{0}^{c}}$. For a suitable positive integer K, we define Ω_{1} as the set of indices of K largest entries of $\mathbf{h}_{\Omega_{0}^{c}}$, Ω_{2} as the set of indices of K largest entries of \mathbf{h} on $(\Omega_{0} \cup \Omega_{1})^{c}$ and so on. Following this argument of [4] gives the proof.

The proof of part (4) follows by choosing $K = 4\left(\frac{4c_2}{c_1}\right)^2 k$. Indeed, (6.9) gives

$$\frac{1-\gamma_{k+K}^{\rm lb}}{\sqrt{2}} - (1+\gamma_K^{\rm ub})\sqrt{\frac{k}{K}} \ge \frac{c_1}{2\sqrt{2}} - 2c_2 \cdot \frac{c_1}{8c_2} = \frac{(\sqrt{2}-1)c_1}{4} = \beta > 0$$

if $n > c'_3(k+K)\log(d^2/(k+K))$, with probability at least $1 - e^{-C_4n}$ for some constant $C_4 > 0$.