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EMPIRICAL BAYES ORACLE UNCERTAINTY QUANTIFICATION FOR REGRESSION

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We propose an empirical Bayes method for high-dimensional linear regression models. Following an oracle approach that quantifies the error locally for each possible value of the parameter, we show that an empirical Bayes posterior contracts at the optimal rate at all parameters and leads to uniform size-optimal credible balls with guaranteed coverage under an “excessive bias restriction” condition. This condition gives rise to a new slicing of the entire space that is suitable for ensuring uniformity in uncertainty quantification. The obtained results immediately lead to optimal contraction and coverage properties for many conceivable classes simultaneously. The results are also extended to high-dimensional additive nonparametric regression models.

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ASYMPTOTIC JOINT DISTRIBUTION OF EXTREME EIGENVALUES AND TRACE OF LARGE SAMPLE COVARIANCE MATRIX IN A GENERALIZED SPIKED POPULATION MODEL

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This paper studies the joint limiting behavior of extreme eigenvalues and trace of large sample covariance matrix in a generalized spiked population model, where the asymptotic regime is such that the dimension and sample size grow proportionally. The form of the joint limiting distribution is applied to conduct Johnson–Graybill-type tests, a family of approaches testing for signals in a statistical model. For this, higher order correction is further made, helping alleviate the impact of finite-sample bias. The proof rests on determining the joint asymptotic behavior of two classes of spectral processes, corresponding to the extreme and linear spectral statistics, respectively.

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SINGULARITY, MISSPECIFICATION AND THE CONVERGENCE RATE OF EM

BY RAAZ DWIVEDI^{1,*}, NHAT HO^{1,†}, KOULIK KHAMARU²,
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A line of recent work has analyzed the behavior of the Expectation-Maximization (EM) algorithm in the well-specified setting, in which the population likelihood is locally strongly concave around its maximizing argument. Examples include suitably separated Gaussian mixture models and mixtures of linear regressions. We consider over-specified settings in which the number of fitted components is larger than the number of components in the true distribution. Such mis-specified settings can lead to singularity in the Fisher information matrix, and moreover, the maximum likelihood estimator based on n i.i.d. samples in d dimensions can have a nonstandard $\mathcal{O}((d/n)^{\frac{1}{4}})$ rate of convergence. Focusing on the simple setting of two-component mixtures fit to a d -dimensional Gaussian distribution, we study the behavior of the EM algorithm both when the mixture weights are different (unbalanced case), and are equal (balanced case). Our analysis reveals a sharp distinction between these two cases: in the former, the EM algorithm converges geometrically to a point at Euclidean distance of $\mathcal{O}((d/n)^{\frac{1}{2}})$ from the true parameter, whereas in the latter case, the convergence rate is exponentially slower, and the fixed point has a much lower $\mathcal{O}((d/n)^{\frac{1}{4}})$ accuracy. Analysis of this singular case requires the introduction of some novel techniques: in particular, we make use of a careful form of localization in the associated empirical process, and develop a recursive argument to progressively sharpen the statistical rate.

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TOWARDS OPTIMAL ESTIMATION OF BIVARIATE ISOTONIC MATRICES WITH UNKNOWN PERMUTATIONS

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Many applications, including rank aggregation, crowd-labeling and graphon estimation, can be modeled in terms of a bivariate isotonic matrix with unknown permutations acting on its rows and/or columns. We consider the problem of estimating an unknown matrix in this class, based on noisy observations of (possibly, a subset of) its entries. We design and analyze polynomial-time algorithms that improve upon the state of the art in two distinct metrics, showing, in particular, that minimax optimal, computationally efficient estimation is achievable in certain settings. Along the way, we prove matching upper and lower bounds on the minimax radii of certain cone testing problems, which may be of independent interest.¹

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HIGH-DIMENSIONAL CONSISTENT INDEPENDENCE TESTING WITH MAXIMA OF RANK CORRELATIONS

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Testing mutual independence for high-dimensional observations is a fundamental statistical challenge. Popular tests based on linear and simple rank correlations are known to be incapable of detecting nonlinear, nonmonotone relationships, calling for methods that can account for such dependences. To address this challenge, we propose a family of tests that are constructed using maxima of pairwise rank correlations that permit consistent assessment of pairwise independence. Built upon a newly developed Cramér-type moderate deviation theorem for degenerate U-statistics, our results cover a variety of rank correlations including Hoeffding’s D , Blum–Kiefer–Rosenblatt’s R and Bergsma–Dassios–Yanagimoto’s τ^* . The proposed tests are distribution-free in the class of multivariate distributions with continuous margins, implementable without the need for permutation, and are shown to be rate-optimal against sparse alternatives under the Gaussian copula model. As a by-product of the study, we reveal an identity between the aforementioned three rank correlation statistics, and hence make a step towards proving a conjecture of Bergsma and Dassios.

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OPTIMAL RATES OF ENTROPY ESTIMATION OVER LIPSCHITZ BALLS

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We consider the problem of minimax estimation of the entropy of a density over Lipschitz balls. Dropping the usual assumption that the density is bounded away from zero, we obtain the minimax rates $(n \ln n)^{-s/(s+d)} + n^{-1/2}$ for $0 < s \leq 2$ for densities supported on $[0, 1]^d$, where s is the smoothness parameter and n is the number of independent samples. We generalize the results to densities with unbounded support: given an Orlicz functions Ψ of rapid growth (such as the subexponential and sub-Gaussian classes), the minimax rates for densities with bounded Ψ -Orlicz norm increase to $(n \ln n)^{-s/(s+d)} (\Psi^{-1}(n))^{d(1-d/p(s+d))} + n^{-1/2}$, where p is the norm parameter in the Lipschitz ball. We also show that the integral-form plug-in estimators with kernel density estimates fail to achieve the minimax rates, and characterize their worst case performances over the Lipschitz ball.

One of the key steps in analyzing the bias relies on a novel application of the Hardy–Littlewood maximal inequality, which also leads to a new inequality on the Fisher information that may be of independent interest.

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LIMIT DISTRIBUTION THEORY FOR BLOCK ESTIMATORS IN MULTIPLE ISOTONIC REGRESSION

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We study limit distributions for the tuning-free max–min block estimator originally proposed in (Fokianos, Leucht and Neumann (2017)) in the problem of multiple isotonic regression, under both fixed lattice design and random design settings. We show that, if the regression function f_0 admits vanishing derivatives up to order α_k along the k th dimension ($k = 1, \dots, d$) at a fixed point $x_0 \in (0, 1)^d$, and the errors have variance σ^2 , then the max–min block estimator $\hat{f}_n$ satisfies

$$(n_*/\sigma^2)^{\frac{1}{2+\sum_{k \in \mathcal{D}_*} \alpha_k - 1}} (\hat{f}_n(x_0) - f_0(x_0)) \rightsquigarrow \mathbb{C}(f_0, x_0).$$

Here, $\mathcal{D}_*$, n_* , depending on $\{\alpha_k\}$ and the design points, are the set of all “effective dimensions” and the size of “effective samples” that drive the asymptotic limiting distribution, respectively. If furthermore either $\{\alpha_k\}$ are relative primes to each other or all mixed derivatives of f_0 of certain critical order vanish at x_0 , then the limiting distribution can be represented as $\mathbb{C}(f_0, x_0) =_d K(f_0, x_0) \cdot \mathbb{D}_\alpha$, where $K(f_0, x_0)$ is a constant depending on the local structure of the regression function f_0 at x_0 , and $\mathbb{D}_\alpha$ is a nonstandard limiting distribution generalizing the well-known Chernoff distribution in univariate problems. The above limit theorem is also shown to be optimal both in terms of the local rate of convergence and the dependence on the unknown regression function whenever such dependence is explicit (i.e., $K(f_0, x_0)$), for the full range of $\{\alpha_k\}$ in a local asymptotic minimax sense.

There are two interesting features in our local theory. First, the max–min block estimator automatically adapts to the local smoothness and the intrinsic dimension of the isotonic regression function at the optimal rate. Second, the optimally adaptive local rates are in general not the same in fixed lattice and random designs. In fact, the local rate in the fixed lattice design case is no slower than that in the random design case, and can be much faster when the local smoothness levels of the isotonic regression function or the sizes of the lattice differ substantially along different dimensions.

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ASSESSMENT OF THE EXTENT OF CORROBORATION OF AN ELABORATE THEORY OF A CAUSAL HYPOTHESIS USING PARTIAL CONJUNCTIONS OF EVIDENCE FACTORS

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An elaborate theory of predictions of a causal hypothesis consists of several falsifiable statements derived from the causal hypothesis. Statistical tests for the various pieces of the elaborate theory help to clarify how much the causal hypothesis is corroborated. In practice, the degree of corroboration of the causal hypothesis has been assessed by a verbal description of which of the several tests provides evidence for which of the several predictions. This verbal approach can miss quantitative patterns. In this paper, we develop a quantitative approach. We first decompose these various tests of the predictions into independent factors with different sources of potential biases. Support for the causal hypothesis is enhanced when many of these evidence factors support the predictions. A sensitivity analysis is used to assess the potential bias that could make the finding of the tests spurious. Along with this multiparameter sensitivity analysis, we consider the partial conjunctions of the tests. These partial conjunctions quantify the evidence supporting various fractions of the collection of predictions. A partial conjunction test involves combining tests of the components in the partial conjunction. We find the asymptotically optimal combination of tests in the context of a sensitivity analysis. Our analysis of an elaborate theory of a causal hypothesis controls for the familywise error rate.

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NONPARAMETRIC DRIFT ESTIMATION FOR I.I.D. PATHS OF STOCHASTIC DIFFERENTIAL EQUATIONS

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We consider N independent stochastic processes $(X_i(t), t \in [0, T])$, $i = 1, \dots, N$, defined by a one-dimensional stochastic differential equation, which are continuously observed throughout a time interval $[0, T]$ where T is fixed. We study nonparametric estimation of the drift function on a given subset A of $\mathbb{R}$. Projection estimators are defined on finite dimensional subsets of $\mathbb{L}^2(A, dx)$. We stress that the set A may be compact or not and the diffusion coefficient may be bounded or not. A data-driven procedure to select the dimension of the projection space is proposed where the dimension is chosen within a random collection of models. Upper bounds of risks are obtained, the assumptions are discussed and simulation experiments are reported.

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DISTANCE-BASED AND RKHS-BASED DEPENDENCE METRICS IN HIGH DIMENSION

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In this paper, we study distance covariance, Hilbert–Schmidt covariance (aka Hilbert–Schmidt independence criterion [In *Advances in Neural Information Processing Systems* (2008) 585–592]) and related independence tests under the high dimensional scenario. We show that the sample distance/Hilbert–Schmidt covariance between two random vectors can be approximated by the sum of squared componentwise sample cross-covariances up to an asymptotically constant factor, which indicates that the standard distance/Hilbert–Schmidt covariance based test can only capture linear dependence in high dimension. Under the assumption that the components within each high dimensional vector are weakly dependent, the distance correlation based t test developed by Székely and Rizzo (*J. Multivariate Anal.* **117** (2013) 193–213) for independence is shown to have trivial limiting power when the two random vectors are nonlinearly dependent but componentwisely uncorrelated. This new and surprising phenomenon, which seems to be discovered and carefully studied for the first time, is further confirmed in our simulation study. As a remedy, we propose tests based on an aggregation of marginal sample distance/Hilbert–Schmidt covariances and show their superior power behavior against their joint counterparts in simulations. We further extend the distance correlation based t test to those based on Hilbert–Schmidt covariance and marginal distance/Hilbert–Schmidt covariance. A novel unified approach is developed to analyze the studentized sample distance/Hilbert–Schmidt covariance as well as the studentized sample marginal distance covariance under both null and alternative hypothesis. Our theoretical and simulation results shed light on the limitation of distance/Hilbert–Schmidt covariance when used jointly in the high dimensional setting and suggest the aggregation of marginal distance/Hilbert–Schmidt covariance as a useful alternative.

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THE DISTANCE STANDARD DEVIATION

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The distance standard deviation, which arises in distance correlation analysis of multivariate data, is studied as a measure of spread. The asymptotic distribution of the empirical distance standard deviation is derived under the assumption of finite second moments. Applications are provided to hypothesis testing on a data set from materials science and to multivariate statistical quality control. The distance standard deviation is compared to classical scale measures for inference on the spread of heavy-tailed distributions. Inequalities for the distance variance are derived, proving that the distance standard deviation is bounded above by the classical standard deviation and by Gini's mean difference. New expressions for the distance standard deviation are obtained in terms of Gini's mean difference and the moments of spacings of order statistics. It is also shown that the distance standard deviation satisfies the axiomatic properties of a measure of spread.

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ROBUST MULTIVARIATE NONPARAMETRIC TESTS VIA PROJECTION AVERAGING

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In this work, we generalize the Cramér–von Mises statistic via projection averaging to obtain a robust test for the multivariate two-sample problem. The proposed test is consistent against all fixed alternatives, robust to heavy-tailed data and minimax rate optimal against a certain class of alternatives. Our test statistic is completely free of tuning parameters and is computationally efficient even in high dimensions. When the dimension tends to infinity, the proposed test is shown to have comparable power to the existing high-dimensional mean tests under certain location models. As a by-product of our approach, we introduce a new metric called *the angular distance* which can be thought of as a robust alternative to the Euclidean distance. Using the angular distance, we connect the proposed method to the reproducing kernel Hilbert space approach. In addition to the Cramér–von Mises statistic, we demonstrate that the projection-averaging technique can be used to define robust multivariate tests in many other problems.

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INFERENCE FOR CONDITIONAL VALUE-AT-RISK OF A PREDICTIVE REGRESSION

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Conditional value-at-risk is a popular risk measure in risk management. We study the inference problem of conditional value-at-risk under a linear predictive regression model. We derive the asymptotic distribution of the least squares estimator for the conditional value-at-risk. Our results relax the model assumptions made in (*Oper. Res.* **60** (2012) 739–756) and correct their mistake in the asymptotic variance expression. We show that the asymptotic variance depends on the quantile density function of the unobserved error and whether the model has a predictor with infinite variance, which makes it challenging to actually quantify the uncertainty of the conditional risk measure. To make the inference feasible, we then propose a smooth empirical likelihood based method for constructing a confidence interval for the conditional value-at-risk based on either independent errors or GARCH errors. Our approach not only bypasses the challenge of directly estimating the asymptotic variance but also does not need to know whether there exists an infinite variance predictor in the predictive model. Furthermore, we apply the same idea to the quantile regression method, which allows infinite variance predictors and generalizes the parameter estimation in (*Econometric Theory* **22** (2006) 173–205) to conditional value-at-risk in the Supplementary Material. We demonstrate the finite sample performance of the derived confidence intervals through numerical studies before applying them to real data.

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SIMULTANEOUS HIGH-PROBABILITY BOUNDS ON THE FALSE DISCOVERY PROPORTION IN STRUCTURED, REGRESSION AND ONLINE SETTINGS

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While traditional multiple testing procedures prohibit adaptive analysis choices made by users, Goeman and Solari (*Statist. Sci.* **26** (2011) 584–597) proposed a simultaneous inference framework that allows users such flexibility while preserving high-probability bounds on the false discovery proportion (FDP) of the chosen set. In this paper, we propose a new class of such simultaneous FDP bounds, tailored for nested sequences of rejection sets. While most existing simultaneous FDP bounds are based on closed testing using global null tests based on sorted p -values, we additionally consider the setting where side information can be leveraged to boost power, the variable selection setting where knockoff statistics can be used to order variables, and the online setting where decisions about rejections must be made as data arrives. Our finite-sample, closed form bounds are based on repurposing the FDP estimates from false discovery rate (FDR) controlling procedures designed for each of the above settings. These results establish a novel connection between the parallel literatures of simultaneous FDP bounds and FDR control methods, and use proof techniques employing martingales and filtrations that are new to both these literatures. We demonstrate the utility of our results by augmenting a recent knockoffs analysis of the UK Biobank dataset.

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CLUSTERING IN BLOCK MARKOV CHAINS

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This paper considers cluster detection in Block Markov Chains (BMCs). These Markov chains are characterized by a block structure in their transition matrix. More precisely, the n possible states are divided into a finite number of K groups or clusters, such that states in the same cluster exhibit the same transition rates to other states. One observes a trajectory of the Markov chain, and the objective is to recover, from this observation only, the (initially unknown) clusters. In this paper, we devise a clustering procedure that accurately, efficiently and provably detects the clusters. We first derive a fundamental information-theoretical lower bound on the detection error rate satisfied under any clustering algorithm. This bound identifies the parameters of the BMC, and trajectory lengths, for which it is possible to accurately detect the clusters. We next develop two clustering algorithms that can together accurately recover the cluster structure from the shortest possible trajectories, whenever the parameters allow detection. These algorithms thus reach the fundamental detectability limit, and are optimal in that sense.

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MODEL SELECTION AND LOCAL GEOMETRY

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We consider problems in model selection caused by the geometry of models close to their points of intersection. In some cases—including common classes of causal or graphical models, as well as time series models—distinct models may nevertheless have identical tangent spaces. This has two immediate consequences: first, in order to obtain constant power to reject one model in favour of another we need local alternative hypotheses that decrease to the null at a slower rate than the usual parametric $n^{-1/2}$ (typically we will require $n^{-1/4}$ or slower); in other words, to distinguish between the models we need large effect sizes or very large sample sizes. Second, we show that under even weaker conditions on their tangent cones, models in these classes cannot be made simultaneously convex by a reparameterization.

This shows that Bayesian network models, amongst others, cannot be learned directly with a convex method similar to the graphical lasso. However, we are able to use our results to suggest methods for model selection that learn the tangent space directly, rather than the model itself. In particular, we give a generic algorithm for learning Bayesian network models.

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IRREDUCIBILITY AND GEOMETRIC ERGODICITY OF HAMILTONIAN MONTE CARLO

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Hamiltonian Monte Carlo (HMC) is currently one of the most popular Markov Chain Monte Carlo algorithms to sample smooth distributions over continuous state space. This paper discusses the irreducibility and geometric ergodicity of the HMC algorithm. We consider cases where the number of steps of the Störmer–Verlet integrator is either fixed or random. Under mild conditions on the potential U associated with target distribution π , we first show that the Markov kernel associated to the HMC algorithm is irreducible and positive recurrent. Under more stringent conditions, we then establish that the Markov kernel is Harris recurrent. We provide verifiable conditions on U under which the HMC sampler is geometrically ergodic. Finally, we illustrate our results on several examples.

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TEST FOR HIGH DIMENSIONAL COVARIANCE MATRICES

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The paper introduces a new test for testing structures of covariances for high dimensional vectors and the data dimension can be much larger than the sample size. Under proper normalization, central and noncentral limit theorems are established. The asymptotic theory is attained without imposing any explicit restriction between data dimension and sample size. To facilitate the related statistical inference, we propose the balanced Rademacher weighted differencing scheme, which is also the delete-half jackknife, to approximate the distribution of the proposed test statistics. We also develop a new testing procedure for substructures of precision matrices. The simulation results show that the tests outperform the existing methods both in terms of size and power. Our test procedure is applied to a colorectal cancer dataset.

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OPTIMAL ESTIMATION OF VARIANCE IN NONPARAMETRIC REGRESSION WITH RANDOM DESIGN

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Consider the heteroscedastic nonparametric regression model with random design

$$Y_i = f(X_i) + V^{1/2}(X_i)\varepsilon_i, \quad i = 1, 2, \dots, n,$$

with $f(\cdot)$ and $V(\cdot)$ α - and β -Hölder smooth, respectively. We show that the minimax rate of estimating $V(\cdot)$ under both local and global squared risks is of the order

$$n^{-\frac{8\alpha\beta}{4\alpha\beta+2\alpha+\beta}} \vee n^{-\frac{2\beta}{2\beta+1}},$$

where $a \vee b := \max\{a, b\}$ for any two real numbers a, b . This result extends the fixed design rate $n^{-4\alpha} \vee n^{-2\beta/(2\beta+1)}$ derived in (*Ann. Statist.* **36** (2008) 646–664) in a nontrivial manner, as indicated by the appearances of both α and β in the first term. In the special case of constant variance, we show that the minimax rate is $n^{-8\alpha/(4\alpha+1)} \vee n^{-1}$ for variance estimation, which further implies the same rate for quadratic functional estimation and thus unifies the minimax rate under the nonparametric regression model with those under the density model and the white noise model. To achieve the minimax rate, we develop a U-statistic-based local polynomial estimator and a lower bound that is constructed over a specified distribution family of randomness designed for both ε_i and X_i .

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ANALYSIS OF A TWO-LAYER NEURAL NETWORK VIA DISPLACEMENT CONVEXITY

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Fitting a function by using linear combinations of a large number N of “simple” components is one of the most fruitful ideas in statistical learning. This idea lies at the core of a variety of methods, from two-layer neural networks to kernel regression, to boosting. In general, the resulting risk minimization problem is nonconvex and is solved by gradient descent or its variants. Unfortunately, little is known about global convergence properties of these approaches.

Here, we consider the problem of learning a concave function f on a compact convex domain $\Omega \subset \mathbb{R}^d$, using linear combinations of “bump-like” components (neurons). The parameters to be fitted are the centers of N bumps, and the resulting empirical risk minimization problem is highly nonconvex. We prove that, in the limit in which the number of neurons diverges, the evolution of gradient descent converges to a Wasserstein gradient flow in the space of probability distributions over Ω . Further, when the bump width δ tends to 0, this gradient flow has a limit which is a viscous porous medium equation. Remarkably, the cost function optimized by this gradient flow exhibits a special property known as *displacement convexity*, which implies exponential convergence rates for $N \rightarrow \infty$, $\delta \rightarrow 0$.

Surprisingly, this asymptotic theory appears to capture well the behavior for moderate values of δ , N . Explaining this phenomenon, and understanding the dependence on δ , N in a quantitative manner remains an outstanding challenge.

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BEYOND GAUSSIAN APPROXIMATION: BOOTSTRAP FOR MAXIMA OF SUMS OF INDEPENDENT RANDOM VECTORS

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The Bonferroni adjustment, or the union bound, is commonly used to study rate optimality properties of statistical methods in high-dimensional problems. However, in practice, the Bonferroni adjustment is overly conservative. The extreme value theory has been proven to provide more accurate multiplicity adjustments in a number of settings, but only on an ad hoc basis. Recently, Gaussian approximation has been used to justify bootstrap adjustments in large scale simultaneous inference in some general settings when $n \gg (\log p)^7$, where p is the multiplicity of the inference problem and n is the sample size. The thrust of this theory is the validity of the Gaussian approximation for maxima of sums of independent random vectors in high dimension. In this paper, we reduce the sample size requirement to $n \gg (\log p)^5$ for the consistency of the empirical bootstrap and the multiplier/wild bootstrap in the Kolmogorov–Smirnov distance, possibly in the regime where the Gaussian approximation is not available. New comparison and anticoncentration theorems, which are of considerable interest in and of themselves, are developed as existing ones interweaved with Gaussian approximation are no longer applicable or strong enough to produce desired results.

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ISOTONIC REGRESSION IN MULTI-DIMENSIONAL SPACES AND GRAPHS

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In this paper, we study minimax and adaptation rates in general isotonic regression. For uniform deterministic and random designs in $[0, 1]^d$ with $d \geq 2$ and $N(0, 1)$ noise, the minimax rate for the ℓ_2 risk is known to be bounded from below by $n^{-1/d}$ when the unknown mean function f is non-decreasing and its range is bounded by a constant, while the least squares estimator (LSE) is known to nearly achieve the minimax rate up to a factor $(\log n)^\gamma$ where n is the sample size, $\gamma = 4$ in the lattice design and $\gamma = \max\{9/2, (d^2 + d + 1)/2\}$ in the random design. Moreover, the LSE is known to achieve the adaptation rate $(K/n)^{-2/d} \{1 \vee \log(n/K)\}^{2\gamma}$ when f is piecewise constant on K hyperrectangles in a partition of $[0, 1]^d$.

Due to the minimax theorem, the LSE is identical on every design point to both the max-min and min-max estimators over all upper and lower sets containing the design point. This motivates our consideration of estimators which lie in-between the max-min and min-max estimators over possibly smaller classes of upper and lower sets, including a subclass of block estimators. Under a q th moment condition on the noise, we develop ℓ_q risk bounds for such general estimators for isotonic regression on graphs. For uniform deterministic and random designs in $[0, 1]^d$ with $d \geq 3$, our ℓ_2 risk bound for the block estimator matches the minimax rate $n^{-1/d}$ when the range of f is bounded and achieves the near parametric adaptation rate $(K/n)\{1 \vee \log(n/K)\}^d$ when f is K -piecewise constant. Furthermore, the block estimator possesses the following oracle property in variable selection: When f depends on only a subset S of variables, the ℓ_2 risk of the block estimator automatically achieves up to a poly-logarithmic factor the minimax rate based on the oracular knowledge of S .

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ROBUST BAYES-LIKE ESTIMATION: RHO-BAYES ESTIMATION

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We observe n independent random variables with joint distribution $\mathbf{P}$ and pretend that they are i.i.d. with some common density s (with respect to a known measure μ) that we wish to estimate. We consider a density model $\bar{\mathcal{S}}$ for s that we endow with a prior distribution π (with support in $\bar{\mathcal{S}}$) and build a robust alternative to the classical Bayes posterior distribution which possesses similar concentration properties around s whenever the data are truly i.i.d. and their density s belongs to the model $\bar{\mathcal{S}}$. Furthermore, in this case, the Hellinger distance between the classical and the robust posterior distributions tends to 0, as the number of observations tends to infinity, under suitable assumptions on the model and the prior. However, unlike what happens with the classical Bayes posterior distribution, we show that the concentration properties of this new posterior distribution are still preserved when the model is misspecified or when the data are not i.i.d. but the marginal densities of their joint distribution are close enough in Hellinger distance to the model $\bar{\mathcal{S}}$.

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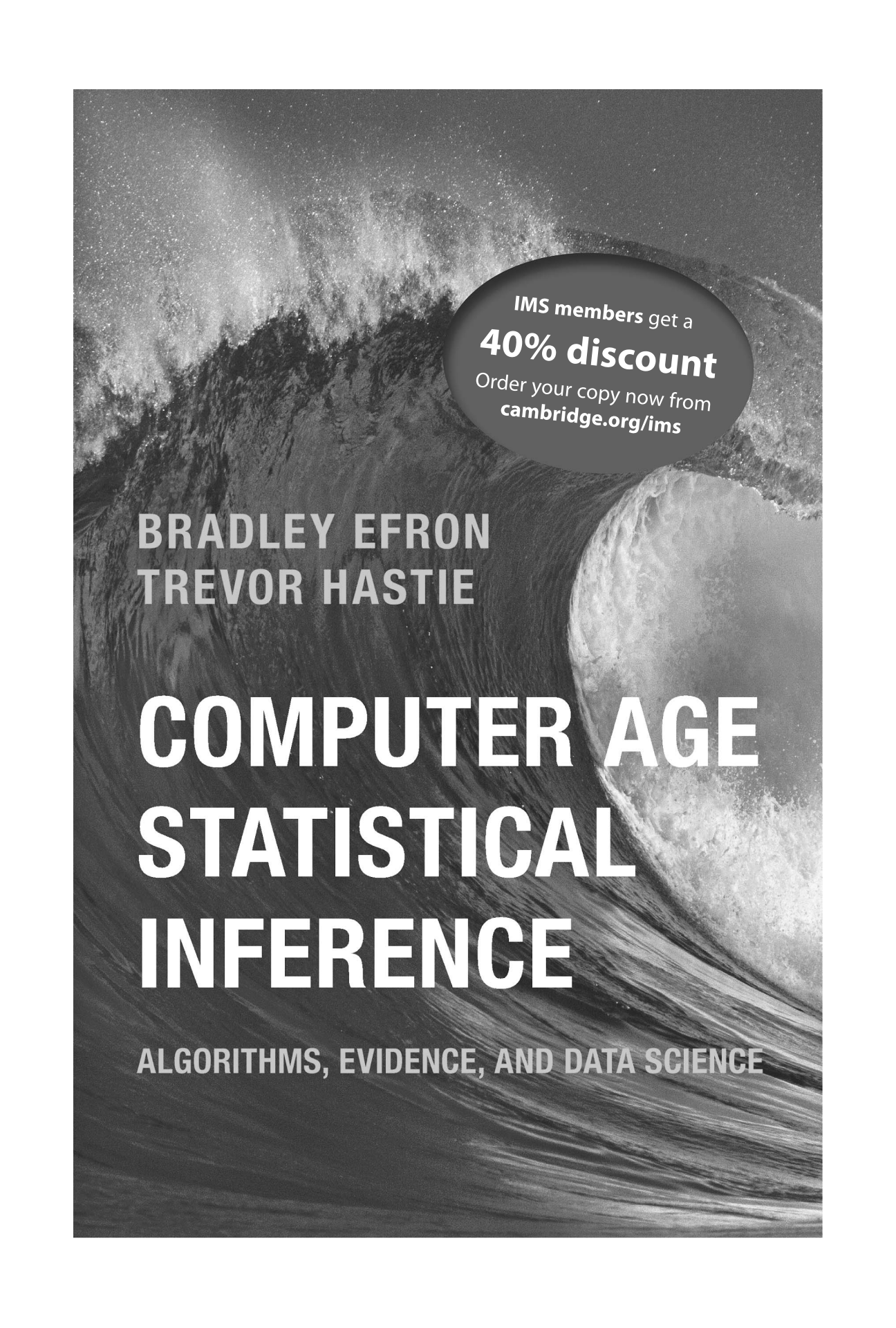
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