

1 Thanks to the reviewers for the insightful and constructive feedback. It will surely improve the manuscript. Due to
 2 space constraints, instead of responding point-by-point, we address points in common with multiple reviews. All
 3 minor comments made by reviewer #1 have been addressed and incorporated into a revised version of the paper.

4
 5 **i.i.d. Gaussian measurement matrix assumption.** While, in general,
 6 AMP theory provides performance guarantees only for i.i.d. sub-Gaussian
 7 data, in practice, favorable performance of AMP seems to be more
 8 universal. For example, in Fig. 1a, we illustrate the performance of AMP
 9 for i.i.d. zero mean, $1/n$ variance design matrices that are *not* Gaussian
 10 (one i.i.d. ± 1 Bernoulli (top) and one i.i.d. shifted exponential (bottom)).
 11 In both cases, AMP converges very fast, thus demonstrating its robustness
 12 to distributional assumptions.

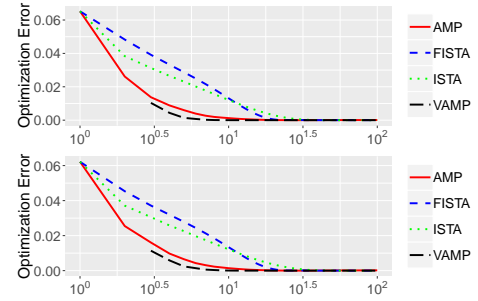
13 Recent work proposes a variant of AMP, called vector-AMP or VAMP,
 14 which is a computationally-efficient algorithm that *provably* works for
 15 a wide range of design matrices, namely, those that are right rotationally-
 16 invariant. We thank reviewer #2 for pointing us to “AMP for convex
 17 optimization with nonseparable penalties” by Manoel et al, which studies
 18 VAMP for a similar setting as SLOPE. However, the type of nonsep-
 19 arability considered in the referenced work requires the penalty to be
 20 separable on subsets of an affine transformation of its input. As such,
 21 the setting does not directly apply to SLOPE, but we have built a hybrid,
 22 “SLOPE VAMP”, based on code generously shared by the authors of
 23 the referenced work, which performs very well in the (non-) i.i.d. (non-)
 24 Gaussian regime (see Fig. 1a and 1b). Motivated by these promising empir-
 25 ical results, we feel that theoretically understanding SLOPE dynamics
 26 with VAMP is an exciting direction that we plan to pursue in other work.

27 **Known signal prior assumption.** We are intrigued by reviewer #3’s
 28 suggestion of using EM- or SURE-based AMP strategies to remove this
 29 assumption. We would like to pursue this within the SLOPE framework,
 30 though we haven’t done so at this time. We believe that developing
 31 such strategies alongside SLOPE VAMP would provide a quite general
 32 framework for recovery of the SLOPE estimator.

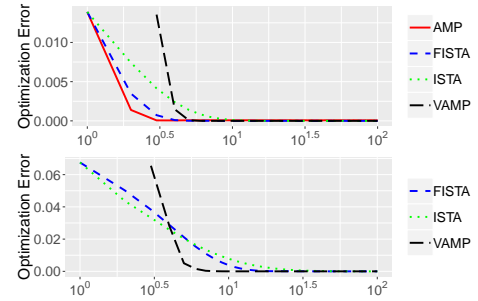
33 **Comparison to MMSE AMP.** In general, the (statistical) motivation for
 34 using methods like LASSO or SLOPE is to perform variable selection,
 35 and in addition, for SLOPE, to control the false discovery rate. Both
 36 methods are therefore biased and, consequently, MMSE AMP strategies
 37 will, by design, outperform *if performance is based on MSE*. To combine
 38 the best of both methods, one could also incorporate a “debiasing” device
 39 in SLOPE AMP, à la “Debiasing the LASSO: optimal sample size for
 40 Gaussian designs” by Javanmard & Montanari, but we will leave this for
 41 future work. Nevertheless, Fig. 1c suggests that SLOPE AMP has MSE
 42 that is not too much worse than MMSE AMP.

43 **Comparison to [20] and [12].** While [20] have the same asymptotic
 44 analysis, we have a clear, rigorous statement of where it applies. That is,
 45 the analysis in [20] applies *if* the state evolution has a unique fixed point,
 46 and our Thm. 1 states precise conditions under which this is true. More-
 47 over, we believe that our algorithmic approach offers a more concrete
 48 connection between the finite-sample behavior of the SLOPE estimator
 49 and its asymptotic distribution. We also agree with reviewer #2 that
 50 a discussion of the main results of [12] to highlight the gap between
 51 optimal estimators and any convex penalty would be useful for readers
 52 and we will add it to the final manuscript.

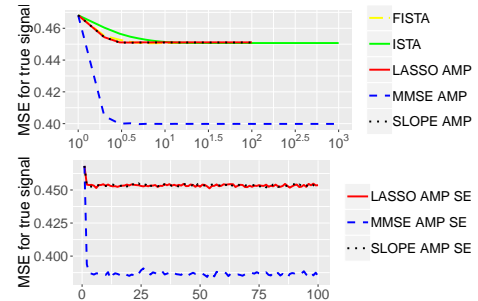
53 **Real-world data.** We agree with reviewer #1 that performing an empirical study on real-world data will significantly
 54 strengthen our results. Previous research has tested SLOPE performance on Genome-Wide Association Studies (GWAS)
 55 data (see “The Northern Finland Birth Cohort of 1966 (NFBC)”). Due to its precedence in the SLOPE literature and its
 56 inherent scientific importance, we intend to test SLOPE AMP on this data, however, there are restrictions on its use and
 57 we are currently undergoing the protocols needed to be granted access to the data by the NIH.



(a) i.i.d. ± 1 Bernoulli design matrix (top) and i.i.d. shifted exponential design matrix (bottom)



(b) i.i.d. Gaussian design matrix (top) and non-i.i.d. right rotationally-invariant design matrix where AMP diverges (bottom)



(c) i.i.d. Gaussian design matrix

Figure 1: Performance of AMP variants in different settings with Bernoulli-Gaussian prior, dimension = 1000, and sample size = 300.