# Fast and reliable confidence intervals for a variance component or proportion

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#### Abstract

We show that confidence intervals for a variance component or proportion, with asymptotically correct uniform coverage probability, can be obtained by inverting certain test-statistics based on the score for the restricted likelihood. The results apply in settings where the variance or proportion is near or at the boundary of the parameter set. Simulations indicate the proposed test-statistics are approximately pivotal and lead to confidence intervals with near-nominal coverage even in small samples. We illustrate our methods' application in spatially-resolved transcriptomics where we compute approximately 15,000 confidence intervals, used for gene ranking, in less than 4 minutes. In the settings we consider, the proposed method is between two and 28,000 times faster than popular alternatives, depending on how many confidence intervals are computed.

## 1 Introduction

Variance component models are widely used in applied statistics. Often, researchers are interested in quantifying how much of the variability in the data is due to a

specific source, possibly after controlling for, or conditioning on, other sources of variability. In modern genetics and genomics, for example, it is common to assume a vector  $Y \in \mathbb{R}^n$  of responses for n sample units satisfies

$$Y \sim \mathcal{N}_n(X\beta, \sigma_q^2 K + \sigma_e^2 I_n), \tag{1}$$

with  $X \in \mathbb{R}^{n \times p}$  a design matrix,  $K \in \mathbb{R}^{n \times n}$  a known positive semi-definite matrix, and parameter  $\theta = (\beta, \sigma_g^2, \sigma_e^2) \in \mathbb{R}^p \times [0, \infty) \times (0, \infty)$  (Purcell, 2002; Kang et al., 2010; Zhang et al., 2010; Lippert et al., 2011; Yang et al., 2011; Zhou and Stephens, 2012; Zhou et al., 2013; Loh et al., 2015; Weissbrod et al., 2016; Moore et al., 2019). In statistical genetics, when elements of Y are phenotypes for n individuals, K can be a genetic relatedness matrix quantifying similarities between individuals' common SNP genotypes (Jiang et al., 2022). Then, the quantity of inferential interest is the heritability (Visscher et al., 2008)

$$h^2 = \frac{\sigma_g^2}{\sigma_a^2 + \sigma_e^2} \in [0, 1). \tag{2}$$

In spatially resolved transcriptomics (Marx, 2021), when elements of Y are a gene's expression at n distinct two- or three-dimensional spatial positions in a tissue sample and K is a matrix encoding distances between positions,  $h^2$  is known as the fraction of spatial variance (Svensson et al., 2018; Kats et al., 2021; Weber et al., 2022; Zeng et al., 2022).

Though (1) may seem a simple model, inference on  $h^2$  can be complicated, especially when it is nearly, or equal to, zero or one, as those are boundary points of the parameter set. It is well-known that common test-statistics, such as Wald and likelihood ratio statistics, have non-standard distributions on boundary points. The distributions can often be obtained by simulation (e.g., Crainiceanu and Ruppert, 2004; Schweiger et al., 2016; Guédon et al., 2023) or approximated, either using asymptotic theory (e.g., Self and Liang, 1987; Geyer, 1994; Stram and Lee, 1994, 1995; Baey et al., 2019), or by other means (e.g. Wood, 2013). Thus, it is often possible to construct tests with (asymptotically) correct pointwise size, and the likelihood ratio

test has recently been suggested for use in related settings (Battey and McCullagh, 2023). However, creating a reliable confidence region by inverting such a test-statistic is more complicated, for at least two reasons.

First, one in general has to compute the statistic for every relevant  $h^2$ . Thus, methods which are designed specifically for testing  $h^2 = 0$  are inapplicable, and simulation-based methods can be prohibitively time-consuming. Secondly, even when inverting a test with asymptotically correct pointwise size is computationally feasible, the resulting confidence region can have poor coverage properties in finite samples. Specifically, the actual coverage probability often depends substantially on how close the true parameter is to the boundary. To illustrate the issues, Fig. 1 shows coverage probabilities for different values of  $h^2$  for Wald, likelihood ratio, and proposed confidence intervals defined in Section 2. For interior points of the parameter set, the confidence intervals are obtained by inverting test-statistics using quantiles of the chi-square distribution with one degree of freedom, as suggested by classical theory. Notably, the coverage probabilities for Wald and likelihood ratio intervals are substantially different from the nominal level when  $h^2$  is near zero or one. The reason is that, when the true parameter is close to the boundary, the distributions of the test-statistics are close to what they are on the boundary. Thus, even when the parameter is in the interior, classical asymptotic theory for interior points provides a poor approximation. Removing the boundary points from the parameter set or using a larger sample size is not sufficient to address the issue, as Fig. 1 illustrates; loosely speaking, for every sample size, there are interior points close enough to the boundary to cause problems. More formally, test-statistics with different asymptotic distributions on boundary and interior points lead to asymptotically incorrect uniform coverage probability (Ekvall and Bottai, 2022, Lemma 2.5).

For reasons to become apparent, we focus on test-statistics based on the score and information of the restricted likelihood. By doing so we are able to construct a  $(1 - \alpha)$  confidence region  $\mathcal{R}_n(\alpha)$  that satisfies, for any compact  $C \subseteq [0, 1]$  and as  $n \to \infty$ ,

$$\sup_{\theta \in C} \left| P_{\theta} \{ h^2 \in \mathcal{R}_n(\alpha) \} - (1 - \alpha) \right| \to 0.$$
 (3)

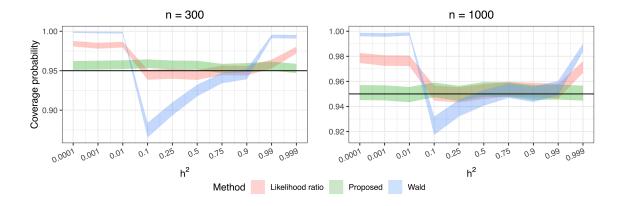


Figure 1: Monte Carlo estimates of coverage probabilities for  $h^2$  in (1) with  $\sigma_g^2 + \sigma_e^2 = 1$ ,  $\beta = 0, X \in \mathbb{R}^{n \times 5}$  a matrix of independent standard normal entries, and  $K_{ij} = 0.95^{|i-j|}$ . The solid horizontal line indicates the nominal level, 0.95. The line widths provide 95% confidence bands for the coverage probability based on  $10^4$  trials.

That is, the proposed confidence region, which is typically an interval, has asymptotically correct uniform coverage probability on compact sets. This property leads to near-nominal coverage of every practically relevant  $h^2$ , including  $h^2$  near or equal to zero, as is illustrated in Fig. 1. We emphasize that when focusing on uniform inference in the presence of boundary points, conclusions about score test-statistics in general do not carry over to Wald and likelihood ratio test-statistics. Indeed, our results show, in a sense to be made precise, that certain score-statistics are asymptotically pivotal, but Wald and likelihood ratio test-statistics are not. The score statistics can therefore be inverted using the same quantiles for every point in the parameter set, leading to a confidence interval that is easy to implement and orders of magnitude faster to compute than competing ones. In a typical example we consider, our confidence intervals take less than a second to compute, whereas the competitors often take 500 seconds, and sometimes much longer.

Previously, the score standardized by expected information has been shown to give confidence regions with asymptotically correct uniform coverage probability in settings with singular Fisher information (Bottai, 2003; Ekvall and Bottai, 2022). We build on ideas from that literature here, but the setting, and hence the arguments required

for the main results, are quite different. In particular, here, the information matrix is positive definite under weak conditions (Theorem 1). There are also two nuisance parameters,  $\beta$  and  $\sigma^2 = \sigma_g^2 + \sigma_e^2$ . While dealing with  $\beta$  is relatively straightforward since  $\beta$  and  $(h^2, \sigma^2)$  are orthogonal in the Fisher information sense, our results are, to the best of our knowledge, the first of their kind with non-orthogonal nuisance parameters. In contrast to classical asymptotic theory for inference in mixed models, we also let the number of predictors grow with the sample size. This is an important development as the restricted likelihood is typically used precisely when the number of predictors is smaller than the number of observations, but large enough to induce substantial bias in the inference on  $h^2$ .

#### 2 Model

Let  $\Sigma = \text{cov}_{\theta}(Y) = \sigma_g^2 K + \sigma_e^2 I_n$ . A log-likelihood corresponding to (1) is

$$l(\beta, \sigma_g^2, \sigma_e^2) = l(\beta, \sigma_g^2, \sigma_e^2; Y, X, K) = -\frac{1}{2} \left\{ \log |\Sigma| + (Y - X\beta)^{\mathrm{T}} \Sigma^{-1} (Y - \beta) \right\}, \quad (4)$$

the domain of which is the parameter set  $\mathbb{R}^p \times [0, \infty) \times (0, \infty)$ . It is important that (4) allows  $\sigma_g^2 = 0$ , which is often of practical interest. When K is positive definite, points with  $\sigma_e^2 = 0$  and  $\sigma_g^2 > 0$  can be added to the parameter set, but with semi-definite K, (4) requires  $\sigma_e^2 > 0$ . Assuming X has full column rank, the restricted log-likelihood, often used for inference on the covariance parameters, is

$$l_R(\sigma_g^2, \sigma_e^2) = -\frac{1}{2} \{ \log |\Sigma| + \log |X^{\mathsf{T}} \Sigma^{-1} X| + (Y - X \tilde{\beta})^{\mathsf{T}} \Sigma^{-1} (Y - X \tilde{\beta}) \},$$

where  $\tilde{\beta} = \tilde{\beta}(\sigma_g^2, \sigma_e^2) = (X^{\mathrm{T}}\Sigma^{-1}X)^{-1}X^{\mathrm{T}}\Sigma^{-1}Y$  is the partial maximizer of  $l(\beta, \sigma_g^2, \sigma_e^2)$  in  $\beta$ , for fixed  $\sigma_g^2$  and  $\sigma_e^2$ . The domain of  $l_R$  is  $[0, \infty) \times (0, \infty)$ . The assumption that X have full column rank is for notational convenience; one could define  $l_R$  similarly as long as the column rank is less than n. Specifically, if the column rank of X is p' < n and  $V \in \mathbb{R}^{n \times (n-p')}$  is a semi-orthogonal matrix satisfying  $V^{\mathrm{T}}X = 0$ , then the restricted likelihood is the likelihood of  $V^{\mathrm{T}}Y$  (Harville, 1974).

By spectral decomposition,  $K = O\Lambda O^{\mathsf{T}}$  for some orthogonal  $O \in \mathbb{R}^{n \times n}$  and  $\Lambda = \operatorname{diag}(\lambda_1, \ldots, \lambda_n)$ ; here and elsewhere, eigenvalues are in decreasing order unless otherwise stated. It follows that  $\Sigma = O(\sigma_q^2 \Lambda + \sigma_e^2 I_n) O^{\mathsf{T}}$  and, hence,

$$l(\beta, \sigma_g^2, \sigma_e^2) = -\frac{1}{2} \sum_{i=1}^n \left\{ \log(\lambda_i \sigma_g^2 + \sigma_e^2) + \frac{(O^{\mathrm{T}} Y - O^{\mathrm{T}} X \beta)_i^2}{\lambda_i \sigma_g^2 + \sigma_e^2} \right\},$$

where  $(\cdot)_i$  means the *i*th element; and

$$l_R(\sigma_g^2, \sigma_e^2) = l(\tilde{\beta}, \sigma_g^2, \sigma_e^2) - \log|(O^{\mathrm{T}}X)^{\mathrm{T}}(\sigma_g^2\Lambda + \sigma_e^2I_n)^{-1}O^{\mathrm{T}}X|/2.$$

Thus, when developing theory, we may upon replacing Y and X by  $O^{\mathrm{T}}Y$  and  $O^{\mathrm{T}}X$ , respectively, assume diagonal covariance matrices  $K = \Lambda$  (equivalently,  $O = I_n$ ) and  $\Sigma = \sigma_g^2 \Lambda + \sigma_e^2 I_n$  without loss of generality. When doing so, the observations are ordered so that the *i*th response  $Y_i$  corresponds to the *i*th eigenvalue  $\lambda_i$ .

At the expense of more notation, the assumption on  $\Sigma$  could be relaxed to  $\Sigma = \sigma_G^2 K + \sigma_e^2 W$  for some known positive definite  $W \in \mathbb{R}^{n \times n}$ . Indeed, in that case Y, X, and K can be replaced by, respectively,  $W^{-1/2}Y$ ,  $W^{-1/2}X$ , and  $W^{-1/2}KW^{-1/2}$  before repeating the above argument to get a diagonal  $\Sigma$ .

In several motivating applications, the parameter of interest is  $h^2 = \sigma_g^2/(\sigma_g^2 + \sigma_e^2)$ . Therefore, we consider a common reparameterization in terms of  $h^2$  and  $\sigma^2 = \sigma_g^2 + \sigma_e^2$ . Then,  $\Sigma = \sigma^2 O\{h^2 \Lambda + (1 - h^2)I_n\}O^{\rm T}$ . Overloading the letter l and assuming  $O = I_n$  without loss of generality, the log-likelihood becomes

$$l(\beta, h^2, \sigma^2) = -\frac{1}{2} \sum_{i=1}^n \left\{ \log(\sigma^2) + \log(\lambda_i h^2 + 1 - h^2) + \frac{(Y_i - X_i^\mathsf{T} \beta)^2}{\sigma^2(\lambda_i h^2 + 1 - h^2)} \right\}, \quad (5)$$

where  $X_i^{\mathsf{T}}$  is the *i*th row of X. The domain of l is the parameter set  $\mathbb{R}^p \times [0,1) \times (0,\infty)$ . The corresponding restricted log-likelihood, with domain  $[0,1) \times (0,\infty)$ , is

$$l_R(h^2, \sigma^2) = l(\tilde{\beta}, h^2, \sigma^2) + \frac{p}{2}\log(\sigma^2) - \frac{1}{2}\log|X^{\mathrm{T}}\{h^2\Lambda + (1 - h^2)I_n\}^{-1}X|.$$

#### 2.1 Test-statistics and confidence regions

There are many potential test-statistics for inference based on  $l(\beta, h^2, \sigma^2)$  and  $l_R(h^2, \sigma^2)$ . However, most common ones, like Wald and likelihood ratio statistics, do not give confidence regions satisfying (3) in boundary settings in general. Therefore, we focus on score statistics standardized to have mean zero and diagonal covariance matrix. Such test-statistics have previously been shown to lead to (3) in settings where, unlike here, the Fisher information is singular (Ekvall and Bottai, 2022). Because our goal is to construct confidence regions for the covariance parameters, effectively treating  $\beta$  as a nuisance parameter, we primarily work with the restricted likelihood.

Let  $V_1 = V_1(h^2, \sigma^2) = \partial \Sigma / \partial h^2 = \sigma^2 O(\Lambda - I_n) O^{\mathrm{T}}$  and  $V_2 = V_2(h^2, \sigma^2) = \partial \Sigma / \partial \sigma^2 = O(h^2 \Lambda + (1 - h^2) I_n) O^{\mathrm{T}}$ . Similarly, let  $U_1(h^2, \sigma^2) = \partial I_R(h^2, \sigma^2) / \partial h^2$  and  $U_2(h^2, \sigma_e^2) = \partial I_R(h^2, \sigma^2) / \partial \sigma^2$ , so the restricted score is  $U(h^2, \sigma^2) = [U_1(h^2, \sigma^2), U_2(h^2, \sigma^2)]^{\mathrm{T}}$ . Routine calculations show, for  $j \in \{1, 2\}$ ,

$$U_{j}(h^{2}, \sigma^{2}) = \frac{1}{2} (Y - X\tilde{\beta})^{\mathrm{T}} \Sigma^{-1} V_{j} \Sigma^{-1} (Y - X\tilde{\beta}) - \frac{1}{2} \operatorname{tr}(\Sigma^{-1} V_{j}) + \frac{1}{2} \operatorname{tr}(\Sigma^{-1} X (X^{\mathrm{T}} \Sigma^{-1} X)^{-1} X^{\mathrm{T}} \Sigma^{-1} V_{j}).$$

Observe  $U_j$  is defined on  $[0,1) \times (0,\infty)$  and has a continuous extension to points where  $h^2 = 1, \sigma^2 > 0$  if K is positive definite. The expression simplifies substantially for j = 2 since  $V_2 = \Sigma/\sigma^2$ . Specifically, assuming  $O = I_n$ ,

$$U_2(h^2, \sigma^2) = -\frac{n-p}{2\sigma^2} + \frac{1}{2\sigma^4} \sum_{i=1}^n \frac{(Y_i - X_i^{\mathrm{T}} \tilde{\beta})^2}{h^2 \lambda_i + 1 - h^2}.$$
 (6)

Let  $\mathcal{I}(h^2, \sigma^2)$  denote the restricted information matrix, that is, the covariance matrix of  $U(h^2, \sigma^2)$  when  $h^2$  and  $\sigma^2$  are the true parameters. We next give an expression for  $\mathcal{I}(h^2, \sigma^2)$  for easy reference. The proof, along with those of other formally stated results, is in Appendix A.

Let 
$$P = \Sigma^{-1/2} X (X^{\mathsf{T}} \Sigma^{-1} X)^{-1} X^{\mathsf{T}} \Sigma^{-1/2}$$
,  $Q = I_n - P$ ,  $D = \{h^2 \Lambda + (1 - h^2) I_n\}^{-1} (\Lambda - I_n)$ , and  $H = ODO^{\mathsf{T}}$ .

**Proposition 1.** If (1) holds for some X with full column rank, then the restricted information matrix has elements

$$\mathcal{I}_{11}(h^2, \sigma^2) = \frac{1}{2}\operatorname{tr}(QHQH); \quad \mathcal{I}_{12}(h^2, \sigma^2) = \frac{1}{2\sigma^2}\operatorname{tr}(QH); \quad \mathcal{I}_{22}(h^2, \sigma^2) = \frac{n-p}{2\sigma^4}.$$

For inference on  $(h^2, \sigma^2)$  jointly, we consider the (quadratic) restricted score standardized by the restricted Fisher information:

$$T_n^R(h^2, \sigma^2) = U(h^2, \sigma^2)^{\mathrm{T}} \mathcal{I}^{-1}(h^2, \sigma^2) U(h^2, \sigma^2), \tag{7}$$

For inference on  $h^2$  only, we use

$$T_n^R(h^2) = U_1(h^2, \tilde{\sigma}^2)^2 \mathcal{I}^{11}(h^2, \tilde{\sigma}^2),$$
 (8)

where  $\mathcal{I}^{11}(h^2, \sigma^2)$  is the leading element of  $\mathcal{I}^{-1}(h^2, \sigma^2)$  and  $\tilde{\sigma}^2 = \tilde{\sigma}^2(h^2)$  is the partial maximizer of  $l_R(h^2, \sigma^2)$  in  $\sigma^2$ , for a fixed  $h^2$ . By (6), assuming  $O = I_n$ ,

$$\tilde{\sigma}^2 = \frac{1}{n-p} \sum_{i=1}^n \frac{(Y_i - X_i^{\mathrm{T}} \tilde{\beta})^2}{h^2 \lambda_i + 1 - h^2} = \frac{1}{n-p} (Y - X \tilde{\beta})^{\mathrm{T}} \{ h^2 \Lambda + (1 - h^2) I_n \}^{-1} (Y - X \tilde{\beta}).$$
(9)

When X has column rank less than n,  $\tilde{\sigma}^2$  is greater than zero with probability one, and hence a valid partial maximizer. Note  $\tilde{\beta}$  does not depend on  $\sigma^2$ , so the last display indeed gives an explicit expression for  $\tilde{\sigma}^2$ . For inference on  $h^2$ , we sometimes also consider  $S_n^R(h) = U_1(h^2, \tilde{\sigma}^2)\mathcal{I}^{11}(h^2, \tilde{\sigma}^2)^{1/2}$ , that is, the signed square root of (8).

The proposed test-statistics require a positive definite  $\mathcal{I}(h^2, \sigma^2)$ , which the following result establishes. Notably,  $\mathcal{I}^{11}(h^2, \sigma^2)$  is positive if  $\mathcal{I}(h^2, \sigma^2)$  is positive definite. To understand the next result, notice that Q has rank n-p, so QHQ has at most n-p non-zero eigenvalues, and recall  $V \in \mathbb{R}^{n \times (n-p)}$  is a semi-orthogonal matrix such that  $V^{\mathrm{T}}X = 0$ .

**Theorem 1.** The information matrix  $\mathcal{I}(h^2, \sigma^2)$  is singular if and only if the following equivalent conditions hold: (i) the n-p possibly non-zero eigenvalues of QHQ are identical; (ii)  $V^{\mathrm{T}}KV = cI_{n-p}$  for some  $c \geq 0$ .

The proof of Theorem 1 is relatively straightforward using the expressions in Proposition 1. The key observation is that the determinant of  $\mathcal{I}(h^2, \sigma^2)$  is

$$\frac{1}{2\sigma^4} \left\{ (n-p) \operatorname{tr}(QHQQHQ) - \operatorname{tr}(QHQ)^2 \right\},\,$$

which is strictly positive by Jensen's inequality applied to the n-p possibly non-zero eigenvalues of QHQ unless they are all identical, in which case the determinant is zero.

Theorem 1 is particularly intuitive if p = 0, so that  $Q = V = I_n$ . Then the eigenvalues of QHQ are the diagonal elements of D, which are identical if and only if the diagonal elements of  $\Lambda$  are; that is, if and only if  $\Lambda$ , and hence K, is proportional to the identity. The parameters are unidentifiable for such K, so we will generally assume K is not proportional to the identity, leading to a positive definite information matrix.

By inverting the proposed test-statistics using quantiles from chi-squared distributions, we get the confidence regions

$$\mathcal{R}_n(\alpha) = \{ h^2 \in [0, 1] : T_n^R(h^2) \le q_{1-\alpha, 1} \};$$

$$\mathcal{R}_n^2(\alpha) = \{ (h^2, \sigma^2) \in [0, 1] \times (0, \infty) : T_n^R(h^2, \sigma^2) \le q_{1-\alpha, 2} \}.$$
(10)

When  $h^2 = 1$  is not in the parameter set, we take  $T_n^R(1) = T_n^R(1, \sigma^2) = \infty$  so the confidence regions are well defined.

## 3 Asymptotic distributions

We assume  $K_n = \Lambda_n = \operatorname{diag}(\lambda_{n1}, \dots, \lambda_{nn})$  in this section without loss of generality, and hence  $\Sigma_n = \sigma_n^2(I_n + \Lambda_n)$  and  $H_n = D_n$  are diagonal. Here, a subscript n has been added to  $\sigma^2$  and other quantities since we will consider a sequence  $\{(\beta_n, h_n^2, \sigma_n^2)\}$  of parameters indexed by the sample size, and a corresponding sequence of models. Similarly, the number of predictors  $p = p_n$  can change with n. By considering a sequence of (true) parameters, we can obtain the asymptotic distribution of  $T_n^R(h_n^2, \sigma_n^2)$ 

and  $T_n^R(h_n^2)$  under that sequence. This is useful because (3) holds for  $\mathcal{R}_n(\alpha)$  in (10) if and only if the asymptotic distribution of  $T_n^R(h_n^2)$  is the same under any convergent sequence of parameters (Ekvall and Bottai, 2022, Lemma 2.5), and similarly for  $\mathcal{R}_n^2(\alpha)$ . This result essentially follows from the equivalence of continuous convergence and uniform convergence on compact sets, applied to the sequence of functions defined by  $\theta \mapsto P_{\theta}\{T_n^R(h^2) \leq q_{1-\alpha,1}\}$ . More generally, examining the distribution under a sequence of parameters tending to a point, can lead to a better understanding of the behavior of the test-statistic near that point. Thus, for our purposes, sequences tending to boundary points are of particular interest. The next result is our first along these lines.

To state the result, let  $\gamma_i(\cdot)$  denote the *i*th eigenvalue in decreasing order.

**Lemma 1.** Assume that, for every  $n \in \{1, 2, ...\}$ , Y satisfies (1) with an X with full column rank,  $K_n = \Lambda_n$ , and parameters  $(\beta_n, h_n^2, \sigma_n^2)$ . If, as  $n \to \infty$ , (i)  $p/n \to 0$ , (ii)  $\sum_{i=1}^p \gamma_i(D_n^2)/\operatorname{tr}(D_n^2) \to 0$ , (iii)  $\limsup_{n\to\infty} \{n^{-1}\operatorname{tr}(D_n)^2/\operatorname{tr}(D_n^2)\} < 1$ ; then, in distribution,

$$T_n^R(h_n^2, \sigma_n^2) \to \chi_2^2. \tag{11}$$

The proof of Lemma 1 establishes (11) by showing asymptotic bivariate normality of  $\mathcal{I}_n(h_n^2, \sigma_n^2)^{-1/2}U_n(h_n^2, \sigma_n^2)$ . The key observation is that the elements of  $U_n(h_n^2, \sigma_n^2)$  are quadratic forms in multivariate normal vectors, centered to have mean zero. Therefore, their distributions are those of linear combinations of independent chisquare random variables. The conditions of the theorem ensure Lyapunov's central limit theorem applies to those linear combinations.

Conditions (ii) and (iii) concern  $\Lambda_n$  and  $h_n^2$ , but neither of  $\beta_n$  and  $\sigma_n^2$  appears in the conditions. Thus, (11) can hold for sequences one may not expect, such as those where  $\sigma_n^2 \to 0$ . This suggests the behavior of our test-statistic is insensitive to the true  $\sigma^2$ , and may perform well even if  $\sigma^2$  is near the boundary, that is, near zero.

Condition (ii) says the sum of the p greatest eigenvalues of  $D_n^2$  should be negligible compared to the sum of all its eigenvalues. An eigenvalue, or diagonal element, of  $D_n^2$  will be relatively large if either (a) the corresponding element of  $\Lambda_n$  is relatively close to zero and  $h^2$  is close to one, or (b) the corresponding element of  $\Lambda_n$  is relatively

large. Thus, condition (ii) suggests the test-statistic may not behave well if a few elements of  $O^{T}Y$  have much larger variance than all other ones, or if some elements have very small variance and most if it is attributable to K.

Condition (iii) essentially says that, when comparing the squared average of the eigenvalues of  $D_n$  to the average of the squared eigenvalues, Jensen's inequality is asymptotically strict; there has to be non-negligible variation in the diagonal elements of  $D_n$ . This is intuitive as, if all the diagonal elements of  $D_n$  are the same, then  $\Lambda_n$  is proportional to the identity and  $h_n^2$  unidentifiable, as noted in the remark following Theorem 1.

Conditions (ii) and (iii) could be replaced by slightly weaker ones by adding assumptions on how X relates to K, but that would substantially increase notational burden.

Our next result shows that the proposed test-statistic for  $h^2$  has the desired asymptotic distribution under the conditions of Lemma 1. Substantial work is needed to establish the result since  $\sigma^2$  is a nuisance parameter for the purposes of inference on  $h^2$  only. In particular, the test-statistic for  $h^2$  cannot depend on or be evaluated at the true parameter, or null hypothesis, which was possible in Lemma 1.

**Lemma 2.** Under the conditions of Lemma 1,  $T_n^R(h_n^2) \to \chi_1^2$  in distribution as  $n \to \infty$ .

Lemma 2 is proven by showing that  $T_n(h_n^2)$  has the same asymptotic distribution as  $\{U_{n1} - \mathcal{I}_{n12}U_{n2}/\mathcal{I}_{n22}\}^2\mathcal{I}_n^{(11)}$ , with all quantities evaluated at the true  $(h_n^2, \sigma_n^2)$  and  $\mathcal{I}_n^{(11)}$  denoting the leading element of  $\mathcal{I}_n^{-1}$ . This result is consistent with what is often obtained for nuisance parameters in classical settings, with fixed interior parameters (Cox and Hinkley, 2000, Section 9.3). This is remarkable because those results in general do not apply under sequences of parameters or with boundary points.

Our next result gives intuitive conditions on model parameters and K which ensure the conditions in Lemmas 1 and 2 hold. The conditions are not necessary in general, but they are more transparent, and easier to assess in practice, than those in the lemmas.

**Theorem 2.** Assume that, for every  $n \in \{1, 2, ...\}$ , Y satisfies (1) with an X with full column rank and parameters  $(\beta_n, h_n^2, \sigma_n^2)$ . If (i)  $\lim_{n\to\infty} p_n/n = 0$ ; (ii\*)  $\limsup_{n\to\infty} \lambda_{n1} < \infty$ ; (iii\*)  $\liminf_{n\to\infty} \lambda_{nn} > 0$  or  $\limsup_{n\to\infty} h_n^2 < 1$ ; (iv\*) there exists an  $\epsilon > 0$  such that for any  $c \in [0, \infty)$ , with  $k_n^c$  denoting the number of eigenvalues of  $\Lambda_n$  in  $[c - \epsilon, c + \epsilon]$ , it holds that  $\limsup_{n\to\infty} k_n^c/n < 1$ , then, in distribution as  $n \to \infty$ ,

$$T_n^R(h_n^2, \sigma_n^2) \to \chi_2^2; \quad T_n^R(h_n^2) \to \chi_1^2.$$
 (12)

Since  $\Lambda_n$  is known, condition (ii\*) can be made to hold, for example by replacing K by  $K/\lambda_{n1}$ . Then, for the conclusions to hold, conditions (iii\*) and (iv\*) would need to hold for the new eigenvalues  $\lambda_{ni}/\lambda_{n1}$ ,  $i \in \{1, ..., n\}$ . Condition (iii\*) illustrates that if K is full rank, then reliable inference on  $h^2$  near one is possible; this is formalized further in Corollary 1. Condition (iv\*) prevents the eigenvalues of K from concentrating in some neighborhood asymptotically. That is, it prevents K from being essentially proportional to the identity asymptotically.

Corollary 1. Under condition (i), (ii\*), and (iv\*) of Theorem 2,  $\mathcal{R}_n(\alpha)$  and  $\mathcal{R}_n^2(\alpha)$  in (10) have asymptotically correct uniform coverage probability on compact subsets of [0,1) and  $[0,1)\times(0,\infty)$ , respectively (c.f. (3)). The conclusion can be strengthened to compact subsets of  $[0,1]\times(0,\infty)$  and [0,1], respectively, if the first part of condition (iii\*) of Theorem 2 also holds.

## 4 Computational considerations

In this section, we consider the computational aspects of our test-statistics. Like popular competing methods, we require a one-time preprocessing step where the eigendecomposition of  $K = O\Lambda O^{\mathrm{T}}$  is used to construct  $Y_O = O^{\mathrm{T}}Y$  and  $X_O = O^{\mathrm{T}}X$ . In practice it is common to use the same K for many different choices of response vector Y, with separate inference on  $h^2$  for each response vector. Then, the pre-processing step is only performed once.

We show that, after the pre-processing, the number of floating point operations required to evaluate the proposed test-statistics grows linearly in n, making our

methods feasible for large-scale applications, such as those described in Section 1. More specifically, we show that, if p is potentially growing with n, the number of operations required is  $O(np^2 + p^3)$ .

Recall  $V_2 = h^2 \Lambda + (1 - h^2) I_n$  and define  $\tilde{P} = V_2^{-1/2} X_O (X_O^{\mathsf{T}} V_2^{-1} X_O)^{-1} X_O^{\mathsf{T}} V_2^{-1/2}$ ,  $\tilde{Q} = I_n - \tilde{P}$ ,  $\tilde{\beta} = (X_O^{\mathsf{T}} V_2^{-1} X)^{-1} X_O^{\mathsf{T}} V_2^{-1} Y_O$ , and  $\tilde{R} = Y_O - X_O \tilde{\beta}$ . Let  $\tilde{R}_i$  denote the *i*th element of  $\tilde{R}$  and  $Q_{ii}$  the *i*th diagonal element of Q. With these definitions the score function is

$$U(h^{2}, \sigma^{2}) = \frac{1}{2} \begin{pmatrix} \sum_{i=1}^{n} D_{ii} \left( \frac{\sigma^{-2} \tilde{R}_{i}^{2}}{h^{2} \lambda_{i} + 1 - h^{2}} - \tilde{Q}_{ii} \right) \\ -\frac{n-p}{\sigma^{2}} + \frac{1}{\sigma^{4}} \sum_{i=1}^{n} \frac{\tilde{R}_{i}^{2}}{h^{2} \lambda_{i} + 1 - h^{2}} \end{pmatrix}.$$
(13)

Both terms in (13) involve the summation of n terms. Construction of  $\tilde{R}$  requires O(np) operations; and computation of all  $\tilde{Q}_{ii}$  requires only  $O(np^2)$  operations. We do not compute or store the entire matrix  $\tilde{Q}$ .

To compute the information efficiently, note  $Q = O\tilde{Q}O^{\top}$ , so by Proposition 1,

$$\mathcal{I}(h^2, \sigma^2) = \frac{1}{2\sigma^4} \begin{pmatrix} \sigma^4 \mathrm{tr}(\tilde{Q}D\tilde{Q}D) & \sigma^2 \mathrm{tr}(\tilde{Q}D) \\ \sigma^2 \mathrm{tr}(\tilde{Q}D) & n-p \end{pmatrix}.$$

Because  $\operatorname{tr}(\tilde{Q}D) = \sum_{i=1}^n \tilde{Q}_{ii}D_{ii}$ ,  $\mathcal{I}_{12}$  can be evaluated in O(n) operations using the  $\tilde{Q}_{ii}$  computed before. Notice  $2\mathcal{I}_{11} = \operatorname{tr}(\tilde{Q}D\tilde{Q}D) = \operatorname{tr}(D^2) - 2\operatorname{tr}(\tilde{P}D^2) + \operatorname{tr}(\tilde{P}D\tilde{P}D)$ . Because D is a diagonal matrix, evaluating  $\operatorname{tr}(D^2)$  requires O(n) operations. Similarly,  $\operatorname{tr}(\tilde{P}D^2) = \sum_{i=1}^n (1-\tilde{Q}_{ii})D_{ii}^2$ , so evaluating this term also requires only O(n) operations. Finally, by the cyclic property of the trace operator,  $\operatorname{tr}(\tilde{P}D\tilde{P}D) = \operatorname{tr}(\tilde{A}\tilde{B}\tilde{A}\tilde{B})$  where  $\tilde{A} = (X_O^{\mathrm{T}}V_2^{-1}X_O)^{-1} \in \mathbb{R}^{p\times p}$  and  $\tilde{B} = X_O^{\mathrm{T}}V_2^{-1/2}DV_2^{-1/2}X_O \in \mathbb{R}^{p\times p}$ . Thus, evaluating  $\operatorname{tr}(\tilde{A}\tilde{B}\tilde{A}\tilde{B})$  requires  $O(np^2 + p^3)$  operations, with the  $p^3$  coming from a Cholesky (or other) decomposition of  $\tilde{A}^{-1}$  to deal with the inverse, and the  $np^2$  is from matrix multiplications.

We have created an R package lmmvar for computing our test-statistics and confidence intervals, available for download at github.com/yqzhang5972/lmmvar.

Our confidence interval for  $h^2$  can be obtained by evaluating  $T_n^R(h^2)$  from at

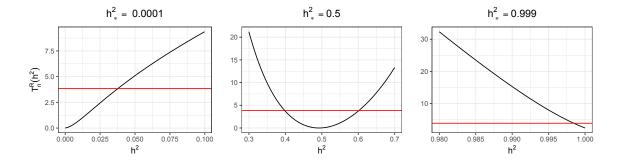


Figure 2: The value  $T_n^R(h^2)$  with n=1000, data generated under the same model as in Fig. 1, and the true  $h^2$ , here denoted  $h_*^2$ , taking values  $\{0.0001, 0.5, 0.999\}$ . Horizontal red lines denote the 95th percentile of the  $\chi_1^2$  distribution.

equally spaced points in [0,1) and comparing the values to quantiles from the chisquare distribution with one degree of freedom. In practice, however, it is typically possible, and faster, to do a search for the points where  $T_n^R(h^2)$  crosses the desired quantile. More specifically, when  $T_n^R(h^2)$  is strictly quasiconvex in  $h^2$ , then  $\mathcal{R}_n(\alpha)$  is an interval, so we need only search for points a such that  $T_n^R(a) = q_{1,1-\alpha}$ . While we have not been able to prove  $T_n^R(h^2)$  is quasiconvex in general due to technical difficulties, we have not encountered an example where it is not. Typically,  $T_n^R(h^2)$  looks roughly like one of the three cases in Fig. 2. Thus, in our software we first find a point a' such that  $T_n^R(a') < q_{1,1-\alpha}$ . More specifically, we start a ternary search for the minimum of  $T_n^R$  and terminate the first time a negative value is encountered. Then, under quasiconvexity, the intervals  $[0, T_n^R(a')]$  and  $[T_n^R(a'), 1)$  contain lower and upper bounds of the interval  $\mathcal{R}_n(\alpha)$ , respectively. We then use a bisection search to find  $b \in [0, T_n^R(a')]$  such that  $T_n^R(b) = q_{1,1-\alpha}$ , and similarly for  $[T_n^R(a'), 1)$ . This procedure often requires less computing time than an exhaustive grid search.

We have also found that, in practice,  $S_n^R(h^2) = \{\mathcal{I}_n^{11}(h^2, \tilde{\sigma}^2)\}^{1/2}U_1(h^2, \tilde{\sigma}^2)$  is typically monotonely decreasing in  $h^2$ , so a search can also be based on that statistic and comparisons to standard normal quantiles; see Section 5.3 for an example with one-sided confidence intervals.

#### 5 Numerical studies

#### 5.1 Finite-sample interval width and coverage probabilities

To supplement our asymptotic theory, we study how  $n, h^2$ , and the eigenvalues of K affect the coverage probability and width of our confidence intervals. We consider  $K_{ij} = \rho^{|i-j|}$  for  $\rho \in (0,1)$  and  $i,j \in \{1,\ldots,n\}$ . When  $\rho$  is close to one, K is approximately  $1_n 1_n^{\mathrm{T}}$ , where  $1_n$  is an n-vector of ones. Thus, approximately,  $\lambda_{n1} = n$  and  $\lambda_{ni} = 0, i \geq 2$ , which is inconsistent with conditions (ii\*) and (iv\*) of Theorem 2. Conversely, when  $\rho$  is close to zero, K is approximately an identity matrix, which is inconsistent with condition (iv\*) of Theorem 2. Thus, we expect our test-statistic to perform best when  $\rho$  is not too close to either zero or one.

In each setting considered, we generate  $X \in \mathbb{R}^{n \times 5}$  with independent and identically distributed standard normal entries, and then generate  $Y \sim \mathcal{N}_n(X\beta, h^2K + (1-h^2)I_n)$  (i.e.,  $\sigma^2 = 1$ ) with  $\beta = 0$ . For every combination of the triplet  $(n, h^2, \rho)$ , we generate  $10^4$  independent realizations of Y and compute the corresponding 95% confidence intervals for  $h^2$ .

The bottom panel of Fig. 3 displays coverage probabilities with  $\rho$  and n varying for four different values of  $h^2$ . For  $\rho \in \{0.1.0.5.0.95\}$ , our interval appears to have the coverage probability 0.95 or higher. Only when  $\rho = 0.999$  is the evidence of under-coverage for some combinations of n and  $h^2$ .

The top panel of Fig. 3 displays interval widths for different values of  $\rho$  and  $h^2$ . When  $\rho = 0.1$ ,  $K \approx I_n$ , which means  $h^2$  is approximately unidentifable. Consequently, interval widths are too large to be practically useful, despite having the nominal coverage level. With  $\rho = 0.5$  or 0.95, the widths are reasonably small and decrease as n increases. When  $\rho = 0.999$ , the eigenvalues of K are approximately  $(n, 0, \ldots, 0)$ , which appear to lead to longer confidence intervals. Nevertheless, as n increases, interval widths decrease as expected. As n approaches 2000, the widths are small enough to be useful in practice.

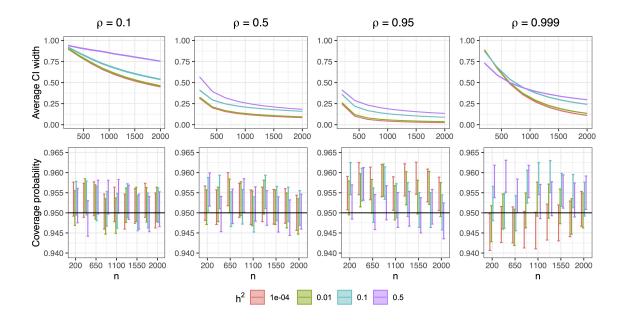


Figure 3: (Top panel) The average width of our confidence interval for  $h^2$  and (bottom panel) the average coverage probability as n varies with  $\rho \in \{0.1, 0.5, 0.95, 0.999\}$ . In the top panels, the widths of the lines provide 95% confidence bands; and in the bottom panel, the bars contain the 95% confidence intervals, each based on  $10^4$  trials.

#### 5.2 Comparison to simulation-based test-statistics

Existing methods for constructing confidence intervals for  $h^2$  often rely on simulation. For example, Schweiger et al. (2016) uses the parametric bootstrap to approximate the distribution of the REML estimator of  $h^2$ . In comparison to a naive implementation of the parametric bootstrap, their implementation is very efficient. Specifically, to estimate the distribution of the REML estimator when the true heritability is  $h^2$ , their method requires (i) drawing N independent samples from (1) with  $\sigma^2 = 1$  and null hypothesis value  $h^2$  given; and (ii) for each sample, finding an interval containing a local maximizer of the restricted log-likelihood. Since each sample is needed for step (ii), Schweiger et al. (2016) show that it is sufficient to draw samples from  $\mathcal{N}_n(0, I_n)$  in step (i) and modify step (ii) accordingly. The modified step (ii) is performed by evaluating the derivative of the restricted log-likelihood at fixed grid points. However,

for a given  $h_2$  the derivative must, in general, be evaluated at many grid points; and this procedure must be repeated for all N samples. Constructing confidence intervals thus requires repeating this procedure many times, which can be computationally burdensome.

In follow-up work, Schweiger et al. (2018) proposed a stochastic approximation to the confidence interval from Schweiger et al. (2016). This approximation estimates the upper and lower bounds of the confidence interval using the Robbins-Monro algorithm. Like our test-statistic, after the same preprocessing step, the operations required for constructing the intervals from Schweiger et al. (2016) and Schweiger et al. (2018) scale linearly with n.

In Fig. 4, we compare the time needed to compute our interval to that of Schweiger et al. (ALBI, 2016) and Schweiger et al. (FIESTA, 2018). The figure indicates our confidence intervals can be computed most efficiently of the three methods. For example, then n=100 and  $h^2=0.01$  our method took on average  $3.7\times10^{-4}$  seconds compared to 3.11 seconds and 45.38 seconds for FIESTA and ALBI, respectively. When n=3000 and  $h^2=0.01$ , our method took 0.07 seconds on average, whereas FIESTA and ALBI took 511.18 seconds and 1908.68 seconds, respectively. ALBI and FIESTA are written in Python, while our software is written in a combination of R and C++. All timings were obtained simultaneously on the University of Florida's high-performance computing cluster, HiPerGator.

We also compared the width of our confidence intervals to those of Schweiger et al. (2018). We found that under a broad range of settings, the distribution of our interval's width was similar to that of Schweiger et al. (2018). In our simulations with n=1000, we recorded the standardized difference in widths (defined as Schweiger et al. (2018)'s width minus our interval's width, divided by our interval's width). On average the standardized differences were 0.24, -0.03, and 0.005 when  $h^2$  was 0.01, 0.1, and 0.5, respectively. Median standardized differences were -0.07, -0.03, and 0.005.

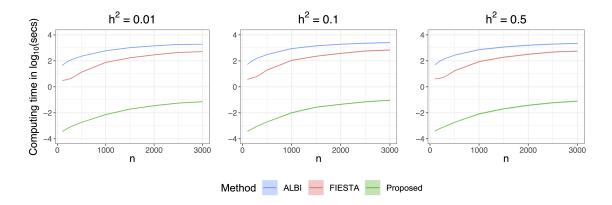


Figure 4: The average computing times for constructing one confidence interval for  $h^2$  using our test-statistic  $T_n^R$  ("Proposed"), the simulation-based interval construction method from Schweiger et al. (2016) ("ABLI"), and the stochastic approximation of the interval from Schweiger et al. (2016) proposed by Schweiger et al. (2018) ("FIESTA"). The shaded regions are 95% confidence bands. All times exclude the required eigendecomposition of K. Data were generated under the same model as in Fig. 1.

#### 5.3 Gene ranking in spatially-resolved transcriptomics

In Section 1, we described one application of our method in the analysis of spatially-resolved transcriptomic data. The rightmost panels of Fig. 5 display data generated in a spatially-resolved RNA sequencing experiment performed on a mouse brain tissue sample slice. In such an experiment, one measures the expression of thousands of genes (the number of RNA reads aligning with each gene) from cells in 50 micron diameter spatial locations (known as spots) across the tissue sample slice. These locations are represented by the colored circles in the rightmost panels of Fig. 5.

In this context, for one gene,  $h^2$  is the fraction of variability in expression attributed to the spatial orientation of the tissue sample. In a typical analysis, one tests  $h^2$  for more than 10,000 genes. Specially, after normalization, it is common to assume that for the  $\ell$ th gene,

$$Y_{\ell} \sim \mathcal{N}_n(X\beta, \sigma^2\{h_{\ell}^2K + (1 - h_{\ell}^2)I_n\}), \quad \ell \in \{1, \dots, d\},$$
 (14)

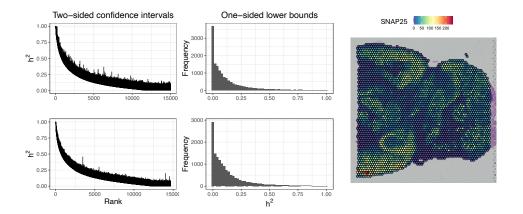


Figure 5: (Left panels) Two-sided 95% confidence intervals for  $h^2$  in anterior (top row) and posterior (bottom row) regions of a mouse's brain tissue, sorted according to their lower bound. (Middle panels) Histograms of the lower bounds of 95% one-sided confidence intervals for  $h^2$  in both anterior and poster regions. (Right panel) Expression counts of SNAP25, one of the most spatially variable genes, across the posterior region. We display counts, but our analysis is performed on log-normalized expression.

where  $h_{\ell}^2$  is the fraction of spatial variability in the  $\ell$ th gene, n is the number of spots in the tissue, d is the number of genes in the study, and  $Y_{\ell}$  and  $Y_{\ell'}$  are independent for all  $\ell \neq \ell'$ . Typically  $K_{ij} = \exp(-\|s_i - s_j\|/a)$  for some a > 0, where  $s_i$  is the two dimensional spatial coordinates of the ith spot and  $\|\cdot\|$  is the Euclidean norm.

To demonstrate the usefulness of our method in this application, we analyzed two slices of a mouse's brain tissue (anterior and posterior regions), which are available as the *stxbrain* data in the **SeuratData** R package. These data were generated on the 10X Visium platform. Assuming (14) with a = 0.02 (chosen based on results from Weber et al. (2022)), after quality control of filtering out low-expressed genes and boundary spots in the tissues, n = 2380 for d = 15117 genes in the anterior slice and n = 3011 for d = 14738 genes in the posterior slice.

Using our test statistics, we can provide a confidence interval for each  $h_{\ell}^2$ . Moreover, in order to rank genes, we can also provide one-sided confidence intervals for  $h_{\ell}^2$  by comparing  $U_1(h^2, \tilde{\sigma}^2) \{ \mathcal{I}^{11}(h^2, \tilde{\sigma}^2) \}^{1/2}$  to quantiles of the standard normal distribution.

Because each of the genes has the same K matrix, we can perform a transcriptomewide analysis very efficiently. After a one-time eigendecomposition of K, the remaining processes took around 3 minutes and 4 minutes for the two tissues, respectively. For comparison, ALBI took approximately 9 minutes and 13 minutes for the two tissues, including the time needed to find the necessary maximum likelihood estimates of each  $h_{\ell}^2$ , which was around 3 minutes. FIESTA, which is generally slower than ALBI when computing many confidence intervals, timed out after 96 hours.

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## A Appendix

Proof of Proposition 1. We first derive a convenient expression for the stochastic part of the restricted score function. Recall  $P = \Sigma^{-1/2} X (X^T \Sigma^{-1} X)^{-1} X^T \Sigma^{-1/2}$  and write  $Y - X \tilde{\beta} = Y - X (X^T \Sigma^{-1} X)^{-1} X^T \Sigma^{-1} Y = (I_n - X (X^T \Sigma^{-1} X)^{-1} X^T \Sigma^{-1}) Y = (I_n - \Sigma^{1/2} P \Sigma^{-1/2}) Y$ . Thus,

$$\begin{split} (Y - X\tilde{\beta})^T \Sigma^{-1} V_j \Sigma^{-1} (Y - X\tilde{\beta}) &= Y^T (I_n - \Sigma^{-1/2} P \Sigma^{1/2}) \Sigma^{-1} V_j \Sigma^{-1} (I_n - \Sigma^{1/2} P \Sigma^{-1/2}) Y \\ &= Y^\mathsf{T} \Sigma^{-1/2} (I_n - P) \Sigma^{-1/2} V_j \Sigma^{-1/2} (I_n - P) \Sigma^{-1/2} Y \\ &= Y^\mathsf{T} \Sigma^{-1/2} Q \Sigma^{-1/2} V_j \Sigma^{-1/2} Q \Sigma^{-1/2} Y. \end{split}$$

Since  $Q\Sigma^{-1/2}X = \Sigma^{-1/2}X - \Sigma^{-1/2}X(X^T\Sigma^{-1}X)^{-1}X^T\Sigma^{-1}X = \Sigma^{-1/2}X - \Sigma^{-1/2}X = 0$ , the preceding display has the same distribution as

$$\xi_n^{\mathsf{T}} Q \Sigma^{-1/2} V_j \Sigma^{-1/2} Q \xi_n,$$

where  $\xi_n \sim N(0, I_n)$  and the distribution holds jointly for  $j \in \{1, 2\}$ . For j = 1 we get

$$\begin{split} Q\Sigma^{-1/2}V_1\Sigma^{-1/2}Q &= Q\Sigma^{-1/2}O\sigma_n^2(\Lambda-I_n)O^T\Sigma^{-1/2}Q \\ &= QO\{h^2\Lambda + (1-h^2)I_n\}^{-1/2}(\Lambda-I_n)\{h^2\Lambda + (1-h^2)I_n\}^{-1/2}OQ \\ &= QHQ, \end{split}$$

and, similarly,  $Q\Sigma^{-1/2}V_2\Sigma^{-1/2}Q = Q\Sigma^{-1/2}(\Sigma/\sigma^2)\Sigma^{-1/2}Q = Q/\sigma^2$ . It follows that

$$\mathcal{I}_{11}(h^2, \sigma^2) = \frac{1}{2} \operatorname{tr}(QHQHQ) = \frac{1}{2} \operatorname{tr}(QHQH)$$

$$\mathcal{I}_{12}(h^2, \sigma^2) = \frac{1}{2} \operatorname{tr}(QHQQ/\sigma^2) = \frac{1}{2\sigma^2} \operatorname{tr}(QH)$$

$$\mathcal{I}_{22}(h^2, \sigma^2) = \frac{1}{2\sigma^4} \operatorname{tr}(Q) = \frac{n-p}{2\sigma^4}.$$

Proof of Theorem 1. We start with some useful observations. First,  $\mathcal{I}_{11}(h^2, \sigma^2) = \text{tr}(QHQQHQ)/2$  and  $\mathcal{I}_{12}(h^2, \sigma^2) = \text{tr}(QHQ)/(2\sigma^2)$  since Q is idempotent. Secondly, the eigenvalues of QHQ are real since QHQ is symmetric, and hence the eigenvalues of QHQQHQ are positive as the squares of real numbers.

Now, suppose QHQ has n-p eigenvalues equal to some  $c \in \mathbb{R}$ , with the remaining p eigenvalues equal to zero. Then the determinant

$$|\mathcal{I}(h^2, \sigma^2)| = \frac{1}{4\sigma^4} \{ (n-p)\operatorname{tr}(QHQQHQ) - \operatorname{tr}(QHQ)^2 \}$$

is zero since  $\operatorname{tr}(QHQQHQ) = (n-p)c^2$  and  $\operatorname{tr}(QHQ) = (n-p)c$ . Thus,  $\mathcal{I}(h^2, \sigma^2)$  is singular.

To establish the other direction, suppose instead that not all of the n-p possibly non-zero eigenvalues of QHQ are identical. Then Jensen's inequality gives

$$\frac{1}{n-p} \sum_{i=1}^{n} \gamma_i (QHQ)^2 > \left\{ \frac{1}{n-p} \sum_{i=1}^{n} \gamma_i (QHQ) \right\}^2,$$

where  $\gamma_i(\cdot)$  is the *i*th largest eigenvalue. Thus,  $|\mathcal{I}(h^2, \sigma^2)| > 0$ . Moreover, at least one eigenvalue of QHQQHQ is strictly positive, so  $\mathcal{I}_{11}(h^2, \sigma^2) > 0$ . Thus,  $\mathcal{I}(h^2, \sigma^2)$  is positive definite by Sylvester's criterion.

**Lemma 3.** If (1) holds for some X with full column rank, then

$$(n-p)\tilde{\sigma}^2(h^2)/\sigma_n^2 \sim \chi_{n-p}^2$$
.

*Proof.* Assuming  $O = I_n$ , by (9) and proof of Proposition 1,

$$\begin{split} (n-p)\tilde{\sigma}^2(h^2)/\sigma_n^2 &= (Y-X\tilde{\beta})^\mathsf{T} \Sigma^{-1} (Y-X\tilde{\beta}) \\ &= Y^\mathsf{T} (I_n - \Sigma^{-1/2} P \Sigma^{1/2}) \Sigma^{-1} (I_n - \Sigma^{1/2} P \Sigma^{-1/2}) Y \\ &= Y^\mathsf{T} \Sigma^{-1/2} (I_n - P) \Sigma^{1/2} \Sigma^{-1} \Sigma^{1/2} (I_n - P) \Sigma^{-1/2} Y \\ &= Y^\mathsf{T} \Sigma^{-1/2} (I_n - P) (I_n - P) \Sigma^{-1/2} Y \\ &= Y^\mathsf{T} \Sigma^{-1/2} Q \Sigma^{-1/2} Y. \end{split}$$

Since  $Q\Sigma^{-1/2}X = 0$  given by proof of Proposition 1,  $Q\Sigma^{-1/2}Y \sim \mathcal{N}(0, Q)$ , then the above quadratic form has the same distribution as  $\xi_n^{\mathsf{T}}Q\xi_n$ , where  $\xi_n \sim \mathcal{N}(0, I_n)$ .

By spectral decomposition,  $Q = C\Omega C^T$  for some orthogonal  $C \in \mathbb{R}^{n \times n}$  and  $\Omega = \operatorname{diag}(1, \ldots, 1, 0, \ldots, 0)$  with rank n - p. Since  $C^{\mathsf{T}} \xi_n \sim \mathcal{N}(0, I_n)$ , we can write

$$\xi_n^{\mathsf{T}} Q \xi_n = \xi_n^{\mathsf{T}} C \Omega C^{\mathsf{T}} \xi_n = (C^{\mathsf{T}} \xi_n)^{\mathsf{T}} \Omega (C^{\mathsf{T}} \xi_n) = \sum_{i=1}^{n-p} (C^{\mathsf{T}} \xi_n)_i^2 \sim \chi_{n-p}^2,$$

where  $(C^{\mathsf{T}}\xi_n)_i$  is the ith element of  $C^{\mathsf{T}}\xi_n$ .

Proof of Lemma 1. We write  $\mathcal{I}_n = \mathcal{I}_n(h_n^2, \sigma_n^2)$  and  $U_n = U(h_n^2, \sigma_n^2)$  for simplicity. By continuous mapping theorem, it suffices to show

$$\mathcal{J}_n^{-1/2} \mathcal{D}_n^{-1} U_n \leadsto \mathcal{N}(0, I_2), \tag{15}$$

where  $\mathcal{I}_n = \mathcal{D}_n \mathcal{J}_n \mathcal{D}_n$ ,  $\mathcal{J}_n$  is the correlation matrix corresponding to  $\mathcal{I}_n$ , and  $\mathcal{D}_n$  is diagonal with positive entries. By definition,  $\mathcal{J}_n$  is the covariance matrix of  $\mathcal{D}_n^{-1}U_n$ .

Also, from Proposition 1,  $\mathcal{D}_{n11} = \|QHQ\|_F/\sqrt{2}$  and  $\mathcal{D}_{n22} = \sqrt{(n-p)}/(\sqrt{2}\sigma_n^2)$ .

To prove (15), it suffices to show that (i) the eigenvalues of  $\mathcal{J}_n$  are bounded away from zero and infinity asymptotically, and (ii) for any unit-length  $w \in \mathbb{R}^2$ ,

$$\frac{w^{\mathsf{T}} \mathcal{D}_n^{-1} U_n}{\text{var}(w^{\mathsf{T}} \mathcal{D}_n^{-1} U_n)^{1/2}} = \frac{w^{\mathsf{T}} \mathcal{D}_n^{-1} U_n}{(w^{\mathsf{T}} \mathcal{J}_n w)^{1/2}} \rightsquigarrow \mathcal{N}(0, 1). \tag{16}$$

We start with (ii). By arguments in the proof of Proposition 1,

$$\sqrt{2}w^{T}\mathcal{D}_{n}^{-1}U_{n} \sim w_{1}\xi_{n}^{\mathsf{T}}(Q_{n}H_{n}Q_{n}/\|Q_{n}H_{n}Q_{n}\|_{F})\xi_{n} - w_{1}\operatorname{tr}(Q_{n}H_{n}Q_{n}/\|Q_{n}H_{n}Q_{n}\|_{F}) 
+ w_{2}\xi_{n}^{\mathsf{T}}Q_{n}\xi_{n}/\sqrt{n-p} - w_{2}\operatorname{tr}(Q_{n}/\sqrt{n-p}) 
= \xi_{n}^{\mathsf{T}}\left(w_{1}\frac{Q_{n}H_{n}Q_{n}}{\|Q_{n}H_{n}Q_{n}\|_{F}} + w_{2}\frac{Q_{n}}{\sqrt{n-p}}\right)\xi_{n} - w_{1}\operatorname{tr}\left(\frac{Q_{n}H_{n}Q_{n}}{\|Q_{n}H_{n}Q_{n}\|_{F}}\right) 
- w_{2}\sqrt{n-p} 
= \xi_{n}^{\mathsf{T}}K_{n}\xi_{n} - \operatorname{tr}(K_{n}),$$

where  $K_n$  is defined by the last equality. Using spectral decomposition of  $K_n$  and rotational invariance of the normal distribution,

$$\xi_n^{\mathsf{T}} K_n \xi_n - \operatorname{tr}(K_n) \sim \sum_{i=1}^{n-p} \gamma_i(K_n) (\xi_{ni}^2 - 1).$$

We establish (16) by verifying Lyapunov's condition (Billingsley, 1995, Theorem 27.3) holds for the right-hand side in the last display. Let

$$a_n = \sum_{i=1}^n \mathsf{E}[\{\gamma_i(K_n)(\xi_{ni}^2 - 1)\}^2] = \mathsf{E}\{(\xi_{11}^2 - 1)^2\} \sum_{i=1}^n \gamma_i(K_n)^2 = \mathsf{E}\{(\xi_{11}^2 - 1)^2\} \|K_n\|_F^2$$

and

$$b_n = \sum_{i=1}^n \mathsf{E}[\{\gamma_i(K_n)(\xi_{ni}^2 - 1)\}^4] = \mathsf{E}\{(\xi_{11}^2 - 1)^4\} \sum_{i=1}^n \gamma_i(K_n)^4 \le \mathsf{E}\{(\xi_{11}^2 - 1)^4\} \|K_n\|^2 \|K_n\|^2_F.$$

Lyapunov's condition says (16) holds if  $b_n/a_n^2 \rightarrow 0$ , which holds if and only if

 $||K_n||/||K_n||_F \to 0$ . We next show  $||K_n|| \to 0$  and  $\liminf_{n \to \infty} ||K_n||_F > 0$ .

For the former we have, by the triangle inequality and submultiplicativity of the spectral norm,

$$||K_n|| \le |w_1| \frac{||Q_n H_n Q_n||}{||Q_n H_n Q_n||_F} + |w_2| \frac{||Q_n||}{\sqrt{n-p}} = |w_1| \frac{||H_n||}{||Q_n H_n Q_n||_F} + |w_2| \frac{1}{\sqrt{n-p}}$$
$$= |w_1| \frac{\gamma_1 (D_n^2)^{1/2}}{||Q_n H_n Q_n||_F} + |w_2| \frac{1}{\sqrt{n-p}},$$

which tends to zero by assumption 2 since, by H.1.h of Marshall et al. (2011),

$$||Q_{n}H_{n}Q_{n}||_{F}^{2} = \operatorname{tr}(Q_{n}H_{n}Q_{n}H_{n})$$

$$= \operatorname{tr}(H_{n}Q_{n}H_{n} - P_{n}H_{n}Q_{n}H_{n})$$

$$= \operatorname{tr}(H_{n}^{2} - H_{n}P_{n}H_{n} - P_{n}H_{n}^{2} + P_{n}H_{n}P_{n}H_{n})$$

$$\geq \operatorname{tr}(H_{n}^{2}) - 2\operatorname{tr}(P_{n}H_{n}^{2})$$

$$\geq \operatorname{tr}(D_{n}^{2}) - 2\sum_{i=1}^{p} \gamma_{i}(D_{n}^{2}).$$

To show  $\liminf_{n\to\infty} ||K_n||_F > 0$  we consider two cases. First, if  $|w_1| \neq |w_2|$ , then by reverse triangle inequality,

$$\left\| w_1 \frac{Q_n H_n Q_n}{\|Q_n H_n Q_n\|_F} + w_2 \frac{Q_n}{\sqrt{n-p}} + \right\|_F \ge \left\| \left\| w_1 \frac{Q_n H_n Q_n}{\|Q_n H_n Q_n\|_F} \right\|_F - \left\| w_2 \frac{Q_n}{\sqrt{n-p}} \right\|_F \right\|_{(17)}$$

$$= ||w_1| - |w_2||,$$

which is greater than zero and does not depend on n. If instead  $|w_1| = |w_2|$ ,

$$\left\| w_{1} \frac{Q_{n} H_{n} Q_{n}}{\|Q_{n} H_{n} Q_{n}\|_{F}} + w_{2} \frac{Q_{n}}{\sqrt{n-p}} \right\|_{F}^{2} = w_{1}^{2} \left\| \frac{Q_{n}}{\sqrt{n-p}} \pm \frac{Q_{n} H_{n} Q_{n}}{\|Q_{n} H_{n} Q_{n}\|_{F}} \right\|_{F}^{2} 
= w_{1}^{2} \operatorname{tr} \left( \frac{Q_{n}^{2}}{n-p} + \frac{(Q_{n} H_{n} Q_{n})^{2}}{\|Q_{n} H_{n} Q_{n}\|_{F}^{2}} \pm 2 \frac{Q_{n}}{\sqrt{n-p}} \frac{Q_{n} H_{n} Q_{n}}{\|Q_{n} H_{n} Q_{n}\|_{F}} \right) 
= 2w_{1}^{2} \left\{ 1 \pm \operatorname{tr} \left( \frac{Q_{n}}{\sqrt{n-p}} \frac{Q_{n} H_{n} Q_{n}}{\|Q_{n} H_{n} Q_{n}\|_{F}} \right) \right\},$$
(18)

so, since  $Q_n$  is idempotent, it suffices to show

$$\limsup_{n \to \infty} \frac{|\operatorname{tr}(Q_n H_n)|}{\sqrt{n-p} ||Q_n H_n Q_n||_F} < 1.$$

We bound the numerator first. By the triangle inequality,

$$|\operatorname{tr}(Q_n H_n)| = |\operatorname{tr}(H_n) - \operatorname{tr}(P_n H_n)| \le |\operatorname{tr}(H_n)| + |\operatorname{tr}(P_n H_n)|.$$

Applying H.1.g and H.1.h of Marshall et al. (2011) (note their comment regarding positive semi-definiteness) to  $tr(P_nH_n)$ , we have

$$\sum_{i=1}^{p} \gamma_{n-p+i}(D_n) \le \operatorname{tr}(P_n H_n) \le \sum_{i=1}^{p} \gamma_i(D_n), \tag{19}$$

which gives an upper bound of  $|tr(P_nH_n)|$ . Using this bound followed by Jensen's inequality gives

$$|\operatorname{tr}(P_n H_n)| \le \max \left\{ \left| \sum_{i=1}^p \gamma_i(D_n) \right|, \left| \sum_{i=1}^p \gamma_{n-p+i}(D_n) \right| \right\}$$

$$\le \sum_{i=1}^p \sqrt{\gamma_i(D_n^2)}$$

$$\le \left( p \sum_{i=1}^p \gamma_i(D_n^2) \right)^{1/2}.$$

We thus have that the ratio to be bounded is no greater than

$$\frac{|\operatorname{tr}(D_n)| + \{p \sum_{i=1}^p \gamma_i(D_n^2)\}^{1/2}}{\sqrt{n-p}\{\operatorname{tr}(D_n^2) - 2 \sum_{i=1}^p \gamma_i(D_n^2)\}^{1/2}} = \frac{|\operatorname{tr}(D_n)|}{\sqrt{n-p}\{\operatorname{tr}(D_n^2) - 2 \sum_{i=1}^p \gamma_i(D_n^2)\}^{1/2}} + o(1).$$

By assumption 2, and since  $n/(n-p) \to 1$  by assumption 1, the last right-hand side

is

$$\{1 + o(1)\} \frac{n^{-1/2} |\operatorname{tr}(D_n)|}{\operatorname{tr}(D_n^2)^{1/2}} + o(1),$$

the upper limit of which is less than one by assumption 3, which establishes (16).

It remains only to establish (i). First, since  $\mathcal{J}_n$  is a correlation matrix, it's entries are no greater than one in absolute value, and hence  $\gamma_1(\mathcal{J}_n) = \|\mathcal{J}_n\| \le \|\mathcal{J}_n\|_F \le 2$ . To bound  $\gamma_2(\mathcal{J}_n)$  away from zero, suppose for contradiction  $\liminf_{n\to\infty} \gamma_2(\mathcal{J}_n) = 0$ . Then we can pick a subsequence along which  $\gamma_2(\mathcal{J}_n) \to 0$ . Since  $\|\mathcal{J}_n\|_F$  is bounded, we may also, by the Bolzano-Weierstrass property, pick the subsequence in such a way that  $\mathcal{J}_n$  converges to some positive semi-definite  $\mathcal{J}$ . Since  $\gamma_2(\mathcal{J}_n) \to 0$ ,  $\mathcal{J}$  has at least one vanishing eigenvalue. Let w be a corresponding eigenvector. Then, along the subsequence,  $w^{\mathsf{T}}\mathcal{J}_n w \to 0$ . But, by the arguments following (16),  $w^{\mathsf{T}}\mathcal{J}_n w = \|K_n\|_F^2$ , and we already proved  $\liminf_{n\to\infty} \|K_n\|_F > 0$ , which gives the desired contradiction. Thus, (i) holds and the proof is complete.

*Proof of Lemma 2.* Define a remainder  $R_{n1}$  to be dealt with later by

$$U_{n1}(h_n^2, \tilde{\sigma}_n^2) = U_{n1}(h_n^2, \sigma_n^2) + \nabla_{12}^2 l_R(h_n^2, \sigma_n^2)(\tilde{\sigma}_n^2 - \sigma_n^2) + R_{n1}.$$

Since

$$U_{n2}(h_n^2, \sigma_n^2) = -\frac{n-p}{2\sigma_n^2} + \frac{(n-p)\tilde{\sigma}_n^2}{2\sigma_{np}^4},$$

we have  $\tilde{\sigma}_n^2 - \sigma_n^2 = 2\sigma_{np}^4 U_{n2}(h_n^2, \sigma_n^2)/(n-p) = U_{n2}/\mathcal{I}_{n22}(h_n^2, \sigma_n^2)$ . For the remainder of the proof, we omit the arguments when they are the true parameters  $(h_n^2, \sigma_n^2)$ . Then

$$U_{n1}(h_n^2, \tilde{\sigma}_n^2) = U_{n1} + (\nabla_{12}^2 l_R) U_{n2} / \mathcal{I}_{n22} + R_{n1} = U_{n1} - \mathcal{I}_{n12} \mathcal{I}_{n22}^{-1} U_{n2} + R_{n1} + R_{n2},$$

where  $R_{n2} = (\nabla_{12}^2 l_R + \mathcal{I}_{n12}) U_{n2} / \mathcal{I}_{n22}$ . We complete the proof by showing (i)  $R_{n1}^2 \mathcal{I}_n^{(11)} \to 0$  in probability, (ii)  $R_{n2}^2 \mathcal{I}_n^{(11)} \to 0$  in probability, and (iii)

$$\{U_{n1} - \mathcal{I}_{n12}U_{n2}/\mathcal{I}_{n22}\}(\mathcal{I}_n^{(11)})^{1/2} \rightsquigarrow \mathcal{N}(0,1).$$

It will be useful to note

$$\mathcal{I}_{n}^{(11)} = (\mathcal{I}_{n11} - \mathcal{I}_{n21}^2 / \mathcal{I}_{n22})^{-1} = \frac{\mathcal{I}_{n22}}{\mathcal{I}_{n11} \mathcal{I}_{n22} - \mathcal{I}_{n21}^2},$$

which in fact does not depend on  $\sigma_n^2$ . We begin with (ii) and have

$$\nabla_{12}^{2} l_{R} = \nabla_{1} U_{n2} = -\frac{1}{2\sigma_{np}^{4}} \sum_{i=1}^{n} \frac{(\lambda_{i} - 1)(Y_{i} - X_{i}^{T} \tilde{\beta})^{2}}{(h_{n}^{2} \lambda_{i} + 1 - h_{n}^{2})^{2}}$$
$$= -\frac{1}{2\sigma_{n}^{2}} (Y - X\tilde{\beta})^{\mathsf{T}} \Sigma_{n}^{-1/2} D_{n} \Sigma_{n}^{-1/2} (Y - X\tilde{\beta})$$

From the proof of Proposition 1,  $Y - X\tilde{\beta} = (I_n - \Sigma_n^{1/2} P \Sigma^{-1/2})Y = \Sigma_n^{1/2} (I_n - P_n) \Sigma_n^{-1/2} Y = \Sigma_n^{1/2} Q_n \Sigma_n^{-1/2} Y$ , and so

$$\nabla_{12}^2 l_R = -\frac{1}{2\sigma_n^2} Y^{\mathsf{T}} \Sigma_n^{-1/2} Q_n D_n Q_n \Sigma_n^{-1/2} Y.$$

Thus,  $-R_{n2}(\mathcal{I}_n^{(11)})^{1/2}$  is

$$\frac{1}{2n^{-1/2}\sigma_n^2(\mathcal{I}_{n11}\mathcal{I}_{n22} - \mathcal{I}_{n12}^2)^{1/2}} \frac{1}{n^{1/2}} \left( Y^\mathsf{T} \Sigma_n^{-1/2} Q_n D_n Q_n \Sigma_n^{-1/2} Y - \operatorname{tr}(Q_n D_n) \right) U_{n2} / \sqrt{\mathcal{I}_{n22}},$$

which we can write as  $I^{-1/2} \times II \times III$  by defining

$$I = 4n^{-1}\sigma_{np}^{4}(\mathcal{I}_{n11}\mathcal{I}_{n22} - \mathcal{I}_{n12}^{2}) = \operatorname{tr}(Q_{n}D_{n}Q_{n}D_{n}) \left\{ 1 - p/n - \frac{n^{-1}\operatorname{tr}(Q_{n}D_{n})^{2}}{\operatorname{tr}(Q_{n}D_{n}Q_{n}D_{n})} \right\};$$

$$II = \frac{1}{n^{1/2}} \left( Y^{\mathsf{T}}\Sigma_{n}^{-1/2}Q_{n}D_{n}Q_{n}\Sigma_{n}^{-1/2}Y - \operatorname{tr}(Q_{n}D_{n}) \right);$$

$$III = \frac{U_{n2}}{\sqrt{\mathcal{I}_{n22}}}.$$

By calculations in the proof of Lemma 1, the term of I in curly brackets is bounded away from zero asymptotically (c.f. condition 3 of that lemma), so  $I \ge c \operatorname{tr}(Q_n D_n Q_n D_n)$  for some c > 0 and all large enough n. Next, II has mean zero and variance  $\operatorname{tr}(Q_n D_n Q_n D_n)/n$ . Thus,  $I^{-1/2} \times II$  has mean zero and variance no greater than

1/(cn), and hence it tends to zero in mean square and probability. By Chebyshev's inequality, III is  $O_P(1)$ , and hence  $I^{-1/2} \times II \times III \to 0$  in probability, which completes the proof of (ii).

To show (i), use that, by Taylor's theorem with Lagrange-form remainder, for some  $\bar{\sigma}_{np}^2$  between  $\sigma_n^2$  and  $\tilde{\sigma}_{np}^2$ ,

$$R_{n1} = \nabla_{122}^3 l_R(h_n^2, \bar{\sigma}_{np}^2) (\tilde{\sigma}_{np}^2 - \sigma_n^2)^2 / 2.$$

Using calculations similar to when deriving  $\nabla_{12}^2 l_R$ , we get

$$\nabla_{122}^{3}l_{R}(h_{n}^{2},\bar{\sigma}_{np}^{6}) = \frac{1}{2\bar{\sigma}_{np}^{6}} \sum_{i=1}^{n} \frac{(\lambda_{i}-1)(Y_{i}-X_{i}^{T}\tilde{\beta})^{2}}{(h_{n}^{2}\lambda_{i}+1-h_{n}^{2})^{2}} = \frac{\sigma_{n}^{2}}{2\bar{\sigma}_{np}^{6}} Y^{\mathsf{T}} \Sigma_{n}^{-1/2} Q_{n} D_{n} Q_{n} \Sigma_{n}^{-1/2} Y.$$

Thus,  $2R_{n1}(\mathcal{I}_n^{(11)})^{1/2}$  is

$$\begin{split} &\frac{\sigma_{n}^{6}}{\bar{\sigma}_{np}^{6}} \frac{1}{2\sigma_{np}^{4}} \frac{\mathcal{I}_{n22}^{1/2}}{(\mathcal{I}_{n11}\mathcal{I}_{n22} - \mathcal{I}_{n12}^{2})^{1/2}} Y^{\mathsf{T}} \Sigma_{n}^{-1/2} Q_{n} D_{n} Q_{n} \Sigma_{n}^{-1/2} Y (U_{n2}/\mathcal{I}_{n22})^{2} \\ &= \frac{\sigma_{n}^{6}}{\bar{\sigma}_{np}^{6}} \frac{n^{-1/2}}{2\sigma_{n}^{2} n^{-1/2} (\mathcal{I}_{n11}\mathcal{I}_{n22} - \mathcal{I}_{n12}^{2})^{1/2}} Y^{\mathsf{T}} \Sigma_{n}^{-1/2} Q_{n} D_{n} Q_{n} \Sigma_{n}^{-1/2} Y \frac{U_{n2}^{2}}{\mathcal{I}_{n22}} \frac{\mathcal{I}_{n22}^{1/2}}{\mathcal{I}_{n22} \sigma_{n}^{2}} \\ &= \frac{\sigma_{n}^{6}}{\bar{\sigma}_{np}^{6}} \frac{n^{-1/2}}{\mathrm{I}^{1/2}} Y^{\mathsf{T}} \Sigma_{n}^{-1/2} Q_{n} D_{n} Q_{n} \Sigma_{n}^{-1/2} Y \times \mathrm{III}^{2} \times \frac{\sqrt{2}}{\sqrt{n-p}} \\ &= \frac{\sigma_{n}^{6}}{\bar{\sigma}_{np}^{6}} \times \mathrm{I}^{-1/2} \times (\mathrm{II} + \mathrm{tr}(Q_{n}D_{n})/\sqrt{n}) \times \mathrm{III}^{2} \times \frac{\sqrt{2}}{\sqrt{n-p}}. \end{split}$$

Note  $|\bar{\sigma}_{np}^2/\sigma_n^2 - 1| \leq |\tilde{\sigma}_n^2/\sigma_n^2 - 1|$  and  $\tilde{\sigma}_n^2/\sigma_n^2 \sim \chi_{n-p}^2/(n-p)$ , which tends to 1 in probability. Thus,  $\bar{\sigma}_{np}^2/\sigma_n^2$  tends to one in probability and, by the arguments used to establish (ii), it suffices to show  $|\operatorname{tr}(Q_nD_n)/n| \times I^{-1/2} \to 0$ . But Jensen's inequality applied to the sum of the n real eigenvalues of QDQ, which are the same as those of QD, gives  $|\operatorname{tr}(Q_nD_n)/n| \leq \operatorname{tr}(Q_nD_nQ_nD_n)^{1/2}n^{-1/2}$  and in the proof of (ii) we argued  $I \geq c \operatorname{tr}(Q_nD_nQ_nD_n)$ , so

$$|\operatorname{tr}(Q_n D_n)/n| \times I^{-1/2} \le (cn)^{-1/2} \to 0.$$

Finally, to show (iii), let  $a_n = (\mathcal{I}_n^{(11)})^{1/2}[1, -\mathcal{I}_{n12}/\mathcal{I}_{n22}]^\mathsf{T}$ , so that what we want to show is  $a_n^\mathsf{T}U_n \rightsquigarrow \mathrm{N}(0,1)$ . A direct calculation shows the variance of  $a_n^\mathsf{T}U_n$  is  $a_n^\mathsf{T}\mathcal{I}_n a_n = 1$ . Let  $b_n = \mathcal{I}_n^{1/2}a_n$ . Then  $||b_n|| = 1$  and  $a_n^\mathsf{T}U_n = b_n^\mathsf{T}\mathcal{I}_n^{-1/2}U_n$ . The proof of Lemma 1 shows  $\mathcal{I}_n^{-1/2}U_n \rightsquigarrow \mathrm{N}(0,I_2)$ . Moroever, by the Bolzano-Weierstrass Theorem and the subsequence principle, we may assume  $b_n \to b$  for some b with ||b|| = 1. The Cramér-Wold Theorem says  $b^\mathsf{T}\mathcal{I}_n^{-1/2}U_n \rightsquigarrow \mathrm{N}(0,1)$ , and therefore  $b_n^\mathsf{T}\mathcal{I}_n^{-1/2}U_n = b^\mathsf{T}\mathcal{I}_n^{-1/2}U_n + (b_n - b)^\mathsf{T}\mathcal{I}_n^{-1/2}U_n = b^\mathsf{T}\mathcal{I}_n^{-1/2}U_n + o_\mathrm{P}(1) \rightsquigarrow \mathrm{N}(0,1)$  by Slutsky's Theorem, which completes the proof.

*Proof of Theorem 2.* Condition 1 is the same as in Lemma 1, so it suffices to verify the remaining conditions of Lemma 1. First, condition 3 ensures

$$\liminf_{n \to \infty} \min_{i \in \{1, \dots, n\}} \{ h_n^2(\lambda_{ni} - 1) + 1 \} > 0,$$

so together with condition 2 we have, for some  $c_{\text{max}} \in (0, \infty)$ ,

$$\limsup_{n \to \infty} \sqrt{\gamma_1(D_n^2)} = \limsup_{n \to \infty} \max_{i \in \{1, \dots, n\}} \left| \frac{\lambda_{ni} - 1}{h_n^2(\lambda_{ni} - 1) + 1} \right| \le c_{\text{max}}. \tag{20}$$

Condition 2 also implies we may upon increasing  $c_{\max}$  if needed have, for any  $\lambda_{ni}$  and  $\lambda_{nj}$ ,

$$0 < (h_n^2(\lambda_{ni} - 1) + 1)(h_n^2(\lambda_{nj} - 1) + 1) \le \max\{1, \lambda_{ni}\} \max\{1, \lambda_{nj}\} \le c_{\max}^2$$

Thus, if  $|\lambda_{ni} - \lambda_{nj}| > \epsilon$ ,

$$\left| \frac{\lambda_{ni} - 1}{h_n^2(\lambda_{ni} - 1) + 1} - \frac{\lambda_{nj} - 1}{h_n^2(\lambda_{nj} - 1) + 1} \right| = \frac{|\lambda_{ni} - \lambda_{nj}|}{(h_n^2(\lambda_{ni} - 1) + 1)(h_n^2(\lambda_{nj} - 1) + 1)} > \frac{\epsilon}{c_{\text{max}}^2}.$$
(21)

Thus, if two diagonal elements of  $\Lambda_n$  are more than  $\epsilon$  apart, then the corresponding diagonal elements of  $D_n$  are more than  $\epsilon/c_{\max}^2$  apart.

Applying (21) with  $\lambda_{nj} = 1$  shows the number of eigenvalues of  $D_n$  in  $[-\epsilon/(c_{\text{max}}^2), \epsilon/(c_{\text{max}}^2)]$  is at most  $k_n^1$ , the number of eigenvalues of  $\Lambda_n$  in  $[1 - \epsilon, 1 + \epsilon]$ . Consequently, together with condition 1 we have

$$\frac{\sum_{i=1}^{p} \gamma_i(D_n^2)}{\sum_{i=1}^{n} \gamma_i(D_n^2)} \le \frac{pc_{\max}^2}{(n-k_n^1)\epsilon^2/c_{\max}^4} \to 0,$$

which verifies condition 2 of Lemma 1.

Let  $\bar{\gamma}_n = \frac{1}{n} \sum_{i=1}^n \gamma_i(D_n)$  and let  $r_n$  be the number of eigenvalues of  $D_n$  in  $[\bar{\gamma}_n - \epsilon/(4c_{\max}^2), \bar{\gamma}_n + \epsilon/(4c_{\max}^2)]$ . Then

$$\sum_{i=1}^{n} \gamma_i(D_n^2) - n\bar{\gamma}_n^2 = \sum_{i=1}^{n} \{\gamma_i(D_n) - \bar{\gamma}_n\}^2 \ge \frac{\epsilon^2}{16c_{\max}^4} (n - r_n).$$

We are done if  $\limsup_{n\to\infty} r_n/n < 1$  since, then, condition 3 of Lemma 1 is satisfied:

$$\limsup_{n \to \infty} \frac{\operatorname{tr}(D)^2}{n \operatorname{tr}(D^2)} = 1 - \liminf_{n \to \infty} \frac{\sum_{i=1}^n \gamma_i(D_n^2) - n \bar{\gamma}_n^2}{\sum_{i=1}^n \gamma_i(D_n^2)} \le 1 - \liminf_{n \to \infty} \frac{\epsilon^2 n}{16c_{\max}^6(n-r_n)} < 1.$$

To see  $r_n/n$  is indeed bounded away from one, observe  $\bar{\gamma}_n$  is bounded by (20), so we may, by the Bolzano–Weierstrass property and passing to a subsequence if necessary, assume  $\bar{\gamma}_n$  converges to some  $\gamma_0 \in \mathbb{R}$ . Thus, for all large enough n, all the eigenvalues of  $D_n$  in  $[\bar{\gamma}_n - \epsilon/(4c_{\max}^2), \bar{\gamma}_n + \epsilon/(4c_{\max}^2)]$  are also in  $[\gamma_0 - \epsilon/(2c_{\max}^2), \gamma_0 + \epsilon/(2c_{\max}^2)]$ . But then, by (21), the corresponding  $\lambda_{ni}$  are in  $[\lambda_0 - \epsilon, \lambda_0 + \epsilon]$  for some fixed  $\lambda_0$ . For example, we can fix any i and n such that  $\gamma_i(D_n) \in [\gamma_0 - \epsilon/(2c_{\max}^2), \gamma_0 + \epsilon/(2c_{\max}^2)]$  and let  $\lambda_0$  be the corresponding  $\lambda_{ni}$ . Thus,  $r_n \leq k_n^{\lambda_0}$  and the proof is completed since  $\lim \sup_{n \to \infty} k_n^{\lambda_0}/n < 1$  by condition 4.