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Numerical implementation of the QuEST function

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ABSTRACT

Certain estimation problems involving the covariance matrix in large dimensions are considered. Due to the breakdown of finite-dimensional asymptotic theory when the dimension is not negligible with respect to the sample size, it is necessary to resort to an alternative framework known as large-dimensional asymptotics. Recently, an estimator of the eigenvalues of the population covariance matrix has been proposed that is consistent according to a mean-squared criterion under large-dimensional asymptotics. It requires numerical inversion of a multivariate nonrandom function called the QuEST function. The numerical implementation of this QuEST function in practice is explained through a series of six successive steps. An algorithm is provided in order to compute the Jacobian of the QuEST function analytically, which is necessary for numerical inversion via a nonlinear optimizer. Monte Carlo simulations document the effectiveness of the code.

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1. Introduction

Many data sets in finance, economics, biostatistics, and electrical engineering, among a host of other fields, contain large numbers of related variables. The estimation of the covariance matrix poses challenging statistical problems when the dimension is not small relative to sample size. Approximations that are valid under traditional asymptotics, that is, when the dimension remains fixed while the sample size goes to infinity, perform poorly. This is why attention has turned to *large-dimensional asymptotics* where the dimension and the sample size go to infinity together, with their ratio converging to a finite, nonzero limit called the *limiting concentration (ratio)*.

Under large-dimensional asymptotics, the sample eigenvalues are not suitable estimators of the population eigenvalues. A new estimator of the population eigenvalues under large-dimensional asymptotics was recently introduced by Ledoit and Wolf (2015); this estimator is consistent in the sense that the mean squared deviation between the estimated eigenvalues and the population eigenvalues converges to zero almost surely. The estimator hinges critically on a multivariate nonrandom function called the QuEST function; this acronym stands for *Quantized Eigenvalues Sampling Transform*. Ledoit and Wolf (2015) provide the mathematical definition of the QuEST function but do not provide any details about the numerical implementation. The problem of numerical implementation is non-trivial because of the complexity of the definition of the QuEST function. A direct application of this method is the optimal estimation of the covariance matrix in the class of rotation-equivariant estimators introduced by Stein (1975, 1986) under various loss functions; see Ledoit and Wolf (2017b).

This paper explains how to numerically implement the QuEST function accurately and efficiently. In addition, given that the estimation of the population eigenvalues requires numerically inverting the QuEST function using a nonlinear optimizer, we also provide the Jacobian analytically.

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The remainder of this paper is organized as follows. Section 2 reviews the literature on this subject. Section 3 gives the definition of the problem that will be solved numerically. Sections 5–10 describe in detail the six steps needed to implement the QuEST function numerically, delineating all the mathematical results that are needed along the way. Section 11 provides extensive Monte Carlo simulations. Section 12 concludes.

2. Literature review

2.1. ‘Fundamental equation’ of random matrix theory

This section serves as a concise primer on relevant facts about covariance-matrix eigenvalues under large-dimensional asymptotics. It lays the background for the subsequent review of the more recent trends in the literature, and also for the definition of the QuEST function in Section 3. The reader looking for a comprehensive monograph-level treatment of the subject should consult Bai and Silverstein (2010). For a more informal, qualitative, and intuitive introduction, see Engle et al. (forthcoming, Section 9).

2.1.1. Spectral distribution

Marčenko and Pastur (1967) were the first to derive a relationship between population and sample covariance matrix eigenvalues in the large-dimensional limit, that is, when the ratio of the matrix dimension $p := p(n)$ to the sample size n converges to some $c \in (0, +\infty)$ called the limiting concentration ratio. Since the number of eigenvalues goes to infinity, they introduced the spectral cumulative distribution functions: $\forall x \in \mathbb{R}, H_n(x) := \sum_{i=1}^p \mathbb{1}_{\{\tau_{n,i} \leq x\}}/p$, and $F_n(x) := \sum_{i=1}^p \mathbb{1}_{\{\lambda_{n,i} \leq x\}}/p$, where $(\tau_{n,1}, \dots, \tau_{n,p})$ and $(\lambda_{n,1}, \dots, \lambda_{n,p})$ are the population and sample eigenvalues, respectively, and $\mathbb{1}$ is the indicator function.

2.1.2. Limiting spectral distribution

Marčenko and Pastur (1967) show that if H_n converges weakly to some limiting population spectral distribution H , and certain other conditions are met, then there exists some nonrandom limiting sample spectral distribution F to which F_n converges weakly in probability. This result was subsequently broadened and strengthened by a number of other authors, including Silverstein (1995), Silverstein and Bai (1995), and Bai and Silverstein (1998, 1999). Of particular note is the formula established by Silverstein and Choi (1995) that gives the mass at zero:

$$F(0) = \max\left\{1 - \frac{1}{c}, H(0)\right\}. \tag{2.1}$$

2.1.3. Stieltjes transform

Marčenko and Pastur (1967) characterize F indirectly through its Stieltjes (1894) transform, which is defined as:

$$\forall z \in \mathbb{C}^+ \quad m_F(z) := \int_{-\infty}^{+\infty} \frac{1}{\lambda - z} dF(\lambda), \tag{2.2}$$

where \mathbb{C}^+ denotes the half-plane of complex numbers with strictly positive imaginary parts. From m_F it is easy to recover F through the well-known inversion formula

$$F(b) - F(a) = \lim_{\eta \rightarrow 0^+} \frac{1}{\pi} \int_a^b \text{Im}[m_F(\xi + i\eta)] d\xi, \tag{2.3}$$

if F is continuous at a and b , where Im denotes the imaginary part of a complex number.

2.1.4. The fundamental equation

In addition, Marčenko and Pastur (1967) provide an equation that links F and H for a given c . It can be reformulated in different ways, and the most convenient version for our purposes is the one initially introduced by Silverstein (1995, Equation (1.4)): for every $z \in \mathbb{C}^+$, $m := m_F(z)$ is the unique solution in the set

$$\left\{ m \in \mathbb{C} : -\frac{1-c}{z} + cm \in \mathbb{C}^+ \right\} \tag{2.4}$$

to the equation

$$m = \int_{-\infty}^{+\infty} \frac{1}{\tau [1 - c - czm] - z} dH(\tau). \tag{2.5}$$

This ‘Fundamental Equation’ essentially goes from H (population eigenvalues) to F (sample eigenvalues); therefore, there has been considerable interest in the question of how to ‘invert’ it, that is, how to recover the population eigenvalues from the sample eigenvalues. The present paper focuses on one specific method to do this, which is through numerical inversion of the QuEST function, a discretized version of the fundamental equation. This method is fully described in Section 3.

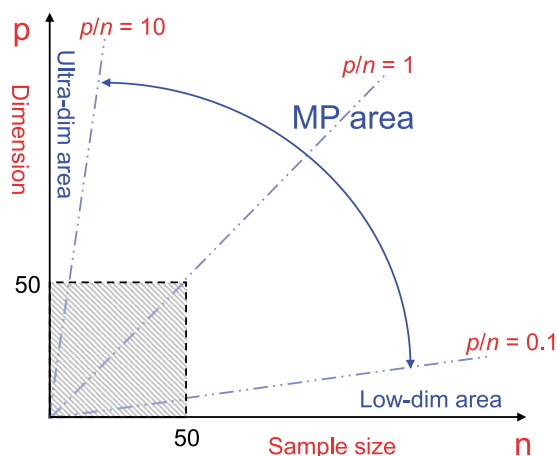


Fig. 1. Heuristic comparison of the area of relevance of Marčenko–Pastur asymptotics versus traditional fixed-dimension asymptotics.

2.1.5. The MP area

Fig. 1, publicized by Yao (2015) in a conference presentation, gives a heuristic view of the area where Marčenko–Pastur asymptotic theory is more useful (labeled “MP area”) versus the area where standard fixed-dimension asymptotic theory applies (labeled “Low-dim area”).

This insight is further developed in the recent book by Yao et al. (2015).

In essence, when $p/n < 0.1$, the amount of effort required to work through the complexities of large-dimensional asymptotics is not rewarded, given that traditional (fixed-dimensional) asymptotics constitute a sufficiently accurate approximation for practical purposes. Obviously, the choice of cutoff is somewhat arbitrary, but we reproduce the picture exactly as it was originally presented. Conversely, when $p/n > 10$, we enter the ultra-high-dimensional area, where the large-dimensional asymptotic theory that assumes c is finite may start to break down; little is known about what happens in that zone, because few have ventured there. Once again, the choice of cutoff is heuristic.

2.2. Estimation of the population covariance matrix eigenvalues

El Karoui (2008) proposes a way to estimate the empirical cumulative distribution function (c.d.f.) of population eigenvalues under large-dimensional asymptotics using a different approach than the QuEST function. However, the code executing his algorithm has not been made available to other researchers in the field, and those who have tried to replicate it have not enjoyed much success. The state of affairs is aptly summarized by Li et al. (2013):

Actually, the general approach in El Karoui (2008) has several implementation issues that seem to be responsible for its relatively low performance as attested by the very simple nature of provided simulation results.

There are three reasons why the same criticisms cannot be leveled against the QuEST function: first, a Matlab code implementing the QuEST function has already been used independently by Welsing (2015), Ito and Kubokawa (2015), Huang and Fryzlewicz (2015), and Lam (2016), among others¹; second, the present paper opens up the code of the QuEST function and its Jacobian to the general public for inspection and potential improvements; and third, Section 11 provides an extensive Monte Carlo study with nearly one third of a million simulations across a variety of challenging scenarios.

Apart from El Karoui (2008), other proposals have been put forward, making this field one of the most active ones in multivariate analysis in recent years.

- Rao et al. (2008) provide a solution when the population spectrum has a staircase structure, typically with half of the eigenvalues equal to one and the rest equal to two. The ability of this approach to handle the general case where there can be up to p distinct population eigenvalues, with p going to infinity, is not established. The Monte Carlo simulations of Rao et al. (2008) assume no more than four distinct population eigenvalues.
- Mestre (2008) provides a solution when the concentration ratio $c_n := p/n$ is sufficiently small and/or the distinct population eigenvalues sufficiently far from one another, that is, when the sample eigenvalues display what is known as ‘spectral separation’. This is a favorable situation where the sample eigenvalues are grouped into easily identifiable clusters, each cluster corresponding to one single population eigenvalue (which can have multiplicity higher than one).

¹ The code can be downloaded at <http://www.econ.uzh.ch/en/people/faculty/wolf/publications.html> under the link “Programming Code”.

Although his algorithm is computationally elegant, the method is severely limited. It cannot be applied when (i) the limiting distribution of population eigenvalues is discrete but spectral separation does not occur or (ii) the limiting distribution of population eigenvalues has a continuous part. Many applications of interest fall under the latter category.

- [Bai et al. \(2010\)](#) propose a solution based on the method of moments when the parametric dimension of the population spectrum is finite. They demonstrate good behavior up to order four.
- [Chen et al. \(2011\)](#) elaborate on the previous paper by providing more rigorous justification of the method when the model order is unknown. But Monte Carlo simulations only go to order three.
- [Yao et al. \(2012\)](#) can be seen as a cross between the papers of [Mestre \(2008\)](#) and [Bai et al. \(2010\)](#), but also requiring a finite number of distinct population eigenvalues. In practice, Monte Carlo simulations provided by the authors do not go above three distinct population eigenvalues.

The common point between all these other methods is that they do not purport to address the general case. They work with a finite number of degrees of freedom (in practice no more than four) in the choice of the population spectral distribution, whereas the real number is p , which goes to infinity. This is why it is important to avoid the criticisms that have been leveled at the only other ostensibly general approach, that of [El Karoui \(2008\)](#), by fully explaining how to numerically implement the QuEST function, and by providing extensive Monte Carlo simulations showing that it works in practice under a wide variety of circumstances.

Finally, [Dobriban \(2015\)](#) also provides a numerical method for solving the so-called ‘Fundamental Equation’ of Random Matrix Theory; see Section 3. He does not compute the QuEST function explicitly, and does not provide the Jacobian analytically; as a result, numerical inversion is very difficult. Though it should be noted that his paper is not focused on the problem of recovering the population eigenvalues.

2.3. Potential applications

The numerical implementation of the QuEST function given in this paper is essential for the estimation of the population eigenvalues, which in turn is essential for nonlinear shrinkage estimation of the covariance matrix under large-dimensional asymptotics; see [Ledoit and Wolf \(2012, 2015\)](#). Many fields are interested in shrinkage estimation of the covariance matrix when the number of variables is large:

Acoustics Optimally removing noise from signals captured from an array of hydrophones ([Zhang et al., 2009](#)).

Cancer Research Mapping out the influence of the Human Papillomavirus (HPV) on gene expression ([Pyeon et al., 2007](#)).

Chemistry Estimating the temporal autocorrelation function (TACF) for fluorescence correlation spectroscopy ([Guo et al., 2012](#)).

Civil Engineering Detecting and identifying vibration-based bridge damage through Random Coefficient Pooled (RCP) models ([Michaelides et al., 2011](#)).

Climatology Detecting trends in average global temperature through the optimal fingerprinting method ([Ribes et al., 2013](#)).

Econometrics Specifying the target covariance matrix in the Dynamic Conditional Correlation (DCC) model to capture time-series effects in the second moments ([Hafner and Reznikova, 2012](#); [Engle et al. forthcoming](#)).

Electromagnetics Studying correlation between reverberation chamber measurements collected at different stirrer positions ([Pirkl et al., 2012](#)).

Entertainment Technology Designing a video game controlled by performing tricks on a skateboard ([Anlauff et al., 2010](#)).

Finance Reducing the risk in large portfolios of stocks ([Ledoit and Wolf, 2003](#)).

Genetics Inferring large-scale covariance matrices from functional genomic data ([Schäfer and Strimmer, 2005](#)).

Geology Modeling multiphase flow in subsurface petroleum reservoirs with the iterative stochastic ensemble method (ISEM) on inverse problems ([Elsheikh et al., 2013](#)).

Image Recognition Detecting anomalous pixels in hyperspectral imagery ([Bachega et al., 2011](#)).

Neuroscience Calibrating brain–computer interfaces ([Lotte and Guan, 2009](#)).

Psychology Modeling co-morbidity patterns among mental disorders ([Markon, 2010](#)).

Road Safety Research Developing an emergency braking assistance system ([Haufe et al., 2011](#)).

Signal Processing Combining data recorded by an array of sensors to minimize the noise ([Chen et al., 2010](#)).

Speech Recognition Automatically transcribing records of phone conversations ([Bell and King, 2009](#)).

Up until now, these fields have had to satisfy themselves with *linear* shrinkage estimation of the covariance matrix ([Ledoit and Wolf, 2003, 2004](#)). However, as explained by [Ledoit and Wolf \(2012, 2015\)](#), this approach is asymptotically suboptimal in the class of rotation-equivariant estimators relative to nonlinear shrinkage, which requires numerical implementation of the QuEST function. The present paper makes this new and improved method universally available in practice.

3. Definition of the QuEST function

The mathematical definition of the QuEST function is given by [Ledoit and Wolf \(2015\)](#); it is reproduced here for convenience. In particular, the large-dimensional asymptotic framework assumes that the number of population eigenvalues, p , goes to infinity together with the sample size, n , with their ratio $c_n := p/n$ converging to a finite, positive constant $c \neq 1$, which is called the *limiting concentration (ratio)*. Note that the case $c = 1$ is excluded because it is not covered by the mathematical treatment, not for algorithmic reasons. Simulation studies indicate that our methodology also can handle the case $c = 1$ in practice; for example, see Section 11.4. The reader is referred to [Ledoit and Wolf \(2015, Section 2.1\)](#) for a detailed description of the asymptotic framework.

For any positive integers n and p , the QuEST function, denoted by $Q_{n,p}$, is the nonrandom multivariate function given by

$$Q_{n,p} : [0, \infty)^p \longrightarrow [0, \infty)^p \tag{3.1}$$

$$\mathbf{t} := (t_1, \dots, t_p)' \longmapsto Q_{n,p}(\mathbf{t}) := (q_{n,p}^1(\mathbf{t}), \dots, q_{n,p}^p(\mathbf{t}))', \tag{3.2}$$

where

$$\forall i = 1, \dots, p \quad q_{n,p}^i(\mathbf{t}) := p \int_{(i-1)/p}^{i/p} (F_{n,p}^{\mathbf{t}})^{-1}(v) dv, \tag{3.3}$$

$$\forall v \in [0, 1] \quad (F_{n,p}^{\mathbf{t}})^{-1}(v) := \sup\{x \in \mathbb{R} : F_{n,p}^{\mathbf{t}}(x) \leq v\}, \tag{3.4}$$

$$\forall x \in \mathbb{R} \quad F_{n,p}^{\mathbf{t}}(x) := \begin{cases} \max\left(1 - \frac{n}{p}, \frac{1}{p} \sum_{i=1}^p \mathbb{1}_{\{t_i=0\}}\right) & \text{if } x = 0, \\ \lim_{\eta \rightarrow 0^+} \frac{1}{\pi} \int_{-\infty}^x \operatorname{Im} [m_{n,p}^{\mathbf{t}}(\xi + i\eta)] d\xi & \text{otherwise,} \end{cases} \tag{3.5}$$

and $\forall z \in \mathbb{C}^+ \quad m := m_{n,p}^{\mathbf{t}}(z)$ is the unique solution in the set

$$\left\{ m \in \mathbb{C} : -\frac{n-p}{nz} + \frac{p}{n} m \in \mathbb{C}^+ \right\} \tag{3.6}$$

to the equation

$$m = \frac{1}{p} \sum_{i=1}^p \frac{1}{t_i (1 - \frac{p}{n} - \frac{p}{n} z m) - z}. \tag{3.7}$$

The basic idea is that p represents the matrix dimension, n the sample size, $\mathbf{t} := (t_1, \dots, t_p)'$ the population eigenvalues, $Q_{n,p}(\mathbf{t}) := (q_{n,p}^1(\mathbf{t}), \dots, q_{n,p}^p(\mathbf{t}))'$ the sample eigenvalues, $F_{n,p}^{\mathbf{t}}$ the estimator of the limiting c.d.f. of the sample eigenvalues, and $m_{n,p}^{\mathbf{t}}$ its Stieltjes transform.

Eqs. (3.3)–(3.4) recuperate the i th smallest sample eigenvalue as the corresponding (smoothed) quantile of the sample spectral distribution $F_{n,p}^{\mathbf{t}}$. Eq. (3.5) is the counterpart of Eqs. (2.1) and (2.3). Eq. (3.6) is a discretization of (2.4), and Eq. (3.7) is a discretization of the Fundamental Equation (2.5). The role of the limiting population spectral distribution H is played by the step function $H_p^{\mathbf{t}}(x) := \sum_{i=1}^p \mathbb{1}_{\{t_i \leq x\}}/p$. The importance of the QuEST function is twofold.

First, inverting it numerically yields an estimator of the population eigenvalues that is consistent under large-dimensional asymptotics. Given the sample eigenvalues $\lambda_{n,1} \leq \lambda_{n,2} \leq \dots \leq \lambda_{n,p}$, we estimate the population eigenvalues $\tau_{n,1} \leq \tau_{n,2} \leq \dots \leq \tau_{n,p}$ by numerically inverting the QuEST function:

$$\hat{\tau}_n := \operatorname{argmin}_{\mathbf{t} \in [0, \infty)^p} \frac{1}{p} \sum_{i=1}^p [q_{n,p}^i(\mathbf{t}) - \lambda_{n,i}]^2.$$

The step function $x \mapsto \sum_{i=1}^p \mathbb{1}_{\{\hat{\tau}_{n,i} \leq x\}}/p$ estimates the object of interest, H , consistently; for further intuition, see Section 11.2.

Second, once this has been achieved, it is possible to use Theorem 2 of [Ledoit and P ech e \(2011\)](#) to construct shrinkage estimators of the covariance matrix that are asymptotically optimal with respect to a given loss function in the p -dimensional space of rotation-equivariant estimators introduced by [Stein \(1975, 1986\)](#). [Ledoit and Wolf \(2017b\)](#) derive the optimal shrinkage formula for five different loss functions, and [Ledoit and Wolf \(forthcoming\)](#) for a sixth.

4. Change of variable

In what follows we omit the subscripts and superscript of $F_{n,p}^{\mathbf{t}}$, $H_p^{\mathbf{t}}$ and c_n in order to simplify the notation. The Fundamental Equation presented in Section 2.1.4 is not convenient for numerical work, since we would need to solve an equation involving complex numbers for every $z \in \mathbb{C}^+$. The first step is to bring everything back to the edge of \mathbb{C}^+ . [Silverstein and Choi \(1995\)](#)

Theorem 1.1) show that $\lim_{z \in \mathbb{C}^+ \rightarrow x} m_F(z) =: \check{m}_F(x)$ exists for all $x \in \mathbb{R} \setminus \{0\}$. For every x in the support of F , which we denote by $\text{Supp}(F)$, $m := \check{m}_F(x)$ is the unique solution in \mathbb{C}^+ to the equation

$$m - \int_{-\infty}^{+\infty} \frac{1}{\tau [1 - c - c x m] - x} dH(\tau) = 0. \tag{4.1}$$

Note that Eq. (4.1) is still inconvenient to solve numerically because we have to zero the real part *and the imaginary part* of the quantity on the left-hand side simultaneously.

There exists a way to rewrite the problem in equivalent form that requires only zeroing a real quantity, which is numerically less demanding. It is detailed in Sections 2.2 and 2.3 of Ledoit and Wolf (2012), and is founded upon the change of variables introduced earlier by Silverstein and Choi (1995). We restate it here succinctly for convenience. We have to work with the modified Stieltjes transform

$$\forall x \in \mathbb{R} \setminus \{0\} \quad \check{m}_F(x) := \frac{c - 1}{x} + c \check{m}_F(x).$$

For every $x \in \text{Supp}(F)$, $m := \check{m}_F(x)$ is the unique solution in \mathbb{C}^+ to the equation

$$m = - \left[x - c \int_{-\infty}^{+\infty} \frac{\tau}{1 + \tau m} dH(\tau) \right]^{-1}. \tag{4.2}$$

Eq. (4.2) appears in slightly different form as equation (1.14) in Marčenko and Pastur (1967). The change of variable $\check{u}_F(x) := -1/\check{m}_F(x)$ enables us to rewrite (4.2) as

$$\forall x \in \text{Supp}(F) \quad x = \check{u}_F(x) - c \check{u}_F(x) \int_{-\infty}^{+\infty} \frac{\tau}{\tau - \check{u}_F(x)} dH(\tau).$$

We now treat $\check{u}_F(x)$ as a variable in $\mathbb{C}^+ \cup \mathbb{R}$, which we call \check{u} , and introduce the function

$$\forall \check{u} \in \mathbb{C}^+ \cup \mathbb{R} \quad \Delta(\check{u}) := \check{u} - c \check{u} \int_{-\infty}^{+\infty} \frac{\tau}{\tau - \check{u}} dH(\tau).$$

For every $x \in \text{Supp}(F)$, there exists some $\check{u} \in \mathbb{C}^+ \cup \mathbb{R}$ such that $\Delta(\check{u}) = x$. Thus, the problem essentially boils down to finding all the \check{u} 's in $\mathbb{C}^+ \cup \mathbb{R}$ such that the imaginary part of $\Delta(\check{u})$ is zero. This is easier than the original formulation (4.1) because now we only have to zero a real quantity, namely, $\text{Im}[\Delta(\check{u})]$.

There is a one-to-one mapping between F -space and u -space, as explained in Ledoit and Wolf (2012, Section 2.2): for every $\check{u} \in \mathbb{C}^+ \cup \mathbb{R}$ such that $\text{Im}[\Delta(\check{u})] = 0$, the limiting sample spectral density evaluated at $x = \Delta(\check{u})$ is equal to

$$\frac{dF}{dx}(x) = \frac{1}{c\pi} \text{Im} \left[-\frac{1}{\check{u}} \right]. \tag{4.3}$$

Remark 4.1. For any given $x \in \text{Supp}(F)$, there exist three mathematically equivalent formulations of the ‘Fundamental Equation’:

$$\begin{aligned} \check{m}_F(x) + \left[x - c \int_{-\infty}^{+\infty} \frac{\tau}{1 + \tau \check{m}_F(x)} dH(\tau) \right]^{-1} &= 0 \\ \check{m}_F(x) - \int_{-\infty}^{+\infty} \frac{1}{\tau [1 - c - c x \check{m}_F(x)] - x} dH(\tau) &= 0 \\ \text{Im} \left[\check{u}_F(x) - c \check{u}_F(x) \int_{-\infty}^{+\infty} \frac{\tau}{\tau - \check{u}_F(x)} dH(\tau) \right] &= 0. \end{aligned} \tag{4.4}$$

The first one is from Marčenko and Pastur (1967, p. 461), and is the most influential for historical reasons because it pioneered this field of research decades before everybody else. The second one is from Silverstein (1995, p. 333), and is the most convenient for definitional purposes because it handles directly the limiting sample spectral distribution. The third one is from Ledoit and Wolf (2012, p. 1030), and is the most convenient for numerical purposes because it zeroes a real number instead of a complex one. We consider all three versions to be *the* ‘Fundamental Equation’. ■

The numerical implementation of the QuEST function consists of a series of six successive operations: (1) finding the support in u -space; (2) choosing a grid that covers the support; (3) solving Eq. (4.4) on the grid; (4) computing the limiting sample spectral density using (4.3); (5) integrating it to obtain the empirical c.d.f. of sample eigenvalues; and (6) interpolating the c.d.f. to compute sample eigenvalues as per Eq. (3.3). Each of these steps is detailed in the following sections.

5. Support in u -space

Decompose the complex variable $\check{u} \in \mathbb{C}^+ \cup \mathbb{R}$ from Section 4 into its real and imaginary parts: $\check{u} := u + iv$, where $u \in \mathbb{R}$ and $v \in [0, +\infty)$. For any given $u \in \mathbb{R}$, there are two and only two mutually exclusive possibilities: either (1) $\text{Im}[\Delta(u + iv)] \neq 0$

for all $v \geq 0$; or (2) there exists a unique $v \geq 0$ such that $\text{Im}[\Delta(u + iv)] = 0$. We call the set of all u 's that fall into the second category the ‘support in u -space’, and denote it by S_U . It maps straight into $\text{Supp}(F)$ through the Δ function.

The motivation for finding S_U is as follows. In theory, we would have to solve Eq. (4.4) for every u in the support. However, doing so is not possible using numerical methods on a digital computer. The best we can do is solve it for a *finite* number of u 's, that is, a grid on the support. To select the grid points, we need to know where the support lies. If, hypothetically, we did not know the location of the support, then some grid points would fall outside, and the corresponding computational expenditures would be wasted.

To determine the boundaries of S_U , we first need to group together the inputs into the QuEST function t_1, \dots, t_p that are equal to one another and, if necessary, discard those that are equal to zero. Let us say that there are K distinct nonzero inputs into the QuEST function $0 < s_1 < \dots < s_K$. We can associate them with their respective weights: if j elements of the vector (t_1, \dots, t_p) are equal to s_k then the corresponding weight is $w_k := j/p$.

5.1. Spectral separation

Now we look for spectral separation between s_k and s_{k+1} ($k = 1, \dots, K - 1$). This is done in two stages. First we run a quick test to see whether we can rule out spectral separation *a priori*. Second, if the test is inconclusive, we carry the full analysis to ascertain whether spectral separation indeed occurs.

5.1.1. Necessary condition for spectral separation

Spectral separation occurs at some $u \in (s_k, s_{k+1})$ if and only if

$$\forall v \in (0, +\infty) \quad \text{Im} \left[\underbrace{(u + iv) - c(u + iv) \sum_{j=1}^K \frac{w_j t_j}{s_j - (u + iv)}}_{\Delta(u+iv)} \right] \neq 0,$$

which is equivalent to

$$\sum_{j=1}^K \frac{w_j s_j^2}{(s_j - u)^2} < \frac{1}{c}. \tag{5.1}$$

Eq. (5.1) holds if and only if the function $x_F(m)$ defined in equation (1.6) of Silverstein and Choi (1995) is strictly increasing at $m = -1/u$. Silverstein and Choi (1995, Section 4) explain how this equivalence enables us to determine the support.

Call $\varphi(u)$ the function on the left-hand side of Eq. (5.1). We can decompose it into

$$\begin{aligned} \varphi(u) &= \theta_k(u) + \psi_k^L(u) + \psi_k^R(u), \\ \text{where } \theta_k(u) &:= \frac{w_k s_k^2}{(s_k - u)^2} + \frac{w_{k+1} s_{k+1}^2}{(s_{k+1} - u)^2}, \\ \psi_k^L(u) &:= \sum_{j=1}^{k-1} \frac{w_j s_j^2}{(s_j - u)^2}, \\ \text{and } \psi_k^R(u) &:= \sum_{j=k+2}^K \frac{w_j s_j^2}{(s_j - u)^2}. \end{aligned}$$

It is easy to see that the function $\theta_k(\cdot)$ is convex over the interval (s_k, s_{k+1}) , diverges to $+\infty$ near s_k and s_{k+1} , and attains its minimum at

$$\widehat{x}_k := (s_k s_{k+1})^{2/3} \frac{w_k^{1/3} s_{k+1}^{1/3} + w_{k+1}^{1/3} s_k^{1/3}}{w_k^{1/3} s_k^{2/3} + w_{k+1}^{1/3} s_{k+1}^{2/3}}; \tag{5.2}$$

therefore, a lower bound for $\theta_k(\cdot)$ on (s_k, s_{k+1}) is $\theta_k(\widehat{x}_k)$.

It is also easy to see that the function $\psi_k^L(\cdot)$ is decreasing over the interval (s_k, s_{k+1}) ; therefore, it attains its minimum at s_{k+1} and is bounded from below by $\psi_k^L(s_{k+1})$. Conversely, the function $\psi_k^R(\cdot)$ is increasing over the interval (s_k, s_{k+1}) , attains its minimum at s_k and is bounded from below by $\psi_k^R(s_k)$. Putting these three results together yields the following lower bound for $\varphi(\cdot)$:

$$\forall u \in (s_k, s_{k+1}) \quad \varphi(u) \geq \frac{w_k s_k^2}{(s_k - \widehat{x}_k)^2} + \frac{w_{k+1} s_{k+1}^2}{(s_{k+1} - \widehat{x}_k)^2} + \sum_{j=1}^{k-1} \frac{w_j s_j^2}{(s_j - s_{k+1})^2} + \sum_{j=k+2}^K \frac{w_j s_j^2}{(s_j - s_k)^2},$$

where \widehat{x}_k is given by Eq. (5.2).

Combining this bound with Eq. (5.1) means that

$$\frac{w_k s_k^2}{(s_k - \widehat{x}_k)^2} + \frac{w_{k+1} s_{k+1}^2}{(s_{k+1} - \widehat{x}_k)^2} + \sum_{j=1}^{k-1} \frac{w_j s_j^2}{(s_j - s_{k+1})^2} + \sum_{j=k+2}^K \frac{w_j s_j^2}{(s_j - s_k)^2} < \frac{1}{c} \tag{5.3}$$

is a necessary (but not sufficient) condition for spectral separation to occur between s_k and s_{k+1} , that is, at some $u \in (s_k, s_{k+1})$. Thus, the numerical procedure can be made more efficient by first computing the quantity on the left-hand side of Eq. (5.3), comparing it with $1/c$, and asserting the absence of spectral separation in the interval (s_k, s_{k+1}) in the case where it is higher than $1/c$. If, on the other hand, it is strictly lower than $1/c$, then further work is needed to ascertain whether spectral separation does indeed occur. In practice, checking this condition seems to save a lot of time by eliminating many intervals (s_k, s_{k+1}) , except perhaps when c is very small and the population eigenvalues are very spread out.

5.1.2. Necessary and sufficient condition for spectral separation

Consider now some $k \in \{1, 2, \dots, K - 1\}$ for which the condition in Eq. (5.3) holds. Given Eq. (5.1), we need to find the minimum of $\varphi(\cdot)$ over (s_k, s_{k+1}) and compare it with $1/c$. It is easy to check that $\varphi(\cdot)$ is strictly convex over (s_k, s_{k+1}) ; therefore, this minimum exists, is unique, and is the only zero in (s_k, s_{k+1}) of the derivative function

$$\varphi'(u) = 2 \sum_{j=1}^K \frac{w_j s_j^2}{(s_j - u)^3}.$$

Most numerical algorithms that find the zero of a function require as inputs two points \underline{x} and \bar{x} such that the sign of the function is different at \underline{x} and at \bar{x} . Finding two such points is the goal of the next step. There are three cases, depending on the sign of $\varphi'(\widehat{x}_k)$:

- $\varphi'(\widehat{x}_k) = 0$: Then the search is immediately over because $\varphi(\cdot)$ attains its minimum at $x_k^* := \widehat{x}_k$. This would not happen generically unless $K = 2$.
- $\varphi'(\widehat{x}_k) < 0$: In this case, given that $\varphi'(\cdot)$ is strictly increasing, the minimizer of $\varphi(\cdot)$ lies in the interval (\widehat{x}_k, s_{k+1}) . We can feed the lower bound $\underline{x} := \widehat{x}_k$ into the numerical procedure that will find the zero of $\varphi'(\cdot)$. It would be also tempting to set $\bar{x} := s_{k+1}$, but unfortunately doing so would not be practicable because $\lim_{u \nearrow s_{k+1}} \varphi'(u) = +\infty$, and most numerical procedures perform poorly near singularity points. Therefore, we need to find some $\bar{x} \in (\widehat{x}_k, s_{k+1})$ such that $\varphi'(\bar{x}) > 0$. Let x_k^* denote the unique value in (\widehat{x}_k, s_{k+1}) such that $\varphi'(x_k^*) = 0$. Then the fact that $w_j s_j^2 / (s_j - u)^3$ is increasing in u for any $j \in \{1, \dots, K\}$ implies that the following inequalities hold:

$$\begin{aligned} \forall u \in (\widehat{x}_k, s_{k+1}) \quad \varphi'(u) &> 2 \frac{w_k s_k^2}{(s_k - \widehat{x}_k)^3} + 2 \frac{w_{k+1} s_{k+1}^2}{(s_{k+1} - u)^3} + \psi_k^{L'}(\widehat{x}_k) + \psi_k^{R'}(\widehat{x}_k) \\ 0 &> 2 \frac{w_k s_k^2}{(s_k - \widehat{x}_k)^3} + 2 \frac{w_{k+1} s_{k+1}^2}{(s_{k+1} - x_k^*)^3} + \psi_k^{L'}(\widehat{x}_k) + \psi_k^{R'}(\widehat{x}_k) \\ -2 \frac{w_k s_k^2}{(s_k - \widehat{x}_k)^3} - \psi_k^{L'}(\widehat{x}_k) - \psi_k^{R'}(\widehat{x}_k) &> 2 \frac{w_{k+1} s_{k+1}^2}{(s_{k+1} - x_k^*)^3} \\ s_{k+1} - x_k^* &> \left(\frac{2w_{k+1} s_{k+1}^2}{-2 \frac{w_k s_k^2}{(s_k - \widehat{x}_k)^3} - \psi_k^{L'}(\widehat{x}_k) - \psi_k^{R'}(\widehat{x}_k)} \right)^{1/3} \\ x_k^* &< s_{k+1} - \left(\frac{2w_{k+1} s_{k+1}^2}{-2 \frac{w_k s_k^2}{(s_k - \widehat{x}_k)^3} - \psi_k^{L'}(\widehat{x}_k) - \psi_k^{R'}(\widehat{x}_k)} \right)^{1/3} \\ x_k^* &< s_{k+1} - \left(\frac{2w_{k+1} s_{k+1}^2}{-2 \frac{w_k s_k^2}{(s_k - \widehat{x}_k)^3} - \varphi'(\widehat{x}_k)} \right)^{1/3}, \end{aligned}$$

where the last inequality follows from $\theta_k'(\widehat{x}_k) = 0$. Thus, if we set

$$\bar{x} := s_{k+1} - \left(\frac{2w_{k+1} s_{k+1}^2}{-2 \frac{w_k s_k^2}{(s_k - \widehat{x}_k)^3} - \varphi'(\widehat{x}_k)} \right)^{1/3},$$

we know that $\varphi'(\bar{x}) > 0$. Launching a zero-finding algorithm for $\varphi'(\cdot)$ on the interval $[\underline{x}, \bar{x}]$ as defined above yields a unique solution x_k^* .

- $\varphi'(\widehat{x}_k) < 0$: A similar line of reasoning points us to

$$\underline{x} := s_k + \left(\frac{2w_k s_k^2}{2 \frac{w_{k+1} s_{k+1}^2}{(s_{k+1} - \widehat{x}_k)^3} + \varphi'(\widehat{x}_k)} \right)^{1/3},$$

$\bar{x} := \widehat{x}_k$, and yields a unique zero x_k^* for $\varphi(\cdot)$ over the interval $[\underline{x}, \bar{x}]$.

Across all three cases, the outcome of this procedure is $x_k^* := \operatorname{argmin}_{u \in (s_k, s_{k+1})} \varphi(u)$. Spectral separation occurs between s_k and s_{k+1} if and only if $\varphi(x_k^*) < 1/c$.

If there is no spectral separation, then we can dismiss the interval (s_k, s_{k+1}) ; otherwise, we need some additional work to compute spectrum boundaries.

5.1.3. Interval boundaries

Consider now some $k \in \{1, 2, \dots, K - 1\}$ for which $x_k^* := \operatorname{argmin}_{u \in (s_k, s_{k+1})} \varphi(u)$ is known and $\varphi(x_k^*) < 1/c$. Spectral separation means that the support ends at some point in (s_k, x_k^*) and starts again at some point in (x_k^*, s_{k+1}) . The equation that characterizes support endpoints is $\varphi(x) = 1/c$. Thus we need to find two zeros of the function $\varphi(\cdot) - 1/c$, one in the interval (s_k, x_k^*) and the other in the interval (x_k^*, s_{k+1}) .

Let us start with the first zero of the function $\varphi(\cdot) - 1/c$, the one that lies in the interval (s_k, x_k^*) . Once again, we employ an off-the-shelf univariate zero-finding routine that takes as inputs two points \underline{x} and \bar{x} such that $\varphi(\underline{x}) > 1/c$ and $\varphi(\bar{x}) < 1/c$. The obvious candidate for \bar{x} is $\bar{x} := x_k^*$. For \underline{x} , however, we cannot use s_k because $\lim_{x \searrow s_k} \varphi(x) = +\infty$. Therefore, we need to find some $\underline{x} \in (s_k, x_k^*)$ that verifies $\varphi(\underline{x}) > 1/c$. Such an \underline{x} can be found by considering the following series of inequalities, which hold for all $x \in (s_k, x_k^*)$:

$$\begin{aligned} \varphi(x) &> \frac{w_k s_k^2}{(s_k - x)^2} + \sum_{j=1}^{k-1} \frac{w_j s_j^2}{(s_j - x_k^*)^2} + \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - s_k)^2} \\ \varphi(x) - \varphi(x_k^*) &> \frac{w_k s_k^2}{(s_k - x)^2} - \frac{w_k s_k^2}{(s_k - x_k^*)^2} + \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - s_k)^2} - \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - x_k^*)^2} \\ \varphi(x) - \frac{1}{c} &> \frac{w_k s_k^2}{(s_k - x)^2} - \frac{w_k s_k^2}{(s_k - x_k^*)^2} + \left[\varphi(x_k^*) - \frac{1}{c} \right] + \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - s_k)^2} - \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - x_k^*)^2}. \end{aligned}$$

Call the function on the right-hand side of the last inequality $g_k(x)$. This function is defined on $(s_k, +\infty)$ and is of the form

$$g_k(x) = \frac{w_k s_k^2}{(s_k - x)^2} + C_k,$$

where C_k is a negative constant. $g_k(\cdot)$ is strictly decreasing, with $\lim_{x \searrow s_k} g_k(x) = +\infty$, and $\lim_{x \nearrow +\infty} g_k(x) = C_k < 0$. Therefore, it admits a unique zero in $(s_k, +\infty)$, given by

$$\underline{x} := s_k + \sqrt{\frac{w_k s_k^2}{\frac{w_k s_k^2}{(s_k - x_k^*)^2} + \left[\frac{1}{c} - \varphi(x_k^*) \right] + \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - x_k^*)^2} - \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - s_k)^2}}.}$$

Notice that

$$g_k(x_k^*) = \left[\varphi(x_k^*) - \frac{1}{c} \right] + \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - s_k)^2} - \sum_{j=k+1}^K \frac{w_j s_j^2}{(s_j - x_k^*)^2} < 0.$$

Combining this fact with the fact that $g_k(\cdot)$ is strictly decreasing implies that $\underline{x} < x_k^*$.

Since $g_k(\underline{x}) = 0$ by construction, $\varphi(\underline{x}) > 1/c$. Feeding (\underline{x}, \bar{x}) thus defined into the zero-finding numerical routine with the function $\varphi(\cdot) - 1/c$ yields an endpoint of the support.

A similar line of reasoning leads to setting $\bar{x} := x_k^*$,

$$\bar{x} := s_{k+1} - \sqrt{\frac{w_{k+1} s_{k+1}^2}{\frac{w_{k+1} s_{k+1}^2}{(s_{k+1} - x_k^*)^2} + \left[\frac{1}{c} - \varphi(x_k^*) \right] + \sum_{j=1}^{k-1} \frac{w_j s_j^2}{(s_j - x_k^*)^2} - \sum_{j=1}^{k-1} \frac{w_j s_j^2}{(s_j - s_{k+1})^2}}.}$$

and running a numerical routine to find a zero of the function $\varphi(\cdot) - 1/c$ on the interval $(\underline{x}, \bar{x}) \subset (x_k^*, s_{k+1})$. This zero will also be a support endpoint.

5.2. Extremities of the support

The procedure described so far identifies all support endpoints lying in the interval $[s_1, s_K]$. In order to complete the determination of the support, we must find the support endpoint that lies in the interval $(-\infty, s_1)$ and the support endpoint that lies in the interval $(s_K, +\infty)$.

5.2.1. Minimum of the support

Let us start with the first support endpoint, the one lying in the interval $(-\infty, s_1)$. The equation that characterizes this point is the same as before: $\varphi(x) = 1/c$. In order to employ the zero-finding numerical routine, we must find two bounds \underline{x} and \bar{x} , both strictly less than s_1 , such that $\varphi(\underline{x}) < 1/c$ and $\varphi(\bar{x}) > 1/c$. The left-hand side bound \underline{x} can be obtained by considering the following inequalities:

$$\forall x \in (-\infty, s_1) \quad \forall j = 1, \dots, K \quad \frac{w_j s_j^2}{(x - s_j)^2} \leq \frac{w_j s_j^2}{(x - s_1)^2}$$

$$\forall x \in (-\infty, s_1) \quad \varphi(x) \leq \frac{\sum_{j=1}^K w_j s_j^2}{(x - s_1)^2}. \tag{5.4}$$

Note that if we set

$$\underline{x} := s_1 - \sqrt{c \sum_{j=1}^K w_j s_j^2} - 1,$$

then

$$\frac{\sum_{j=1}^K w_j s_j^2}{(\underline{x} - s_1)^2} < \frac{1}{c},$$

which in turn implies by Eq. (5.4) that $\varphi(\underline{x}) < 1/c$, as desired.

The right-hand side bound \bar{x} can be found by considering a different inequality:

$$\forall x \in (-\infty, s_1) \quad \varphi(x) \geq \frac{w_1 s_1^2}{(x - s_1)^2}. \tag{5.5}$$

Note that if we set

$$\bar{x} := s_1 - \frac{\sqrt{c w_1 s_1^2}}{2},$$

then

$$\frac{w_1 s_1^2}{(\bar{x} - s_1)^2} > \frac{1}{c},$$

which in turn implies by Eq. (5.5) that $\varphi(\bar{x}) > 1/c$, as desired. Launching the numerical routine to find a zero of the function $\varphi(\cdot) - 1/c$ over the interval (\underline{x}, \bar{x}) thus defined yields the first endpoint of the support.

5.2.2. Maximum of the support

For the last endpoint of the support, the one that lies in the interval $(s_K, +\infty)$, a similar line of reasoning leads us to define

$$\underline{x} := s_K + \frac{\sqrt{c w_K s_K^2}}{2} \quad \text{and} \quad \bar{x} := s_K + \sqrt{c \sum_{j=1}^K w_j s_j^2} + 1.$$

Launching the numerical routine to find a zero of the function $\varphi(\cdot) - 1/c$ over the interval (\underline{x}, \bar{x}) thus defined yields the last endpoint of the support.

5.3. Output

The main outputs of this procedure are $\nu \geq 1$, the number of distinct intervals that constitute the support, and $u_1, \dots, u_{2\nu}$, the support endpoints. The support in u -space is $S_U = [u_1, u_2] \cup \dots \cup [u_{2\nu-1}, u_{2\nu}]$.

Another output of this procedure is a set of positive integers $\omega_1, \dots, \omega_\nu$ summing up to p that tell us how many population eigenvalues correspond to each support interval. If $\nu = 1$, then there is no spectral separation and $\omega_1 = p$. If $\nu \geq 2$ and the

first spectral separation occurs between s_k and s_{k+1} for some $k = 1, \dots, K - 1$, then $\omega_1 = p \sum_{j=1}^k w_j$. If some population eigenvalues are equal to zero, then ω_1 needs to be augmented accordingly; see the discussion below Lemma 4 of [Bai and Silverstein \(1999\)](#) for details.

If $\nu \geq 2$ and the last spectral separation occurs between $s_{k'}$ and $s_{k'+1}$ for some $k' = 1, \dots, K - 1$, then $\omega_\nu = p \sum_{j=k'+1}^K w_j$. If $\nu \geq 3$ and the i th support interval (for $i = 2, \dots, \nu - 1$) is delimited on the left-hand side by spectral separation occurring between s_k and s_{k+1} , and on the right-hand side by spectral separation occurring between $s_{k'}$ and $s_{k'+1}$ (where $1 \leq k < k' \leq K$), then $\omega_i = p \sum_{j=k+1}^{k'} w_j$. This information will turn out to be useful in subsequent operations.

5.4. Derivative of the support endpoints

If the QuEST function defined by Eqs. (3.1)–(3.7) is to be used efficiently in an optimization algorithm, it is desirable to be able to compute its derivative analytically. Since this function is constructed as a chain of six successive operations, the first of which is the determination of support endpoints, its derivative can be computed in the same way, provided that we start by computing analytically the derivative of support endpoints with respect to t_k for all $k = 1, \dots, K$.

Every u_i for $i = 1, \dots, 2\nu$ is a zero of the function

$$\tilde{\varphi}(u; t_1, \dots, t_p) := \frac{1}{p} \sum_{j=1}^p \frac{t_j^2}{(t_j - u)^2} - \frac{1}{c}.$$

By differentiating the equation $\tilde{\varphi}(u; t_1, \dots, t_p) = 0$ we get:

$$\begin{aligned} \frac{\partial \tilde{\varphi}}{\partial u} \cdot du + \frac{\partial \tilde{\varphi}}{\partial t_k} \cdot dt_k &= 0, \\ \frac{\partial u}{\partial t_k} &= - \frac{\frac{\partial \tilde{\varphi}}{\partial t_k}}{\frac{\partial \tilde{\varphi}}{\partial u}}. \end{aligned}$$

The partial derivatives of the function $\tilde{\varphi}(\cdot)$ are as follows:

$$\begin{aligned} \frac{\partial \tilde{\varphi}}{\partial u}(u; t_1, \dots, t_p) &= \frac{2}{p} \sum_{j=1}^p \frac{t_j^2}{(t_j - u)^3} \\ \frac{\partial \tilde{\varphi}}{\partial t_k}(u; t_1, \dots, t_p) &= - \frac{2}{p} \frac{t_k u}{(t_k - u)^3}; \end{aligned}$$

therefore,

$$\forall i = 1, \dots, 2\nu \quad \forall k = 1, \dots, p \quad \frac{\partial u_i}{\partial t_k} = \frac{\frac{t_k u_i}{(t_k - u_i)^3}}{\sum_{j=1}^p \frac{t_j^2}{(t_j - u_i)^3}}. \tag{5.6}$$

6. Grid

The first operation generated the support in u -space $S_U = [u_1, u_2] \cup \dots \cup [u_{2\nu-1}, u_{2\nu}]$ and the number of population eigenvalues corresponding to each interval: $\omega_1, \dots, \omega_\nu$. The goal of the second operation is to produce a grid that covers this support. This problem can be broken down by considering each interval $i = 1, \dots, \nu$ separately.

6.1. Formula for the grid points

Take some $i \in \{1, \dots, \nu\}$. How shall we determine a grid that covers the interval $[u_{2i-1}, u_{2i}]$? The number of points on the grid will be a function of ω_i . Specifically, we shall take ω_i points in the open interval (u_{2i-1}, u_{2i}) , plus the two endpoints u_{2i-1} and u_{2i} . Thus, the total number of points covering the closed interval $[u_{2i-1}, u_{2i}]$ will be $\omega_i + 2$. Let us call these points $\xi_0^i, \dots, \xi_{\omega_i+1}^i$, with the convention that $\xi_0^i := u_{2i-1}$ and $\xi_{\omega_i+1}^i := u_{2i}$. Thus, what is left is to define $\xi_1^i, \dots, \xi_{\omega_i}^i$.

There are many ways to choose such a grid, depending on how densely we want to cover the various parts of the interval. The simplest idea would be to have uniform coverage through a linearly spaced grid. But it is more judicious to increase coverage density near the edges of the interval because this is where a lot of the action is taking place. [Silverstein and Choi \(1995\)](#) demonstrate that the limiting density of sample eigenvalues has ‘square root’-type behavior near boundary points. This fact points us towards the inverse c.d.f. function of the beta distribution with parameters (0.5, 0.5), also known as the arcsine distribution:

$$\forall j \in \{0, \dots, \omega_i + 1\} \quad \xi_j^i := u_{2i-1} + (u_{2i} - u_{2i-1}) \sin^2 \left[\frac{\pi j}{2(\omega_i + 1)} \right]. \tag{6.1}$$

Compared to the beta distribution with parameters (1, 1), which is the uniform distribution on the interval [0, 1], reducing both parameters from 1 to 0.5 increases coverage density near the edges of the interval. Note that the density of the arcsine distribution goes to infinity at the edges of the interval (as does the derivative of the square root function), but the c.d.f., its inverse, and the grid all remain well-behaved. The goal here is to enhance numerical accuracy.

6.2. Derivative of the grid points

In keeping with our earlier stated objective (see Section 5.4) of building towards an analytical formula for the partial derivative of λ_i with respect to t_k for all $i, k \in \{1, \dots, p\}$, at this stage we need to compute $\partial \xi_j^i / \partial t_k$ for all $j \in \{1, \dots, \omega_i\}$. From Eq. (6.1), one can see immediately that it is

$$\frac{\partial \xi_j^i}{\partial t_k} = \left\{ 1 - \sin^2 \left[\frac{\pi j}{2(\omega_i + 1)} \right] \right\} \frac{\partial u_{2i-1}}{\partial t_k} + \sin^2 \left[\frac{\pi j}{2(\omega_i + 1)} \right] \frac{\partial u_{2i}}{\partial t_k}, \tag{6.2}$$

where $\partial u_{2i-1} / \partial t_k$ and $\partial u_{2i} / \partial t_k$ are given by Eq. (5.6).

7. Solving the fundamental equation in u -space

In this section, we will assume that the interval index $i \in \{1, \dots, \nu\}$ is fixed.

7.1. Statement of the problem

Given a grid coverage $(\xi_j^i)_{j=0, \dots, \omega_i}$ of the i th support interval, the third operation solves the Fundamental Equation at ξ_j^i . For every $j = 0, \dots, \omega_i + 1$, define the function

$$\forall y \in [0, +\infty) \quad \Gamma_j^i(y) := \frac{1}{p} \sum_{k=1}^p \frac{t_k^2}{(t_k - \xi_j^i)^2 + y^2} - \frac{1}{c}.$$

Since by definition of the grid ξ_j^i lies in the support S_U , Eq. (5.1) implies that $\Gamma_j^i(0) \geq 0$. In addition, it is easy to verify that $\Gamma_j^i(\cdot)$ is strictly decreasing on $[0, +\infty)$ and that $\lim_{y \rightarrow +\infty} \Gamma_j^i(y) = -1/c$.

The solution to the Fundamental Equation at ξ_j^i is the thus unique $y \in [0, +\infty)$ such that

$$\Gamma_j^i(y) = 0. \tag{7.1}$$

Call this solution y_j^i . This line of attack is directly inspired by Ledoit and Wolf (2012, Section 2.3).

From the definition of the ξ_j^i 's in Section 6.1, it is obvious that $y_0^i = y_{\omega_i+1}^i = 0$. What remains to be determined is $(y_j^i)_{j=1, \dots, \omega_i}$. In the remainder of this section, we will assume that j is fixed in the set $\{1, \dots, \omega_i\}$.

The solution y to the equation $\Gamma_j^i(y) = 0$ is computed by some standard numerical routine that finds the zero of a real univariate function. As usual, we need to input into this routine a lower bound $\underline{y}_j^i \in [0, +\infty)$ such that $\Gamma_j^i(\underline{y}_j^i) > 0$ and an upper bound $\bar{y}_j^i \in (0, +\infty)$ such that $\Gamma_j^i(\bar{y}_j^i) < 0$.

7.2. Lower bound

From Section 5, (s_1, \dots, s_K) is the vector of unique nonzero population eigenvalues, with corresponding weights (w_1, \dots, w_K) . Let $\delta_j^i := \min_{k \in \{1, \dots, K\}} (s_k - \xi_j^i)^2$ and $\Omega_j^i := \{k \in \{1, \dots, K\} : (s_k - \xi_j^i)^2 = \delta_j^i\}$. Then we have

$$\Gamma_j^i(y) \geq \frac{\sum_{k \in \Omega_j^i} w_k s_k^2}{\delta_j^i + y^2} - \frac{1}{c}. \tag{7.2}$$

Looking at the right-hand side of Eq. (7.2), one can see that

$$\frac{\sum_{k \in \Omega_j^i} w_k s_k^2}{\delta_j^i + y^2} - \frac{1}{c} \geq 0 \iff y^2 \leq c \sum_{k \in \Omega_j^i} w_k s_k^2 - \delta_j^i.$$

Therefore, if we set

$$y_j^i := \sqrt{\frac{\max\left(0, c \sum_{k \in \Omega_j^i} w_k s_k^2 - \delta_j^i\right)}{2}},$$

then $\Gamma(y_j^i) > 0$, as desired.

7.3. Upper bound

We use the inequalities

$$\begin{aligned} \forall k \in \{1, \dots, K\} \quad \frac{1}{(s_k - \xi_j^i)^2 + y^2} &\leq \frac{1}{\delta_j^i + y^2} \\ \Gamma_j^i(y) &\leq \frac{\sum_{k=1}^K w_k s_k^2}{\delta_j^i + y^2} - \frac{1}{c}. \end{aligned} \tag{7.3}$$

Note that if we set

$$\bar{y}_j^i := \sqrt{c \sum_{k=1}^K w_k s_k^2 - \delta_j^i + 1},$$

then

$$\frac{\sum_{k=1}^K w_k s_k^2}{\delta_j^i + (\bar{y}_j^i)^2} - \frac{1}{c} < 0;$$

therefore, by Eq. (7.3), $\Gamma_j^i(\bar{y}_j^i) < 0$, as desired.

7.4. Output

Launching a standard numerical routine to find the zero of the function $\Gamma_j^i(\cdot)$ over the interval (y_j^i, \bar{y}_j^i) yields y_j^i , the solution to the Fundamental Equation at ξ_j^i . The output of this operation is more conveniently expressed as the complex number $z_j^i := \xi_j^i + \sqrt{-1}y_j^i$.

7.5. Derivative

The derivative of the real part of z_j^i with respect to t_k has been computed in Section 6.2. As for the derivative of the imaginary part, y_j^i , consider the function

$$\tilde{\Gamma}_j^i(y; t_1, \dots, t_p) := \frac{1}{p} \sum_{k=1}^p \frac{t_k^2}{(t_k - \xi_j^i)^2 + y^2} - \frac{1}{c}.$$

One can view y_j^i as a function of (t_1, \dots, t_p) : $y_j^i = \tilde{y}_j^i(t_1, \dots, t_p)$. Then the manner in which y_j^i is obtained in Section 7.1 can be expressed through the equation

$$\tilde{\Gamma}_j^i(\tilde{y}_j^i(t_1, \dots, t_p); t_1, \dots, t_p) = 0.$$

Taking the partial derivative with respect to t_k while holding the other population eigenvalues constant yields

$$\begin{aligned} \frac{\partial \tilde{\Gamma}_j^i}{\partial y} \cdot \frac{\partial \tilde{y}_j^i}{\partial t_k} + \frac{\partial \tilde{\Gamma}_j^i}{\partial t_k} &= 0, \\ \frac{\partial \tilde{y}_j^i}{\partial t_k} &= -\frac{\frac{\partial \tilde{\Gamma}_j^i}{\partial t_k}}{\frac{\partial \tilde{\Gamma}_j^i}{\partial y}}. \end{aligned}$$

The partial derivatives of the function \tilde{F}_j^i are

$$\frac{\partial \tilde{F}_j^i}{\partial t_k}(y; t_1, \dots, t_p) = \frac{2t_k}{(t_k - \xi_j^i)^2 + y^2} - \frac{2t_k^2(t_k - \xi_j^i)}{[(t_k - \xi_j^i)^2 + y^2]^2}$$

$$\frac{\partial \tilde{F}_j^i}{\partial y}(y; t_1, \dots, t_p) = -2 \sum_{l=1}^p \frac{t_l^2 y}{[(t_l - \xi_j^i)^2 + y^2]^2}.$$

Therefore,

$$\frac{\partial \tilde{y}_j^i}{\partial t_k}(t_1, \dots, t_p) = \frac{\frac{t_k}{(t_k - \xi_j^i)^2 + (y_j^i)^2} - \frac{t_k^2(t_k - \xi_j^i)}{[(t_k - \xi_j^i)^2 + (y_j^i)^2]^2}}{\sum_{l=1}^p \frac{t_l^2 y_j^i}{[(t_l - \xi_j^i)^2 + (y_j^i)^2]^2}}. \tag{7.4}$$

Now this is only part of the answer because in this analysis we held ξ_j^i constant, whereas in reality it is also a function of the population eigenvalues. Thus, the partial derivative of y_j^i with respect to t_k is given by the formula

$$\frac{\partial y_j^i}{\partial t_k} = \frac{\partial \tilde{y}_j^i}{\partial t_k} + \frac{\partial y_j^i}{\partial \xi_j^i} \cdot \frac{\partial \xi_j^i}{\partial t_k}, \tag{7.5}$$

where $\partial \tilde{y}_j^i / \partial t_k$ is given by Eq. (7.4) and $\partial \xi_j^i / \partial t_k$ is given by Eq. (6.2). All that remains to be computed is $\partial y_j^i / \partial \xi_j^i$. This is done by temporarily ignoring the direct dependency on the population eigenvalues and setting up the function

$$\hat{\Gamma}(y; \xi) := \frac{1}{p} \sum_{l=1}^p \frac{t_l^2}{(t_l - \xi)^2 + y^2} - \frac{1}{c}.$$

Differentiating the equation $\hat{\Gamma}(y; \xi) = 0$ yields:

$$\frac{\partial \hat{\Gamma}}{\partial y} dy + \frac{\partial \hat{\Gamma}}{\partial \xi} d\xi = 0 \implies \frac{\partial y}{\partial \xi} = -\frac{\frac{\partial \hat{\Gamma}}{\partial \xi}}{\frac{\partial \hat{\Gamma}}{\partial y}}.$$

The partial derivatives of the function $\hat{\Gamma}$ are

$$\frac{\partial \hat{\Gamma}}{\partial \xi}(y; \xi) = 2 \sum_{l=1}^p \frac{t_l^2(t_l - \xi)}{[(t_l - \xi)^2 + y^2]^2}$$

and

$$\frac{\partial \hat{\Gamma}}{\partial y}(y; \xi) = -2 \sum_{l=1}^p \frac{t_l^2 y}{[(t_l - \xi)^2 + y^2]^2};$$

therefore,

$$\frac{\partial y_j^i}{\partial \xi_j^i} = \frac{\sum_{l=1}^p \frac{t_l^2(t_l - \xi_j^i)}{[(t_l - \xi_j^i)^2 + (y_j^i)^2]^2}}{\sum_{l=1}^p \frac{t_l^2 y_j^i}{[(t_l - \xi_j^i)^2 + (y_j^i)^2]^2}}.$$

Plugging this formula into Eq. (7.5) yields the partial derivative of y_j^i with respect to t_k .

8. Density of the limiting distribution of the sample eigenvalues

8.1. Mapping

This is the operation where we leave u -space and map back to $(x, F(x))$, where F is the limiting distribution of sample eigenvalues. The underlying mathematics for this mapping can be found in equations (2.7)–(2.8) of Ledoit and Wolf (2012). The mapping can be expressed with the notation of the present paper as

$$x = \Delta(\check{u}) = \check{u} - c \check{u} \frac{1}{p} \sum_{k=1}^p \frac{t_k}{t_k - \check{u}}.$$

In the remainder of this section, we will assume that the interval index $i \in \{1, \dots, \nu\}$ is fixed. For every $j \in \{0, 1, \dots, \omega_i + 1\}$, map z_j^i into

$$x_j^i = z_j^i - c z_j^i \frac{1}{p} \sum_{k=1}^p \frac{t_k}{t_k - z_j^i}. \tag{8.1}$$

Even though z_j^i is generally a complex number, Eq. (7.1) guarantees that x_j^i is real.

Using Eq. (4.3), we can also obtain the value of the limiting sample spectral density F' evaluated at x_j^i as $F'(x_j^i) =: f_j^i$ where

$$f_j^i = \frac{1}{c\pi} \operatorname{Im} \left[-\frac{1}{z_j^i} \right] = \frac{1}{c\pi} \frac{y_j^i}{(\xi_j^i)^2 + (y_j^i)^2}. \tag{8.2}$$

Note that $f_1^i = f_{\omega_i+1}^i = 0$.

The output of this operation is $(x_j^i, f_j^i)_{j=0,1,\dots,\omega_i+1}$, for every $i \in \{1, \dots, \nu\}$.

8.2. Derivative

From Eq. (8.2), it is easy to compute the partial derivative of f_j^i with respect to τ_k as

$$\frac{\partial f_j^i}{\partial t_k} = \frac{1}{c\pi} \operatorname{Im} \left[\frac{\partial z_j^i}{\partial t_k} \cdot \frac{1}{(z_j^i)^2} \right], \tag{8.3}$$

where
$$\frac{\partial z_j^i}{\partial t_k} = \frac{\partial \xi_j^i}{\partial t_k} + \sqrt{-1} \frac{\partial y_j^i}{\partial t_k}, \tag{8.4}$$

$\partial \xi_j^i / \partial \tau_k$ is given by Eq. (6.2), and $\partial y_j^i / \partial \tau_k$ is given by Eq. (7.5).

In order to differentiate Eq. (8.1) more easily, introduce the function $m_{LH}(\cdot)$ defined as per Ledoit and Wolf (2012, Section 2.2):

$$\forall z \in \mathbb{C}^+ \quad m_{LH}(z; t_1, \dots, t_p) := \frac{1}{p} \sum_{l=1}^p \frac{t_l}{t_l - z} = 1 + z \frac{1}{p} \sum_{l=1}^p \frac{1}{t_l - z}.$$

This enables us to rewrite Eq. (8.1) as

$$x_j^i = z_j^i - c z_j^i m_{LH}(z_j^i; t_1, \dots, t_p). \tag{8.5}$$

The full derivative of $m_{LH}(z_j^i; t_1, \dots, t_p)$ with respect to t_k is

$$\frac{dm_{LH}}{dt_k} = \frac{\partial m_{LH}}{\partial t_k} + \frac{\partial m_{LH}}{\partial z_j^i} \cdot \frac{\partial z_j^i}{\partial t_k},$$

where the last term is given by Eq. (8.4). The partial derivatives of m_{LH} are

$$\frac{\partial m_{LH}}{\partial t_k} = -z_j^i \times \frac{1}{p} \frac{1}{(t_k - z_j^i)^2}$$

and
$$\frac{\partial m_{LH}}{\partial z_j^i} = \frac{1}{p} \sum_{l=1}^p \frac{t_l}{(t_l - z_j^i)^2};$$

therefore,

$$\frac{dm_{LH}}{dt_k} = -z_j^i \times \frac{1}{p} \frac{1}{(t_k - z_j^i)^2} + \frac{\partial z_j^i}{\partial t_k} \times \frac{1}{p} \sum_{l=1}^p \frac{t_l}{(t_l - z_j^i)^2}.$$

Finally, differentiating Eq. (8.5) enables us to compute the partial derivative of x_j^i with respect to t_k as follows:

$$\frac{\partial x_j^i}{\partial t_k} = \frac{\partial z_j^i}{\partial t_k} \times [1 - c \cdot m_{LH}(z_j^i; t_1, \dots, t_p)] - c \cdot z_j^i \frac{dm_{LH}}{dt_k}(z_j^i; t_1, \dots, t_p). \tag{8.6}$$

9. Cumulative distribution function

9.1. Numerical integration of the density

The objective is to compute $F_j^i := F(x_j^i)$. It holds that

$$F(0) = F_0^1 = \max\left(0, 1 - \frac{1}{c}\right). \tag{9.1}$$

Since the support of F is $\cup_{i=1}^v [x_0^i, x_{\omega_i+1}^i]$, with the possible addition of $\{0\}$ if $p > n$, as soon as v is greater than or equal to two, $F_0^{i+1} = F_{\omega_i+1}^i$, for $i = 1, \dots, v - 1$. Bai and Silverstein (1999) show that

$$\forall i = 1, \dots, v \quad F_{\omega_i+1}^i = \frac{1}{p} \sum_{j=1}^i \omega_j. \tag{9.2}$$

All that remains is to compute F_j^i for $j \in \{1, \dots, \omega_i\}$. First, we will get an approximation of F_j^i by using the trapezoidal integration formula over $[x_0^i, x_j^i]$. Then we will refine this approximation using the fact stated in Eq. (9.2). The trapezoidal method yields the approximation:

$$\forall j = 1, \dots, \omega_i + 1 \quad \tilde{F}_j^i := F_0^i + \frac{1}{2} \sum_{l=1}^j (x_l^i - x_{l-1}^i)(f_l^i + f_{l-1}^i). \tag{9.3}$$

Now the problem is that $\tilde{F}_{\omega_i+1}^i$ thus defined would generally differ from $\sum_{j=1}^i \omega_j/p$ due to numerical error in the integration formula. This is why, in a second step, we refine the approximation by computing

$$F_j^i := F_0^i + (\tilde{F}_j^i - F_0^i) \frac{F_{\omega_i+1}^i - F_0^i}{\tilde{F}_{\omega_i+1}^i - F_0^i} \quad \text{for } j = 1, \dots, \omega_i. \tag{9.4}$$

9.2. Derivatives with respect to QuEST function inputs

The computation of these derivatives is subdivided into two steps that mirror the ones performed in Section 9.1. First, by differentiating Eq. (9.3) with respect to x_l^i and f_l^i we obtain:

$$\begin{aligned} \forall j = 1, \dots, \omega_i + 1 \quad \frac{\partial \tilde{F}_j^i}{\partial t_k} &= \frac{1}{2} \sum_{l=1}^j \left(\frac{\partial x_l^i}{\partial t_k} - \frac{\partial x_{l-1}^i}{\partial t_k} \right) (f_l^i + f_{l-1}^i) \\ &\quad + \frac{1}{2} \sum_{l=1}^j (x_l^i - x_{l-1}^i) \left(\frac{\partial f_l^i}{\partial t_k} + \frac{\partial f_{l-1}^i}{\partial t_k} \right), \end{aligned} \tag{9.5}$$

where the partial derivatives of x_l^i and f_l^i with respect to t_k are given by Eqs. (8.6) and (8.3), respectively. Second, differentiating Eq. (9.4) with respect to \tilde{F}_j^i and $\tilde{F}_{\omega_i+1}^i$ yields:

$$\frac{\partial F_j^i}{\partial t_k} = (F_{\omega_i+1}^i - F_0^i) \frac{\frac{\partial \tilde{F}_j^i}{\partial t_k} - F_0^i}{\tilde{F}_{\omega_i+1}^i - F_0^i} - (F_{\omega_i+1}^i - F_0^i) \frac{\partial \tilde{F}_{\omega_i+1}^i}{\partial t_k} \cdot \frac{\tilde{F}_j^i - F_0^i}{(\tilde{F}_{\omega_i+1}^i - F_0^i)^2}, \tag{9.6}$$

where the partial derivatives of \tilde{F}_j^i and $\tilde{F}_{\omega_i+1}^i$ with respect to t_k are given by Eq. (9.5).

10. Discretization of the sample spectral C.D.F.

10.1. Estimated sample eigenvalues

The final operation involves extracting from F a set of p estimated sample eigenvalues $(\lambda_1, \dots, \lambda_p)$. First, we take care of zero eigenvalues when $c > 1$. By Eq. (9.1),

if $p < n$ then $\lambda_1, \dots, \lambda_{p-n} = 0$.

In what follows, we will assume that we have fixed an interval index i in the set $\{1, \dots, \nu\}$.

Let the function $X^i(\alpha)$ denote the approximation to $\int_{F_0^i}^\alpha F^{-1}(x)dx$ that is obtained by fitting a piecewise linear function to $F^{-1}(\cdot)$ over the interval $[F_0^i, F_{\omega_i+1}^i]$. This piecewise linear function passes through every point $(F_j^i, x_j^i)_{j=0, \dots, \omega_i+1}$. Using once again the trapezoidal integration formula, we obtain:

$$\forall j = 0, \dots, \omega_i \quad \int_{F_j^i}^{F_{j+1}^i} F^{-1}(x) dx \approx X^i(F_{j+1}^i) - X^i(F_j^i) = (F_{j+1}^i - F_j^i) \frac{x_j^i + x_{j+1}^i}{2}. \tag{10.1}$$

For every integer κ such that $pF_0^i \leq \kappa < pF_{\omega_i+1}^i$, define $j(\kappa)$ as the unique integer in $\{0, \dots, \omega_i\}$ such that $F_{j(\kappa)}^i \leq \kappa < F_{j(\kappa)+1}^i$. Then it holds that:

$$\begin{aligned} \int_{F_{j(\kappa)}^i}^{\kappa/p} F^{-1}(x) dx &\approx X^i(\kappa/p) - X^i(F_{j(\kappa)}^i), \quad \text{where} \\ X^i(\kappa/p) - X^i(F_{j(\kappa)}^i) &= \left(\frac{\kappa}{p} - F_{j(\kappa)}^i\right) \left[x_{j(\kappa)}^i + \frac{\frac{\kappa}{p} - F_{j(\kappa)}^i}{2(F_{j(\kappa)+1}^i - F_{j(\kappa)}^i)} (x_{j(\kappa)+1}^i - x_{j(\kappa)}^i) \right] \\ &= \left(\frac{\kappa}{p} - F_{j(\kappa)}^i\right) x_{j(\kappa)}^i + \frac{\left(\frac{\kappa}{p} - F_{j(\kappa)}^i\right)^2}{2(F_{j(\kappa)+1}^i - F_{j(\kappa)}^i)} (x_{j(\kappa)+1}^i - x_{j(\kappa)}^i). \end{aligned} \tag{10.2}$$

Putting together Eqs. (10.1)–(10.2) yields:

$$\begin{aligned} X^i(\kappa/p) &= \sum_{l=0}^{j(\kappa)-1} (F_{l+1}^i - F_l^i) \frac{x_l^i + x_{l+1}^i}{2} \\ &\quad + \left(\frac{\kappa}{p} - F_{j(\kappa)}^i\right) x_{j(\kappa)}^i + \frac{\left(\frac{\kappa}{p} - F_{j(\kappa)}^i\right)^2}{2(F_{j(\kappa)+1}^i - F_{j(\kappa)}^i)} (x_{j(\kappa)+1}^i - x_{j(\kappa)}^i). \end{aligned}$$

Finally, we can define the estimated sample eigenvalues that belong to the i th support interval as:

$$\forall \kappa \in \{pF_0^i + 1, pF_0^i + 2, \dots, pF_{\omega_i+1}^i\} \quad \lambda_\kappa := X^i\left(\frac{\kappa}{p}\right) - X^i\left(\frac{\kappa - 1}{p}\right). \tag{10.3}$$

10.2. Partial derivatives of estimated sample eigenvalues with respect to the inputs of the QuEST function

As in Section 10.1, we handle separately the zero eigenvalues when the sample covariance matrix is singular:

$$\text{if } p < n \text{ then } \forall \kappa = 1, \dots, p - n \quad \frac{\partial \lambda_\kappa}{\partial t_k} = 0.$$

In the remainder of this section, we will assume that we have fixed an interval index i in the set $\{1, \dots, \nu\}$. Differentiating Eq. (10.1) with respect to F_j^i and x_j^i yields:

$$\begin{aligned} \forall j = 0, \dots, \omega_i \quad \frac{\partial X^i}{\partial t_k}(F_{j+1}^i) - \frac{\partial X^i}{\partial t_k}(F_j^i) &= \frac{1}{2} \left(\frac{\partial F_{j+1}^i}{\partial t_k} - \frac{\partial F_j^i}{\partial t_k} \right) (x_j^i + x_{j+1}^i) \\ &\quad + \frac{1}{2} (F_{j+1}^i - F_j^i) \left(\frac{\partial x_j^i}{\partial t_k} + \frac{\partial x_{j+1}^i}{\partial t_k} \right), \end{aligned} \tag{10.4}$$

where the partial derivatives of F_j^i and x_j^i with respect to t_k are given by Eqs. (9.6) and (8.6), respectively. Similarly, differentiating Eq. (10.2) yields:

$$\begin{aligned} \frac{\partial X^i}{\partial t_k} \left(\frac{\kappa}{p} \right) - \frac{\partial X^i}{\partial t_k} (F_{j(\kappa)}^i) &= \left(\frac{\kappa}{p} - F_{j(\kappa)}^i \right) \frac{\partial x_{j(\kappa)}^i}{\partial t_k} - \frac{\partial F_{j(\kappa)}^i}{\partial t_k} x_{j(\kappa)}^i \\ &\quad - \frac{\partial F_{j(\kappa)}^i}{\partial t_k} \times \frac{\left(\frac{\kappa}{p} - F_{j(\kappa)}^i \right)^2}{F_{j(\kappa)+1}^i - F_{j(\kappa)}^i} (x_{j(\kappa)+1}^i - x_{j(\kappa)}^i) \\ &\quad + \frac{\left(\frac{\kappa}{p} - F_{j(\kappa)}^i \right)^2}{2(F_{j(\kappa)+1}^i - F_{j(\kappa)}^i)} \left(\frac{\partial x_{j(\kappa)+1}^i}{\partial t_k} - \frac{\partial x_{j(\kappa)}^i}{\partial t_k} \right) \\ &\quad - \left(\frac{\partial F_{j(\kappa)+1}^i}{\partial t_k} - \frac{\partial F_{j(\kappa)}^i}{\partial t_k} \right) \frac{\left(\frac{\kappa}{p} - F_{j(\kappa)}^i \right)^2}{2(F_{j(\kappa)+1}^i - F_{j(\kappa)}^i)^2} (x_{j(\kappa)+1}^i - x_{j(\kappa)}^i). \end{aligned} \tag{10.5}$$

We obtain the partial derivative of X^i with respect to t_k evaluated at κ/p from Eqs. (10.4)–(10.5) in the following way:

$$\frac{\partial X^i}{\partial t_k} \left(\frac{\kappa}{p} \right) = \sum_{l=0}^{j(\kappa)-1} \left[\frac{\partial X^i}{\partial t_k} (F_{l+1}^i) - \frac{\partial X^i}{\partial t_k} (F_l^i) \right] + \left[\frac{\partial X^i}{\partial t_k} \left(\frac{\kappa}{p} \right) - \frac{\partial X^i}{\partial t_k} (F_{j(\kappa)}^i) \right],$$

which enables us to compute the partial derivatives of the sample eigenvalues that belong to the i th support interval with respect to the population eigenvalues as:

$$\forall \kappa \in \{pF_0^i + 1, pF_0^i + 2, \dots, pF_{o_i+1}^i\} \quad \forall k = 1, \dots, p \quad \frac{\partial \lambda_\kappa}{\partial t_k} = \frac{\partial X^i}{\partial t_k} \left(\frac{\kappa}{p} \right) - \frac{\partial X^i}{\partial t_k} \left(\frac{\kappa - 1}{p} \right). \tag{10.6}$$

This derivation concludes the description of the numerical implementation of the QuEST function and its analytical Jacobian.

11. Monte Carlo simulations

Ledoit and Wolf (2015, Section 5.1.1) already provide some preliminary evidence for the accuracy of the estimator of the population eigenvalues obtained by numerically inverting the QuEST function. The simulations presented below are more extensive and highlight the convergence rate in log–log scale for various shapes of the population spectrum.

11.1. Population spectrum

The population eigenvalues are taken from the distribution of $1 + (\kappa - 1)X$, where κ is the condition number and X is a random variable whose support is the compact interval $[0, 1]$. Throughout the whole simulation study, we carry four different shapes for the distribution of X .

1. The original shape is left-skewed: it is the Kumaraswamy (1980) distribution with parameters $(3, 1/3)$. The Kumaraswamy family is similar in spirit to the Beta family, but more tractable: the density, the c.d.f. and the quantile function are all available in closed form. For reference, the c.d.f. of Kumaraswamy $(3, 1/3)$ is

$$\forall x \in [0, 1] \quad H_1(x) = 1 - (1 - x^3)^{1/3}. \tag{11.1}$$

All the other shapes are derived from this one.

2. The next shape is right-skewed, obtained by taking the mirror image of the density about the midpoint of the support. Jones (2009, p. 73) observes that there is “a pleasing symmetry” in this case: it is equivalent to taking the mirror image of the c.d.f. about the 45 degrees line, that is, replacing it with its inverse, the quantile function:

$$\forall x \in [0, 1] \quad H_2(x) = [1 - (1 - x)^3]^{1/3}. \tag{11.2}$$

3. A symmetric bimodal distribution is generated by combining right-skewness on $[0, 1/2]$ with left-skewness on $[1/2, 1]$:

$$\forall x \in [0, 1] \quad H_3(x) = \begin{cases} \frac{1}{2} [1 - (1 - 2x)^3]^{1/3} & \text{if } x \in [0, 1/2], \\ 1 - \frac{[1 - (2x - 1)^3]^{1/3}}{2} & \text{if } x \in [1/2, 1]. \end{cases} \tag{11.3}$$

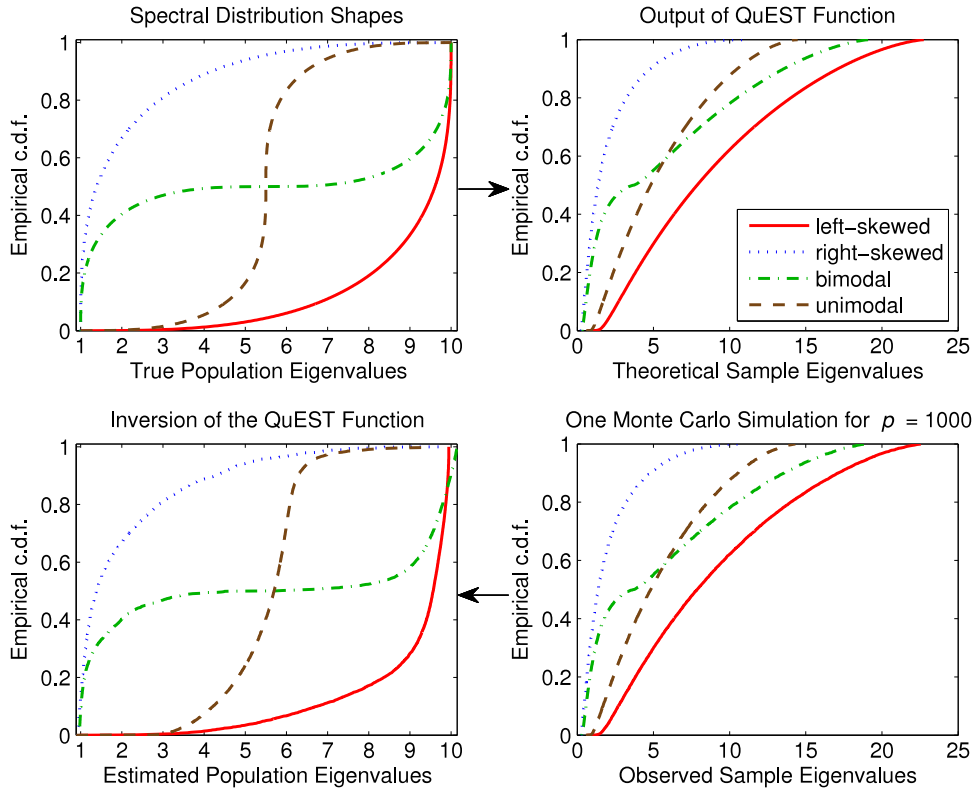


Fig. 2. Population vs. sample spectrum. Top panel is the direct QuEST function, bottom panel the inverse of the QuEST function for estimation purposes.

4. Finally a symmetric unimodal distribution is generated by combining left-skewness on $[0, 1/2]$ with right-skewness on $[1/2, 1]$:

$$\forall x \in [0, 1] \quad H_4(x) = \begin{cases} \frac{1 - [1 - (2x)^3]^{1/3}}{2} & \text{if } x \in [0, 1/2], \\ \frac{1 + [1 - (2 - 2x)^3]^{1/3}}{2} & \text{if } x \in [1/2, 1]. \end{cases} \quad (11.4)$$

All four densities diverge to infinity, so the set of shapes chosen is a challenging one.

11.2. Intuition

As mentioned in Section 3, given the sample eigenvalues $\lambda_{n,1} \leq \lambda_{n,2} \leq \dots \leq \lambda_{n,p}$, we estimate the population eigenvalues $\tau_{n,1} \leq \tau_{n,2} \leq \dots \leq \tau_{n,p}$ by numerically inverting the QuEST function:

$$\hat{\tau}_n := \operatorname{argmin}_{\mathbf{t} \in [0, \infty)^p} \frac{1}{p} \sum_{i=1}^p [q_{n,p}^i(\mathbf{t}) - \lambda_{n,i}]^2. \quad (11.5)$$

The simulation study presented below centers on the base-case scenario where the condition number is $\kappa = 10$, variates are normally distributed, and the concentration ratio is $c = 1/3$. For dimension $p = 1,000$, Fig. 2 provides a side-by-side comparison of the population spectra specified in Section 11.1 with their sample counterparts.

Let us start with the top panel. It shows the effect of the QuEST function. For each of the four distribution shapes, the population eigenvalues in the top left graph get mapped into the limiting sample spectra shown in the top right graph. This mapping illustrates the transformation coming out of the Fundamental Equation. There is a lot of distortion, but the relative positions of the four color-coded c.d.f.'s have been preserved. Therefore, the information has not been destroyed: it is just waiting to be deciphered by a suitable method. We are essentially facing a severe nonlinear bias-correction problem.

The bottom panel goes in the opposite direction: the QuEST function gets inverted. At the bottom right are the sample eigenvalues generated in one Monte Carlo simulation. Observe how closely they match the nonrandom distributions in the top right. This is because, as mentioned above, in the large-dimensional asymptotic limit randomness vanishes. Then

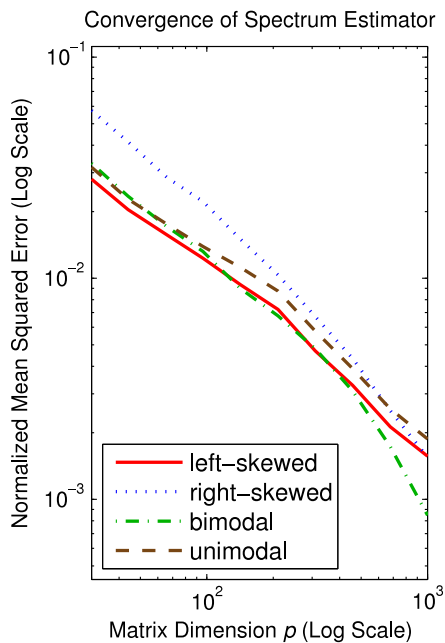


Fig. 3. Consistency of the estimator of population eigenvalues in the base case scenario.

numerically inverting the QuEST function yields the estimated population eigenvalues shown in the bottom left graph. They closely match the truth (shown top left); the distortion has been undone, and the original shapes of the spectral distributions have been restored. The bottom panel is our estimation procedure in a nutshell.

11.3. Base-case scenario

Ledoit and Wolf (2015, Theorem 2.2) prove that the mean squared deviation between the estimated eigenvalues and the population eigenvalues

$$\frac{1}{p} \sum_{i=1}^p [\hat{\tau}_{n,i} - \tau_{n,i}]^2 \tag{11.6}$$

converges almost surely to zero under large-dimensional asymptotics. This quantity is scale-sensitive, whereas the problem is scale-invariant. Therefore, we study in the Monte Carlo simulations the scale-adjusted quantity

$$\frac{\frac{1}{p} \sum_{i=1}^p [\hat{\tau}_{n,i} - \tau_{n,i}]^2}{\left(\frac{1}{p} \sum_{i=1}^p \tau_{n,i}\right)^2} \tag{11.7}$$

instead, called the (empirical) *normalized mean squared error*. This change in performance measure does not affect the strong consistency result, given that Ledoit and Wolf (2015) assume that the population eigenvalues are bounded away from zero and infinity: The quantity (11.6) converges to zero almost surely if and only if the quantity (11.7) converges to zero almost surely. Our preference for the performance measure (11.7) is due to the fact that we do not want to give the visual impression that covariance matrices with a larger trace are estimated less accurately, since on a relative basis they are not.

The matrix dimension ranges from $p = 30$ to $p = 1,000$. Convergence of the scale-adjusted mean squared deviation defined by Eq. (11.7) is displayed in Fig. 3 on a log–log scale for the four distribution shapes.

In all log–log graphs presented in this paper, including this one, the scales of the x- and y-axes have been equalized, so that the -45° line corresponds to a convergence rate of p . Each point in the curves corresponds to the average across 1,000 Monte Carlo simulations.

In terms of speed, Fig. 4 shows that the numerical recipe presented in this paper for the implementation of the QuEST function is sufficiently fast for practical purposes. (These numbers were run using Matlab R2016a on an Apple Mac Pro with a 3.5 GHz Intel Xeon E5 processor.)

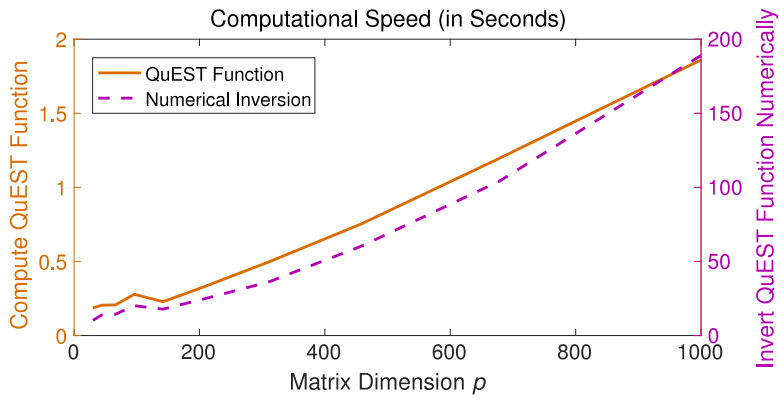


Fig. 4. Speed benchmark for computing the QuEST function and estimating population eigenvalues. The QuEST function relates to the vertical axis on the left-hand side, and its numerical inversion to the one on the right-hand side.

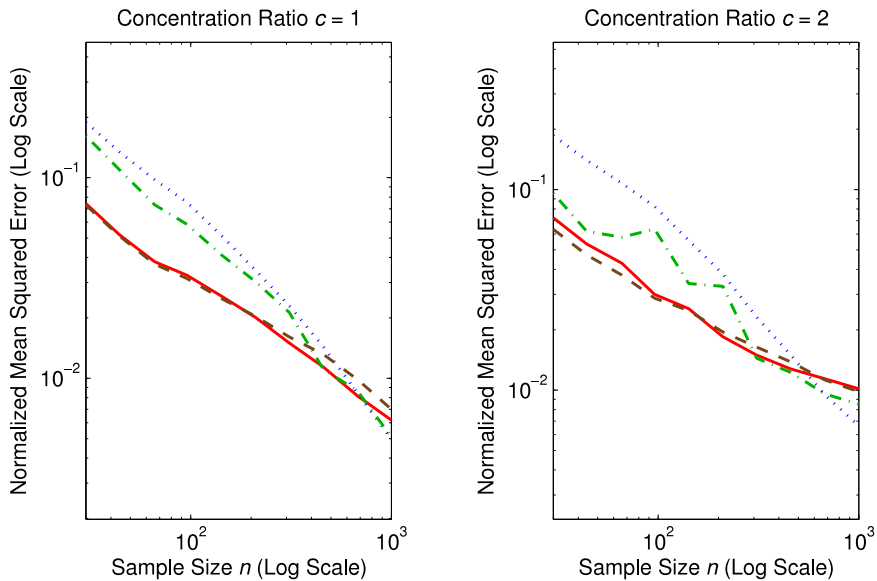


Fig. 5. Consistency of the estimator of population eigenvalues for higher concentration ratios. Color and line-style codes are as in Figs. 2 and 3.

The remainder of Section 11 is dedicated to demonstrating the robustness of the base-case convergence pattern in three directions: different concentration ratios $c = p/n$, condition numbers κ , and variate distributions D .

11.4. Concentration ratio

First, we increase the concentration ratio; we pick two values: $c = 1$ and $c = 2$. The first case is not covered by the mathematical theory of Ledoit and Wolf (2015), but the numerical results displayed on the left panel of Fig. 5 seem to indicate that satisfactory convergence is achieved nonetheless. In the second case, we manage to consistently estimate p eigenvalues, in spite of the fact that the sample covariance matrix has only $n = p/2$ nontrivial eigenvalues. Note that this is the only graph where we let n (instead of p) range from 30 to 1,000, because of $n \leq p$.

11.5. Condition number

The second axis of deviation from the baseline case is to look at condition numbers other than $\kappa = 10$. We consider a smaller condition number, $\kappa = 2$, and a larger one, $\kappa = 100$. The results are displayed in Fig. 6.

These results show that we still obtain convergence in spite of changes in the condition number.

11.6. Distribution of the variates

Finally, we deviate from the base-case scenario in the direction of having other distributions than Gaussian for the random variates. First, we take a fat-tailed distribution: the “Student” t -distribution with 5 degrees of freedom; and second, the

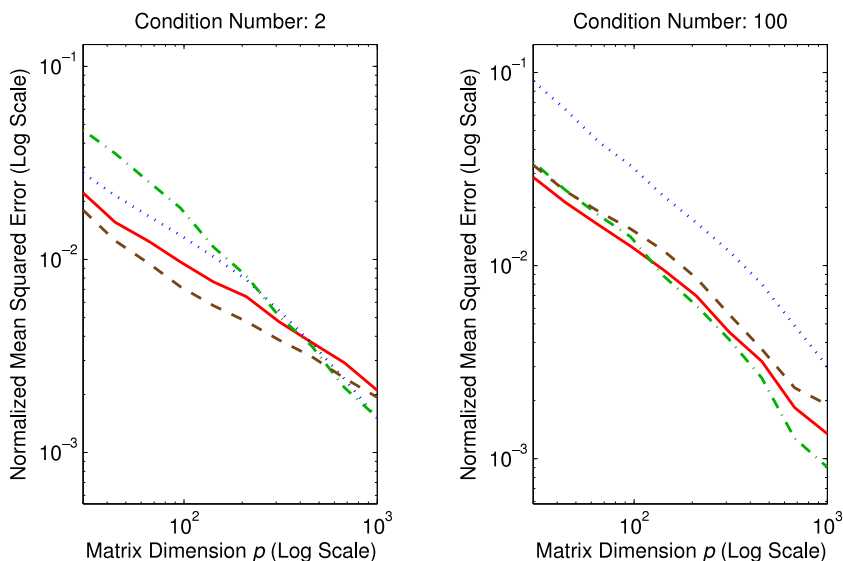


Fig. 6. Consistency of the estimator of population eigenvalues for various condition numbers. Color and line-style codes are as in Figs. 2 and 3.

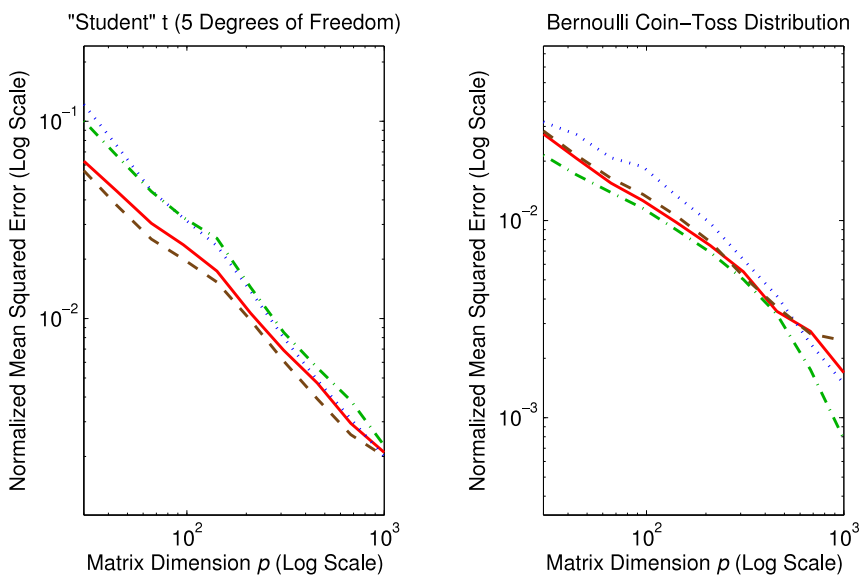


Fig. 7. Consistency of the estimator of population eigenvalues when the variates have thick or thin tails. Color and line-style codes are as in Figs. 2 and 3.

most thin-tailed of all distributions: the Bernoulli coin-toss distribution with probability 1/2. The results are displayed in Fig. 7.

We also consider a skewed distribution: the exponential with parameter one. The results are displayed in Fig. 8.

These results show that we obtain convergence across a variety of variate distributions.

11.7. Overview of the simulation results

The Monte Carlo simulations presented above illustrate the ability of the estimator of population eigenvalues constructed by numerically inverting the QuEST function to get closer to the truth as the matrix dimension and the sample size go to infinity together. This exercise has been extensive, involving a grand total of 320,000 Monte Carlo simulations. The point was to build practical comfort around the theoretical result. Best-fit lines in log–log space have slopes that vary in the range from -0.70 to -1.10 , giving some empirical indication about the exponent of the convergence rate of the mean squared deviation between true and estimated population eigenvalues.

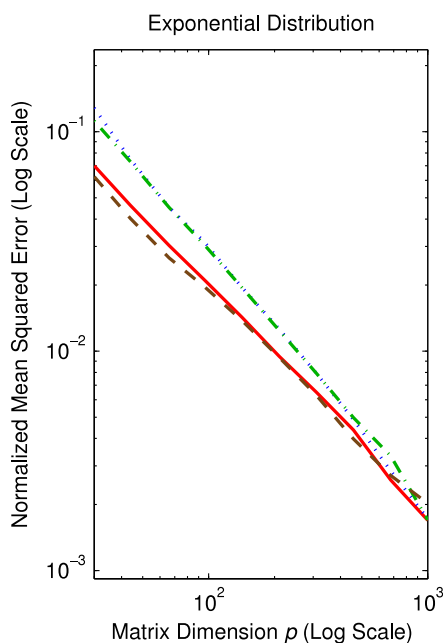


Fig. 8. Consistency of the estimator of population eigenvalues when the variates are skewed. Color and line-style codes are as in Figs. 2 and 3.

Remark 11.1 (*Further Numerical Results*). To keep the exposition compact, we only present simulation results for our estimator of the population eigenvalues in this paper. Important applications of this estimator include nonlinear shrinkage estimation covariance matrices and principal component analysis (PCA) in large dimensions. For corresponding numerical results, in terms of both simulation studies and empirical applications to real-life data, the interested reader is referred to Ledoit and Wolf (2015, forthcoming, 2017b) and Engle et al. (forthcoming). Note that the simulation results in Ledoit and Wolf (2012) do not fall in this category, since they are based on a different estimator of the population eigenvalues. ■

12. Conclusion

When the matrix dimension is not negligible with respect to the sample size, finite-dimensional asymptotic approximations are no longer close to the truth: We enter the Marčenko and Pastur (1967) zone instead. In this zone, the sample eigenvalues are a very distorted version of their population counterparts. Only after the works of El Karoui (2008) and Mestre (2008), did researchers in the field of large-dimensional multivariate statistics start to harbor any hope of unwinding this distortion.

Ledoit and Wolf (2015) put forward a natural discretization of the Fundamental Equation of Random Matrix Theory that can be inverted numerically. Even though the sample eigenvalues are far from their population counterparts, the distortion can be inverted through this particular procedure. The present paper describes in great detail how to discretize the Marčenko–Pastur equation, also called the Fundamental Equation. In addition, we provide extensive Monte Carlo simulations demonstrating the practical effectiveness of the method in terms of recovering the population eigenvalues. There are many applications in the field of large-dimensional multivariate statistics, including nonlinear shrinkage estimation of covariance matrices, as proposed by Ledoit and Wolf (2012, 2015, 2017b) and Engle et al. (forthcoming), and PCA, as proposed by Ledoit and Wolf (2015).

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