

ON THE ACCURACY OF SOLVING CONFLUENT PRONY SYSTEMS*

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Abstract. In this paper we consider several nonlinear systems of algebraic equations which can be called “Prony-type.” These systems arise in various reconstruction problems in several branches of theoretical and applied mathematics, such as frequency estimation and nonlinear Fourier inversion. Consequently, the question of stability of solution with respect to errors in the right-hand side becomes critical for the success of any particular application. We investigate the question of “maximal possible accuracy” of solving Prony-type systems, putting stress on the “local” behavior which approximates situations with low absolute measurement error. The accuracy estimates are formulated in very simple geometric terms, shedding some light on the structure of the problem. Numerical tests suggest that “global” solution techniques such as Prony’s algorithm and the ESPRIT method are suboptimal when compared to this theoretical “best local” behavior.

Key words. confluent Prony system, Prony method, algebraic sampling, Jacobian determinant, confluent Vandermonde matrix, Hankel matrix, PACE model, ESPRIT, frequency estimation

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1. Introduction.

1.1. Problem definition. Consider the following system of algebraic equations:

$$(1.1) \quad \sum_{i=1}^{\kappa} a_i \xi_i^k = m_k,$$

where $a_i, \xi_i \in \mathbb{C}$ are unknown parameters and the measurements $\{m_k\}_{k=0,1,\dots}$ are given. This “exponential fitting” system, or “Prony system,” appears in several branches of theoretical and applied mathematics, such as frequency estimation, Padé approximation, array processing, statistics, interpolation, quadrature, radar signal detection, and error correction codes. The literature on this subject is huge (for instance, the bibliography on Prony’s method from [3] is some 50+ pages long). Our interest in this system (and other, more general, systems of this kind, to be specified below) is motivated by its central role in algebraic sampling—a recent approach to reconstruction of nonlinear parametric models from measurements. There, it arises as the problem of reconstructing a signal modeled by a linear combination of Dirac δ -distributions,

$$(1.2) \quad f(x) = \sum_{i=1}^{\kappa} a_i \delta(x - \xi_i), \quad a_i, \xi_i \in \mathbb{R},$$

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from the measurements given by the power moments

$$(1.3) \quad m_k(f) \stackrel{\text{def}}{=} \int_0^1 x^k f(x) \, dx.$$

While the above problem may be considered mainly of theoretical interest, it is actually one of the most basic problems in algebraic sampling. On one hand, if $s(x)$ is a piecewise-constant signal with jump discontinuities at the locations $\xi_1, \dots, \xi_{\mathcal{K}}$, then $s'(x) = f(x)$ as in (1.2). Thus, the “signal” $f(x)$ essentially captures the nonsmooth nature of $s(x)$. On the other hand, the moments (1.3) are convenient to consider because of the respective simplicity of the arising algebraic equations, while other types of measurements (e.g., Fourier coefficients) may be recast into moments after a change of variables.

An important generalization of the Prony system, which is of great interest to us, arises when the simple model (1.2) is extended to include higher-order derivatives (see [8, 46] for examples of such constructions):

$$(1.4) \quad f(x) = \sum_{i=1}^{\mathcal{K}} \sum_{j=0}^{l_i-1} a_{i,j} \delta^{(j)}(x - \xi_i), \quad a_{i,j}, \xi_j \in \mathbb{R},$$

where $\delta^{(j)}$ is the j th derivative of the Dirac delta (in the sense of distributions).

From now on, we denote the number of unknown coefficients $a_{i,j}$ by $C \stackrel{\text{def}}{=} \sum_{i=1}^{\mathcal{K}} l_i$ and the overall number of unknown parameters by $R \stackrel{\text{def}}{=} C + \mathcal{K}$. Taking moments of $f(x)$ in (1.4), we arrive¹ at the following “confluent Prony” system:

$$(1.5) \quad \sum_{i=1}^{\mathcal{K}} \sum_{j=0}^{l_i-1} a_{i,j} (k)_j \xi_i^{k-j} = m_k, \quad a_{i,j}, \xi_i, m_k \in \mathbb{C},$$

where the Pochhammer symbol $(i)_j$ denotes the falling factorial

$$(i)_j = i(i-1) \cdots (i-j+1), \quad i \in \mathbb{R}, \quad j \in \mathbb{N},$$

and the expression $(k)_j \xi_i^{k-j}$ is defined to be zero for $k > j$.

The Prony-type systems appear in various recent reconstruction methods of signals with discontinuities; see [7, 8, 9, 10, 11, 14, 18, 20, 21, 23, 24, 28, 30, 32]. In particular, finite rate of innovation (FRI) techniques [19, 31, 46] have spawned a rather extensive literature (see, e.g., a recent addition [44]). Usually, the ξ_i represent “location” parameters of the problem, such as discontinuity locations or complex frequencies $\xi_j = e^{i\omega_j}$. These variables enter the equations in a nonlinear way, and we call them “nodes.” The coefficients $a_{i,j}$, on the other hand, enter the equations linearly, and we call them “magnitudes.”

While algebraic sampling provides exact reconstruction for noise-free data in many cases mentioned above, a critical issue remains, namely, stability, or accuracy of solution. Stable solution of Prony-type systems is generally considered to be a difficult problem, and in recent years many algorithms have been devised for this task (e.g., [6, 25, 26, 33, 34, 36, 38, 42, 45]). Perhaps the simplest version of the stability problem can be formulated as follows (cf. Definitions 3.1 and 4.1 and subsection 1.4).

¹Strictly speaking, this will result in a “real” confluent Prony system.

Assume that the measurements $\{m_k\}_{k=0,\dots,S-1}$ are known with some error: $m_k + \varepsilon_k$. Given an estimate $\varepsilon = \max_k |\varepsilon_k|$, how large can the error in the reconstructed model parameters (i.e., $|\Delta\xi_j| \stackrel{\text{def}}{=} |\tilde{\xi}_j - \xi_j|$ and $|\Delta a_{i,j}| \stackrel{\text{def}}{=} |\widetilde{a_{i,j}} - a_{i,j}|$) be in the worst case in terms of ε , the number of measurements S , and the true parameters $\{\xi_j\}, \{a_{i,j}\}$?

In more detail, our ultimate goal may be described as follows:

1. determine the qualitative dependence of the accuracy on the values of the parameters;
2. quantify this dependence as precisely as possible; and
3. determine how (and if at all) increasing the number of measurements (i.e., oversampling) improves accuracy.

1.2. Related work. Matching the ubiquity of Prony-type systems is the impressive body of literature devoted to both designing methods of solution and analyzing the accuracy/robustness of these methods; see references cited above. Although there appears to be no simple answer to the above question of “maximal possible accuracy,” several important results in this direction are available in the literature, which we now briefly discuss.

Methods of solution can be roughly divided into three categories (see, e.g., [41], [43, section 4]): direct nonlinear minimization (nonlinear least squares), recurrence-based methods (such as the original Prony method; see section 2), and subspace methods (such as Pisarenko’s method, MUSIC, ESPRIT, and matrix pencils; see, e.g., [38]).

In the framework of statistical signal estimation [27], the subspace methods are known to be more efficient and robust to noise, mainly due to the fact that the noise is assumed to have certain statistical properties. The confluent Prony system (1.5) is also known as the “polynomial amplitude complex exponential” (PACE) model. A standard measure of estimator performance is Cramer–Rao bound and related lower bounds (CRB). These have been recently established for the PACE model in [5] (see also related results for FRI models [15]). Furthermore, it has been demonstrated that the performance of the generalized ESPRIT algorithm ([4, 6] and subsection 5.2) is close to the optimal CRB; therefore, we consider it to represent the state of the art in the subspace methods.

We do not assume any particular statistical model or other structure for either the error terms ε_k or the estimation algorithm (such as white noise or unbiasedness). Therefore, the CRB and related lower bounds cannot provide the full answer to the stability problem *as is*. Still, it turns out that the stability bounds developed in this paper resemble the CRB as established in [5]; see subsection 5.1 below for details.

Recent papers of Tasche, Peter, and Potts [34, 36] contain some uniform error bounds for solving Prony systems. In particular, the authors develop the so-called approximate Prony method, analyze its worst-case error, and numerically compare it with the ESPRIT method (showing similar performance). Although they consider the nonconfluent version of the Prony system (1.1) and analyze only the error in recovering the magnitudes a_j , we believe these results to be an important step toward answering the stability problem as posed above. See subsection 5.3 below for details.

Very recently, Candes and Fernandez-Granda [18] investigated stable solution of Prony systems by total variation minimization under assumptions of minimal node separation, in the context of superresolution.

Considering all of the above, we believe that a full answer to our somewhat rigid l^∞ formulation of the stability problem may contribute to the understanding of limitations of using Prony systems and methods both in signal processing applications

and in function approximation, in particular compressed sensing, nonlinear Fourier inversion, FRI techniques, and related problems.

1.3. Notation. In what follows we use the infinity norm distance

$$\forall \mathbf{x}, \mathbf{y} \in \mathbb{C}^n : \quad \text{dist}(\mathbf{x}, \mathbf{y}) \stackrel{\text{def}}{=} \max_{1 \leq i \leq n} |x_i - y_i|$$

and denote by $B(\mathbf{a}, \varepsilon)$ the ε -ball around a point $\mathbf{a} \in \mathbb{C}^n$ in this norm.

1.4. Summary of results. In section 3 we define “best possible pointwise accuracy” as follows. We consider the “Prony map” $\mathcal{P}_S : \mathbb{C}^R \rightarrow \mathbb{C}^S$ which associates to any parameter vector $\mathbf{x} = \{\{a_{ij}\}, \{\xi_i\}\} \in \mathbb{C}^R$ its corresponding measurement vector $\mathbf{y} = (m_0, \dots, m_{S-1}) \in \mathbb{C}^S$ (where the m_k are given by (1.5)).

Now if instead of \mathbf{y} we are given a noisy $\tilde{\mathbf{y}} \in B(\mathbf{y}, \varepsilon)$, then this $\tilde{\mathbf{y}}$ can correspond to any parameter vector $\tilde{\mathbf{x}} \in \mathbb{C}^R$ for which $\mathcal{P}_S(\tilde{\mathbf{x}}) \in B(\tilde{\mathbf{y}}, \varepsilon)$. Therefore, we define the best possible accuracy at a point \mathbf{x} to be equal to the maximal (over all $\tilde{\mathbf{y}}$) spread of the preimage of this $B(\tilde{\mathbf{y}}, \varepsilon)$, that is (see Definition 3.1),

$$\sup_{\tilde{\mathbf{y}} \in B(\mathbf{y}, \varepsilon)} \frac{1}{2} \text{diam} \mathcal{P}_S^{-1}(B(\tilde{\mathbf{y}}, \varepsilon)).$$

We then simplify the setting by assuming that the number of measurements S equals the number of unknowns R , and looking at the (local) linear approximation to the Prony map \mathcal{P}_S . Then the solution error at some (noncritical) point in the parameter space can be estimated by the local Lipschitz constant of the (regular) inverse map \mathcal{P}_S^{-1} . We derive such simple estimates in section 4 and compare them to the “global” accuracy of the original Prony method (derived for completeness in section 2).

Our main result (Theorem 4.5) can be summarized as follows (all statements are valid for small ε):

1. The stability of recovering a node ξ_i depends on the separation of the nodes, is inversely proportional to the magnitude of the highest coefficient corresponding to this node ($|a_{i,l_i-1}|$), and does not depend on any other magnitude.
2. For $1 \leq j \leq l_i - 1$, the stability of recovering a magnitude $a_{i,j}$ depends on the separation of the nodes, is proportional to $1 + \frac{|a_{i,j-1}|}{|a_{i,l_i-1}|}$, and does not depend on any other magnitude. Note that in fact every magnitude influences only the next highest magnitude corresponding to the same node.
3. The stability of recovering the lowest magnitudes $a_{i,0}$ is the same for all nodes and depends only on the separation of the nodes.

The separation of the nodes is specified in terms of norms of inverse confluent Vandermonde matrices on the nodes, which is roughly of the same order as some finite power of $\prod_{1 \leq i < j \leq \kappa} |\xi_j - \xi_i|^{-1}$.

Our numerical experiments (section 6) confirm the above theoretical estimates. We also test the performance of two well-known solution methods—namely, the recurrence-based Prony method (section 2) and the generalized ESPRIT (subsection 5.2)—in the same setting as above (i.e., high SNR). The results suggest the following.

1. The recurrence-based global Prony method does not achieve the above theoretical limits, and so it is not optimal even in the case of small data perturbations.
2. The subspace methods (in particular, the ESPRIT algorithm) behave better than the Prony method, but they are still not optimal for small perturbations and small sample size.

The ‘‘Prony map’’ approach can in principle be generalized to obtain both global accuracy bounds as well as to study effects of oversampling by considering the case $S > R$ and taking into account second-order terms in the Taylor expansion of \mathcal{P}_S . We discuss these directions in section 7.

2. The Prony method. In this section we describe the most basic solution method for the system (1.5), which is in fact a slight generalization of the (historically earliest) method due to Prony [37]. By factorizing the so-called *data matrix*, one immediately obtains necessary and sufficient conditions for a unique solution as well as an estimate of the numerical stability of the method.

Most of the results of section 2 are not new and are scattered throughout the literature. Nevertheless, we believe that our presentation can be useful for further study of the various singular situations, such as collision of two nodes.

2.1. The description of the method. The nontrivial part is the recovery of the nodes ξ_j . Note that the case of a priori known nodes has been extensively treated in the literature (see, e.g., [1, 35] for the most recent results). Using the framework of finite difference calculus, one can easily prove the following result (see [8, Theorem 2.8]).

PROPOSITION 2.1. *Let the sequence $\{m_k\}$ be given by (1.5). Then this sequence satisfies the recurrence relation (of length at most $C + 1$)*

$$\left(\prod_{i=1}^{\mathcal{K}} (\mathbf{E} - \xi_i \mathbf{I})^{l_i} \right) \{m_k\} = 0,$$

where \mathbf{E} is the forward shift operator in k and \mathbf{I} is the identity operator.

COROLLARY 2.2. *For all $k \in \mathbb{N}$ we have the recurrence relation $\sum_{j=0}^C q_j m_{k+j} = 0$, where q_0, q_1, \dots, q_C are the coefficients of the polynomial $q(x) \stackrel{\text{def}}{=} \prod_{i=1}^{\mathcal{K}} (x - \xi_i)^{l_i}$.*

This suggests the following reconstruction procedure.²

Algorithm 1 The Prony method

Let there be given $\{m_k\}_{k=0}^{2C-1}$ (where $C = \sum_{i=1}^{\mathcal{K}} l_i$).

1. Solve the linear system (here we set $q_C = 1$ for normalization)

$$(2.1) \quad \underbrace{\begin{pmatrix} m_0 & m_1 & \cdots & m_{C-1} \\ m_1 & m_2 & \cdots & m_C \\ \vdots & \vdots & \ddots & \vdots \\ m_{C-1} & m_C & \cdots & m_{2C-2} \end{pmatrix}}_{\stackrel{\text{def}}{=} M_C} \begin{pmatrix} q_0 \\ q_1 \\ \vdots \\ q_{C-1} \end{pmatrix} = - \begin{pmatrix} m_C \\ m_{C+1} \\ \vdots \\ m_{2C-1} \end{pmatrix}$$

for the unknown coefficients q_0, \dots, q_{C-1} .

2. Find all the roots of $q(x) = x^C + \sum_{j=0}^{C-1} q_j x^j$. These roots, with appropriate multiplicities, are the unknowns $\xi_1, \dots, \xi_{\mathcal{K}}$ (use, e.g., arithmetic means to estimate multiple roots which are scattered by the noise into clusters).
 3. Substitute the recovered ξ_i 's back into the original (1.5). Solve the resulting overdetermined linear system (C unknowns and $2C$ equations) with respect to the magnitudes $\{a_{i,j}\}$ by the least squares method.
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²Equivalent derivation of the method is based on Padé approximation to the function $I(z) = \sum_{k=0}^{\infty} m_k z^k$; see [37] and, for instance, [39].

Several comments are in order.

1. The number of measurements used in step 1 equals $2C$, which can be greater than the number of unknowns $R = C + \mathcal{K}$ (equality for the order zero Prony system). If more measurements are available, the linear system (2.1) can be modified in a straightforward way to be overdetermined and subsequently solved by, say, the least squares method.
2. The linear system for the magnitudes has a special “Vandermonde-like” structure (see below), and so certain efficient algorithms can be used to solve it (e.g., [16, 29]).

The remainder of this section is organized as follows. The Hankel matrix M_C is shown to factor into the product of a generalized “Vandermonde-type” matrix which depends only on the nodes ξ_j , with an upper triangular matrix depending only on the amplitudes $a_{i,j}$. We also explicitly write down the linear system for the $a_{i,j}$ (see step 3 in Algorithm 1 above). These calculations lead to simple nondegeneracy conditions and stability estimates for the Prony method.

2.2. Factorization of the data matrix. Let us start by recalling a well-known type of matrix.

DEFINITION 2.3. For every $j = 1, \dots, \mathcal{K}$ and $k \in \mathbb{N}$ let the symbol $\mathbf{u}_{j,k}$ denote the following $1 \times l_j$ row vector:

$$(2.2) \quad \mathbf{u}_{j,k} \stackrel{\text{def}}{=} \left[\xi_j^k, \quad k\xi_j^{k-1}, \quad \dots, \quad (k)_{l_j-1} \xi_j^{k-l_j+1} \right].$$

DEFINITION 2.4. Let $U = U(\xi_1, l_1, \dots, \xi_{\mathcal{K}}, l_{\mathcal{K}})$ denote the matrix

$$(2.3) \quad U = \begin{bmatrix} \mathbf{u}_{1,0} & \mathbf{u}_{2,0} & \dots & \mathbf{u}_{\mathcal{K},0} \\ \mathbf{u}_{1,1} & \mathbf{u}_{2,1} & \dots & \mathbf{u}_{\mathcal{K},1} \\ & & \dots & \\ \mathbf{u}_{1,C-1} & \mathbf{u}_{2,C-1} & \dots & \mathbf{u}_{\mathcal{K},C-1} \end{bmatrix}.$$

This matrix is called the “confluent Vandermonde” [16, 22] matrix. It has been long known in numerical analysis due to its central role in Hermite polynomial interpolation. Its determinant is [40, p. 30]

$$(2.4) \quad \det U = \prod_{1 \leq i < j \leq \mathcal{K}} (\xi_j - \xi_i)^{l_j l_i} \prod_{\mu=1}^{\mathcal{K}} \prod_{\nu=1}^{l_{\mu}-1} \nu!.$$

It is straightforward to see that the matrix U defines the linear system for the jump magnitudes $a_{i,j}$.

PROPOSITION 2.5. Let \mathbf{a} be the column vector containing all the magnitudes $\{a_{i,j}\}$, i.e.,

$$\mathbf{a} \stackrel{\text{def}}{=} [a_{1,0}, \dots, a_{1,l_1-1}, a_{2,0}, \dots, a_{2,l_2-1}, \dots, a_{\mathcal{K},0}, a_{\mathcal{K},l_{\mathcal{K}}-1}]^T,$$

and $\mathbf{m} \stackrel{\text{def}}{=} [m_0, \dots, m_{C-1}]^T$. Then we have

$$(2.5) \quad U(\xi_1, l_1, \dots, \xi_{\mathcal{K}}, l_{\mathcal{K}}) \mathbf{a} = \mathbf{m}.$$

It is known that every Hankel matrix H admits a factorization $H = UDU^T$, where U is given by (2.3) and D is a block diagonal matrix; see [17]. Using different notation, such a factorization is proved in [4, Proposition III.7] for the Hankel matrix M_C .

LEMMA 2.6. For the system (1.5), the matrix M_C admits the following factorization:

$$(2.6) \quad M_C = UBU^T,$$

where $U = U(\xi_1, l_1, \dots, \xi_{\mathcal{K}}, l_{\mathcal{K}})$ is the confluent Vandermonde matrix (2.3) and B is the $C \times C$ block diagonal matrix $B = \text{diag}\{B_1, \dots, B_{\mathcal{K}}\}$ with each block of size $l_i \times l_i$ given by

$$(2.7) \quad B_i \stackrel{\text{def}}{=} \begin{bmatrix} a_{i0} & a_{i1} & \cdots & \cdots & a_{i, l_i-1} \\ a_{i1} & & & \binom{l_i-1}{l_i-2} a_{i, l_i-1} & 0 \\ \cdots & & & \cdots & 0 \\ & \binom{l_i-1}{2} a_{i, l_i-1} & 0 & \cdots & 0 \\ a_{i, l_i-1} & 0 & \cdots & \cdots & 0 \end{bmatrix}.$$

In other words, B_i is a “flipped” upper triangular matrix whose j th antidiagonal equals

$$a_{ij} \cdot \begin{bmatrix} 1 & \binom{j}{2} & \cdots & \binom{j}{j-1} & 1 \end{bmatrix}$$

for $j = 0, \dots, l_i - 1$.

The formula (2.6) is useful because it separates the jump locations $\{\xi_i\}$ from the magnitudes $\{a_{i,j}\}$, simplifying the analysis considerably.

THEOREM 2.7. The system (1.5) for $k = 0, 1, \dots, 2C$ has a unique solution if and only if all the $\{\xi_i\}$'s are pairwise different and all the $\{a_{i, l_i-1}\}$'s (just the highest coefficients) are nonzero.

Proof. Existence of a unique solution to the system (2.1) is equivalent to the nondegeneracy of $M_C = UBU^T$. Furthermore, the system for the jump magnitudes is given by (2.5). Therefore, existence of a unique solution to (1.5) is equivalent to the conditions $\det U \neq 0$ and $\det B \neq 0$. The proof is completed by (2.4) and (2.7). \square

2.3. Stability estimates. The stability of the Prony method can be estimated by the condition numbers of the matrices B and U . In particular, we have the following well-known result (e.g., [47]) from numerical linear algebra.

LEMMA 2.8. Consider the linear system $A\mathbf{x} = \mathbf{b}$, and let \mathbf{x}_0 be the exact solution. Let this system be perturbed,

$$(A + \Delta A)\mathbf{x} = \mathbf{b} + \Delta \mathbf{b},$$

and let $\mathbf{x}_0 + \Delta \mathbf{x}$ denote the exact solution of this perturbed system. Denote $\delta \mathbf{x} = \frac{\|\Delta \mathbf{x}\|}{\|\mathbf{x}_0\|}$, $\delta A = \frac{\|\Delta A\|}{\|A\|}$, $\delta \mathbf{b} = \frac{\|\Delta \mathbf{b}\|}{\|\mathbf{b}\|}$, and the condition number $\kappa = \|A\| \|A^{-1}\|$ for some vector norm $\|\cdot\|$ and the induced matrix norm. Then

$$(2.8) \quad \delta \mathbf{x} \leq \frac{\kappa}{1 - \kappa \cdot \delta A} (\delta A + \delta \mathbf{b}).$$

Now we can easily estimate the stability of the Prony method (compare with similar estimates in [4, eq. (19)]).

COROLLARY 2.9. Let the measurements $\{m_k\}$ be given with an error bounded by ε . Denote $u = \kappa(U)$, $b = \kappa(B)$. Assume that $|\xi_i| \leq \Xi$ for all $i = 1, \dots, \mathcal{K}$. Then the Prony method recovers the parameters $\{\xi_j, a_{i,j}\}$ with the following accuracy as $\varepsilon \rightarrow 0$:

$$|\Delta \xi_j| \sim (u^2 b \varepsilon)^{\frac{1}{l_j}} + O\left(\varepsilon^{\frac{2}{l_j}}\right),$$

$$|\Delta a_{i,j}| \sim C(\Xi) u (u^2 b \varepsilon)^{\frac{1}{\max_j l_j}} + L.O.T.,$$

where $C(\Xi)$ is a constant depending on the number Ξ .

Proof. Using the factorization of Lemma 2.6, we obtain that $\kappa(M_C) \leq u^2b$. Therefore, according to (2.8), the coefficient vector $\mathbf{q} = (q_0, \dots, q_{C-1})$ is recovered with the accuracy

$$\begin{aligned} \|\delta\mathbf{q}\| &\sim \frac{\kappa(M_C)}{1 - \kappa(M_C)\delta M_C} \cdot (\delta M_C + \delta\mathbf{m}) \\ &\leq \frac{u^2b\varepsilon}{1 - u^2b\varepsilon} \sim u^2b\varepsilon + O(\varepsilon^2). \end{aligned}$$

The parameters ξ_1, \dots, ξ_K are the roots of the polynomial with coefficient vector \mathbf{q} , with multiplicities l_1, \dots, l_K . Therefore, by the general theory of stability of polynomial roots (see, e.g., [47]) it is known that $\Delta\xi_j \sim (\delta\mathbf{q})^{\frac{1}{l_j}}$. The first part of the claim is thus proved.

Now consider the linear system (2.5) for recovering the jump magnitudes. Note that the matrix U is known only approximately. Again, by (2.8) we have

$$(2.9) \quad \delta\mathbf{a} \sim \frac{\kappa(U)}{1 - \kappa(U)\delta U} (\delta U + \delta\mathbf{m}).$$

Assuming that $|\xi_j| \leq \Xi$, it is easy to see that $\delta U \sim C(\Xi) (u^2b\varepsilon)^{\frac{1}{\max_j l_j}}$. Plugging this value into (2.9), we get the desired result. \square

Inverses of confluent Vandermonde matrices and their condition numbers are extensively studied in numerical linear algebra (e.g., [12, 13, 22]).³ In general, $\kappa(U)$ will grow exponentially with K and will also depend on the ‘‘node separation’’ $\prod_{i \neq j} |\xi_j - \xi_i|^{-1}$. As for $\kappa(B)$, we are not aware of a general formula except for the simplest cases.⁴

Finally, notice that the stability estimates of Corollary 2.9 suggest that when the Prony method is used, the parameters of the problem are ‘‘coupled’’ to each other,

³In particular, the paper [22, Theorem 3] contains the following estimate for the norm of $\{U(\xi_1, 1, \dots, \xi_K, 1)\}^{-1}$ when the nodes are arbitrary complex numbers:

$$\|U^{-1}\|_\infty \leq \max_{1 \leq i \leq K} b_i \prod_{j=1, j \neq i}^K \left(\frac{1 + |\xi_j|}{|\xi_i - \xi_j|} \right)^2,$$

where

$$b_i \stackrel{\text{def}}{=} \max \left(1 + |\xi_i|, 1 + 2(1 + |\xi_i|) \sum_{j \neq i} \frac{1}{|\xi_j - \xi_i|} \right).$$

⁴The following are estimates of the spectral condition numbers:

- For the standard Prony system we have

$$\kappa(B) = \frac{\max_j |a_{j,0}|}{\min_j |a_{j,0}|}.$$

- For multiplicity 1 confluent system, assuming $a_{j,1} \neq 0$ and denoting $\mu_j \stackrel{\text{def}}{=} \frac{a_{j,0}}{a_{j,1}}$, brute force calculation gives

$$\kappa(B) = \frac{\max_j \sqrt{\frac{\mu_j^2 + 2 + \mu_j \sqrt{\mu_j^2 + 4}}{\mu_j^2 + 2 - \mu_j \sqrt{\mu_j^2 + 4}}}}{\min_j \sqrt{\frac{\mu_j^2 + 2 + \mu_j \sqrt{\mu_j^2 + 4}}{\mu_j^2 + 2 - \mu_j \sqrt{\mu_j^2 + 4}}}}.$$

in the sense that the accuracy of recovering either a node ξ_i or a magnitude $a_{i,j}$ will depend on the values of *all the parameters at once*. This undesired behavior is confirmed by our numerical experiments in section 6.

3. Measurement set and the Prony map. Assume that the number of measurements is $S \geq R$ (where R is the overall number of parameters in the confluent Prony system). Then we define $\mathcal{M}_{R,S}$ to be the set⁵ of all possible exact measurements, i.e.,

$$\mathcal{M}_{R,S} \stackrel{\text{def}}{=} \left\{ (m_0, m_1, \dots, m_{S-1}) : m_k = \sum_{i=1}^{\mathcal{K}} \sum_{j=0}^{l_i-1} a_{i,j}(k)_j \xi_i^{k-j}, a_{i,j} \in \mathbb{C}, \xi_j \in \mathbb{C} \right\} \\ \subset \mathbb{C}^S.$$

This $\mathcal{M}_{R,S}$ is the image of \mathbb{C}^R under the ‘‘Prony map’’ $\mathcal{P}_S : \mathbb{C}^R \rightarrow \mathbb{C}^S$ defined as

$$(3.1) \quad \mathcal{P}_S(\{a_{ij}\}, \{\xi_i\}) = (m_0, m_1, \dots, m_{S-1}) : m_k = \sum_{i=1}^{\mathcal{K}} \sum_{j=0}^{l_i-1} a_{i,j}(k)_j \xi_i^{k-j}.$$

Now let $\mathbf{x} = \{\{a_{ij}\}, \{\xi_i\}\} \in \mathbb{C}^R$ be an unknown parameter vector and $\mathbf{y} = \mathcal{P}_S(\mathbf{x}) \in \mathcal{M}_{R,S}$ its corresponding exact measurement vector. The absolute error in each measurement is bounded from above by ε ; therefore, the actual measurement satisfies $\tilde{\mathbf{y}} \in B(\mathbf{y}, \varepsilon)$. Now consider the set

$$T_{\tilde{\mathbf{y}}, \varepsilon} \stackrel{\text{def}}{=} \mathcal{M}_{R,S} \cap B(\tilde{\mathbf{y}}, \varepsilon)$$

of all possible noise-free measurements corresponding to the given noisy one $\tilde{\mathbf{y}}$. Any algorithm which receives this $\tilde{\mathbf{y}}$ as input will therefore produce worst-case error, which is at least

$$\frac{1}{2} \text{diam } \mathcal{P}_S^{-1}(T_{\tilde{\mathbf{y}}, \varepsilon}),$$

where \mathcal{P}_S^{-1} denotes the full preimage set.

This prompts us to make the following definition.

DEFINITION 3.1. Assign to each one of the parameters $\{a_{ij}\}, \{\xi_i\}$ a unique index $1 \leq p \leq R$. The best possible pointwise accuracy of solving the noisy confluent Prony system (1.5) with each noise component bounded above by ε at the point $\mathbf{x} = (\{a_{ij}\}, \{\xi_i\}) \in \mathbb{C}^R$ with respect to the parameter p is defined to be

$$\text{ACC}(\mathbf{x}, \varepsilon, p) \stackrel{\text{def}}{=} \sup_{\tilde{\mathbf{y}} \in B(\mathcal{P}_S(\mathbf{x}), \varepsilon)} \frac{1}{2} \text{diam}_p \mathcal{P}_S^{-1}(\mathcal{M}_{R,S} \cap B(\tilde{\mathbf{y}}, \varepsilon)),$$

where $\text{diam}_p A$ is the diameter of the set A along the dimension p .

Obviously, $\text{ACC}(\mathbf{x}, \varepsilon)$ will depend on the point $\mathbf{x} \in \mathbb{C}^R$ in a nontrivial way because the chart \mathcal{P}_S is nonlinear. Calculation of the function ACC may be considered as one possible answer to the stability problem posed in the introduction.

⁵Formally, $\mathcal{M}_{R,S}$ is a projection of the complex algebraic variety defined by the set of the S confluent Prony equations onto the corresponding S coordinate axes. If all parameters are real-valued, this is a semialgebraic set.

4. Local accuracy. Having given the general definition of accuracy, in the remainder of this paper we restrict ourselves to the “local” setting in the following sense: we assume that ε is small enough so that the set $\mathcal{M}_{R,S}$ can be approximated by the linear part of the Prony map, and furthermore we take $S = R$ so that the preimage will be given by the usual inverse function. For such an analysis to be valid, it should be done at noncritical points of \mathcal{P}_S so that this map is locally invertible. By definition, the point \mathbf{x} is a critical point of \mathcal{P}_S if the Jacobian determinant of \mathcal{P}_S vanishes at \mathbf{x} .

To summarize, let us give the following definition of the local accuracy which is nothing more than the first-order Taylor approximation to the inverse function $\mathcal{N} = \mathcal{P}_S^{-1}$ at a regular point of \mathcal{P}_S .

DEFINITION 4.1. Assume $S = R$. Let $\mathbf{x} = (\{a_{ij}\}, \{\xi_i\}) \in \mathbb{C}^R$ be a regular point of \mathcal{P}_S , and assume ε to be small enough so that that the inverse function $\mathcal{N} = \mathcal{P}_S^{-1}$ exists in the ε -neighborhood of $\mathbf{y} = \mathcal{P}_S(\mathbf{x})$. Assign, as before, to each one of the parameters $\{a_{ij}\}, \{\xi_i\}$ a unique index $1 \leq p \leq R$. The best possible local pointwise accuracy of solving the noisy confluent Prony system (1.5) with each noise component bounded above by ε at the point \mathbf{x} with respect to the parameter p is

$$ACC_{LOC}(\mathbf{x}, \varepsilon, p) \stackrel{\text{def}}{=} \sup_{\tilde{\mathbf{y}} \in B(\mathbf{y}, \varepsilon)} \left| [\mathcal{J}_{\mathcal{N}}(\mathbf{y})(\tilde{\mathbf{y}} - \mathbf{y})]_p \right|,$$

where $\mathcal{J}_{\mathcal{N}}(\mathbf{y})$ is the Jacobian of \mathcal{N} at the point \mathbf{y} and $[v]_p$ is the p th component of the vector \mathbf{v} .

In Theorem 4.5 below we estimate the function ACC_{LOC} . The key technical tool is the following factorization of the Jacobian of \mathcal{P}_S , which separates the nonlinear part depending on the nodes $\{\xi_j\}$ from the linear part, which depends on the magnitudes $\{a_{i,j}\}$.

LEMMA 4.2. Let $\mathbf{x} = (\{a_{ij}\}, \{\xi_i\}) \in \mathbb{C}^R$. Then

$$(4.1) \quad \mathcal{J}_{\mathcal{P}_S}(\mathbf{x}) = U(\xi_1, l_1 + 1, \dots, \xi_{\mathcal{K}}, l_{\mathcal{K}} + 1) \cdot \text{diag}\{D_1, \dots, D_{\mathcal{K}}\},$$

where $U(\dots)$ is the confluent Vandermonde matrix (2.3), and D_i is the $(l_i+1) \times (l_i+1)$ block

$$(4.2) \quad D_i \stackrel{\text{def}}{=} \begin{bmatrix} 1 & 0 & 0 & \cdots & 0 \\ 0 & 1 & 0 & \cdots & a_{i,0} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \cdots & a_{i,l_i-1} \end{bmatrix}.$$

Proof. We have by (3.1)

$$\begin{aligned} \frac{\partial m_k}{\partial a_{ij}} &= (k)_j \xi_i^{k-j}, \\ \frac{\partial m_k}{\partial \xi_i} &= \sum_{j=0}^{l_i-1} a_{ij} (k)_j (k-j) \xi_i^{k-(j+1)} = \sum_{j=1}^{l_i} a_{i,j-1} (k)_j \xi_i^{k-j}. \end{aligned}$$

The rest of the proof is just a straightforward calculation. \square

COROLLARY 4.3. $\mathbf{x} = (\{a_{ij}\}, \{\xi_i\}) \in \mathbb{C}^R$ is a critical point of \mathcal{P}_S if and only if at least one of the following conditions is satisfied:

1. $\xi_i = \xi_j$ for any pair of indices $i \neq j$.
2. $a_{i,l_i-1} = 0$ for any $1 \leq i \leq \mathcal{K}$.

COROLLARY 4.4. Let $\mathbf{x} \in \mathbb{C}^R$ be a regular point of \mathcal{P}_S . Then the Jacobian matrix of the inverse function $\mathcal{N} = \mathcal{P}_S^{-1}$ at $\mathbf{y} = \mathcal{P}_S(\mathbf{x})$ is equal to

$$\begin{aligned} \mathcal{J}_{\mathcal{N}}(\mathbf{y}) &= \{\mathcal{J}_{\mathcal{P}_S}(\mathbf{x})\}^{-1} = \frac{\partial(a_{10}, \dots, a_{1, l_1-1}, \xi_1, \dots, a_{\mathcal{K}, 0}, \dots, a_{\mathcal{K}, l_{\mathcal{K}}-1}, \xi_{\mathcal{K}})}{\partial(m_0, \dots, m_{R-1})} \\ &= \text{diag}\{D_1^{-1}, \dots, D_{\mathcal{K}}^{-1}\} \cdot U^{-1}(\xi_1, l_1 + 1, \dots, \xi_{\mathcal{K}}, l_{\mathcal{K}} + 1), \end{aligned}$$

where

$$(4.3) \quad D_i^{-1} = \begin{bmatrix} 1 & 0 & 0 & \cdots & 0 \\ 0 & 1 & 0 & \cdots & (-1)^{l_i-1} \frac{a_{i,0}}{a_{i, l_i-1}} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \cdots & \frac{1}{a_{i, l_i-1}} \end{bmatrix}.$$

Now we are ready to formulate and prove our local stability result.

THEOREM 4.5. Assume $S = R$. Let $\mathbf{x} = (\{a_{ij}\}, \{\xi_i\}) \in \mathbb{C}^n$ be a regular point of \mathcal{P}_S , and assume ε to be small enough so that the inverse function $\mathcal{N} = \mathcal{P}_S^{-1}$ exists in the ε -neighborhood of $\mathbf{y} = \mathcal{P}_S(\mathbf{x})$.

Then there exists a positive constant C_1 depending only on $\xi_1, \dots, \xi_{\mathcal{K}}$ and $l_1, \dots, l_{\mathcal{K}}$ such that for all $i = 1, \dots, \mathcal{K}$

$$\begin{aligned} \text{ACC}_{\text{LOC}}(\mathbf{x}, \varepsilon, a_{ij}) &= \begin{cases} C_1 \varepsilon, & j = 0, \\ C_1 \varepsilon \left(1 + \frac{|a_{i, j-1}|}{|a_{i, l_i-1}|}\right), & 1 \leq j \leq l_i - 1, \end{cases} \\ \text{ACC}_{\text{LOC}}(\mathbf{x}, \varepsilon, \xi_i) &= C_1 \varepsilon \frac{1}{|a_{i, l_i-1}|}. \end{aligned}$$

Proof. Express the Jacobian matrix $\mathcal{J}_{\mathcal{N}}(\mathbf{y})$ as

$$\mathcal{J}_{\mathcal{N}}(\mathbf{y}) = [\mathbf{s}_{10}^T \quad \cdots \quad \mathbf{s}_{1, l_1-1}^T \quad \mathbf{t}_1^T \quad \cdots \quad \mathbf{s}_{n0}^T \quad \cdots \quad \mathbf{s}_{\mathcal{K}, l_{\mathcal{K}}-1}^T \quad \mathbf{t}_{\mathcal{K}}^T]^T,$$

where

$$\begin{aligned} \mathbf{s}_{ij} &\stackrel{\text{def}}{=} \left[\frac{\partial a_{ij}}{\partial m_0} \quad \frac{\partial a_{ij}}{\partial m_1} \quad \cdots \quad \frac{\partial a_{ij}}{\partial m_{S-1}} \right], \\ \mathbf{t}_i &\stackrel{\text{def}}{=} \left[\frac{\partial \xi_i}{\partial m_0} \quad \frac{\partial \xi_i}{\partial m_1} \quad \cdots \quad \frac{\partial \xi_i}{\partial m_{S-1}} \right]. \end{aligned}$$

Let $\tilde{\mathbf{y}} = (m_0 + \Delta m_0, \dots, m_{S-1} + \Delta m_{S-1})$, where each $|\Delta m_k| < \varepsilon$. Denote by $\|\cdot\|_1$ the l_1 vector norm; i.e., if $\mathbf{v} = (v_i)$ is an n -vector, then $\|\mathbf{v}\|_1 \stackrel{\text{def}}{=} \sum_{i=1}^n |v_i|$. Then

$$\begin{aligned} [\mathcal{J}_{\mathcal{N}}(\mathbf{y})(\tilde{\mathbf{y}} - \mathbf{y})]_{a_{ij}} &= \left| \sum_{k=0}^{P-1} \frac{\partial a_{ij}}{\partial m_k} \Delta m_k \right| \leq \varepsilon \|\mathbf{s}_{ij}\|_1, \\ [\mathcal{J}_{\mathcal{N}}(\mathbf{y})(\tilde{\mathbf{y}} - \mathbf{y})]_{\xi_i} &= \left| \sum_{k=0}^{P-1} \frac{\partial \xi_i}{\partial m_k} \Delta m_k \right| \leq \varepsilon \|\mathbf{t}_i\|_1. \end{aligned}$$

By Corollary 4.4, the matrix $\mathcal{J}_{\mathcal{N}}$ is the product of the block diagonal matrix $D^* \stackrel{\text{def}}{=} \text{diag}\{D_1^{-1}, \dots, D_{\mathcal{K}}^{-1}\}$ with the matrix $U^* \stackrel{\text{def}}{=} (U(\xi_1, l_1 + 1, \dots, \xi_{\mathcal{K}}, l_{\mathcal{K}} + 1))^{-1}$. Therefore,

\mathbf{s}_{ij} and \mathbf{t}_i are the products of the corresponding rows of D_i^{-1} with U^* . Let $D_i^{-1} = (d_{k,l}^{(i)})$ and $U^* = (u_{k,l})$. Then

$$\|\mathbf{s}_{ij}\|_1 = \sum_{k=1}^P \left| \sum_{l=1}^{l_i+1} d_{j,l}^{(i)} u_{l,k} \right| \leq \sum_{l=1}^{l_i+1} |d_{j,l}^{(i)}| \sum_{k=1}^P |u_{l,k}|,$$

and likewise

$$\|\mathbf{t}_i\|_1 \leq \sum_{l=1}^{l_i+1} |d_{l_i+1,l}^{(i)}| \sum_{k=1}^P |u_{l,k}|.$$

Let $\|\cdot\|_\infty$ denote the “maximal row sum” matrix norm; i.e., for any $n \times n$ matrix $C = (c_{ij})$ we have $\|C\|_\infty \stackrel{\text{def}}{=} \max_{i=1,\dots,n} \sum_{j=1}^n |c_{ij}|$.

Denote $C_1 \stackrel{\text{def}}{=} \|U^*\|_\infty$. Then substitute for $d_{l_i,k}^{(i)}$ the actual entries of D_i^{-1} from (4.3) into the above, and get the desired result. \square

5. Comparison with known results.

5.1. CRB for PACE model. The confluent Prony system (1.5) is equivalent to the PACE model [4, 5]. The Cramer–Rao bound (CRB) (which gives a lower bound for the variance of any unbiased estimator) of the PACE model in colored Gaussian noise is as follows (note that the original expressions have been appropriately modified to match the notation of this paper).

THEOREM 5.1 (see [5, Proposition III.1]). *Let the noise variance be σ^2 ; then⁶*

$$\begin{aligned} \text{CRB}\{\xi_i\} &= C_2 \frac{\sigma^2}{|\xi_i|^2 |a_{i,l_i-1}|^2}, \\ \text{CRB}\{a_{i,0}\} &= C_3 \sigma^2, \\ \text{CRB}\{a_{i,j}\} &= C_4 \sigma^2 \left(C_5 \left| \frac{a_{i,j-1}}{a_{i,l_i-1}} \right|^2 + C_6 \Re \left\{ \frac{a_{i,j-1}}{a_{i,l_i-1}} \right\} + 1 \right), \quad j = 1, 2, \dots, l_i - 1, \end{aligned}$$

where C_2, \dots, C_6 are constants depending on the configuration of the nodes $\{\xi_i\}$, while in addition C_4, C_5, C_6 depend on the index j .

As mentioned in subsection 1.2, there exist several essential differences between our setting and the statistical signal estimation framework; in particular,

1. no a priori statistical model of the noise is available;
2. no assumptions on the reconstruction algorithm (estimator) such as unbiasedness are made; and
3. the measure of performance is the worst-case error rather than estimator variance.

The expressions for the CRB in Theorem 5.1 are very similar to the local pointwise accuracy bounds of Theorem 4.5. The reason for such similarity is not a priori clear (although it could be partially attributed to the fact that both methods require calculation of the partial derivatives of the measurements with respect to the parameters), and it certainly warrants further investigation.

⁶Here $\Re(\cdot)$ denotes the real part.

5.2. ESPRIT method. The ESPRIT algorithm is one of the best performing subspace methods for estimating parameters of the Prony systems with white Gaussian noise. Originally developed in the context of frequency estimation [43, section 4.7], it has been generalized to the full PACE model [4], and its performance has been shown to approach the CRB in the case of high signal-to-noise ratio (SNR) and infinite observation length.

In essence, the ESPRIT (and other subspace methods) relies on the following observations:

1. The range (column space) of both the data matrix M_C (2.1) and the confluent Vandermonde matrix U (2.3) are the same (follows directly from (2.6)).
2. The matrix U has the so-called *rotational invariance property* [4]:

$$U^\uparrow = U_\downarrow J,$$

where U^\uparrow denotes U without the first row, U_\downarrow denotes U without the last row, and J is a block diagonal matrix whose i th block is the $l_i \times l_i$ Jordan block with the number ξ_i on the diagonal.

Suppose we know U ; then the matrix J could be found by

$$J = U_\downarrow \# U^\uparrow$$

(where $\#$ denotes the Moore–Penrose pseudoinverse), and then the nodes ξ_j could be recovered as the eigenvalues of J .

Unfortunately, U is unknown in advance, but suppose we had at our disposal a matrix W whose column space was identical to that of U . In that case, we would have $W = UG$ for an invertible G , and consequently

$$W^\uparrow = W_\downarrow \Phi,$$

where

$$\Phi = G^{-1} J G,$$

which means that the eigenvalues of Φ are also $\{\xi_i\}$. Such a matrix W can be obtained, for example, from the singular value decomposition (SVD) of the data matrix/covariance matrix. To summarize, the ESPRIT method for estimating $\{\xi_i\}$, as used in our experiments below, is as follows.

Algorithm 2 ESPRIT method for recovering the nodes $\{\xi_i\}$.

Let M_S be a rectangular $n \times l$ Hankel matrix built from the measurements.

1. Compute the SVD $M_S = W \Sigma V^T$.
 2. Calculate $\Phi = W_\downarrow \# W^\uparrow$.
 3. Set $\{\xi_i\}$ to be the eigenvalues of Φ with appropriate multiplicities (use, e.g., arithmetic means to estimate multiple nodes which are scattered by the noise).
-

Note that the dimensions n, l are not fixed a priori, but in [6] it is shown that taking $n = 2l$ or $l = 2n$ results in optimal performance for nonconfluent Prony system (1.1).

Since the performance of the ESPRIT method is close to the CRB, which, in turn, resembles our local bounds, we regard the ESPRIT as the best candidate among the “global” solution methods of the confluent Prony system. It should be noted, however, that the analysis of ESPRIT as presented in [6] suggests a relatively complicated dependence of the estimator performance on the model parameters for a small number of measurements S .

5.3. Approximate Prony method. In [36] the authors develop the approximate Prony method for solving the system (1.1) (restricting ξ_j to unit length) and analyze its performance for small measurement errors. In more detail, the model is defined as

$$h(x) = \sum_{j=1}^M c_j e^{if_j x}, \quad x \in \mathbb{R}, c_j \in \mathbb{C}, f_j \in (-\pi, \pi).$$

The measurements are given with errors

$$\tilde{h}(k) = h(k) + e_k, \quad k = 0, \dots, 2N,$$

where the number of measurements N satisfies $N \geq 2M + 1$. Finally, the coefficients c_j are assumed to be large with respect to the noise level, i.e.,

$$|e_k| \leq \varepsilon_1 \ll |c_j|.$$

The proposed solution method is as follows.

Algorithm 3 Approximate Prony method.

1. Build the Hankel matrix $\tilde{H} \in \mathbb{C}^{2N-L, L}$ from the measurements where L is an upper bound on the number of nodes. Compute the SVD of \tilde{H} , and take the smallest nonzero singular value and its singular vector $\mathbf{v} = (v_i)$. Finally, compute the roots of the polynomial $p(z) = \sum_{i=0}^L v_i z^i$. These are the approximations of $\{f_j\}$.
 2. Find $\{c_j\}$ by solving an overdetermined Vandermonde linear system.
-

The stability analysis of the approximate Prony method is performed only for step 2 above, assuming that the frequencies $\{f_j\}$ have been recovered with high accuracy. Potts and Tasche [36, Theorem 5.2] give the following estimate:

$$(5.1) \quad |c_j - \tilde{c}_j| \sim \sqrt{NM} \left| f_j - \tilde{f}_j \right| \max_k |h_k| + \max_k |\Delta h_k|.$$

While missing the explicit analysis of step 1 above (however, the actual numerical accuracy of this step was shown in [34] to be comparable to the performance of the ESPRIT method) and dealing with single poles only, these results may provide an important insight as to the dependence of the accuracy on the number of measurements N , as well as to the applicability of the Vandermonde inversion for recovering the magnitudes (the errors in fact *increase* with $N!$). In addition, the authors notice that the accurate recovery of the magnitudes depends greatly on a sufficient accuracy of recovering the nodes, and this fact is also reflected in our numerical experiments (section 6).

6. Numerical experiments. In our numerical experiments we had two distinct goals:

1. Numerically investigate the “best possible local accuracy” of inverting (1.5) as a function of the various parameters of the problem, and compare the results with the predictions of Theorem 4.5.
2. Ascertain whether there exist some regular patterns in the behavior of the global solution methods (Prony and ESPRIT) in a similar “local” setting, and compare their performance to the optimal one.

6.1. Experimental setup.

1. Given \mathcal{K}, d , choose the jumps $\xi_1, \dots, \xi_{\mathcal{K}} \in [0, 1]$ and the magnitudes $a_{1,0}, \dots, a_{\mathcal{K},d-1} \in [-1, 1]$.
2. Change one or more of the parameters according to a particular experiment.
3. Calculate the perturbed moments $\tilde{m}_k = m_k + \varepsilon_k$, where m_k is given by (1.5) and $\varepsilon_k \ll 1$ (on the order of 10^{-10}) are randomly chosen.
4. Invert (1.5) with the right-hand side given by \tilde{m}_k by one of the following three methods:
 - (a) Nonlinear least squares minimization (using MATLAB `lsqnonlin` routine) with the initial guess being very close to the true parameter values. This is our simulation of the “local” setting.
 - (b) Global Prony method—Algorithm 1.
 - (c) ESPRIT method—Algorithm 2.
5. Calculate the absolute errors $|\Delta\xi_j| = |\xi_j - \tilde{\xi}_j|$ and $|\Delta a_{i,j}| = |a_{i,j} - \tilde{a}_{i,j}|$.

In all the experiments we took $\mathcal{K} = 2$. All solution methods were applied to the same moment sequence $\{m_k\}$. The number of measurements is the minimal necessary for exact inversion, namely, R for least squares and $2C$ both for Prony and ESPRIT.

6.2. Results.

6.2.1. Changing the highest coefficient. In the first set of experiments, we checked how the reconstruction errors $|\Delta\xi_i|, |\Delta a_{i,j}|$ depend on the magnitude of the highest coefficient $|a_{i,l_i-1}|$. The results are presented in Figure 6.1(a)–(c).

For both least squares and ESPRIT (but not for Prony), the inverse proportionality $|\Delta\xi_i| \sim \frac{1}{|a_{i,l_i-1}|}$ is seen in Figure 6.1(a), (c), matching the theoretical predictions of Theorem 4.5.

For LS and ESPRIT, the errors $|\Delta a_{i,j}|$ seem to be unaffected by the increase in $|a_{i,l_i-1}|$. This can be explained very well by the formula $|\Delta a_{i,j}| \sim 1 + \frac{|a_{i,j-1}|}{|a_{i,l_i-1}|}$ so that indeed $|\Delta a_{i,j}|$ should remain close to constant as $|a_{i,l_i-1}| \rightarrow \infty$.

The Prony method’s performance with respect to the recovery of the magnitudes actually *degrades* with the increase in $|a_{i,l_i-1}|$. Although both Prony and ESPRIT use the same method for the recovery of the magnitudes, it appears that the initial error in recovering the nodes, which is significantly smaller in ESPRIT (see subsection 6.2.3 below), influences this step greatly—in accordance with the predictions of [36, 34] (see also the discussion in subsection 5.3).

In addition, the Prony method fails to separate recovery of a node and its magnitudes (say, $\Delta\xi_1, \Delta a_{1,j}$) from the highest magnitude associated with *another* node (e.g., $|a_{2,l_2-1}|$); these results are not shown for the purpose of saving space.

6.2.2. Changing coefficient other than the highest. In the second set of experiments, we changed the magnitude of some coefficient other than the highest, i.e., $a_{i,j}$ for $j < l_i - 1$. The results are presented in Figure 6.1(d)–(f).

For the least squares method, the dependence of $|\Delta a_{i,j}|$ on the “previous” magnitude $|a_{i,j-1}|$ for $j \neq 0$ is consistent with the formula $|\Delta a_{i,j}| \sim 1 + \frac{|a_{i,j-1}|}{|a_{i,l_i-1}|}$; such behavior should be visible when $|a_{i,j-1}| \gg |a_{i,l_i-1}|$, as can indeed be noticed in Figure 6.1(d). In addition, the other magnitudes and the jumps are unaffected, as predicted.

On the contrary, neither Prony nor ESPRIT succeeds in confining the influence of $|a_{i,j-1}|$ only to the recovery of the next magnitude $|\Delta a_{i,j}|$. In particular, $|\Delta\xi_1|$ increases with $|a_{1,0}|$ in both of them. The error in *all the magnitudes* grows with $|a_{1,0}|$, as opposed to the least squares, where only $|\Delta a_{1,1}|$ is increased.

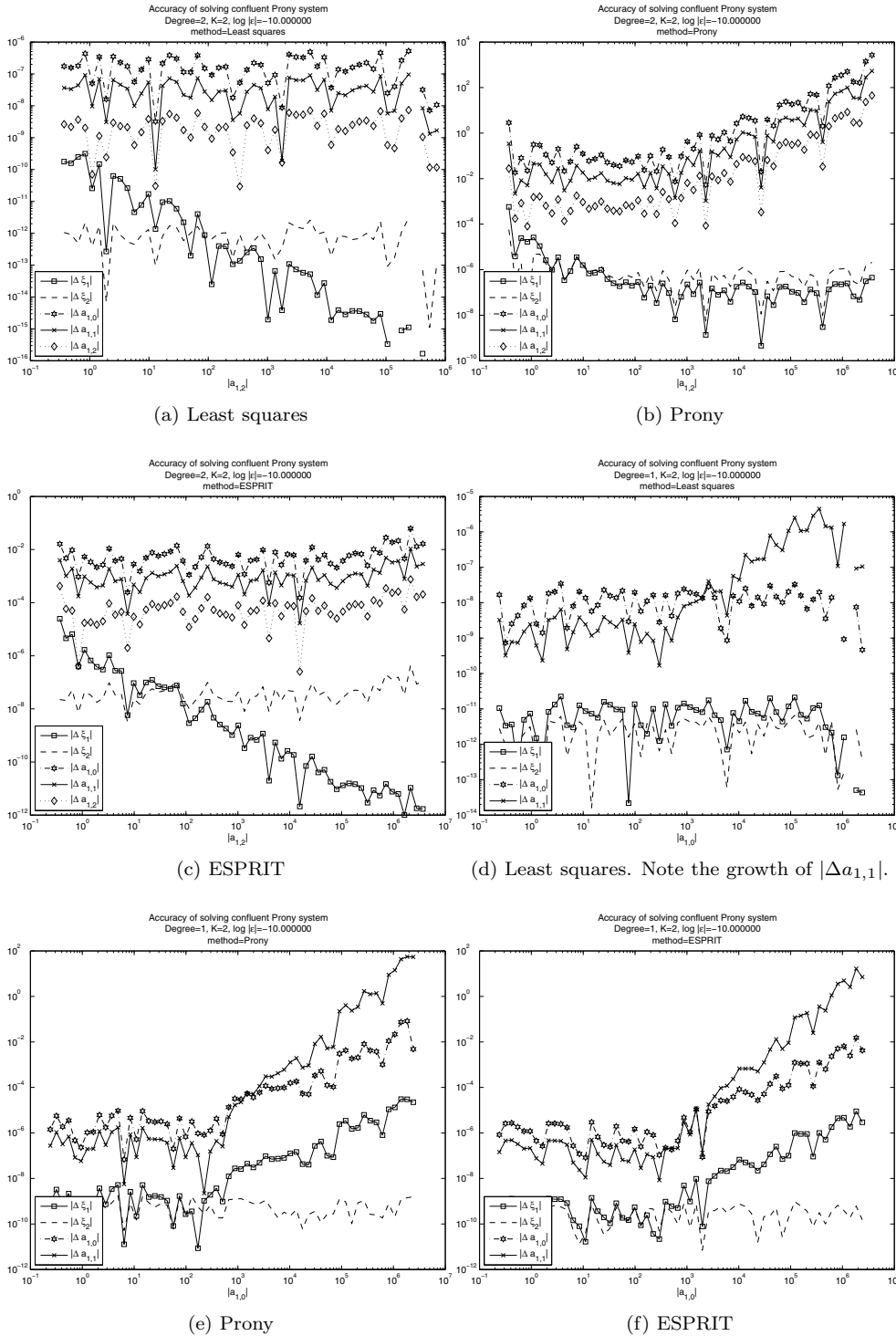


FIG. 6.1. (a)–(c): Dependence of the reconstruction error on the magnitude of the highest coefficient; degree = 2. (d)–(f): Dependence of the reconstruction error on the magnitude of the “previous” coefficient; degree = 1.

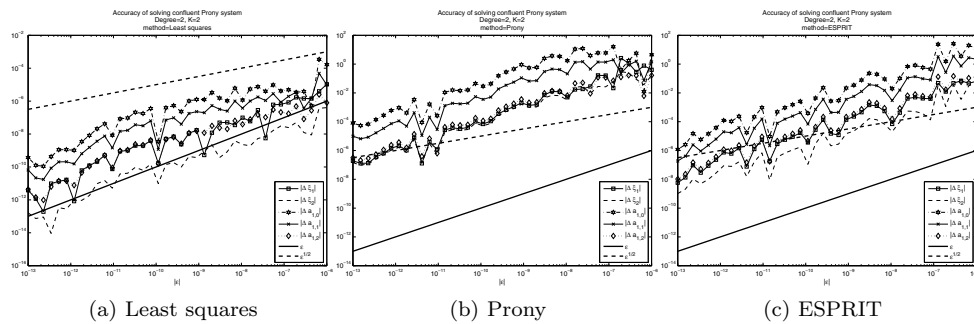
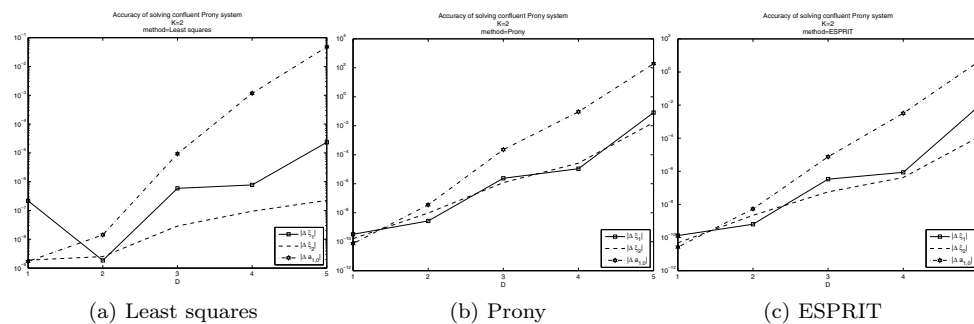
FIG. 6.2. Reconstruction error as $\varepsilon \rightarrow 0$, degree = 2.

FIG. 6.3. Dependence of the reconstruction error on the order of the model.

6.2.3. Dependence on the measurement error. In the next experiment, we kept all the parameters constant and changed the magnitude of the error $\max_k \varepsilon_k$. The results are presented in Figure 6.2. The ESPRIT performs slightly better than Prony, but both of them are worse than the optimal least squares. Note, however, that the asymptotic error (the slope) is $O(\varepsilon)$ in spite of the fact that both algorithms involve extraction of multiple roots, which should decrease the accuracy to $O(\varepsilon^{\frac{1}{d}})$, where d is the order of the pole. This phenomenon can be explained by the effect of averaging the clustered roots (see [4, Proposition V.3]).

6.2.4. Dependence on the model order. Next, we checked the dependence of the reconstruction error on the model order $D \stackrel{\text{def}}{=} \max_{i=1, \dots, \kappa} l_i$. The results are presented in Figure 6.3. The reconstruction error for all the parameters grows exponentially in D for all the methods.

6.2.5. Dependence on the node separation. Finally, we checked the dependence of the reconstruction error on the distance between the two nodes $|\xi_2 - \xi_1|$. The results are presented in Figure 6.4. For all the three methods, the results are consistent with

$$|\Delta \xi_i|, |\Delta a_{i,j}| \sim |\xi_2 - \xi_1|^{-D}.$$

6.3. Conclusions. In the numerical experiments we have investigated the “best possible local accuracy” via the least squares method, comparing it both with the theoretical results of Theorem 4.5 and with the performance of two “global” solution

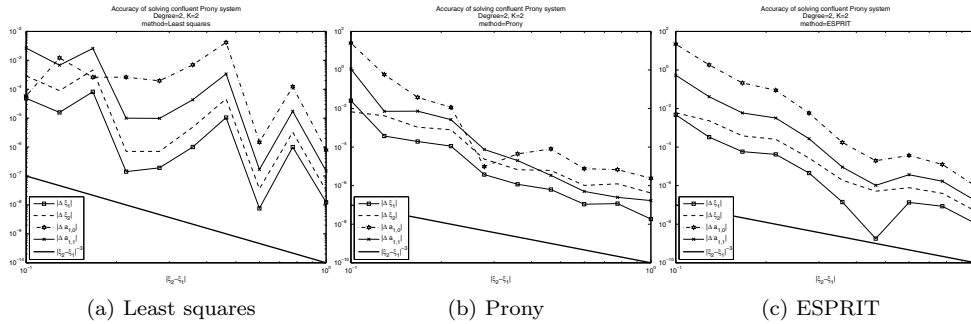


FIG. 6.4. Dependence of the reconstruction error on the node separation.

techniques, namely, Prony and ESPRIT methods, for small perturbations (high SNR). Our results suggest the following.

1. The numerical behavior of the solution in the case of small data perturbations indeed exhibits the patterns predicted by Theorem 4.5, in particular the qualitative dependence of the reconstruction error on the values of the parameters of the problem.
2. The Prony solution method largely fails to separate the parameters which could be separated in theory. Furthermore, its performance actually *degrades* when the highest coefficient $|a_{i,l_i-1}|$ is increased. ESPRIT separates the parameters better than Prony but is still worse than optimal.
3. In terms of absolute reconstruction error, ESPRIT is better than Prony but still worse than the optimal least squares.
4. In terms of dependence of the reconstruction error on the model order and the node separation, both Prony and ESPRIT behave close to the predicted law, namely, exponential increase in the order and polynomial increase in the separation distance.

7. Discussion. We believe that the analytical approach of this paper has the potential to provide relatively complete answers to several important questions related to stable solution of Prony-type systems, as briefly discussed below.

The numerical experiments suggest that the least squares method approximates the optimal “local” behavior very well. However, it is well known that a very accurate initial approximation is required in order to find the global minima. It is customary to use one of the global solution methods to obtain such an initial value. Further analysis of the Prony sets $\mathcal{M}_{R,S}$ may provide explicit conditions for such an initialization to be sufficiently close to the true solution.

The general case $S \geq R$ should be well understood in order to estimate the feasibility of taking more measurements than strictly needed (oversampling). Without assumptions on the noise, it is not a priori obvious that averaging should improve the accuracy in any way. Again, such an understanding is hopefully achievable via the investigation of $\mathcal{M}_{R,S}$ with $S \gg R$.

In practice it is often the case that neither the number of nodes \mathcal{K} nor the numbers $\{l_i\}$ are known a priori, but only *their upper* bounds are. In this case, given a noisy measurement vector, more than one “explanation” is possible for this data, in which case a good reconstruction algorithm needs somehow to select the “optimal” configuration. One possible way to achieve this goal is to characterize, for each configuration

of the system (i.e., $\{\mathcal{K}, \{l_i\}_{i=1}^{\mathcal{K}}\}$), the “stable regions” of the corresponding measurement sets $\mathcal{M}_{R,S}$ for which the accuracy function \mathcal{ACC} does not exceed a predefined upper bound. Based on the initial measurement $\tilde{\mathbf{y}} \in \mathbb{C}^S$ and the error bound ε , an algorithm would choose the closest “stable measurement set,” i.e., select a configuration for which the local accuracy is optimal. Using this approach, collision of two nodes ξ_i, ξ_j can in principle be handled in a stable way by substituting the configuration $\{\mathcal{K}, \{l_i\}_{i=1}^{\mathcal{K}}\}$ with $\{\mathcal{K} - 1, \{l_1, \dots, l_i + l_j, \dots, l_{\mathcal{K}}\}\}$ once the measurement vector leaves the stability region associated with the former configuration. In this regard, we note that such a singular behavior has been studied in [48] (see also [33]), where it is shown that if the solution is represented in the basis of divided differences, then the inverse operator is uniformly bounded with respect to the corresponding expansion coefficients. Analogous developments for extraction of multiple roots of polynomials [49] might be very relevant as well.

In order to achieve the above goals, we propose computing the function \mathcal{ACC} as accurately as possible. For that purpose, more detailed analysis of the Prony map⁷ is necessary. In particular, its essential nonlinearity should be quantified using the second-order terms in the Taylor expansion.

In addition to (1.5), other generalizations of the basic Prony system (1.1) appear in applications. One such extension arises in Eckhoff’s method [21] for reconstructing piecewise-smooth functions from Fourier coefficients. There, an additional parameter appears: namely, the measurements m_k are given starting from some large index $k = M$. In [11], we presented an algorithm for solving this system with high accuracy (in the sense of asymptotic rate of convergence as $M \rightarrow \infty$). However, the question of “maximal possible accuracy” for this problem is still open. It will be most desirable to reinterpret those results in the sense of global stability bounds for Prony-like systems.

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⁷Its nonconfluent version appears in a paper by Arnol’d [2] under the name “Vandermonde map.”

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