Computation of Electromagnetic Fields Scattered From Objects With Uncertain Shapes Using Multilevel Monte Carlo Method

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Abstract—Computational tools for characterizing electromagnetic scattering from objects with uncertain shapes are needed in various applications ranging from remote sensing at microwave frequencies to Raman spectroscopy at optical frequencies. Often, such computational tools use the Monte Carlo (MC) method to sample a parametric space describing geometric uncertainties. For each sample, which corresponds to a realization of the geometry, a deterministic electromagnetic solver computes the scattered fields. However, for an accurate statistical characterization, the number of MC samples has to be large. In this paper, to address this challenge, the continuation multilevel Monte Carlo (CMLMC) method is used together with a surface integral equation solver. The CMLMC method optimally balances statistical errors due to sampling of the parametric space, and numerical errors due to the discretization of the geometry using a hierarchy of discretizations, from coarse to fine. The number of realizations of finer discretizations can be kept low, with most samples computed on coarser discretizations to minimize computational cost. Consequently, the total execution time is significantly reduced, in comparison to the standard MC scheme.

Index Terms—Fast Fourier transform (FFT), fast multipole method (FMM), integral equation, multilevel Monte Carlo method (MLMC), numerical methods, uncertain geometry, uncertainty quantification.

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I. INTRODUCTION

N UMERICAL methods for predicting radar and scattering cross sections (RCS and SCS) of complex targets find engineering applications ranging from microwave remote sensing soil/ocean surface and vegetation [1], [2] to enhancing Raman spectroscopy using metallic nanoparticles [3], [4]. When the target size is comparable to or larger than the wavelength at the operation frequency, the scattered field is a strong function of the target shape. However, in many of the applications, whether the target is a (vegetated) rough surface or a nanoparticle, its exact shape may not be known and has to be parameterized using a stochastic/statistical model. Consequently, computational tools, which are capable of generating statistics of a quantity of interest (QoI) (RCS or SCS in this case) given a geometry described using random variables/parameters, are required.

These computational tools often use the Monte Carlo (MC) method [5]–[11]. The MC method is nonintrusive and straightforward to implement; therefore, its incorporation with an existing deterministic EM solver is rather trivial. However, the traditional MC method has an error convergence rate of $\mathcal{O}(N^{-1/2})$ [12], where N is the number of samples in the parametric space used for describing the geometry. Provided more regularity of the QoI w.r.t. the geometry parameters, quasi-MC methods may have a better convergence rate, $\mathcal{O}(N^{-1})$ with a multiplicative log-term that depends on the number of parameters [12], [13]. Both the traditional MC and quasi-MC methods require large N to yield accurate statistics of the QoI and are computationally expensive since the function evaluation at each sampling point, which corresponds to the computation of the QoI for one realization of the deterministic problem, requires the execution of a simulation. For practical real-life scattering scenarios, each of these simulations may take a few hours, if not a few days.

In recent years, schemes that make use of surrogate models have received significant attention as potential alternatives to the MC method [14]–[20]. The surrogate model is generated using the values of the QoI that are computed by the simulator at a small number of "carefully selected" (collocation) points in the parametric space. The surrogate model is then probed using the MC method to obtain statistics of the QoI. Consequently, the surrogate model-based schemes are more efficient than the MC method that operates directly on the computationally expensive

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simulator. The efficiency and accuracy of these schemes leveraging generalized polynomial chaos expansion (gPCE) [21]–[23] can be improved using a refinement strategy that adaptively divides the parametric space into smaller domains, in each of which a separate collocation scheme is used [16], [17]. Additionally, if the QoI can be approximated in terms of low-order contributions from only a few of the parameters, one can use high-dimensional model representation expansions to accelerate the generation of the surrogate model [18]. The surrogate model-based schemes and their accelerated variants have been successfully applied to certain stochastic EM problems [24]– [32]. On the other hand, for the problem of scattering from geometries with uncertain shapes, all parameters and their highorder combinations contribute significantly to the QoI limiting the efficient use of a surrogate model [33]–[35].

Recently, the multilevel Monte Carlo (MLMC) methods have seen increasing use due to their efficiency, robustness, and simplicity [36]-[42]. The MLMC methods operate on a hierarchy of meshes and perform most of the simulations using low-fidelity models (coarser meshes) and only a few simulations using highfidelity models (finer meshes). By doing so, their cost becomes significantly smaller than the cost of the traditional MC methods using only high-fidelity models. The continuation multilevel Monte Carlo (CMLMC) algorithm [42] uses the MLMC method to calibrate the average cost per sampling point and the corresponding variance using Bayesian estimation, taking particular notice of the deepest levels of the mesh hierarchy, to minimize the computational cost. To balance discretization and statistical errors, the CMLMC method estimates how the discretization error and the computational cost depend on the mesh level and uses this information to select optimal numbers of levels and samples on each level.

Surrogate-model-based schemes leveraging gPCE [14], [15], [43] (as briefly described above) could be an alternative to the CMLMC method in general. If the response/surrogate model is smooth and simple, then the collocation scheme used for computing/finding the coefficients of the gPCE requires only a few deterministic solver executions. However, gPCE-based surrogate models suffer from several drawbacks: it is not clear how to determine the number of terms in the gPCE, how to avoid aliasing effects/errors by computing higher order polynomials, and how to control the accuracy of the integration needed for computing the gPCE coefficients. In addition, as mentioned above, specifically for the scattering problem, surrogate models lose their efficiency. Ideas from CMLMC and gPCE methods could be combined, but it goes out of the scope of this paper.

In this paper, a computational framework, which makes use of the CMLMC method to efficiently and accurately characterize EM wave scattering from dielectric objects with uncertain shapes, is proposed. The deterministic simulations required by the CMLMC method to compute the samples at different levels are carried out using the Poggio–Miller–Chan–Harrington– Wu–Tsai surface integral equation (PMCHWT-SIE) solver [44]. The PMCHWT-SIE is discretized using the method of moments (MoM) and the iterative solution of the resulting matrix system is accelerated using a (parallelized) fast multipole method (FMM)–fast Fourier transform (FFT) scheme [45]–[51].



Fig. 1. Description of the scattering problem.

Numerical results, which demonstrate the accuracy, efficiency, and convergence of the proposed computational framework, are presented. The developed computational framework is proven effective not only in studying the effects of uncertainty in the geometry on scattered fields but also in increasing the robustness of the FMM–FFT accelerated PMCHWT-SIE solver by testing its convergence for a large set of scenarios with deformed geometries and varying mesh densities, quadrature rules, iterative solver tolerances, and FMM parameters.

Remark 1: Reasons for this specific choice of SIE formulation (namely PMCHWT) can be explained as follows. One can define two types of "convergence" for the SIE formulation used within the CMLMC framework: 1) convergence of the iterative solver, and 2) convergence of the solution as mesh is made denser. The PMCHWT formulation produces a first-kind SIE, and it is well known that there are other formulations that produce second-kind SIEs, which are better behaved under convergence type (2). However, results obtained by solving these SIEs are less accurate than those of the PMCHWT formulation [52]. Additionally, PMCHWT-SIE is better behaved under convergence type (1), i.e., its solution gets more accurately consistently as mesh is made denser. Having said that, it is clear that the convergence type (1) might have a significant effect on the efficiency of the CMLMC. Use of second-kind SIEs within the CMLMC framework will be investigated elsewhere.

II. FORMULATION

This section describes a computational framework for characterizing scattering from dielectric objects with uncertain shapes (Fig. 1). It is assumed that the shape of the scatterer can be parameterized by means of random variables. The QoI is the SCS over a user-defined solid angle (i.e., a measure of far-field scattered power in a cone). Section II-A describes the MLMC and CMLMC methods, Section II-B formulates a scheme to parameterize and generate objects with uncertain shapes, and Section II-C outlines the FMM–FFT accelerated PMCHWT-SIE solver used for computing the SCS of a given object.

A. MLMC and CMLMC Methods

This section describes elements of the MLMC and CMLMC methods relevant to the characterization of scattering from objects with uncertain shapes. A more indepth description of these techniques can be found in [42].

Let $\boldsymbol{\xi}$ and $g(\boldsymbol{\xi})$ represent the vector of random parameters/variables and the QoI, respectively. The goal of the MLMC method is to approximate the expected value, E[g], to a guaranteed tolerance TOL with predefined probability, and minimal computational cost. To achieve this, the method constructs a telescoping sum, defined over a sequence of mesh levels $\ell = 0, \dots, L$ as described next.

Let $\{P_\ell\}_{\ell=0}^L$ and $\{h_\ell\}_{\ell=0}^L$ be sequences of meshes that discretize the randomly generated object's surface and their average element edge sizes, respectively. It is assumed $\{P_\ell\}_{\ell=0}^L$ are generated hierarchically with $h_\ell = h_0 \beta^{-\ell}$ for $h_0 > 0$ and a constant $\beta > 1$. A method to such meshes for the purpose of illustrating the proposed technique is described in Section II-B.

Let $g_{\ell}(\boldsymbol{\xi})$ represent the approximation to $g(\boldsymbol{\xi})$ computed using mesh P_{ℓ} . The MLMC method expresses $E[g_L]$, the expected value of the most accurate approximation g_L , using

$$\mathbf{E}\left[g_L\right] = \sum_{\ell=0}^{L} \mathbf{E}\left[G_\ell\right] \tag{1}$$

where G_{ℓ} is defined as

$$G_{\ell} = \begin{cases} g_0 & \text{if } \ell = 0\\ g_{\ell} - g_{\ell-1} & \text{if } \ell > 0. \end{cases}$$
(2)

Note that g_{ℓ} and $g_{\ell-1}$ are computed using the same input random parameter $\boldsymbol{\xi}$.

In the telescoping sum (1), the expected values in practice are replaced by sample averages, i.e., $E[G_{\ell}] \approx \tilde{G}_{\ell} = M_{\ell}^{-1} \sum_{m=1}^{M_{\ell}} G_{\ell,m}$, where random variable $G_{\ell,m}$ have the same distribution as G_{ℓ} and are independent identically distributed samples. $G_{\ell,m}$ represents the *m*th random sample of the random output G_{ℓ} . G_{ℓ} and $G_{\ell,m}$ are random variables because they depend on random (input) parameters $\boldsymbol{\xi}$, which is defined in (17).

As ℓ increases, the variance of G_{ℓ} decreases. As a result, the total computational cost can be reduced by approximating $E[G_{\ell}]$ with a smaller number of samples.

The CMLMC algorithm is an improved version of the MLMC method in that it approximates E[g] with a sequence of decreasing tolerances [42]. In doing so, it continuously improves estimates of several problem-dependent parameters, while solving relatively inexpensive problems that by themselves would yield large tolerances. The CMLMC algorithm assumes that the convergence rates for the mean (weak convergence) and variance (strong convergence) follow

$$\mathbf{E}\left[g - g_{\ell}\right] \approx Q_W h_{\ell}^{q_1} \tag{3a}$$

$$\operatorname{Var}\left[g_{\ell} - g_{\ell-1}\right] \approx Q_{S} h_{\ell-1}^{q_{2}} \tag{3b}$$

for $Q_W \neq 0$, $Q_S > 0$, $q_1 > 0$, and $0 < q_2 \le 2q_1$ [42]. Note that variables q_1 and q_2 are solver- and problem-dependent.

For example, the CMLMC algorithm estimates $q_1 \approx 2$ and $q_2 \approx 4$, and $q_1 \approx 3$ and $q_2 \approx 5$ for the examples in Sections III-B and III-C, respectively. These parameter estimates are crucial to optimally distribute the computational effort, as shown below. The total computational cost of the adaptive algorithm is close to that of the MLMC method with correct values of parameters given *a priori*.

For the sake of completeness, the main ingredients of the CMLMC algorithm (described in full in [42]) are stated here. To estimate the number of samples, the algorithm invokes of *cost per sample* and *total cost* as described next.

The CMLMC estimator for the QoI, \mathcal{A} , can be written as $\mathcal{A} = \sum_{\ell=0}^{L} \widetilde{G}_{\ell}$. Let the average cost of generating one sample of G_{ℓ} (cost of one deterministic simulation for one random realization) be

$$W_{\ell} \propto h_{\ell}^{-d\gamma} = h_0^{-d\gamma} \beta^{\ell d\gamma} \tag{4}$$

where d is the spatial dimension and γ is determined by the computational complexity of the deterministic solver. For the FMM–FFT-accelerated PMCHWT-SIE solver used here d = 2 (only surfaces are discretized) and $\gamma \approx 1$ (Sections II-C and III-B). Note that this solver calls for an iterative solution of the MoM system of equations (Section II-C), and that the cost of computing a sample of G_{ℓ} may fluctuate for different realizations depending on the number of iterations required. Finally, for the method used for generating $\{P_{\ell}\}_{\ell=0}^{L}$ (Section II-B), $\beta = 2$.

The total CMLMC computational cost is

$$W = \sum_{\ell=0}^{L} M_{\ell} W_{\ell}.$$
 (5)

The estimator A satisfies a tolerance with a prescribed failure probability $0 < \nu \leq 1$, i.e.,

$$P[|E[g] - \mathcal{A}| \le TOL] \ge 1 - \nu \tag{6}$$

while minimizing W. The total error is split into bias and statistical error

$$|\mathbf{E}\left[g\right] - \mathcal{A}| \leq \underbrace{|\mathbf{E}\left[g - \mathcal{A}\right]|}_{\text{Bias}} + \underbrace{|\mathbf{E}\left[\mathcal{A}\right] - \mathcal{A}|}_{\text{Statistical error}}$$

where $\theta \in (0, 1)$ is a splitting parameter, so that

$$TOL = \underbrace{(1-\theta)TOL}_{\text{Bias tolerance}} + \underbrace{\theta TOL}_{\text{Statistical error tolerance}}.$$
 (7)

The CMLMC algorithm bounds the bias, B = |E[g - A]|, and the statistical error as

$$B = |\mathbf{E}[g - \mathcal{A}]| \le (1 - \theta) \mathrm{TOL}$$
(8)

$$|\mathbf{E}[\mathcal{A}] - \mathcal{A}| \le \theta \mathrm{TOL} \tag{9}$$

where the latter bound holds with probability $1 - \nu$. Note that θ itself can be a variable [42].

To satisfy condition in (9), we require

$$\operatorname{Var}\left[\mathcal{A}\right] \le \left(\frac{\theta \mathrm{TOL}}{C_{\nu}}\right)^2 \tag{10}$$

for some given confidence parameter, C_{ν} , such that $\Phi(C_{\nu}) = 1 - \frac{\nu}{2}$, (see more in [53]); here, Φ is the cumulative distribution function of a standard normal random variable.

By construction of the MLMC estimator, $E[\mathcal{A}] = E[g_L]$, and by independence, $Var[\mathcal{A}] = \sum_{\ell=0}^{L} V_{\ell} M_{\ell}^{-1}$, where $V_{\ell} = Var[G_{\ell}]$. Given *L*, TOL, and $0 < \theta < 1$, and by minimizing *W* subject to the statistical constraint (10) for $\{M_\ell\}_{\ell=0}^L$ gives the following optimal number of samples per level ℓ (apply ceiling function to M_ℓ if necessary):

$$M_{\ell} = \left(\frac{C_{\nu}}{\theta \text{TOL}}\right)^2 \sqrt{\frac{V_{\ell}}{W_{\ell}}} \left(\sum_{\ell=0}^{L} \sqrt{V_{\ell} W_{\ell}}\right).$$
(11)

Summing the optimal numbers of samples over all levels yields the following expression for the total optimal computational cost in terms of TOL:

$$W(\text{TOL}, L) = \left(\frac{C_{\nu}}{\theta \text{TOL}}\right)^2 \left(\sum_{\ell=0}^{L} \sqrt{V_{\ell} W_{\ell}}\right)^2.$$
(12)

The total cost of the CMLMC algorithm can be estimated using Theorem 1 as follows [36], [39], [40], [54], [55].

Theorem 1: Let $d = \{1, 2, 3\}$ denote the problem dimension. Suppose there exist positive constants $q_1, q_2, \gamma > 0$ such that $q_1 \ge \frac{1}{2} \min(q_2, \gamma d)$, and $|\mathbf{E}[g_{\ell} - g]| = \mathcal{O}(h_{\ell}^{q_1})$, $\operatorname{Var}[g_{\ell} - g_{\ell-1}] = \mathcal{O}(h_{\ell}^{q_2})$, and $W_{\ell} = \mathcal{O}(h_{\ell}^{-d\gamma})$. Then for any accuracy TOL and confidence level ν , $0 < \nu \le 1$, there exist a deepest level L(TOL) and number of realizations $\{M_{\ell} \text{ (TOL)}\}$ such that

$$\lim_{\text{TOL}\to 0} \inf P\left[\left(|\mathbb{E}\left(g\right) - \mathcal{A}| \le \text{TOL} \right) \right] \ge 1 - \nu$$
 (13)

with cost

$$W(\text{TOL}) \leq \begin{cases} \mathcal{O}(\text{TOL}^{-2}), & q_2 > d\gamma \\ \mathcal{O}\left(\text{TOL}^{-2}\left(\log(\text{TOL}^{-1})\right)^2\right), & q_2 = d\gamma \\ \mathcal{O}\left(\text{TOL}^{-\left(2 + \frac{d\gamma - q_2}{q_1}\right)}\right), & q_2 < d\gamma. \end{cases}$$
(14)

This theorem shows that even in the worst case scenario, the CMLMC algorithm has a lower computational cost than that of the traditional (single level) MC method, which scales as $\mathcal{O}(\text{TOL}^{-2-d\gamma/q_1})$. Furthermore, in the best case scenario presented above, the computational cost of the CMLMC algorithm scales as $\mathcal{O}(\text{TOL}^{-2})$, i.e., identical to that of the MC method assuming that the cost per sample is $\mathcal{O}(1)$. In other words, for this case, the CMLMC algorithm can in effect remove the computational cost required by the discretization, namely $\mathcal{O}(\text{TOL}^{-d\gamma/q_1})$.

Remark 2: Note that the numerical errors relevant to the deterministic solver, such as discretization error, error due to the numerical integration used for computing the MoM matrix entries, and error due to the FMM used for accelerating the matrixvector product, do not have a significant effect on the statistical error, but they contribute to the bias error. Convergence of bias and the statistical errors are essential for CMLMC.

B. Random Shape Generation

This section describes a method to generate the sequence of meshes $\{P_\ell\}_{\ell=0}^L$ that are used in the numerical experiments (Section III) to demonstrate the properties of the CMLMC scheme. Alternative ways to generating random perturbations

can be found in [22], [56]–[59]. The method used here relies on perturbing an initial mesh generated on a sphere and refining it as level ℓ is increased as described next.

First, the unit sphere is discretized using a mesh P_0 with triangular elements, then the perturbations are generated by moving the nodes of this mesh using

$$v(\vartheta_m, \varphi_m) \approx \tilde{v}(\vartheta_m, \varphi_m) + \sum_{k=1}^{K} a_k \kappa_k(\vartheta_m, \varphi_m)$$
 (15)

where ϑ_m and φ_m are angular coordinates of node m, $v(\vartheta_m, \varphi_m)$ is its (perturbed) radial coordinate, and $\tilde{v}(\vartheta_m, \varphi_m) = 1$ m is its (unperturbed) radial coordinate on the unit sphere (all in spherical coordinate system). Here, $\kappa_k(\vartheta, \varphi)$ are obtained from spherical harmonics by rescaling their arguments and a_k , which satisfy $\sum_{k=1}^{K} a_k < 0.5$, are uncorrelated random variables.

For the numerical experiments considered in this paper, K = 2, and $\kappa_1(\vartheta, \varphi) = \cos(\alpha_1 \vartheta)$ and $\kappa_2(\vartheta, \varphi) = \sin(\alpha_2 \vartheta) \sin(\alpha_3 \varphi)$, where α_1, α_2 , and α_3 are positive constants. If a $\kappa_k(\vartheta, \varphi)$ depends on φ , then its dependence on ϑ must vanish at the poles $\vartheta = \{0, \pi\}$. Therefore, α_2 must be an integer. Additionally, α_3 must be an integer to generate a smooth perturbation.

Mesh P_0 , which is now after the application of (15), is also rotated and scaled using the simple transformation

$$\begin{bmatrix} x'_m \\ y'_m \\ z'_m \end{bmatrix} = \bar{L}(l_x, l_y, l_z) \bar{R}_x(\varphi_x) \bar{R}_y(\varphi_y) \bar{R}_z(\varphi_z) \begin{bmatrix} x_m \\ y_m \\ z_m \end{bmatrix}.$$
 (16)

Here, (x_m, y_m, z_m) and (x'_m, y'_m, z'_m) are the coordinates of node *m* before and after the transformation, matrices $\bar{R}_x(\varphi_x)$, $\bar{R}_y(\varphi_y)$, and $\bar{R}_z(\varphi_z)$ perform rotations around *x*, *y*, and *z* axes by angles φ_x , φ_y , and φ_z , and matrix $\bar{L}(l_x, l_y, l_z)$ implements (down) scaling along *x*, *y*, and *z* axes by l_x , l_y , and l_z , respectively.

The random variables used in generating the final version of P_0 are the perturbation weights a_k , k = 1, ..., K, the rotation angles φ_x, φ_y , and φ_z , and the scaling factors l_x, l_y , and l_z , making the dimension of the stochastic space K + 6, i.e., random parameter vector

$$\boldsymbol{\xi} = \{a_1, \dots, a_K, \varphi_x, \varphi_y, \varphi_z, l_x, l_y, l_z\}.$$
 (17)

Note that P_0 is the coarsest mesh used in CMLMC ($\ell = 0$) [see Fig. 2(a) for an example]. The mesh of the next level ($\ell = 1$), P_1 , is generated by refining each triangle of the perturbed P_0 into four (by halving all three edges and connecting midpoints) [Fig. 2(b)]. The mesh at level $\ell = 2$, P_2 , is generated in the same way from P_1 [Fig. 2(c)], and so on. All meshes P_ℓ at all levels $\ell = 1, \ldots, L$ are nested discretizations of P_0 . This method of refinement results in $\beta = 2$ in (4). Note that no uncertainties are added on meshes P_ℓ , $\ell > 0$; the uncertainty is introduced only at level $\ell = 0$. It is assumed that P_0 is fine enough to accurately represent the variations of the highest harmonic in the expansion used in (15).



Fig. 2. Example of four nested meshes with (a) 320, (b) 1280, (c) 5120, and (d) 20 480 triangular elements, which are generated for a perturbed shape with $\alpha_1 = 2$, $\alpha_2 = 3$, $\alpha_3 = 2$, $a_1 = 0.04$ m, $a_2 = 0.048$ m, $\varphi_x = 0.32$ rad, $\varphi_y = 0.88$ rad, $\varphi_z = 0.81$ rad, $l_x = 1.06$, $l_y = 1.08$, and $l_z = 1.07$.

C. Electromagnetic Solver

This section briefly describes the PMCHWT-SIE solver used for the RCS and SCS of a dielectric scatterer with surface S. It is assumed that sources and fields are time harmonic, i.e., their time dependence varies as $e^{j\omega t}$, where t and ω are the time and angular frequency, respectively.

Let V_1 and V_0 represent the space internal and external to S, respectively. The permittivity, permeability, and characteristic impedance in V_i , $i \in \{0, 1\}$ are ε_i , μ_i , and $\eta_i = \sqrt{\mu_i/\varepsilon_i}$, respectively. It is assumed that the scatterer is excited by an external electromagnetic field with electric and magnetic components $\mathbf{E}^{\text{inc}}(\mathbf{r})$ and $\mathbf{H}^{\text{inc}}(\mathbf{r})$. Using the surface equivalence theorem and enforcing the tangential continuity of total electromagnetic fields on S yield the PMCHWT-SIE [44].

$$\hat{\mathbf{n}}(\mathbf{r}) \times \mathbf{E}^{\text{inc}}(\mathbf{r}) = \hat{\mathbf{n}}(\mathbf{r}) \times \Big\{ \mathcal{L}_0[\mathbf{J}](\mathbf{r}) + \mathcal{L}_1[\mathbf{J}](\mathbf{r}) \\ - \mathcal{K}_0[\mathbf{M}](\mathbf{r}) - \mathcal{K}_1[\mathbf{M}](\mathbf{r}) \Big\}, \mathbf{r} \in S$$
(18)

$$\hat{\mathbf{n}}(\mathbf{r}) \times \mathbf{H}^{\text{inc}}(\mathbf{r}) = \hat{\mathbf{n}}(\mathbf{r}) \times \Big\{ \mathcal{K}_0[\mathbf{J}](\mathbf{r}) + \mathcal{K}_1[\mathbf{J}](\mathbf{r}) \\ + \eta_0^{-2} \mathcal{L}_0[\mathbf{M}](\mathbf{r}) + \eta_1^{-2} \mathcal{L}_1[\mathbf{M}](\mathbf{r}) \Big\}, \mathbf{r} \in S.$$
(19)

Here, $\hat{\mathbf{n}}(\mathbf{r})$ is the outward pointing unit normal vector, and $\mathbf{J}(\mathbf{r})$ and $\mathbf{M}(\mathbf{r})$ represent equivalent electric and magnetic surface current densities on S. The integral operators $\mathcal{L}_i[.](\mathbf{r})$ and

 $\mathcal{K}_{i}[.](\mathbf{r})$ in (18) and (19) are

$$\mathcal{L}_{i}[\mathbf{X}](\mathbf{r}) = j\omega\mu_{i}\int_{S}\left[\overline{\mathbf{I}} + \frac{\nabla\nabla'}{k_{i}^{2}}\right] \cdot \mathbf{X}(\mathbf{r}')g_{i}(\mathbf{r},\mathbf{r}')ds'$$
$$\mathcal{K}_{i}[\mathbf{X}](\mathbf{r}) = \nabla \times \int_{S}\mathbf{X}(\mathbf{r}')g_{i}(\mathbf{r},\mathbf{r}')ds'$$

where $g_i(\mathbf{r}, \mathbf{r}') = e^{-jk_i|\mathbf{r}-\mathbf{r}'|}/(4\pi|\mathbf{r}-\mathbf{r}'|)$ is the Green function of the Helmholtz equation in the unbounded medium with wave number $k_i = \omega \sqrt{\varepsilon_i \mu_i}$.

To numerically solve (18) and (19) for the unknowns $\mathbf{J}(\mathbf{r})$ and $\mathbf{M}(\mathbf{r})$, S is discretized by triangular elements, and $\mathbf{J}(\mathbf{r})$ and $\mathbf{M}(\mathbf{r})$ are approximated as

$$\mathbf{J}(\mathbf{r}) = \sum_{n=1}^{N} \bar{I}_n \mathbf{f}_n(\mathbf{r})$$
(20)

$$\mathbf{M}(\mathbf{r}) = \sum_{n=N+1}^{2N} \bar{I}_n \mathbf{f}_{n-N}(\mathbf{r})$$
(21)

where $\mathbf{f}_n(\mathbf{r})$, n = 1, ..., N, represent Rao–Wilton–Glisson basis functions [60], and $\overline{I} = [I_1, ..., I_{2N}]^T$ is the vector of unknown coefficients. Substituting (20) and (21) into (18) and (19), and testing the resulting equations with $\mathbf{f}_m(\mathbf{r})$, m = 1, ..., N, yield the MoM system of equations

$$\bar{V} = \bar{Z}\bar{I} \tag{22}$$

where the entries of the vector \bar{V} and the MOM matrix \bar{Z} are

$$\bar{V}_{m} = \begin{cases} \langle \mathbf{f}_{m}, \mathbf{E}^{\mathrm{inc}} \rangle, & 1 \leq m \leq N \\ \langle \mathbf{f}_{m-N}, \mathbf{H}^{\mathrm{inc}} \rangle, & N+1 \leq m \leq 2N \end{cases}$$
(23)
$$\bar{Z}_{m,n} = \begin{cases} \langle \mathbf{f}_{m}, \mathcal{L}_{0}[\mathbf{f}_{n}] + \mathcal{L}_{1}[\mathbf{f}_{n}] \rangle \\ 1 \leq m \leq N, 1 \leq n \leq N \\ - \langle \mathbf{f}_{m-N}, \mathcal{K}_{0}[\mathbf{f}_{n}] + \mathcal{K}_{1}[\mathbf{f}_{n}] \rangle \\ N+1 \leq m \leq 2N, 1 \leq n \leq N \\ \langle \mathbf{f}_{m}, \mathcal{K}_{0}[\mathbf{f}_{n-N}] + \mathcal{K}_{1}[\mathbf{f}_{n-N}] \rangle \\ 1 \leq n \leq N, N+1 \leq n \leq 2N \\ \langle \mathbf{f}_{m-N}, \eta_{0}^{-2} \mathcal{L}_{0}[\mathbf{f}_{n}] + \eta_{1}^{-2} \mathcal{L}_{1}[\mathbf{f}_{n}] \rangle \\ N+1 \leq m \leq 2N, N+1 \leq n \leq 2N. \end{cases}$$
(24)

Here, the inner product is $\langle \mathbf{f}_m, \mathbf{a} \rangle = \int_S \mathbf{f}_m(\mathbf{r}) \cdot \mathbf{a}(\mathbf{r}) d\mathbf{r}$.

Matrix equation (22) is solved iteratively for \overline{I} . The computational cost of multiplying \overline{Z} with a trial solution vector scales as $\mathcal{O}(N^{\text{iter}}N^2)$, where N^{iter} is the number of iterations required for the residual error to reach the desired level: typically $N^{\text{iter}} \ll N$. Likewise, the storage costs of the unaccelerated/classical solution scale as $\mathcal{O}(N^2)$.

To minimize the computational and storage cost while executing the MLMC algorithm, an FMM–FFT scheme is used. A detailed formulation of FMM and its extension FMM–FTT can be found in [45]–[51]. The scheme encloses the scatterer in a fictitious box that is embedded into a uniform grid of smaller boxes. Two nonempty boxes (i.e., boxes containing at least a pair of patches) constitute a far-field pair if there is at least one box between them. Otherwise, they form a near-field pair. Interactions between basis and test functions in near-field pairs (and same boxes) are computed using (24) and stored in matrix \overline{Z}^{near} . The contribution of near- and self-interactions to the matrix-vector multiplication $\overline{Z}\overline{I}$ is calculated by simply multiplying \overline{Z}^{near} with I. The interactions between basis and test functions in farfield pairs are represented using the radiation patterns of basis and test functions (sampled over a solid angle of a sphere) and a translation operator and their contributions to ZI are computed using the FMM-FFT scheme. The computational cost and memory requirement of the FMM-FFT accelerated iterative solution scale as $\mathcal{O}(N^{\text{iter}}N^{4/3}\log^{2/3}N)$ and $\mathcal{O}(N^{4/3}\log^{2/3}N)$, respectively [50], [51]. Note that these estimates are higher than those of the multilevel FMM [61], [62]; the FMM-FFT scheme is preferred here since its parallel implementation is significantly simpler [47]–[51], [63]–[69]. The parallelization strategy implemented here uses a hybrid message passing interface/open multiprocessing (MPI/OpenMP) standard to uniformly distribute the memory and computational load among processors [47]-[51], [63]–[69].

Note that the computational cost estimate provided above is obtained under the assumption that the ratio of the wavelength to the (average) element edge length stays the same as N is decreased/increased [47]-[51]. Within the CMLMC algorithm, the mesh is refined from one level to the next, while the frequency is kept constant. This means that the ratio of the wavelength to the edge length increases and the computational cost of the FFT-FMM-accelerated solver is expected to scale differently. Indeed, numerical experiments in Section III show that the CMLMC algorithm estimates the computational cost parameter as $\gamma \approx 1$ for objects comparable to the wavelength in size.

To compute the RCS and SCS, the scatterer is excited by a plane wave: $\mathbf{E}^{\text{inc}}(\mathbf{r}) = \mathbf{E}_0 e^{-jk_0 \hat{\mathbf{u}}^{\text{inc}} \cdot \mathbf{r}}$ and $\mathbf{H}^{\text{inc}}(\mathbf{r}) = (\hat{\mathbf{u}}^{\text{inc}} \times$ \mathbf{E}_0 / $\eta_0 e^{-jk_0 \hat{\mathbf{u}}^{\text{inc.}} \mathbf{r}}$, where $\hat{\mathbf{u}}^{\text{inc}}$ is the direction of propagation. The unknown vector \overline{I} is solved for under this excitation and the RCS $\sigma^{\rm rcs}(\vartheta,\varphi)$ along direction $\hat{\mathbf{u}}(\vartheta,\varphi) = \hat{\mathbf{x}}\sin\vartheta\cos\varphi + \hat{\mathbf{y}}\cos\varphi$ $\hat{\mathbf{y}}\sin\vartheta\sin\varphi + \hat{\mathbf{z}}\cos\vartheta$ is computed using [70]

$$\sigma^{\rm rcs}(\vartheta,\varphi) = \frac{\left|\mathbf{F}(\vartheta,\varphi)\right|^2}{4\pi \left|\mathbf{E}_0\right|^2}.$$
(25)

Here, $\mathbf{F}(\vartheta, \varphi)$ is the scattered electric field pattern in the far field and computed using

^ ()

$$\begin{split} \mathbf{F}(\vartheta,\varphi) &= -j\omega\mu_0 \hat{\mathbf{u}}(\vartheta,\varphi) \\ &\times \hat{\mathbf{u}}(\vartheta,\varphi) \times \sum_{n=1}^{N} \bar{I}_n \int_{S_n} \mathbf{f}_n(\mathbf{r}') e^{jk_0 \hat{\mathbf{u}}(\vartheta,\varphi) \cdot \mathbf{r}'} d\mathbf{r}' \\ &- jk_0 \hat{\mathbf{u}}(\vartheta,\varphi) \times \sum_{n=N+1}^{2N} \bar{I}_n \int_{S_{n-N}} \mathbf{f}_{n-N}(\mathbf{r}') e^{jk_0 \hat{\mathbf{u}}(\vartheta,\varphi) \cdot \mathbf{r}'} d\mathbf{r}'. \end{split}$$

The SCS $C^{\text{sca}}(\Omega)$ is obtained by integrating $\sigma^{\text{rcs}}(\vartheta,\varphi)$ over the solid angle Ω [70].

$$C^{\rm sca}(\Omega) = \frac{1}{4\pi} \int_{\Omega} \sigma^{\rm rcs}(\vartheta, \varphi) \sin \vartheta d\vartheta d\varphi.$$
 (26)

The integral in (26) is efficiently computed using the exact quadrature rule in [71].

III. NUMERICAL RESULTS

In all numerical experiments considered in this section, the scatterer resides in free space (vacuum) with $\mu_0 = 4\pi \times$ 10^{-7} H/m and $\varepsilon_0 = c_0^2/\mu_0$ F/m, where c_0 is the speed of light in vacuum, and the scatterer's permittivity and permeability are $\mu_1 = \mu_0$ and $\varepsilon_1 = 4\varepsilon_0$, respectively. For the plane wave excitation, $\mathbf{E}_0 = \mathbf{\hat{x}}, \mathbf{\hat{u}}^{\text{inc}} = -\mathbf{\hat{z}}$, and $\omega = 2\pi f$ with f = 300 MHz. The QoI is the SCS $C^{\text{sca}}(\Omega)$ computed over the cone $\Omega = [\vartheta_0, \vartheta_1] \times$ $[\varphi_0, \varphi_1] = [1/6, 11/36]\pi \operatorname{rad} \times [5/12, 19/36]\pi \operatorname{rad}$. For all random perturbations, the parameters of the quasi-spherical harmonics are $\alpha_1 = 2$, $\alpha_2 = 3$, and $\alpha_3 = 2$. Meshes P_{ℓ} at levels $\ell = 0, 1, 2, 3, 4$, which are generated using the method described in Section II-B, have {320, 1280, 5120, 20480, 81920} triangles, respectively. Note that not all five mesh levels are required in every experiment. Since the CMLMC algorithm is stochastic, 15 independent CMLMC runs are executed for each experiment to statistically characterize the performance of the method.

The matrix system in (22) is solved iteratively using the transpose-free quasi-minimal residual method [72] with a tolerance (residual) of $\epsilon_{\ell} = \epsilon_0 \beta^{-2\ell}$, where $\ell = 0, 1, 2, 3, 4$ represent mesh levels, $\epsilon_0 = 6.0 \times 10^{-4}$, and $\beta = 2$ (Section II-B). Using a level-dependent tolerance is consistent with the dependence of the discretization error in the QoI, which scales at a (heuristically determined) rate $\mathcal{O}(h_{\ell}^2) \propto \mathcal{O}(\beta^{-2\ell})$ and ensures that the number of iterations is kept in check when analyzing coarser meshes. The integrals in (23) and (24) are computed using the Gaussian quadrature rules; the accuracy of the rule, i.e., number of quadrature points, adjusts with the mesh level: it is seven for $\ell = 0, 1, 2$ and six for $\ell = 3, 4$. For all levels of meshes, the parameters of the FMM-FFT scheme are selected carefully to ensure that it has six digits of accuracy [61], [62]. All simulations are executed on Intel(R) Xeon(R) CPU E5-2680 v2 @ 2.80-GHz DELL workstations with 40 cores and 128-GB RAM.

For the examples in Sections III-B and III-C, the performance of the CMLMC is compared to that of the traditional MC method for varying values of TOL. Note that the MC runs are not actually executed, but their execution time is predicted using the results of the CMLMC runs. More specifically, the variance $Var(q_L)$ computed using the CMLMC method is used to predict the number of required MC samples as $N_{\rm MC} \sim {\rm Var}(g_L) {\rm TOL}^{-2}$. Also, note that, since 15 independent CMLMC runs are executed for each experiment, 15 MC curves are obtained and confidence intervals are used on plots to visualize the results obtained from these 15 curves.

A. Single Realization of Random Variables

This section demonstrates that the scattered fields strongly depend on the shape of the object for the scenarios considered in the numerical experiments, i.e., f = 300 MHz, $\varepsilon_1 = 4\varepsilon_0$, and the size of the object is roughly 2 m. More specifically, J(r) and M(r) induced on a (rotated and scaled) perturbed geometry and its RCS are compared to the same quantities of the unit sphere. The rotated, scaled, and perturbed surface is generated using



Fig. 3. Amplitudes of (a) $\mathbf{J}(\mathbf{r})$ and (b) $\mathbf{M}(\mathbf{r})$ induced on the unit sphere under excitation by an $\hat{\mathbf{x}}$ -polarized plane wave propagating in $-\hat{\mathbf{z}}$ direction at 300 MHz. Amplitudes of (c) $\mathbf{J}(\mathbf{r})$ and (d) $\mathbf{M}(\mathbf{r})$ induced on the perturbed shape shown in Fig. 2(c) under excitation by the same plane wave. For all figures, amplitudes are normalized to 1 and plotted in *dB* scale.

 $a_1 = 0.04$ m, $a_2 = 0.048$ m, $\varphi_x = 0.32$ rad, $\varphi_y = 0.88$ rad, $\varphi_z = 0.81$ rad, $l_x = 1.06$, $l_y = 1.08$, and $l_z = 1.07$. The discretization is refined twice, resulting in the mesh with 5120 triangles [Fig. 2(c)]. A mesh with the same number of triangles is generated on the unit sphere. Fig. 3(a) and (b) shows normalized amplitudes of $\mathbf{J}(\mathbf{r})$ and $\mathbf{M}(\mathbf{r})$ induced on this unit sphere. Fig. 3(c) and (d) plots the normalized amplitudes of $\mathbf{J}(\mathbf{r})$ and $\mathbf{M}(\mathbf{r})$ on the perturbed shape. It is clear that there is a significant difference between the current distributions induced on the sphere and the perturbed shaped. Fig. 4(a) and (b) compares $\sigma^{\mathrm{rcs}}(\vartheta,\varphi)$ of the unit sphere and the perturbed shaped the perturbed shaped computed on the $xz - (\vartheta \in [0, \pi] \operatorname{rad}, \varphi = 0, \operatorname{and} \varphi = \pi \operatorname{rad})$ and $yz - (\vartheta \in [0, \pi] \operatorname{rad}, \varphi = \pi/2 \operatorname{rad}, \operatorname{and} \varphi = 3\pi/2 \operatorname{rad})$ planes, respectively.

As expected, the RCS of the perturbed geometry is significantly different than that of the unit sphere. The results presented in Figs. 3 and 4 clearly demonstrate the need for a computational tool that can predict EM fields scattered from objects with uncertain shapes parameterized using random variables.

Next, convergence of the error in the SCS of the perturbed object is demonstrated. The reference SCS value used in the computation of the relative norm error is obtained using the finest mesh at the highest level. Fig. 5 plots this error versus N. A good convergence is observed. Having said that, the error convergence depends on the specific random perturbation, i.e., the realization of the random input vector $\boldsymbol{\xi}$, defined in (17). For some realizations that result surfaces with large perturba-



Fig. 4. RCS of the unit sphere and the perturbed shape shown in Fig. 2(c) computed on (a) xz and (b) yz planes under excitation by an \hat{x} -polarized plane wave propagating in $-\hat{z}$ direction at 300 MHz. For (a), $\varphi = 0$ and $\varphi = \pi$ rad in the first and second halves of the horizontal axis, respectively. For (b), $\varphi = \pi/2$ rad and $\varphi = 3\pi/2$ rad in the first and second halves of the horizontal axis, respectively.



Fig. 5. Relative norm error in SCS versus N.

tions, the convergence of the SCS error might be poor. Large number of such realizations results in nonoptimal identification of CMLMC parameters q_1 and q_2 , and may slow down the CMLMC method. A possible remedy is not to allow very large perturbations.



Fig. 6. (a) Computation times required by the CMLMC and MC methods versus TOL. (b) Value of the computational cost estimate W [given by (5)] for the CMLMC and MC methods versus TOL. Computation times are averaged over 15 repetitions of the experiment.

B. High Variability

In this example, the CMLMC algorithm is executed for random variables a_1 , $a_2 \sim U[-0.14, 0.14]$ m, φ_x , φ_y , $\varphi_z \sim U[0.2, 3]$ rad, and l_x , l_y , $l_z \sim U[0.9, 1.1]$; here, U[a, b] is the uniform distribution between a and b. The CMLMC algorithm is run for TOL values ranging from 0.2 to 0.008. At the lowest value of TOL, the CMLMC algorithm requires five mesh levels, i.e., P_{ℓ} , $\ell = 0, 1, 2, 3, 4$.

Fig. 6(a) compares computation times for the CMLMC and MC algorithms as a function of TOL. The figure reveals three convergence regimes (TOL zones).

The first zone, $0.1 \lesssim \text{TOL} \lesssim 0.3$, covers the range where TOL is larger than the sum of bias and statistical error. No additional samples and no additional mesh refinements are required by the CMLMC algorithm. The MC and CMLMC methods consume similar computational resources.

The second zone, $0.012 \leq \text{TOL} \leq 0.1$, corresponds to a preasymptotic regime. Both convergence rates are approxi-

mately 2. The statistical error is dominant, and many new samples on existing meshes P_{ℓ} , $\ell = 0, 1, 2$ (by default) are required. The MC and CMLMC methods again consume similar computational resources. To achieve higher accuracy, both the bias and the statistical error should be reduced. The statistical error could be reduced by taking more samples, and the bias by using finer meshes (i.e., increasing ℓ).

The third zone, $0.008 \lesssim \text{TOL} \lesssim 0.012$, corresponds to the start of the asymptotic regime. The bias becomes important and finer meshes are required. MC computation time increases rapidly and for the smallest TOL used, the CMLMC algorithm becomes almost 10 times faster than the MC method. Note that in the first and second regimes, the MC method may outperform the CMLMC algorithm since the latter carries some initial overhead.

For example, for TOL ≥ 0.012 , only P_{ℓ} , $\ell = 0, 1, 2, 3$ are required, but for TOL ≤ 0.012 an additional finer mesh, P_{ℓ} , $\ell = 4$, is required. Sampling on level $\ell = 4$ is more expensive; therefore, the MC computation time increases rapidly for TOL ≤ 0.012 . On the other hand, the CMLMC algorithm requires only very few samples on level $\ell = 4$, and is, therefore, significantly faster.

The computational cost estimate W is an indicator of computation time. It depends on how the computational cost of the deterministic solver changes from level $\ell - 1$ to ℓ [as indicated by parameters γ and β in (4)] and on the order of decay for the mean and the variance [parameters q_1, q_2 in (3a)]. Fig. 6(b) plots the values of W versus TOL. The curve is similar to that of the CMLMC computation time given in Fig. 6(a) demonstrating that $\gamma \approx 1, q_1 \approx 3$, and $q_2 \approx 5$ are reasonably accurate.

Fig. 7(a) shows the time required to compute $G_{\ell} = g_{\ell} - g_{\ell-1}$ versus ℓ . Computation times vary roughly as $2^{2\ell}$, which verifies that $\gamma \approx 1$ [since d = 2 and $\beta = 2$ in (4)]. Fig. 7(b) shows $E_{\ell} = E[G_{\ell}]$ versus ℓ , revealing that $E_{\ell} \sim 2^{-3\ell}$ (assumed weak convergence obtained with $q_1 = 3$). Fig. 7(c) shows $V_{\ell} = \text{Var}[G_{\ell}]$ versus ℓ , demonstrating that $V_{\ell} \sim 2^{-5\ell}$ (assumed strong convergence with $q_2 = 5$).

The results presented in Fig. 6(a) and (b), and Fig. 7(a)–(c) confirm the assumptions stated in Section II-A as well as the CMLMC scheme's quasi-optimality.

Fig. 8 shows θ versus TOL and demonstrates the complex relationship between TOL and the bias and statistical error. θ decreases from 1 to ≈ 0.7 . $\theta = 1$ implies that the ratio of bias to the total error is negligible, i.e., that the ratio of of statistical error to the total error is 1. Such variability in θ is one of the differences between the CMLMC algorithm and the MLMC method where $\theta = 0.5$. The blue dots show that for TOL ≈ 0.2 , $\theta \approx 1$, meaning that the impact of the bias is negligible and that there is no need to further extend the mesh hierarchy by adding finer mesh levels.

A higher accuracy can be achieved by decreasing either the bias or the statistical error (i.e., smaller TOL). $\theta \approx 1$ means that the bias is negligible and that the statistical error should be decreased to achieve a smaller TOL. Only when introducing a sufficient number of new samples will the statistical error decrease and the ratio of bias to TOL becomes higher. The bias can be decreased by including an additional mesh level. After that, the ratio of bias to TOL is dropping again, and the



 $\begin{array}{c} 1.0 \\ 0.8 \\ 0.6 \\ 0.4 \\ 0.2 \\ 0.0 \\$

Fig. 8. Value of θ used by the CMLMC algorithm versus TOL. Computation time is averaged over 15 repetitions of the experiment, i.e., there are 15 values of θ for a given value of TOL.

TOL



Fig. 7. (a) Time required to compute G_{ℓ} versus ℓ . (b) $E_{\ell} = E[G_{\ell}]$ versus ℓ and assumed weak convergence curve $2^{-3\ell}$ $(q_1 = 3)$. (c) $V_{\ell} = \operatorname{Var}[G_{\ell}]$ versus ℓ and assumed strong convergence curve $2^{-5\ell}$ $(q_2 = 5)$. The experiment is repeated 15 times independently and the obtained values are shown as error bars on the curves.

statistical error becomes dominant and should be decreased, etc. For TOL ≈ 0.01 , the present mesh hierarchy is not sufficient anymore, and the CMLMC algorithm adds one more mesh level.

Another way to study the behavior of G_{ℓ} is to look at its probability density function (pdf). Fig. 9(a) and (b) plots empirical pdfs of G_{ℓ} from {2000, 400, 50} samples for $\ell = 1$

Fig. 9. CMLMC pdfs of $g_{\ell} - g_{\ell-1}$ for (a) $\ell = 1$ and (b) $\ell = \{2, 3\}$.

and $\ell = \{2,3\}$, respectively. Fig. 9(a) shows that $\operatorname{Var}[G_1] = \operatorname{Var}[g_1 - g_0]$, where g_0 and g_1 are computed using the meshes P_0 and P_1 , varies roughly in the range (0, 2.5), i.e., this variance is large. Note that $\operatorname{E}[g_1 - g_0] \approx 1.0$, but the following $\operatorname{E}[g_\ell - g_{\ell-1}]$ are close to zero, that is why we show two plots. Fig. 9(b) shows that $G_2 = g_2 - g_1$, where g_2 are g_1 are computed using the meshes P_2 and P_1 , varies in the interval (-0.1, 0.1) and $\operatorname{E}[g_2 - g_1] \approx 0.02$. Finally, $G_3 = g_3 - g_2$, where g_3 are g_2 are computed using the meshes P_3 and P_2 , varies in the interval (-0.02, 0.02) and $\operatorname{E}[G_3] \approx 0$. The pdfs of G_ℓ concentrate more and more around zero [with the rates shown in Fig. 7(b) and (c)] as ℓ increases.





Fig. 10. (a) Computation times required by the CMLMC and MC methods versus TOL. (b) Value of the computational cost estimate W [given by (4)] for the CMLMC and MC methods versus TOL. Computations times are averaged over 15 repetitions of the experiment.

C. Low Variability

In this example, the CMLMC algorithm is executed for random variables a_1 , $a_2 \sim U[0.014, 0.49]$ m, φ_x , φ_y , $\varphi_z \sim U[0.2, 1.0]$ rad, and l_x , l_y , $l_z \sim U[0.8, 1.2]$. The variability of these random variables is lower than that of the variables used in the previous example. The value of TOL is changed from 0.2 to 0.02. For the lowest TOL value, the CMLMC algorithm requires four mesh levels, i.e., P_{ℓ} , $\ell = 0, 1, 2, 3$ are the only meshes levels used for this experiment.

Fig. 10(a) compares the computation time of the CMLMC and MC methods as a function of TOL. There are two zones in the figure. The first zone, $0.07 \leq \text{TOL} \leq 0.15$, describes the regime when TOL is higher than the sum of the bias and statistical error. No additional samples or refinements are needed.

The second zone, TOL ≤ 0.07 , describes the preasymptotic regime. As TOL gets smaller, the CMLMC algorithm becomes more efficient than the MC method. For values of TOL close to 0.02, the CMLMC algorithm is roughly 10 times faster than

Fig. 11. (a) $E_{\ell} = E[G_{\ell}]$ versus ℓ and assumed weak convergence curve $2^{-2\ell}$ $(q_1 = 2)$. (b) $V_{\ell} = \text{Var}[G_{\ell}]$ versus ℓ and assumed strong convergence curve $2^{-4\ell}$ $(q_2 = 4)$. The experiment is repeated 15 times and error bars are shown on the curves.

MC. Fig. 10(b) shows the values of the computational cost estimate W versus TOL. The curve is similar to that of the CMLMC computation time given in Fig. 10(a) validating the model developed for the CMLMC algorithm. Fig. 11(a) and (b) shows $E_{\ell} = \mathbb{E}[G_{\ell}]$ and $V_{\ell} = \operatorname{Var}[G_{\ell}]$ versus ℓ , respectively, revealing dependencies on ℓ that vary as $2^{-2\ell}$ (assuming weak convergence obtained with $q_1 = 2$) and $2^{-4\ell}$ (assuming strong convergence obtained with $q_2 = 4$). Note that in the previous example these parameters are 3 and 5, respectively, demonstrating that the dependence of q_1 and q_2 on the variability of the random variables used to described the geometry. Convergence in Fig. 11(a) and (b) is faster than the predicted rates of $2^{-2\ell}$ and $2^{-4\ell}$, respectively. A possible reason is that we are still in a preasymptotic regime and we were excessively pessimistic with factors $q_1 = 2$ and $q_2 = 4$. Although we tried to avoid "outliers" in generating the random sphere-like geometries, we could not completely avoid them. So some shape samples which were used for "training" of the CMLMC method, were "bad" and the deterministic solver produced solutions which were very different from others. As a result, we were forced to give pessimistic *a priori* estimates for q_1 and q_2 (we remind that the CMCLMC method computes 10–20 samples on each of the first three nested meshes). Later samples were better than ones used for "training" the CMLMC method. The remedy to get a better fitting of curves in Fig. 11(a) and (b) is a more accurate preliminary estimates of q_1 and q_2 . For this, we need either to take more samples for training or to generate "smoother" perturbations of the original shape.

IV. CONCLUSION

A computational framework is developed to efficiently and accurately characterize EM wave scattering from dielectric objects with uncertain shapes. To this end, the framework uses the CMLMC algorithm, which reduces the computational cost of the traditional MC method by performing most of the simulations with lower accuracy and lower cost (using coarser meshes) and smaller number of simulations with higher accuracy and higher cost (using finer meshes). To increase the efficiency further, each of the simulations is carried out using the FMM-FFT accelerated PMCHWT-SIE solver. Numerical results demonstrate that the CMLMC algorithm can be 10 times faster than the traditional MC method depending on the amplitude of the perturbations used for representing the uncertainties in the scatterer's shape. This paper confirms that the known advantages of the CMLMC algorithm can be observed when it is applied to EM wave scattering: nonintrusiveness, dimension independence, better convergence rates compared to the classical MC method, and higher immunity to irregularity w.r.t. uncertain parameters, than, for example, sparse grid methods.

For optimal performance (for a given value of accuracy parameter TOL), the CMLMC algorithm requires the mean and the variance to have reliable convergence rates (i.e., one should be able estimate q_1 and q_2 without much difference from one level to next). However, some random perturbations may affect the convergence rates. With difficult-to-predict convergence rates, it is hard for the CMLMC algorithm to estimate the computational cost W, the number of levels L, the number of samples on each level M_{ℓ} , the computation time, and the parameter θ , and the variance in QoI. All these may result in a suboptimal performance. Indeed, numerical results demonstrate that there is a significant preasymptotic regime where the performance is not optimal. Additionally, it is observed that the settings of the FMM-FFT accelerated PMCHWT-SIE solver, which regulate the computation time and the accuracy (such as the iterative solver threshold, the number of quadrature points, and the FMM-FFT parameters), have a significant effect on the performance of the CMLMC algorithm.

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