Parameter Estimation in SAR Imagery using Stochastic Distances and Asymmetric Kernels

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Abstract—In this paper we analyze several strategies for the estimation of the roughness parameter of the $G_0^0$ distribution. It has been shown that this distribution is able to characterize a large number of targets in monopolarized SAR imagery, deserving the denomination of “Universal Model.” It is indexed by three parameters: the number of looks (which can be estimated in the whole image), a scale parameter, and the roughness or texture parameter. The latter is closely related to the number of elementary backscatter in each pixel, one of the reasons for receiving attention in the literature. Although there are efforts in providing improved and robust estimates for such quantity, its dependable estimation still poses numerical problems in practice. We discuss estimators based on the minimization of stochastic distances between empirical and theoretical densities, and argue in favor of using an estimator based on the Triangular distance and kurtosis. We also provide new results regarding the heavytailedness of this distribution.

Index Terms—Feature extraction, image texture analysis, statistics, synthetic aperture radar, speckle

I. INTRODUCTION

THE statistical modeling of the data is essential in order to interpret SAR images. Speckled data have been described under the multiplicative model using the $G$ family of distributions which is able to describe rough and extremely rough areas better than the $K$ distribution [1], [2]. The survey article [3] discusses in detail several statistical models for this kind of data.

Under the $G$ model different degrees of roughness are associated to different parameter values, therefore it is of paramount importance to have high quality estimators. Several works have been devoted to the subject of improving estimation with two main venues of research, namely, analytic and resampling procedures.

The analytic approach was taken by Vasconcellos et al. [4] who quantified the bias in the estimation of the roughness parameter of the $G_0^0$ distribution by maximum likelihood (ML). They proposed an analytic change for improved performance with respect to bias and mean squared error. Also, Silva et al. [5] computed analytic improvements for that estimator. Such approaches reduce both the bias and the mean squared error of the estimation, at the expense of computing somewhat cumbersome correction terms, and yielding estimators whose robustness is largely unknown.

Cribari-Neto et al. [6] compared several numerical improvements for that estimator using bootstrap. Again, the improvement comes at the expense of intensive computing and with no known properties under contamination. Allende et al. [7] and Bustos et al. [8] also sought for improved estimation, but keeping the robustness of the procedure. The new estimators are resistant to contamination, and in some cases they also improve the mean squared error, but they require dealing with influence functions and asymptotic properties not always immediately available to remote sensing practitioners.

A common issue in all the aforementioned estimation procedures, including ML, and to those based on fractional moments [1] and log-cumulants [9]–[11] is the need of iterative algorithms for which there is no granted convergence to global solutions. Such lack of convergence usually arises with small samples, precluding the use of such techniques in, e.g., statistical filters. Frery et al. [12] and Pianto and Cribari-Neto [13] proposed techniques which aim at alleviating such issue, at the cost of additional computational load.

The main idea of this work is to develop an estimation method for the $G_0^0$ model with good properties (as measured by its bias, the mean squared error and its ability to resist contamination) even with samples of small and moderate size, and low computational cost. In order to achieve this task, we propose minimizing a stochastic distance between the fixed empirical evidence and the estimated model.

Shannon proposed a divergence between two density functions as a measure of the relative information between the two distributions. Divergences were studied by Kullback and Leibler and by Rényi [14], among others. These divergences have multiple applications in signal and image processing [15], medical image analysis diagnosis [16], and automatic region detection in SAR imagery [17]–[20]. Liese and Vajda [21] provide a detailed theoretical analysis of divergence measures.

Cassetti et al. [22] compared estimators based on the Hellinger, Bhattacharyya, Rényi and Triangular distance with the ML estimator. They presented evidence that the Triangular distance is the best choice for this application, but noticed that histograms led to many numerical instabilities. This work presents improvements with respect to those results in the following regards: we assess the impact of contamination in the estimation, we employ kernels rather than histograms and
we compare estimators based on the Triangular distance with the ML, Fractional Moments and Log-Cumulants estimators. Among the possibilities for such estimate, we opted for employing asymmetric kernels.

Kernels have been extensively applied with success to image processing problems as, for instance, object tracking [23]. Kernels with positive support, which are of particular interest for our work, were employed in [24], [25]. The general problem of using asymmetric kernels was studied in [26].

In [27], the authors demonstrate, through extensive modeling of real data, that SAR images are best described by families of heavy-tailed distributions. We provide new results regarding the heaviness-tailedness of the $\mathcal{G}_j^0$ distribution; in particular, we show that it has heavy tails with tail index $1 - \alpha$.

The paper unfolds as follows. Section II recalls the main properties of the $\mathcal{G}_j^0$ model and the Maximum Likelihood, the $\frac{1}{4}$-moment and the log-cumulants methods for parameter estimation. Estimation with stochastic distances is presented in Section III including different ways of calculating the estimate of the underlying density function and the contamination models employed to assess the robustness of the procedure. Section IV presents the main results. Section V discusses the conclusions.

II. THE $\mathcal{G}_j^0$ MODEL

The return in monopolarized SAR images can be modeled as the product of two independent random variables, one corresponding to the backscatter $X$ and the other to the speckle noise $Y$. In this manner $Z = XY$ represents the return in each pixel under the multiplicative model.

For monopolarized data, speckle is modeled as a Gamma distributed random variable, with unitary mean and shape parameter $L \geq 1$, the number of looks, while the backscatter is considered to obey a reciprocal of Gamma law. This gives rise to the $\mathcal{G}_j^0$ distribution for the return. Given the mathematical tractability and descriptive power of the $\mathcal{G}_j^0$ distribution for intensity data [2], [28] it represents an attractive choice for SAR data modeling.

The density function for intensity data is given by

$$f_{\mathcal{G}_j^0}(z) = \frac{L^L \Gamma(L - \alpha)}{\Gamma(L) \Gamma(L - \alpha) \Gamma(\gamma z L)^{\alpha - 1}} \frac{z^{L-1}}{\gamma},$$

where $-\alpha, \gamma, z > 0$ and $L \geq 1$. The $r$-order moments are

$$E[Z^r] = \left(\frac{\gamma}{L}\right)^r \frac{\Gamma(L + r) \Gamma(-\alpha - r)}{\Gamma(-\alpha) \Gamma(L + r)} \frac{\Gamma(-\alpha - r)}{\Gamma(-\alpha)},$$

provided $\alpha < -r$, and infinite otherwise.

With the double purpose of simplifying the calculations and making the results comparable, in the following we choose the scale parameter such that $E(Z) = 1$, which is given by $\gamma^* = -\alpha - 1$.

One of the most important features of the $\mathcal{G}_j^0$ distribution is the interpretation of the $\alpha$ parameter, which is related to the roughness of the target. Values close to zero (typically above $-3$) suggest extreme textured targets, as urban zones. As the value decreases, it indicates regions with moderate texture (usually $\alpha \in [-6, -3]$), as forest zones. Textureless targets, e.g. pasture, usually produce $\alpha \in (-\infty, -6)$. This is the reason why the accuracy in the estimation of $\alpha$ is so important.

Let $z = (z_1, \ldots, z_n)$ be a random sample of $n$ independent draws from the $\mathcal{G}_j^0$ model. Assuming $\gamma^* = -\alpha - 1$, the ML estimator for the parameter $\alpha$, namely $\hat{\alpha}_{\text{ML}}$ is the solution of the following nonlinear equation:

$$\Psi^0(\hat{\alpha}_{\text{ML}}) - \Psi^0\left(\frac{L}{1 - \hat{\alpha}_{\text{ML}}} - \log(1 - \hat{\alpha}_{\text{ML}})\right) + \frac{\hat{\alpha}_{\text{ML}}}{1 - \hat{\alpha}_{\text{ML}}} + \frac{1}{n} \sum_{i=1}^{n} \log(1 - \hat{\alpha}_{\text{ML}} + L z_i) - \frac{\hat{\alpha}_{\text{ML}} - L}{n} \sum_{i=1}^{n} \frac{1}{1 - \hat{\alpha}_{\text{ML}} + L z_i} = 0,$$

where $\Psi^0(\cdot)$ is the digamma function. Some of the issues posed by the solution of this equation have been discussed, and partly solved, in [12].

Fractional moments estimators have been widely used with success [1], [29]. Using $r = 1/2$ in (2) one has to solve

$$\frac{1}{n} \sum_{i=1}^{n} \sqrt{z_i} - \sqrt{-\hat{\alpha}_{\text{Mom12}} - 1} \frac{\Gamma(-\hat{\alpha}_{\text{Mom12}} - \frac{r}{2})}{\Gamma(-\alpha_{\text{Mom12}})} \frac{\Gamma\left(\frac{r}{2}ight)}{\Gamma(L) \Gamma(\gamma z_i)} = 0.$$

(3)

Estimation based on log-cumulants is gaining space in the literature due to its nice properties and good performance [9]–[11]. Following Tison et al. [30] the main second kind statistics can be defined as:

- First second kind characteristic function: $\phi_x(s) = \int_{0}^{\infty} u^{s-1} p_x(u)du$
- Second second kind characteristic function: $\psi_x(s) = \log \phi_x(s)$
- First order second kind characteristic moment:

$$\tilde{m}_r = \frac{d\phi_x(s)}{ds} \bigg|_{s=1}.$$  

(4)

- First order second kind characteristic cumulant (log-cumulant)

$$\tilde{k}_r = \frac{d\psi_x(s)}{ds} \bigg|_{s=1}.$$  

(5)

If $p_x(u) = f_{\mathcal{G}_j^0}(u)$ then we have:

- $\phi_x(s) = \left(\frac{\gamma}{L}\right)^s \Gamma(1 + L - s) \Gamma(1 - \alpha) \Gamma(1 + s - \alpha)$
- $\tilde{m}_1 = \frac{d\phi_x(s)}{ds} \bigg|_{s=1} = -\log\left(\frac{\gamma}{L}\right) + \Psi^0(L) - \Psi^0(-\alpha) - \alpha$

Using the developments presented in [30], we have that $\tilde{k}_1 = \tilde{m}_1$ and the empirical expression for the first log-cumulant estimator for $n$ samples $z_i$ is $\tilde{k}_1 = n^{-1} \sum_{i=1}^{n} \log z_i$. Therefore

$$\tilde{k}_1 = -\log \frac{L}{\gamma} + \Psi^0(L) - \Psi^0(-\alpha).$$

(6)

Assuming $\gamma^* = -\alpha - 1$, the Log-Cumulant estimator of $\alpha$, denoted by $\hat{\alpha}_{\text{LCum}}$ is then the solution of $\tilde{k}_1 = -\log \frac{L}{1 - \hat{\alpha}_{\text{LCum}}} + \Psi^0(L) - \Psi^0(-\hat{\alpha}_{\text{LCum}})$, that is, the solution of

$$\frac{1}{n} \sum_{i=1}^{n} \log z_i = -\log \frac{L}{1 - \hat{\alpha}_{\text{LCum}}} + \Psi^0(L) + \Psi^0(-\hat{\alpha}_{\text{LCum}}).$$

(7)

The ability of these estimators to resist outliers has not still been assessed.
In the following we provide new results regarding the heavy-tailedness of the $G^0_T$ distribution. This partly explains the numerical issues faced when seeking for estimators for its parameters: the distribution is prone to producing extreme values. The main concepts are from \[31\]–\[33\].

**Definition 1:** The function $\ell : \mathbb{R} \to \mathbb{R}$ is slow-varying in the infinite if for every $t > 0$ holds that
\[
\lim_{x \to +\infty} \frac{\ell(tx)}{\ell(x)} = 1.
\]

**Definition 2:** A probability density function $f(x)$ has heavy tails if for any $\eta > 0$ holds that
\[
f(x) = \ell(x)x^{-\eta},
\]
where $\ell$ is a slow-varying function in the infinite, and $\eta$ is the tail index. The smaller the tail index is, the more prone to producing extreme observations the distribution is.

**Proposition 1:** The $G^0_T$ distribution has heavy tails with tail index $1 - \alpha$.

**Proof:** Defining $\ell(x) = f_{G^0_T}(x)x^{-\alpha+1}$ we have that
\[
\lim_{x \to +\infty} \frac{\ell(tx)}{\ell(x)} = \lim_{x \to +\infty} \frac{(tx)^{-L-1}(Ltx + \gamma)^{\alpha-L}(tx)^{1-\alpha}}{x^{-L-1}(Lx + \gamma)^{\alpha-L}x^{1-\alpha}} = \lim_{x \to +\infty} t^{-\alpha}\left(\frac{Lx + \gamma}{Ltx + \gamma}\right)^{L-\alpha} = 1.
\]
This holds for every $t > 0$, so $\ell$ is as in Def. 1.

As expected, the tail index is a decreasing function on $\alpha$, then the $G^0_T$ distribution is more prone to producing extreme observations when the roughness parameter is bigger.

The remainder of this section is devoted to proving that the $G^0_T$ model is outlier-prone. Consider the random variables $z_1, \ldots, z_n$ and the corresponding order statistics $Z_{1:n} \leq \cdots \leq Z_{n:n}$.

**Definition 3:** The distribution is absolutely outlier-prone if there are positive constants $\varepsilon, \delta$ and an integer $n_0$ such that
\[
\Pr(Z_{n:n} - Z_{n-1:n} > \varepsilon) \geq \delta
\]
holds for every integer $n \geq n_0$.

A sufficient condition for being absolutely outlier-prone is that there are positive constants $\varepsilon, \delta, x_0$ such that the density of the distribution satisfies, for every $x \geq x_0$, that
\[
\left.\frac{f(x + \varepsilon)}{f(x)} \geq \delta\right.\quad (8)
\]

Since
\[
\lim_{x \to +\infty} \frac{f_{G^0_T}(x + \varepsilon)}{f_{G^0_T}(x)} = \lim_{x \to +\infty} \frac{(x + \varepsilon)^{-L-1}((Lx + \gamma)^{\alpha-L}(x + \varepsilon)^{1-\alpha}}{x^{-L-1}(Lx + \gamma)^{\alpha-L}x^{1-\alpha}} = 1,
\]
we are in the presence of an absolutely outlier-prone model.

### III. Estimation by the Minimization of Stochastic Distances

Information Theory provides divergences for comparing two distributions; in particular, we are interested in distances between densities. Our proposal consists of, given the sample $z$, computing $\hat{\alpha}$, estimator of $\alpha$, as the point which minimizes the distance between the density $f_{G^0_T}$ and an estimate of the underlying density function.

Cassetti et al. [22] assessed the Hellinger, Bhattacharyya, Rényi and Triangular distances, and they concluded that the latter outperforms the other ones in a variety of situations. The Triangular distance between the densities $f_V$ and $f_W$ with common support $S$ is given by
\[
d_T(f_V, f_W) = \int_S \frac{(f_V - f_W)^2}{f_V + f_W}. \quad (9)
\]

Let $z = (z_1, \ldots, z_n)$ be a random sample of $n$ independent $G^0_T(a_0, \gamma_0, L_0)$-distributed observations. An estimate of the underlying density function of $z$, denoted $\hat{f}$, is used to define the objective function to be minimized as a function of $\alpha$. The estimator for the $\alpha$ parameter based on the minimization of the Triangular distance between an estimate of the underlying density $f$ and the model $G^0_T$, denoted by $\hat{\alpha}_T$, is given by
\[
\hat{\alpha}_T = \arg \min_{-20 \leq \alpha \leq 1} d_T(f_{G^0_T}(\alpha, \gamma_0^*, L_0), \hat{f}(z)), \quad (10)
\]
where $\gamma_0^*$ and $L_0$ are known and $d_T$ is given in (9). Solving (10) requires two steps: the integration of (9) using the $G^0_T$ density and the density estimate $\hat{f}$, and optimizing with respect to $\alpha$. To the best of the authors’ knowledge, there are no explicit analytic results for either problem, so we rely on numerical procedures. The range of the search is established to avoid numerical instabilities.

We also calculate numerically the ML, the $\frac{1}{2}$-moment and Log-Cumulants based estimators using the same established range of the search and compare all the methods through the bias and the mean squared error by simulation.

### A. Estimate of the Underlying Distribution

The choice of the way in which the underlying density function is computed is very important in our proposal. In this section we describe several possibilities for computing it, and we justify our choice.

Histograms are the simplest empirical densities, but they lack unicity and they are not smooth functions.

Given that the model we are interested is asymmetric, another possibility is to use asymmetric kernels [24], [26]. Let $z = (z_1, \ldots, z_n)$ be a random sample of size $n$, with an unknown density probability function $f$, an estimate of its density function using kernels is given by
\[
\hat{f}_b(t; z) = \frac{1}{n} \sum_{i=1}^{n} K(t; z_i, b),
\]
where $b$ is the bandwidth of the kernel $K$.

Among the many available asymmetric kernels, we worked with two: the densities of the Gamma and Inverse Gaussian
These kernels are given by
\[ K_1(t; z_i, b) = \frac{t^{z_i/b} \exp\{-t/b\}}{b^{z_i/b+1} \Gamma(z_i/b+1)}, \]
and
\[ K_{IG}(t; z_i, b) = \frac{1}{\sqrt{2\pi}b t^3} \exp\left\{ -\frac{1}{2b} \left( \frac{t}{z_i} + \frac{z_i}{t} - 2 \right) \right\}, \]
respectively, for every \( t > 0 \). Empirical studies led us to employ \( b = n^{-1/2}/5 \).

As an example, Figure 1 shows the \( G_{IG}^0(-3, 2, 1) \) density and three estimates of the underlying density function obtained with \( n = 30 \) samples: those produced by the Gamma and the Inverse Gaussian kernels, and the histogram computed with the Freedman-Diaconis method.

![Gamma Kernel IG Kernel Truth](image)

**Fig. 1.** Fitting the \( G_{IG}^0(-3, 2, 1) \) density of thirty \( G_{IG}^0(-3, 2, 1) \) observations in different ways, along with the histogram.

As it can be seen, fitting the underlying density function using kernels is better than using a histogram. After extensive numerical studies, we opted for the \( K_{IG} \) kernel due to its low computational cost, its ability to describe observations with large variance, and its good numerical stability. In agreement with what Bouezmarni and Scaillet [26] reported, the Gamma kernel is also susceptible to numerical instabilities.

### B. Contamination

Estimators in signal and image processing are often used in a wide variety of situations, so their robustness is of highest importance. Robustness is the ability to perform well when the data obey the assumed model, and to not provide completely useless results when the observations do not follow it exactly. Robustness, in this sense, is essential when designing image processing and analysis applications. Filters, for instance, employ estimators based, typically, in small samples which, more often than not, receive samples from more than one class. Supervised classification relies on samples, and the effect of contamination (often referred to as “training errors”) on the results has been attested, among other works, in [34].

In order to assess the robustness of the estimators, we propose three contamination models able to describe realistic departures from the hypothetical “independent identically distributed sample” assumption.

One of the sources of contamination in SAR imagery is the phenomenon of double bounce which results in some pixels having a high return value. The presence of such outliers may provoke big errors in the estimation.

In order to assess the robustness of the proposal, we generate contaminated random samples using three types (cases) of contamination, with \( 0 < \epsilon \ll 1 \) the proportion of contamination. Let the Bernoulli random variable \( B \) with probability of success \( \epsilon \) model the occurrence of contamination. Let \( C \in \mathbb{R}_+ \) be a large value.

- **Case 1:** Let \( W \) and \( U \) be such that \( W \sim G_{IG}^0(\alpha_1, \gamma_1^*, L) \), and \( U \sim G_{IG}^0(\alpha_2, \gamma_2^*, L) \). Define \( Z = BU + (1 - B)W \), then we generate \( \{z_1, \ldots, z_n\} \) identically distributed random variables with cumulative distribution function
  \[ (1 - \epsilon)F_{G_{IG}^0(\alpha_1, \gamma_1^*, L)}(z) + \epsilon F_{G_{IG}^0(\alpha_2, \gamma_2^*, L)}(z), \]
  where \( F_{G_{IG}^0(\alpha, \gamma, L)} \) is the cumulative distribution function of a \( G_{IG}^0(\alpha, \gamma, L) \) random variable.

- **Case 2:** Consider \( W \sim G_{IG}^0(\alpha_1, \gamma_1^*, L) \); return \( Z = BC + (1 - B)W \).

- **Case 3:** Consider \( W \sim G_{IG}^0(\alpha, \gamma^*, L) \) and \( U \sim G_{IG}^0(\alpha, 10^k \gamma^*, L) \) with \( k \in \mathbb{N} \). Return \( Z = BU + (1 - B)W \), then \( \{z_1, \ldots, z_n\} \) are identically distributed random variables with cumulative distribution function
  \[ (1 - \epsilon)F_{G_{IG}^0(\alpha, \gamma^*, L)}(z) + \epsilon F_{G_{IG}^0(\alpha, 10^k \gamma^*, L)}(z), \]

All these models consider departures from the hypothesized distribution \( G_{IG}^0(\alpha, \gamma^*, L) \). The first type of contamination assumes that, with probability \( \epsilon \), instead of observing outcomes from the “right” model, a sample from a different one will be observed; notice that the outlier may be close to the other observations. The second type returns a fixed and typically large value, \( C \), with probability \( \epsilon \). The third type is a particular case of the first, where the contamination assumes the form of a distribution whose scale is \( k \) orders of magnitude larger than the hypothesized model. We use the three cases of contamination models in our assessment.

### IV. RESULTS

A Monte Carlo experiment was set to assess the performance of each estimation procedure. The parameter space consists of the grid formed by (i) three values of roughness: \( \alpha = \{ -1.5, -3, -5 \} \), which are representative of areas with extreme and moderate texture; (ii) three usual levels of signal-to-noise processing, through \( L = \{ 1, 3, 8 \} \); (iii) sample sizes \( n = \{ 9, 25, 49, 81, 121, 1000 \} \), related to squared windows of side 3, 5, 7, 9 and 11, and to a large sample; (iv) each of the three cases of contamination, with \( \epsilon = \{ 0.001, 0.005, 0.01 \} \), \( \alpha_2 = \{ -4, -15 \} \), \( C = 100 \) and \( k = 2 \).

One thousand samples were drawn for each point of the parameter space, producing \( \{\hat{\alpha}_1, \ldots, \hat{\alpha}_{1000}\} \) estimates of each kind. Estimates of the mean \( \hat{\alpha} = \{ 1000 \}^{-1}\sum_{i=1}^{1000} \hat{\alpha}_i \), bias \( \hat{B}(\hat{\alpha}) = \hat{\alpha} - \alpha \) and mean squared error (MSE) are then computed and compared.

In the following figures, “ML”, “T”, “Mom12” and “LCum” denote the estimator based on the Maximum likelihood, Triangular distance, \( \frac{1}{2} \)-moment and Log-Cumulant, respectively.
Sample sizes are in the abscissas, which are presented in logarithmic scale. The estimates of the mean are presented with error bars which show the Gaussian confidence interval at the 95% level of confidence. Only a few points of this parameter space are presented for brevity, but the remaining results are consistent with what is discussed here.

Figure 2 shows the mean of the estimators $\hat{\alpha}$ in uncontaminated data ($\epsilon = 0$) with different values of $n$ and $L$. Only two of the four estimators lie, in mean, very close to the true value: $\hat{\alpha}_{\text{MV}}$ and $\hat{\alpha}_{\text{T}}$; it is noticeable how far from the true value lie $\hat{\alpha}_{\text{LCum}}$ and $\hat{\alpha}_{\text{Mom12}}$ when $\alpha = -5$. It is noticeable that $\hat{\alpha}_{\text{ML}}$ has a systematical tendency to underestimate the true value of $\alpha$. Vasconcellos et al. [4] computed a first order approximation of such bias for a closely related model, and our results are in agreement with those. The estimator based on the Triangular distance $\hat{\alpha}_{\text{T}}$ compensates this bias.

Figure 3 shows the sample mean squared error of the estimates under the same situation, i.e., uncontaminated data. In most cases, all estimators have very similar mse, not being possible to say that one is systematically the best one. This is encouraging, since it provides evidence that $\hat{\alpha}_{\text{T}}$ exhibits the first property of a good robust estimator, i.e., not being unacceptable under the true model.

The mean times of processing, measured in seconds, for each method and each case, were computed, as an example, the mean times of processing for $L = 1$ and $n = 81$ are presented in Table I. It can be seen that the new method has a higher computational cost. The other cases are consistent with this table. The details of the computer platform are presented in the appendix.

### Table I
**Mean times for simulated data without contamination, $L = 1$, $n = 81$**

|        | MV   | DT   | Moment $\frac{1}{2}$ | Log Cumulant |
|--------|------|------|----------------------|--------------|
| Time   | 0.003| 2.223| 0.0001               | 0.003        |

Figures 4 and 5 show, respectively, the sample mean and mean squared error of the estimates under Case 1 contamination with $\alpha_2 = -15$, $\epsilon = 0.01$, and varying $n$ and $L$. This type of contamination injects, with probability $\epsilon = 0.01$, observations with almost no texture in the sample under analysis. As expected, the influence of such perturbation is more noticeable in those situations where the underlying model is further away from the contamination, i.e., for larger values of $\alpha$. This is particularly clear in Fig. 5 which shows that the mean squared errors of $\hat{\alpha}_{\text{ML}}$, $\hat{\alpha}_{\text{Mom12}}$ and $\hat{\alpha}_{\text{LCum}}$ are larger than that of $\hat{\alpha}_{\text{T}}$ for $L = 3, 8$, with not a clear distinction for $L = 1$ except that $\hat{\alpha}_{\text{T}}$ is at least very competitive in the $\alpha = -3, -5$ cases.

Figures 6 and 7 present, respectively, the sample mean and sample mean squared error of the estimates under Case 2 contamination with $\epsilon = 0.001$ and $C = 100$. This type of contamination injects a constant value ($C = 100$) with probability $\epsilon = 0.001$ instead of each observation from the $G^0_I$ distribution. Since we are considering samples with unitary mean, this is a large contamination. In this case, $\hat{\alpha}_{\text{T}}$ is, in mean, closer to the true value than the other methods, and its mean square error the smallest.

Figures 8 and 9 show, respectively, the sample mean and mean squared error of the estimates under Case 3 with $\epsilon = 0.005$ with $k = 2$. This kind of contamination draws, with probability $\epsilon = 0.005$, an observation from a $G^0_I$ distribution with a scale one hundred times larger than that of the “correct” model. The behavior of the estimators follows the same pattern for $L = 3, 8$: $\hat{\alpha}_{\text{T}}$ produces the closest estimates to the true value with reduced mean squared error. There is no good estimator for the single-look case with this case of contamination.

As previously mentioned, the iterative algorithms which implement the $\frac{1}{2}$-moment and Log-Cumulant estimators often fail to converge. In the Monte Carlo experiment, if $\frac{1}{2}$-moment or Log-Cumulant fail, we eliminate the estimators computed...
with the other methods in the corresponding iteration, so the amount of elements for calculating the mean $\bar{\alpha}$, the bias and the mean squared error is lower than 1000. As an example, Table II informs the number of such situations for Case 1 and $\alpha_2 = -15, \epsilon = 0.01$. These results are consistent with other situations under contamination, and they suggest that these methods are progressively more prone to fail in more heterogeneous areas.

Data from a single-look L-band HH polarization in intensity format E-SAR [35] image was used in the following. Figure 10 shows the regions used for estimating the texture parameter. Table III shows the results of estimating the $\alpha$ parameter for

| $L$ | $\frac{1}{2}$-Moment | Log-Cumulant |
|-----|----------------------|--------------|
| 1   | 22.87                | 21.56        |
| 3   | 11.71                | 11.87        |
| 8   | 5.81                 | 6.04         |

Fig. 3. Sample mean squared error of estimates under uncontaminated data.
each rectangular region, where \(NA\) means that the corresponding estimator is not available.

The Kolmogorov-Smirnov test (KS-test) is applied to two samples: \(x\), from the image, and \(y\), simulated. Babu and Feigelson [36] warn about the use of the same sample for parameter estimation and for performing a KS test between the data and the estimated cumulative distribution function. We then took the real sample \(x\), used to estimate the parameters with the four methods under assessment, and then samples of the same size \(y\) were drawn from the \(G_0^I\) law with those parameters. The KS test was then performed between \(x\) and \(y\) for the null hypothesis \(H_0\) “both samples come from the same distribution,” and the complementary alternative hypothesis.

Table [IV] shows the sample \(p\)-values. It is observed that the null hypothesis is not rejected with a significance level of 5% in any of the cases. This result justifies the adequacy of the model for the data.

Fig. [I] shows the regions used for estimating the texture parameter under the influence of a corner reflector. Table [V] shows the estimates of \(\alpha\) for each rectangular region in the image of Fig. [II(b)]. The Maximum Likelihood, \(\frac{1}{2}\)-moment and Log-Cumulant estimators are unable to produce any estimate in small samples. They only produce sensible values when the sample is of at least 90 observations. The Log-Cumulant method is the one which requires the largest sample size to produce an acceptable estimate. The estimator based on the Triangular distance yields plausible values under contamination even with very small samples.

Table [VI] presents the \(p\)-values of the KS test, applied as previously described to the samples of Fig. [II(b)]. Estimators based on the \(\frac{1}{2}\)-moments and on Log-Cumulants failed to produce estimates in two samples and, therefore, it was not
possible to apply the procedure. The other samples did not fail to pass the KS test, except for the Blue sample and the Maximum Likelihood estimator. These results lead us to conclude that the underlying model is a safe choice regardless the presence of contamination and the estimation procedure.

V. CONCLUSIONS

We proposed a new estimator for the texture parameter of the $G_0^I$ distribution based on the minimization of a stochastic distance between the model (which is indexed by the estimator) and an estimate of the probability density function built with asymmetric kernels. We defined three models of contamination inspired in real situations in order to assess the impact of outliers in the performance of the estimators. The difficulties of estimating the textured parameter of the $G_0^I$ distribution were further investigated and justified throughout new theoretical results regarding its heavytailedness.

Regarding the impact of the contamination on the performance of estimators, we observed the following. Only Fractional Moment and Log-Cumulant estimators fail to converge.

- Case 1: Regardless the intensity of the contamination, the

![Graphs showing MSE for different sample sizes and contamination levels.](image)

Fig. 5. Sample mean squared error of estimates, Case 1 with $\alpha_2 = -15$ and $\epsilon = 0.01$. 
bigger the number of looks, the smaller the percentage of situations for which no convergence if achieved.

- Cases 2 and 3: the percentage of situations for which there is no convergence increases with the level of contamination, and reduces with $\alpha$.

In the single-case look the proposed estimator does not present excellent results, but it never fails to converge whereas the others are prone to produce useless output.

The new estimator presents good properties as measured by its bias and mean squared error. It is competitive with the Maximum Likelihood, Fractional Moment and Log-Cumulant estimators in situations without contamination, and outperforms the other techniques even in the presence of small levels of contamination.

For this reason, it would be advisable to use $\hat{\alpha}_T$ in every situation, specially when small samples are used and/or when there is the possibility of having contaminated data. The extra computational cost incurred in using this estimator is, at most, the twenty times the required to compute $\hat{\alpha}_{ML}$, but its advantages outnumber these extra computer cycles.

**APPENDIX**

Simulations were performed using the R language and environment for statistical computing [37] version 3.0.2. The adaptIntegrate function from the cubature package was used to perform the numerical integration required to evaluate the Triangular distance, the algorithm utilized is an adaptive multidimensional integration over hypercubes.

In order to numerically find $\hat{\alpha}_{LCum}$ we used the function uniroot implemented in R to solve (7).

The computer platform is Intel(R) Core i7, with 8 GB of memory and 64 bits Windows 7. Codes and data are available upon request from the corresponding author.

![Sample mean of estimates, Case 2 with $C = 100, \epsilon = 0.001$.](image)
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Fig. 7. Sample mean squared error of estimates, Case 2 with \( C = 100 \), \( \epsilon = 0.001 \).
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![Fig. 8](image-url)  
Fig. 8. Sample mean of estimates, Case 3 with $\epsilon = 0.005$ and $k = 2$. 
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Fig. 9. Sample mean squared error of estimates, Case 3 with $\epsilon = 0.005$ and $k = 2$.  

$$
\begin{array}{|c|c|c|c|c|}
\hline
L & \text{Case} & sample size & MSE \hline
1 & 3 & & \hline
3 & 3 & & \hline
8 & 3 & & \hline
\end{array}
$$
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Fig. 10. Real SAR image and the regions used to estimate the $\alpha$-parameter.

(a) Single-look E-SAR image with a corner reflector.

(b) Regions of interest.

Fig. 11. Samples of several sizes in a real SAR image with a corner reflector, used to estimate the $\alpha$-parameter.
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| TABLE III |
| --- |
| **Estimations of the \( \alpha \)-Parameter Using the Samples Shown in Fig. 10** |
| Color | size | \( \hat{\alpha}_{MV} \) | \( \hat{\alpha}_T \) | \( \hat{\alpha}_{Mom12} \) | \( \hat{\alpha}_{LCum} \) | time MV | time DT | time \( \hat{\alpha}_{Mom} \) | time \( \hat{\alpha}_{LCum} \) |
| Magenta | 100 | -1.9 | -2.7 | -1.9 | -1.7 | 0.03 | 5.85 | 0.03 | 0.02 |
| Green | 48 | -2.5 | -2.5 | -2.9 | -3.1 | 0.00 | 3.31 | 0.00 | 0.00 |
| Blue | 25 | -4.9 | -3.0 | N.A | N.A | 0.00 | 2.08 | 0.00 | 0.00 |
| Yellow | 90 | -6.2 | -5.1 | -6.6 | -6.8 | 0.00 | 5.16 | 0.00 | 0.00 |
| Red | 64 | -1.8 | -1.9 | -1.9 | -1.8 | 0.00 | 4.17 | 0.00 | 0.00 |

| TABLE IV |
| --- |
| **Sample \( p \)-Values of the Kolmogorov-Smirnov Test with Samples from the Image in Fig. 10** |
| Color | TestMV | TestDT | TestMom | TestLogCum |
| Magenta | 0.38 | 0.93 | N.A | N.A |
| Green | 0.11 | 0.93 | N.A | N.A |
| Blue | 0.01 | 0.11 | 0.400 | 0.11 |
| Yellow | 0.19 | 0.31 | 0.460 | 0.15 |
| Red | 0.23 | 0.12 | 0.008 | 0.02 |

| TABLE V |
| --- |
| **Estimations of the \( \alpha \)-Parameter Using the Samples Shown in Fig. 11(b)** |
| Color | size | \( \hat{\alpha}_{MV} \) | \( \hat{\alpha}_T \) | \( \hat{\alpha}_{Mom12} \) | \( \hat{\alpha}_{LCum} \) | time MV | time DT | time \( \hat{\alpha}_{Mom12} \) | time \( \hat{\alpha}_{LCum} \) |
| Magenta | 15 | -20.0 | -4.1 | N.A | N.A | 0.03 | 1.95 | 0.03 | 0.03 |
| Green | 42 | -9.2 | -5.0 | N.A | N.A | 0.00 | 4.04 | 0.00 | 0.00 |
| Blue | 90 | -3.5 | -2.7 | -4.7 | -14.2 | 0.02 | 4.85 | 0.00 | 0.00 |
| Yellow | 156 | -2.2 | -1.8 | -2.6 | -3.4 | 0.01 | 8.35 | 0.00 | 0.00 |
| Red | 225 | -1.9 | -1.7 | -2.1 | -2.5 | 0.02 | 10.97 | 0.00 | 0.00 |

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