#### **ORIGINAL ARTICLE**



# **Percentage porosity computation of three‑dimensional non‑convex porous geometries using the direct Monte Carlo simulation**

**Mauricio Campillo1 · Pablo Pérez1 · Jorge Daher1 · Luis Pérez<sup>1</sup>**

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#### **Abstract**

The pursuit of more representative numerical models for open-cell metallic foams requires the computation of volume and percentage porosity of geometries containing randomly distributed interconnected pores, which is one of the main characteristics that determines its mechanical properties. From a mathematical standpoint, the analytical defnition of foam geometries forms a three-dimensional non-convex set. It is known that the volume computation of *n*-dimensional polytopes and sets is a P-hard problem. A common way to approach this problem is using the Monte Carlo techniques; however, eforts are oriented toward the treatment of convex polytopes and polyhedrons. In this article, the Direct Monte Carlo Simulation (DMCS) is used to compute the percentage porosity of three-dimensional non-convex sets. A single-thread Python code was implemented, and tests were run to estimate the percentage porosity of three-dimensional open-cell porous geometries. Measurements of percentage porosity and runtime requirements over cubical and cylindrical geometries containing from 100 to 4000 overlapping spherical pores showed high accuracy and consistency in non-convex three-dimensional sets, while the proposed algorithm achieved a signifcant reduction in computing time with respect to the currently available method. In the same manner, results from the proposed algorithm were compared with a similar software available, showing a gain in both performance time and accuracy.

**Keywords** Direct Monte Carlo simulation · Percentage porosity · Non-convex sets · Metallic foams · Python

### **1 Introduction**

Metallic foams have gained importance in recent years which can be verifed by considerable number of investigations in the literature on this subject. They are composed of a metal-based structure with internal cavities called pores, similar to a conventional Metal Matrix Composite (MMC) but containing a void secondary phase. Percentage porosity is measured as the volume fraction of the void phase to the overall volume, which is the complementary percentage of the relative density of the solid phase. Porosity can be introduced in metal foams using a wide variety of fabrication methods, including processes using metals in solid, liquid, and gaseous states [\[1](#page-21-0)]. The main characteristic that determines the mechanical properties of metallic foams is the

 $\boxtimes$  Mauricio Campillo mauricio.campillo@postgrado.usm.cl percentage porosity. This characteristic is mainly infuenced by the random distribution of pores and the manufacturing process which establishes the internal structure of the foams. When pores exhibit a structure where a membrane bounds each one independently (e.g., honeycombs), it is said to be a closed-cell structure, while if interconnection exists between pores (e.g., sponges), it is said to be an open-cell structure [[2\]](#page-21-1).

The Finite-Element Analysis (FEA) has been a powerful and feasible numerical technique to model the mechanical behavior of both open-cell and closed-cell foams in the aid to identify plausible fabrication routes, depending on its desired microstructure and macroscopic mechanical properties. However, several challengers remain to be overcome to achieve better accuracy in the representation of its three-dimensional internal microstructure and macroscopic mechanical behavior, as its properties are highly dependent of the quality of the Computer-Aided Design (CAD) geometrical models [\[3](#page-21-2)]. FEA of heterogenous and porous media is commonly carried out by a Representative Volume Element (RVE) analysis [\[4](#page-21-3)], where the mechanical properties

<sup>1</sup> Department of Mechanical Engineering, Universidad Técnica Federico Santa María, Av. España 1680, Casilla 110-V, Valparaíso, Chile

of the medium are obtained by solving a Boundary-Value Problem (BVP) over a small and rather simple domain from which the equivalent mechanical properties are transferred to a more complex one assumed as a continuum [\[5](#page-21-4)[–7\]](#page-21-5).

One approach to improve the representativeness of the numerical results obtained in FEA-based models is by the inclusion of randomly distributed spherical pores, mimicking the natural distribution that can be observed in the microstructure of open-cell metallic foams obtained using the Space Holders Phase (SHP) technique, either by conventional powder metallurgy (PM) [\[8](#page-21-6)] or by infltration of liquid metal [\[9](#page-21-7)]. These models must represent adequately the topological parameters of the metallic foams (e.g., pore distribution and sizes, interconnection, porosity percentage, etc.). To generate these metallic foam models, one of the most used mechanisms is using a CAD software package and executing a series of sequential command operations contained in a script [\[10](#page-21-8)[–12](#page-21-9)], examples of these models can be seen in Fig. [1](#page-1-0).

However, despite good results for the mechanical behavior of foams are reported in literature, the script commands' processing time is considerably excessive in relation to the FEA total time, which poses the necessity of developing an alternative to overcome this shortcoming. Figure [2](#page-1-1) shows an example of the average time required to import a geometry into FEA preprocessor ANSYS v18 Design Modeler module and a quadratic relation with the number of cutting operations required to include pores.

In three dimensions, RVEs of porous media with open interconnected pores can be defned by the intersection of the interior of an external body (e.g., a cube or a cylinder) and the exterior of the union of spheres, which are allowed to intersect the exterior surface of the external body, to produce surface porosity seen in Fig. [1.](#page-1-0) Due to the overlapping pores and the manner in which the set is defned, both the geometry of the RVE and the union of the spheres of the void phase are non-convex sets. Hereafter, the concept of non-convex set may be referred to either of the prior.

In this article, an algorithm based on the Direct Monte Carlo Simulation (DMCS), implemented in Python language, is developed to estimate the percentage porosity of CAD generated metallic foam geometries. A grid cell-based linked list strategy and a Latin Hypercube Sampling (LHS) approach are used to achieve better computational efficiency and convergence [[13](#page-21-10)]. A well-defned data set composed of 40 individual and distinct non-convex sets (i.e., 3D foam geometries) was used to evaluate the proposed method. This implementation has shown a time reduction between 80 and 90% with respect to the execution of the complete extruding and cutting operations by a CAD software package, as well as better accuracy and runtimes when compared to McVol [[14\]](#page-21-11), which is a Monte Carlo-based software intended to



<span id="page-1-0"></span>**Fig. 1** 3D foam geometries of solids containing spherical overlapping pores of 350–450 µm in diameter: **a** porous cylinder of diameter 2.25 mm and height 1.8 mm with 100 pores and 75.03% porosity and **b** porous cubes of size 3.6 mm with 1320 pores and 70.08% porosity



<span id="page-1-1"></span>**Fig. 2** Average loading time measured for cube-based foam geometries of size 3.6 mm into FEA Preprocessor Ansys v18 Design Modeler containing diferent number of spherical overlapping pores of 350–450 µm in diameter via JavaScript scripts execution

compute molecular volume of proteins, developed by Till et al.

### **2 Methodology**

### **2.1 CAD geometries' generation**

The process to generate the CAD geometries of the foams is based in a basic three-step procedure, which is schematically shown in Fig. [3](#page-2-0), for a two-dimensional case. First, a Discrete-Element Method (DEM) [[15\]](#page-21-12) simulation is carried out, from which the instantaneous element positions are used as the center position distribution of the pores (Fig. [3a](#page-2-0)). Then, using the defned pores center distribution, each pore

diameter is redefned to meet the required criteria to ft the intender microstructure, such as pore interconnectivity and size distribution (Fig. [3b](#page-2-0)). Finally, a CAD geometry is completely defned as a script, which after being imported to an FEA Preprocessor or CAD package is seen as the remaining solid phase of the intersection between the interior of the external body and the exterior of the union of spheres, as shown in Fig. [3](#page-2-0)c.

Two sets of 3D CAD geometries containing randomly distributed spherical pores have been generated using DEM, where one set corresponds to cylinder-based geometries, while the second set is cube-based, similar to those shown in Fig. [1.](#page-1-0) These two general shapes have been selected due to their relevance on both numerical and experimental studies of mechanical behavior of metallic foams. While experimental tests use cylindrical specimens for compressive testing [\[16](#page-21-13), [17\]](#page-21-14), numerical simulations are oriented toward the characterization of mechanical behavior based on a reduced RVE in association to a multiscale scheme  $[18, 19]$  $[18, 19]$  $[18, 19]$  $[18, 19]$ , and hence, the cubic-shaped geometry is more suited for orthogonal load testing [[20–](#page-21-17)[22\]](#page-21-18).

Originally presented to model the behavior of granular media, the DEM is a powerful numerical tool to model the random distribution of pores generated in foam fabrication methods such as SHP due to the solid mixing process [\[17](#page-21-14)] or in liquid metal infltration process [\[23](#page-21-19), [24](#page-21-20)]. The DEM represents the medium as a collection of material particles exhibiting independent rigid body motion behavior where the total external force acting over each particle is determined by the interaction with neighboring particles in which each pair interaction force is governed by a low-range contact force law [[15\]](#page-21-12). In this work, a Hertz-type law with dampening has been considered for this purpose, where the total force has a normal component and a tangential component contributing to the linear and angular momentum Newton's conservation law correspondently. This type of contact law is found to be usual on the simulation of granular media [[25](#page-21-21)[–27](#page-21-22)].

Geometries based on cylinders of 6.75 mm in diameter (*D*) and 5.4 mm in height (*H*), and cubes of size (*A*) 6.75 mm containing 100, 500, 1000, 2000, and 4000 pores

with diameters (*d*) ranging from 350 to 450 µm have been created using DEM open software LIGGGHTS [[28](#page-21-23)]. An *H*/*D* ratio of 0.8 has been chosen to avoid flexural deflection and inelastic buckling, in accordance with geometric recommendations for short specimens in ASTM E9-09 standard [[29\]](#page-21-24). A ratio of characteristic length to maximum pore size of 15 (i.e., *D/d* for cylinders and *A/d* for cubes) was chosen, to allow the inclusion of a wide range of number of pores throughout the tests. The maximum number of pores was set to 4000, as loading times into the FEA preprocessor becomes excessive.

Cylinder-based geometries have been set to have its revolution axis coaxial with the *Z*-axis, while the base is contained in the  $X-Y$  plane protruding through the  $Z > 0$ subspace. On the other hand, cube-shaped geometries were oriented to have three of its faces aligned with the *X*–*Y*, *Y*–*Z,* and *Z*–*X* planes, while its remaining three faces contained in the  $X, Y, Z > 0$  subspace.

DEM special domain used considered 1 time the maximum pore size (450 µm) for each side, being the radius, lower and upper height limits for cylinders and for each lower and upper direction for cubes, to promote the presence of pores on the surface of the resulting geometry and particle size was set to 0.25 times the maximum pore size  $(112 \mu m)$  to achieve interconnection between pores on the generated CAD geometries.

For both geometry types, the conditions for particles insertion were initial velocity (0.5 mm/s on each direction), high Young's modulus (5.0e8 MPa), and coefficient of restitution (0.95) and near zero gravity with the intention of generating high level of interaction between particles and maintaining high total kinetic energy in the system during simulation, according to Pérez et al. [\[9\]](#page-21-7).

Results from DEM simulations were post-processed using a purposely developed Python script to generate ANSYS v18 Design Modeler input JavaScript scripts, using the corresponding center of the particles and assigning a uniformly distributed random value for each pore diameter between 350 and 450 µm, similar to Ref. [[9\]](#page-21-7). The aforementioned script follows the following pseudo-code structure.

<span id="page-2-0"></span>**Fig. 3** Two-dimensional representation of the RVE generation of porous media with opencell porosity: **a** Pores' center distribution obtained by DEM; **b** Uniform random definition of pores diameter; and **c** resulting foam geometry



```
1. #Define Type of Geometry
2. (Geo.Type) = {CUBE, CYLINDER}3. #Define Characteristic Lengths
4. Switch Case (Geo.Type):<br>5. CUBE: (Geo.charleng
        CUBE: (Geo(charlength) = [(A)]6. CYLINDER: (Geo.charlength) = [(H), (D)]
7. #Define Pore Minimum and Maximum Diameter
8. (pore diam) = [(min diam), (max diam)]9. #Retrieve List of Centers
10. input_file.open()
11. for each line in (input_file):
12. Append info to (X_vector) and (Y_vector) and (Z_vector)
13. input_file.close()
14. #Build JavaScript file
15. output_file.open()
16. #Set Global Dimensions<br>17. Switch Case (Geo.T
17. Switch Case (Geo.Type):<br>18. CUBE: output file.
            CUBE: output_file.write((A))
19. CYLINDER: output_file.write((H), (D))
20. #Write Instructions for Basic Extrusion
21. Switch Case (Geo.Type):
22. CUBE:<br>23. 0
            output file.write(Build Square of Edge (A))
24. output_file.write(Extrude Square (A) Units)<br>25. CYLINDER:
25. CYLINDER:<br>26. outpu
            output file.write(Build Circle of Diameter (D))
27. output_file.write(Extrude Circle (H) Units)
28. #Write instructions for Pores as Cutting Operations
29. for i in range(list_of_pores):
        #Define Pore Center and Diameter
31. (diameter) = random.uniform((pore_diam.min),(pore_diam.max))
32. (center)=(X_vector[i],Y_vector[i],Z_vector[i])
33. #Write Instructions
34. output file.write(Create Sphere Cut with (center), (diameter))
35. output_file.close()
```
### **2.2 The Monte Carlo simulation**

Volume computation of polytopes is a problem which has been studied for at least 3 decades [[30](#page-21-25)[–32](#page-21-26)], and since then, author has been using diferent approaches to provide polynomial solutions to this problem, at least for low-dimensional cases. One of the most common approaches is based on the principle of divide-and-conquer, and the polyhedron is divided into smaller and simpler instances where the volume can be computed after by addition. Algorithms such as volume decomposition [[33](#page-21-27)] and triangulation [[34\]](#page-21-28) are examples of this approach. Another commonly used approach is based on the approximation of volume by means of the Monte Carlo (MC) simulation. Here, various algorithm implementations have been developed for the volume estimation of convex *n*-dimensional polytopes for which validation is restricted to low-dimension geometries with known volume, whether it is analytically or not [[35,](#page-22-4) [36](#page-22-5)]. These algorithms are typically based under a 'hit-and-run' technique and random walks used for a uniform sampling throughout the geometry surface of the *n*-dimensional volume, providing a polynomial time solution for the treatment of convex polytopes [[37\]](#page-22-0). It is worth noticing that a general non-convex set can be approximated by a non-convex polytope and the approximating non-convex polytope can be rewritten as a union of convex and simpler polytopes. Hence, in principle, the volume of non-convex sets can be computed using algorithms and techniques intended for the treatment of convex polytopes, but the convex representation of non-convex polytopes can be extremely expensive and algorithms efficiency can be severely affected by the number of instances needed to represent the original set [\[38](#page-22-1)].

Although spheres from which pores are generated correspond to convex three-dimensional bodies, the union of two or more pores, which is called a macro-pore, is not guaranteed to hold this property [[39](#page-22-2)]. In the case of the CAD geometries treated in this work, the existing overlap between spheres in relation to the sphere diameters is mainly unknown, which supposes that the generated volume is highly likely to form a non-convex set. Due to this nonconvexity, the application of the previously described algorithms is restricted and a diferent technique, such as simple quadrature, may be applicable [[40\]](#page-22-3). This problem has been tackled in felds such as biochemistry, where the volume of proteins, represented as a collection of overlapping spheres, can be computed using algorithms based on volume decomposition [\[41\]](#page-22-6) or the MC simulation [[14\]](#page-21-11).

#### **2.2.1 Direct Monte Carlo simulation (DMCS)**

In this work, the MC simulation is used as a mechanism to gather information about an analytic 3D CAD geometry. In this approach, a set of *N* independent Bernoulli random variables,  $X_j^N$ , are evaluated, each one according to a probability distribution function such as:

$$
f(X) = \begin{cases} 1 & X \in solid \\ 0 & \text{Elsewise} \end{cases}
$$
 (1)

where each  $X_j^N$  corresponds to an independent three-dimensional vector in the subspaces enclosed by the exterior surface of the geometry. As all the information relative to the pores is known a priori (being its centers and radii), an alternative which poses an equivalent but less expensive to solve problem is the one on estimating the void volume inside the geometry and then to compute the solid portion as the diference between the solid volume and the estimated void volume. In this manner, the probability distribution function proposed in Eq. [1,](#page-4-0) can be rewritten as:

$$
f(X) = \begin{cases} 1 & X \in \text{a void} \\ 0 & X \notin \text{any void} \end{cases}
$$
 (2)

With this, for each Bernoulli random variable, if the vector  $X_j^N$  happens to reside inside a pore, it considers a 'hit', while if it does not reside inside any void, it is considered a 'miss'. For any given number *N* of Bernoulli random variables, it is known that they follow a Binomial distribution, where its estimation,  $\lambda_N$ , of the expected value,  $\lambda$ , is determined as the product between the probability of success (or 'hit') and the total number of independent variables, *N*. As the expected value of any given distribution is known from the evaluation of every variable, the probability of 'hit' on any Binomial distribution,  $p_N$ , may be determined as a function of the number of independent variables as:

$$
p_N = \frac{\lambda_N}{N}.\tag{3}
$$

With this, as the number of independent Bernoulli variables increases, the estimator,  $\lambda_N$ , tends to the real expected value,  $\lambda$ , of the Binomial distribution  $[42]$  $[42]$  and, as a consequence, the real probability of success, *p*, is found. This is:

$$
\lim_{N \to \infty} p_N = \lim_{N \to \infty} \frac{\lambda_N}{N} = p. \tag{4}
$$

In terms of the DMCS, each Bernoulli random variable evaluation of the binary probability function in Eq. [2](#page-4-1) is named a *membership oracle call*, since there is an algorithm established a priori to determine whether the returned for an arbitrary input value is 0 or 1. Later, the wanted percentage porosity of a CAD geometry foam is associated with the probability of success of the related Binomial distribution followed by the random points submitted to the membership oracle.

#### **2.2.2 Random point generation**

<span id="page-4-0"></span>A Latin Hypercube Sampling (LHS) [\[34](#page-21-28)] strategy was used to generate the sampling points to be submitted to the membership oracle. To implement the LHS, each dimension of the domain is stratifed in *m* equiprobable strata. Later, each stratum is randomly sampled one time. For cubes, each dimension has a span from 0 to A; therefore, each stratum has equal length  $l = A/m$ . Whether for cylinders, to achieve the equiprobability condition in cylindrical coordinates, it is required that each stratum has a diferent length in the radial direction (*R*), so the volume of each stratum can remain constant, and hence, the *j*th radial stratum will be of length  $l_j^R$ :

$$
l_j^R = \sqrt{\frac{(j-1)}{k}} \left(\frac{D}{2}\right). \tag{5}
$$

<span id="page-4-5"></span><span id="page-4-2"></span>Later, each stratum,  $S_{i,j,k}$ , is defined as:

<span id="page-4-1"></span>
$$
\mathbf{S}_{i,j,k}^{CUBE} = \begin{bmatrix} il, (i+1)l \\ il, (j+1)l \\ kl, (l+1)l \end{bmatrix} \mid \{i,j,k\} \in [0, 1, ..., (m-1)], \quad (6)
$$

whether for cylinders, each stratum is defned by:

$$
S_{i,j,k}^{CYL} = \begin{bmatrix} \sqrt{i}l^R, \sqrt{i+1}l^R \\ jl^{\theta}, (j+1)l^{\theta} \\ kl^z, (k+1)l^z \end{bmatrix} | \{i,j,k\} \in [0, 1, ..., (m-1)],
$$
\n(7)

<span id="page-4-4"></span>where the lengths of the strata in the other directions  $l_{\theta} = 2\pi/m$  and  $l_z = H/m$  are constant. Although, for arbitrary parallelepipeds, individual lengths,  $l_i = A_i/m$  shall be used for the *i*th dimension in Eq. [6,](#page-4-2) the proposed implementation does not feature this option.

Finally, for each stratum,  $S_n^N = S_{i,j,k}$ , a uniform random point  $(X_n^N)$  is generated where:

<span id="page-4-3"></span>
$$
X_n^N = X_{i,j,k} = \{ X \in S_{i,j,k} \subset \mathbb{R}^3 \forall i,j,k \in \{0,1,\ldots,(m-1)\} \},\tag{8}
$$

where  $n = [0, (m^3 - 1)]$  or in terms of the total number of Bernoulli random variables,  $n = [0, (N - 1)]$ .

#### **2.2.3 Porosity estimation**

The expected value,  $\lambda_k$ , for an arbitrary number, *k*, of random measurements of the random variable *X* distributed Binomial with parameters *n* and *p* is known to tend to the expected value of the Binomial distribution,  $\mu$ , this is:

$$
\lambda(\bar{X}) = \sum_{i=1}^{k} \frac{1}{k} \mu_i = n\left(\frac{\mu}{n}\right) = \mu \cdot |k \to \infty \,.
$$
 (9)

The prior is based on the premise that the random variable *X*, which represents the number of successes in any random sampling of size *n*, is distributed Normal with mean  $\mu = np$  and variance  $\sigma^2 = np(1 - p)$ . This is:

$$
X = N(\mu, \sigma^2) = N(np, np(1-p)).
$$
\n(10)

A standardization of the later Normal distribution results in a new random variable *Z* also distributed Normal defned by:

$$
Z = \frac{X_i - n_i p}{n_i p (1 - p)} = \frac{\frac{X_i}{n_i} - p}{p (1 - p)}.
$$
\n(11)

The above random variable *Z*, measured by  $X_i/n_i$  associated with the previously defned Binomial distribution, is then distributed Normal with mean value *p* and variance  $p(1-p)$ . As mean value of a collection of *k* random samples *Z̄* is known to be

$$
E(\bar{Z}) = \mu(z) = \frac{1}{k} \sum_{i=1}^{k} Z_i = 0 | k \to \infty,
$$
 (12)

it is necessary that the summation term in Eq. [10](#page-5-0) satisfes:

$$
\frac{1}{k}\sum_{i=1}^{k}\left(\frac{X_i}{n_i}-p\right)=0|k\to\infty.\tag{13}
$$

From Eq. [11,](#page-5-1) given that

$$
\frac{1}{k} \sum_{i=1}^{k} p = \frac{1}{k} (kp) = p, |k \to \infty,
$$
\n(14)

it follows consequently that

$$
p = \frac{1}{k} \sum_{i=1}^{k} \frac{X_i}{n_i} = \frac{1}{k} \sum_{i=1}^{k} p_i = \bar{p} |k \to \infty.
$$
 (15)

Hence, for a random sample of measurements of the probability of success,  $p_i = X_i/n_i$ , of the Binomial distribution done by arbitrary samples over the distribution, it is guaranteed that its mean value,  $\bar{p}$ , represents an estimator of the unknown probability of success, *p* [[43\]](#page-22-8).

#### **2.2.4 Convergence**

As it is known that the sample mean,  $\bar{p}$ , in general will not coincide with the random distribution mean, *p*, it is a common practice to evaluate the accuracy of the results from MC simulations based on the distribution of the independently computed sample mean. Therefore, rather than computing the exact value of the distribution,  $p$ , a confidence interval is established to limit the error on the estimator of the mean, *p̄* [[44](#page-22-9), [45](#page-22-10)]. For normally distributed variables with unknown standard deviation and small sample sizes, *n*, it is well known that a confdent interval for the mean, with a confidence level of  $(1 - \alpha)$ , can be established based on the estimator of both the mean,  $\bar{p}_n$ , and the standard deviation, *S<sub>n</sub>*, or rather the standard error,  $S_n / \sqrt{n}$ , by:

<span id="page-5-2"></span><span id="page-5-0"></span>
$$
\Pr\left[\bar{p}_n - t_{n-1}\left(1 - \frac{\alpha}{2}\right)\frac{S_n}{\sqrt{n}} \le p \le \bar{p}_n + t_{n-1}\left(1 - \frac{\alpha}{2}\right)\frac{S_n}{\sqrt{n}}\right] = 1 - \alpha,
$$
\n(16)

where  $t_{n-1}$  is the Student's distribution of  $(n - 1)$  degrees of freedom. Furthermore, when the sample size is large enough (e.g.,  $n \ge 30$ ), the Student's distribution can be approximated by the Standard's Normal distribution (*Z*). Hence, Eq. [16](#page-5-2) can be restated as:

<span id="page-5-3"></span><span id="page-5-1"></span>
$$
\Pr\left[\bar{p} - Z\left(1 - \frac{\alpha}{2}\right)\frac{S_n}{\sqrt{n}} \le p \le \bar{p} + Z\left(1 - \frac{\alpha}{2}\right)\frac{S_n}{\sqrt{n}}\right] = 1 - \alpha.
$$
\n(17)

When the standard normal percentile  $Z(1 - \alpha/2) = 3$ , Eq. [17](#page-5-3) takes the form of the well-known three-sigma rule, which established a confdence interval with a confdence level of 99.97%. Hence, for any sample size, an uncertainty level, ε, can be established, although its confdence level will increase as the sample size increases. With this, a stopping rule for the sampling algorithm can be established based on the sample size and its standard deviation as:

<span id="page-5-5"></span>
$$
a\frac{S_n}{\sqrt{n}} \le \varepsilon \Rightarrow \frac{S_n}{\sqrt{n}} \le \frac{\varepsilon}{a},\tag{18}
$$

where the parameter *a* is the number of standard deviations allowed in the uncertainty and the corresponding confdent level is related to the confdence level of either the student or the standard normal distribution percentile as:

<span id="page-5-4"></span>
$$
a = \begin{cases} t_{n-1} \left( 1 - \frac{a}{2} \right) & n < 30 \\ Z \left( 1 - \frac{a}{2} \right) & n \ge 30. \end{cases} \tag{19}
$$

In this article, the three-sigma rule and a minimum sample size of 4 MC experiments have been adopted for the stopping rule of the MC simulations, which establishes a minimum confdence level of 94%. This is:

$$
\Pr\left[\bar{p}_n - 3\frac{S_n}{\sqrt{n}} \le p \le \bar{p}_n + 3\frac{S_n}{\sqrt{n}}\right] \ge 0.939 |n \ge 4. \tag{20}
$$

#### **2.2.5 Performance**

For each MC experiment, *N* calls to the membership oracle are needed. In addition, each call to the oracle is resolved in

a maximum of *m* independent operations, given by the evaluation of the distance between the random sample point and the center of the pores contained in all the neighbor cells, defned by the linked lists; this means that, in the worst-case scenario, the membership oracle will evaluate the pores contained in a maximum of 27 cells (i.e., the central cell and all the adjacent cells that shares one face or vertex with it) as every other pore is discarded since they are too far from the point so that it is possible for it to be inside them. With this, if each independent operation made by the oracle is considered equal to one FLOP, it is expected for the algorithm to perform a complete experiment in a total amount of FLOP with an upper bound given by:

$$
FLOP_{total}^{N} = N \cdot m. \tag{21}
$$

Hence, the total FLOPs required to perform an arbitrary MC experiment for any given geometry are determined by the number of pores contained in the geometry and the number of calls made to the oracle. If an iterative process is considered, where a total of *k* MC independent experiments are performed, the total FLOP is determined by the summation of the times required to execute each individual MC experiment. Assuming a process where each MC experiment is preformed using *N* calls to the oracle, it is expected for the algorithm to require a total number of computing operations of:

$$
FLOP_{total} = k \cdot N \cdot m. \tag{22}
$$

From Eq. [22,](#page-6-0) the implemented algorithm is expected to exhibit a time performance linear with both the number of pores, given by the expected number of pores, *m*, contained in the neighboring cell space and the base number of calls to the oracle, *N*. In this work, the implemented algorithm considers a constant sampling size, similar to the test procedure presented by Liu et al. [[36](#page-22-5)], although an LHS strategy was adopted to improve sampling efficiency. A total of 50 strata per dimension, which translates to a sample size of 125 thousand points per MC experiment, were used. Using the LHS strategy and linked lists to identify neighboring pores, it is expected to achieve faster convergence than uniform random sampling as generated samples are non-collapsing in space and linked lists allow to dismiss verifcation of very distant pores and, therefore, unnecessarily to check.

#### **2.3 Implementation**

An algorithm to compute porosity percentage based on the DMCS introduced in the previous section was implemented. A complete version of the Python code can be found in ["Appendix A](#page-15-0)". For the execution of the code, the usage of Python libraries numpy for scientifc computing and numba, through its jit decorator, for computational optimization, along with the built-in modules math, random, and sys for fle handling and arithmetic operations was required. When a geometry is under analysis, independent MC experiments are performed using a given number of samples size, *N*, from which the success probability,  $p_i$ , is estimated. MC experiments are performed until the standard error,  $S_n / \sqrt{n}$ , of the estimated success probability,  $\bar{p}$ , given by the samples mean, reaches a prescribed error,  $\epsilon/3$ .

### **2.3.1 Function for retrieving geometry information contained in FEA preprocessor script**

<span id="page-6-0"></span>The frst step in the code execution consists in gathering the information regarding the geometry under analysis which is retrieved from the corresponding input JavaScript scripts for ANSYS preprocessor. This is achieved by the sequential reading of the script fle, from which general dimensions of the geometry (i.e., diameter, *D*, and height, *H*, for cylinders and the side, *A*, for cubes) as well as the complete lists for the pores (i.e., center position in directions *x*, *y*, *z* and radii). Once all the relevant information is retrieved, the fle is closed and dismissed.

1. def retrieve file info(inp file): 2. # File is read line by line<br>3 with onen(inn file) as with open(inp\_file) as ifile: 4. n\_void = 0 radii, x, y, z =  $[]$ ,  $[]$ ,  $[]$ 6. **for** line **in** ifile: 7. **if** 'var H=' **in** line: # Cylinder he ight 8.  $l_H = float(linestrip("var H=").strip("j\n");$ <br>9. if 'var D=' in line: 9. **if** 'var D=' **in** line: # Cylinder di ameter 10.  $1_D = float(line.strip("var D=").strip("; \n',\n')$ 11. **if** 'var A=' **in** line: # Cube edge s ize 12. l A = float(line.strip("var A=").strip(";\n")) 13. **if** ('radio=0;' **not in** line **and** 'radio=' **in** line): # Sphere radi us 14. **n\_void += 1**<br>15. **n\_void += 1** radii.append(float(line.strip("radio=").strip(";\n"))) 16. **if** ('posicionx=0;' **not in** line **and** "posicionx=" **in** line): # Sphere cent er X-coordinate 17. <br> **17. x.append(float(line.strip("posicionx=").strip("; \n")))**<br> **18. if ('posiciony=0:' not in line and "posiciony=" in line):** 18. **if** ('posiciony=0;' **not in** line **and** "posiciony=" **in** line): # Sphere cent er Y-coordinate 19. y.append(float(line.strip("posiciony=").strip("; \n")))<br>20. if ('posicionz=0:' not in line and "posicionz=" in line): 20. **if** ('posicionz=0;' **not in** line **and** "posicionz=" **in** line): # Sphere cent er Z-coordinate 21. z.append(float(line.strip("posicionz=").strip("; \n"))) 22. ifile.close() 23. # Checks retrieved type of geometry and sets zero the unused variables<br>24. **if not 'l\_A' in locals():** # Checks if i  $if not 'l A' in locals():$ s not a Cube 25.  $1_A = 0$ 26. **elif not** ('l\_D' **in** locals() **and** 'l\_H' **in** locals()): # Checks if i s not a Cylinder 27.  $1_H$ ,  $1_D = 0$ , 0<br>28. else: 28. **else**: # Acts if doe s not detect neither a cube nor a cylinder 29.  $1_A$ ,  $1_D$ ,  $1_H = 0$ ,  $0$ ,  $0$ <br>30.  $\frac{1}{2}$  print('Frror retrieving 30. **print**('Error retrieving file info. Not a Cylinder nor a Cube was found.') 31. sys.exit(2) 32. **return** l\_H,l\_D,l\_A,np.asarray(x),np.asarray(y),np.asarray(z),np.asarray(radii),n\_ void

#### **2.3.2 Function to build the linked lists**

The process of building the linked lists was capsuled in a function called build\_lists. This function takes as input the information of the pores (i.e., pores' center and radii) and returns as output the linked lists head and list and two arrays containing the values of the corresponding grid lengths and the point from which the grid is deployed, respectively. First, temporal arrays are built to capture the extreme most coordinates that the spheres will have in the *n*-dimensional space (three-dimensional for this application). This is achieved by adding and subtracting the corresponding radii to each sphere center and selecting the minimum and maximum values (lines 3, 4, and 5). Then, the corresponding number of cells in each dimension is defned (line 8) by dividing the diference between the two extreme values by the biggest sphere diameter (line 6), and by the *math.ceil()* function, the closest upper integer is selected. With this, it is clear that no pore residing more than one cell away can be considered for evaluation, later by the membership oracle, as it will be away from the random point by, at least, two times the maximum radius. Later, the corresponding grid lengths are computed by dividing the corresponding dimensional span by the recently defned number of cells (line 9). Finally, for each sphere in the array, its corresponding *cell* value is computed and stored in the linked lists by flling the corresponding cell with its index in the *head* list and pushing the already stored information into the *list* list under the corresponding index.



### **2.3.3 Function for the execution of independent MC experiments**

Once all the information that defnes the geometry to be analyzed is retrieved, an iterative cycle is generated to perform each MC experiment. For this purpose, the process was encapsulated as a function, monte carlo exp. Each independent simulation consists in a single uniform random sampling of each domain strata; therefore, a total of  $N = \prod_{i=1}^{3} n_i$ random points are generated throughout the entire domain, in a non-collapsing way. Each random 3D point is defned as a vector in Cartesian coordinates (i.e., *dart[i]*) and then is parsed to the membership oracle to resolve whether it corresponds to a *'hit'* or a *'miss'*. As for the cylinder-based geometries, cylindrical coordinates are used to stratify the domain; each generated random point (*temp*) must be transformed to its corresponding Cartesian coordinates (*dart*) prior to submission to the membership oracle. These random points correspond to those defned in Eq. [8](#page-4-3), where each stratum is defned in Eq. [6](#page-4-2) for cube-based and in Eq. [7](#page-4-4) cylinderbased geometries.

Once the *N* random points are evaluated, the quotient of the summation of hits over every dimension and the total generated points is returned to the main program as the computed success probability of the experiment,  $p_i$ , according to Eq. [15.](#page-5-4)

```
1. def monte_carlo_exp(H, D, A, x_vec, y_vec, z_vec, r_vec, seeds):<br>2. # Allocates list for sampling results
        # Allocates list for sampling results
3. sampling_list = np.zeros([nx,ny,nz])
4. for k in range(nz):
5. for j in range(ny):
6. for i in range(nx):
                    # Takes a smple from the [i,j,k] stratum
8. if A!=0: # CUBE
9. dart = [random.uniform(dx*(i), dx*(i+1)), random.uniform(dy*(j), dy)*(j+1)), random.uniform(dz*(k),dz*(k+1))]10. if (D != 0 and H != 0): # CYLINDER
11. tmp = [random.uniform(dr*math.sqrt(i),dr*math.sqrt(i+1)), random.
   uniform(dt*(j),dt*(j+1)), random.uniform(dz*(k),dz*(k+1))]
12. dart = [tmp[0]*np.cos(tmp[1]), tmp[0]*np.sin(tmp[1]), tmp[2]]<br>13. # Evaluates the sampling point
                    # Evaluates the sampling point
14. sampling_list[i,j,k] = member_oracle(np.asarray(dart), head, list, np
.asarray([x_vec, y_vec, z_vec]), r<sub>_</sub>vec, vecDelta, vecMin)<br>15. # # Computes expected value E[x]# Computes expected value E[x]
16. hits_num = sum(sum(sum(sampling_list)))<br>17. # Returns the computed probability p(x,# Returns the computed probability p(x,n) = E[x]/n18. return float(hits_num/seeds)
```
Results in this article were obtained by stratifying the three-dimensional space in 50 strata per spatial dimension, giving a sample size of 125 thousand points for each MC experiment.

### **2.3.4 Function for the evaluation of membership oracle calls**

When any random sampling point, defned by its coordinates (i.e.,  $vecP[i]$ ), is parsed to the membership oracle to be evaluated, the function member\_oracle was generated to answer based on the information given a priori regarding the list of voids, defned by the center positions, *vecPores*, and its radii,

*vecRadii*. This function identifes the corresponding grid cell associated with the coordinates of the random point, and then, it iterates looking for the surrounding cells checking if any of the spheres that lies in the neighborhood of the corresponding cell will satisfy the *'hit'* condition, using the linked lists. This *'hit'* condition, as stated in Eq. [2](#page-4-1), means that if the distance from the point to the center of a sphere is less than its radius, then the point lays inside the sphere, returning a 1 as a result, hence *'hit'*, stopping the iteration. In the case of after checking all the relevant cells, no *'hit'* is found, the function returns a 0, hence *'miss'*. The numba decorator jit for Just-In-Time compilation is used to speed up the code.





<span id="page-9-0"></span>**Fig. 4** Computed percentage porosity and standard error versus number of generated random points for a cube-based geometry of size 6.75 [mm] with 4000 pores (CUBE\_4000\_1) using 125,000 random points per iteration

#### **2.3.5 Main program**

To run the previously showed functions, a short main program is coded at the end of the Python script fle, where the geometry information (i.e., flename with its extension) is provided as system argument and a few computation parameters (i.e., LHS grid size, uncertainty tolerance, and minimum MC iterations) are defned in lines 2 through 4. After that, the function retrieve\_fle\_info is called in line 17 and all the information regarding the geometry is gathered according to the already explained structure. Subsequently, in lines 19 and 20, the corresponding domain strata are defned based on the retrieved type of geometry. Then, in line 23, the *build\_lists* function is called to generate all the required data to speed up the membership oracle response. With this, from line 26 through 31, an iterative procedure is defned to perform subsequent MC experiments by calling the monte\_carlo\_exp function, which is only exit when both conditions (i.e., minimum number of MC experiments and standard error threshold) are achieved. Finally, in lines 34 and 35, relevant results such as the required number of MC experiments (or iterations) and computed (average) percentage porosity are printed in screen.

```
1. # Input Parameters
2. n_grid = 50 \# LHS Grid size per dimension<br>3. c int = 5.0*10**(-4) \# Confidence Interval Size
                                # Confidence Interval Size
4. min iter = 4 \# Minimum MC iterations
5. # Variables
6. inppath = sys.path[0]+"/" # Work Path
7. c iter = 1    # Iterations Counter
8. usum = 0 <br># Sum of computed porosities
9. sum2 = \theta \neq Sum of the square of computed porosities<br>10. stderr = 1 \neq Std Error of computed porosities
10. stderr = 1 # Std Error of computed porosities
                                # Latin Hypercube Sampling Grid Size
12. ni = n_grid**3 # Monte Carlo Sampling Size (1x each stratum)
13. try:
14. file = sys.argv[1]
15. print("File: "+file)
16. # Retrieve File Information
17. H,D,A,Xvec,Yvec,Zvec,Rvec,n_sph = retrieve_file_info(inppath + file)<br>18. # Definition of Strata Sizes
        # Definition of Strata Sizes
19. if A != 0: [dx, dy, dz] = [A/n_grid]*3
               # Strata sizes for CUBE
20. if D !=0 and H != 0: [dr, dt, dz] = [D/(2*math.sqrt(n_grid)), (2*np.pi)/n_grid, H
/n_grid] # Strata sizes for CYLINDER
       # Linked Lists Building
22. print('Building Neighbour Lists')
23. head, list, vecDelta, vecMin = build_lists(Xvec, Yvec, Zvec, Rvec)<br>24. # MC Independent Simulations
        # MC Independent Simulations
25. print("Computing Volume Using "+str(ni)+" MC Points.")
26. while stderr >= c_int/3 or c_iter <= min_iter:
27. inst_porosity = monte_carlo_exp(H,D,A,Xvec,Yvec,Zvec,Rvec,ni)
28. [usum, sum2] = [usum+inst_porosity, sum2+inst_porosity**2]
29. [mean_val, std_dev] = [usum/c_iter, math.sqrt((sum2-
    (usum**2/c_iter))/c_iter)] # Mean Value and Std Deviation of Porosity
30. stderr = std_dev/math.sqrt(c_iter)
               # Standard Error for current iteration
31. c_iter += 1<br>32. print("Porosity
        32. print("Porosity Computation: DONE")
33. # Prints results in screen
34. print('Iterations needed: '+str(c_iter-1))<br>35. print("Computed percentage porosity: "+str
        35. print("Computed percentage porosity: "+str(round(mean_val*100,2))+"%")
36. except:
37. print("Error: Unexpected Exit")
38. sys.exit(2)
```
### **3 Results**

Algorithm implementations proposed in previous works are focus in using the MC method for the computation of volume in *n*-dimensional convex polytopes [[35,](#page-22-4) [36](#page-22-5)]. Algorithms such as the one presented by Liu et al. [\[36](#page-22-5)], which is based on a Markov chain method, or the one presented by Emeris and Fisikopoulos [[37\]](#page-22-0), based on a 'hit-and-run' random walk method are simple and efficient but limited to convex geometries. On the other hand, regarding non-convex sets, algorithms such the one presented by Cazals et al. [[41\]](#page-22-6) and Till et al. [[14\]](#page-21-11) give solutions, based on decomposition and the uniformly random sample MC simulations, respectively, to a union of spheres.

In this work, a membership oracle approach has been followed to implement a simple yet efficient algorithm to estimate the actual percentage porosity of three-dimensional non-convex geometries, represented by the intersection of the exterior of a union of spheres and the interior of a bounding volume. This algorithm has been particularized for the case of porous three-dimensional cubic-based and cylinder-based geometries. Percentage porosity, for diferent instances containing a wide range of number of pores, has been computed using a confdence interval stopping procedure and results have been compared with the reference value obtained by importing the corresponding geometries into ANSYS v18 Design Modeler module to quantify its accuracy. Both computed percentage porosity and runtime have been compared with available program McVol, an MCbased software for protein volume computation developed by Till et al. [[14\]](#page-21-11), as a benchmark.

A data set with a total of 40 geometries was used. The data set is composed by 20 cube-based and 20 cylinder-based geometries, all generated using DEM software LIGGGHTS. A total of eight subsets generated by running DEM simulation using 100, 500, 1000, 2000, and 4000 elements, contained in two basic geometry types (i.e., cube and cylinder), were used. For each subset, 200 unique distributions of pores per number of elements used per basic geometry type were generated, from which 200 distinct geometries were then



<span id="page-11-0"></span>**Fig. 5** Average runtime to compute the percentage porosity for geometries containing diferent number of pores using DMCS (this work) and McVol [\[14\]](#page-21-11) after 20 independent runs for **a** cube-based geometries and **b** cylinder-based geometries



<span id="page-11-1"></span>**Fig. 6** Average MC iterations needed to achieve convergence of 0.5% uncertainty in standard error versus number of pores of cubebased size 6.75 mm and cylinder-based diameter 6.75 mm and height 5.4 mm foam geometries

created. Later, four geometries were randomly selected for each subset, to obtain a workable size data set. Each geometry was individually identifed based on the basic geometry type, the total number of spherical pores, and a correlative (e.g. CUBE\_100\_1, CYL\_2000\_3). Results using the proposed method were obtained using an LHS grid of 50 strata per dimension, representing a sample size of 125 thousand random points per iteration and an uncertainty of 0.5% was established as convergence criteria, under the three-sigma rule.

For each geometry in the data set, a total of 20 independent analysis were run using both the proposed method and McVol and loaded into ANSYS v18 Design Modeler module to retrieve the reference value of percentage porosity. To use McVol, all 40 geometries of the data set were transformed to the corresponding input fle *.pqr* and scaled, so the maximum pore radius did not exceed 5 [Å], as required by the Till

<span id="page-11-2"></span>**Table 1** Average runtime of analysis of 4 diferent geometries after 20 independent runs using the proposed method and McVol for cubebased size 6.75 mm and cylinder-based diameter 6.75 mm and height 5.4 mm geometries containing pores of 350–450 µm in diameter

<b>Subset</b>	This work		McVol	
	Avg $[s]$	Std Dev [s]	Avg $[s]$	Std Dev [s]
CUBE 100	9.1	0.1	12.3	0.1
<b>CUBE 500</b>	10.7	0.1	42.7	0.1
<b>CUBE 1000</b>	17.3	0.1	76.1	0.1
<b>CUBE 2000</b>	31.3	0.1	121.9	0.1
<b>CUBE 4000</b>	54.5	0.6	119.2	12.7
<b>CYL 100</b>	12.8	3.7	10.7	5.2
<b>CYL 500</b>	21.2	2.3	41.1	5.2
CYL 1000	37.8	2.7	65.4	5.1
<b>CYL 2000</b>	69.0	4.4	78.4	10.2
<b>CYL 4000</b>	102.3	8.9	70.5	10.3

et al. All other setup parameters for McVol were reused from the example case provided in the documentation. For each analysis, both percentage porosity and runtime ware registered and results between the proposed method and McVol were compared. Runtimes were measured using the standard C function *time()*.

To establish the convergence character of the proposed implementation, the instantaneous percentage porosity and the standard error were registered for each iteration of the MC experiment. In Fig. [4,](#page-9-0) results from the analysis of the cube-based geometry containing 4000 pores, CUBE  $4000$  1, are shown, where samples of  $125,000$ random points per iteration were used. As it can be seen, as the number of random points increases, the corresponding mean percentage porosity stabilizes, while the standard error decreases asymptotically. This behavior is expected and predicted in Eq. [18](#page-5-5), as the standard error normalized the standard deviation of the measured values by the square root of the number of measurements. Also, as the standard error decreases, the stabilizing behavior of the mean percentage porosity tends to the expected value of the corresponding distribution, and the uncertainty of the confidence interval decreases for a constant confdence level (see Eq. [17](#page-5-3)).

Regarding runtime requirements, the currently available method to compute the percentage porosity in this type of geometries is to import the geometry into the FEA preprocessor. This process is expensive, especially when geometries contain a large number of pores, as it can be seen in Fig. [2.](#page-1-1) As an alternative, the DMCS poses a much faster way to estimate the percentage porosity. In Fig. [5](#page-11-0), the average runtime for the proposed implementation, for each subset, is provided and compared to that obtained by analyzing the same geometries using McVol. The average runtime was measured for all four geometries in each subset after 20 independent runs.

For the average case, based on the analyzed subsets, the runtime of the proposed algorithm scales linear with respect to the number of contained pores ( $R^2$  = 0.995 and  $R^2$ =0.978). In the case of cube-based geometries, Fig. [5a](#page-11-0) shows that for all the analyzed range, the proposed algorithm is consistently faster than McVol, although, for cylinder-based geometries, in Fig. [5b](#page-11-0), results for the proposed implementation show to be faster for geometries containing 2000 pores or less. As suspected, when comparing the average runtime of the proposed implementation for both cubebased and cylinder-based geometries, cube-based geometries exhibit faster average runtimes, of nearly half, than its cylindrical counterparts. Either way, for the analyzed range, the proposed method showed an average convergence time of order  $O(n)$ , where *n* is the number of contained pores.

The diference in runtime between cube-based and cylinder-based geometries is believed to be based on two main factors: (a) the fact that cylinder-based geometries require on average more iterations to achieve convergence and (b) each iteration is slower than for cube-based geometries as the equiprobable space is more complex and a transformation from cylindrical to cartesian coordinates must be done after each random point is generated, to be evaluated by the membership oracle. Regarding required number of iterations to achieve convergence, Fig. [6](#page-11-1) shows the average number of iterations needed to achieve the established convergence criterion as a function of the number of pores. The average has been measured as the average of the four geometries of each subset. Examination of Fig. [6](#page-11-1) shows that for any given number of pores, the number of iterations needed to achieve convergence is consistently higher for cylinder-based geometries than for the cube-based ones. Measurement of the cumulative time required to execute the corresponding lines to generate the 125 thousand random points per iteration have

shown a consistent average time of 2.4 µs per point for the cube-based geometries versus 7.84 µs per point for cylinderbased geometries, which, in addition to the larger number of iteration needed, are supporting evidence of both assumptions. More details regarding average and standard deviation in runtime for the proposed method and McVol can be seen in Table [1](#page-11-2). Alternatively, this diference in behavior may be infuenced by a dependence of the runtime with respect to the percentage porosity, although this relation has not been addressed at this point.

The main objective of the propose method is to estimate the percentage porosity of non-convex sets, and the tested implementation is oriented toward foam geometries, represented by the intersection between the exterior of a union of spheres and the interior of a surrounding primitive geometry such as a cube or a cylinder. The computed percentage porosity obtained by the proposed method, and by McVol, was compared by measuring the absolute error respect to the reference value, which is obtained by loading the corresponding geometries into ANSYS v18 Design Modeler module and retrieving the solid volume from there. Figure [7](#page-13-0) shows the results for the cube-based subsets, while Fig. [8](#page-14-0) shows the results obtained for the cylinder-based subsets.

More detailed information regarding the minimum, maximum, and average computed percentage porosity, as well as the absolute error, for each of the geometries contained in the data set is provided in Table [2](#page-19-0) of ["Appendix B"](#page-18-0). Also, in ["Appendix C"](#page-20-0), Table [3](#page-20-1) provides a more detailed information regarding the minimum, maximum, and average runtime for each of the tested geometries using both the proposed implementation and McVol.

Examination of Figs. [7](#page-13-0) and [8](#page-14-0) shows that although the proposed method tends to produce less precise results; in general, these results are more accurate than those produced by McVol. This difference is based on a key aspect that differentiates the proposed method form it, which is the introduction of the exterior bound. While McVol defnes a bounding box, which encloses the complete union of spheres, in our method, the domain is bounded by the exterior primitive volume and allows the spheres to intersect this boundary. Further examination of Fig. [7](#page-13-0) shows that the computed results using the proposed method are bounded by 0.35%, which is lower than the 0.5% percent limit established a priori for the uncertainty. On the other hand, detailed examination of Fig. [8](#page-14-0) shows that the absolute error distribution for cylinderbased subsets was less predictable than for cube-based ones. Figure [8d](#page-14-0) shows the widest distribution of error obtained in computing the percentage porosity of cylinder containing 2000 pores, as high as 1%, while Fig. [8a](#page-14-0)–c, e shows an excellent prediction capability for all the other cases, less than 0.2% in all cases which is, again, less than the prescribed 0.5% uncertainty.



<span id="page-13-0"></span>**Fig. 7** Comparison of absolute error in percentage porosity computation of cube-based geometries of size 6.5 mm by the proposed method and McVol [\[14\]](#page-21-11) for geometries containing **a** 100, **b** 500, **c** 1000, **d** 2000, and **e** 4000 distributed pores



<span id="page-14-0"></span>**Fig. 8** Comparison of absolute error in percentage porosity computation of cylinder-based geometries of diameter 6.75 mm and height 5.4 mm by the proposed method and McVol [[14](#page-21-11)] for geometries containing **a** 100, **b** 500, **c** 1000, **d** 2000, and **e** 4000 distributed pores

Measurements of the diference between the computed percentage porosity and its reference value for geometries containing a larger number of pores (i.e., greater than 4000 pores) could not be considered for this work due to the excessive time required to import each geometry into the FEA preprocessor, though similar results are expected. On the other hand, the proposed implementation considers just the use of a single-thread algorithm and, due to the potential for parallelization of the independent MC experiments, further optimizations can be done to achieve better time performance. It is known by the authors that Python libraries such as Joblib and Numba provide fast and easy alternatives for that matter.

### **4 Conclusions**

A single-thread algorithm for DMCS, partially based on the one presented by Liu et al. [[36\]](#page-22-5), has been implemented in the Python language to estimate percentage porosity in cylindrical and cubical geometries containing interconnected spherical voids analytically defned with spatial distribution obtained by DEM simulation and uniformly random radii, as those required by Perez et al. [\[10\]](#page-21-8) in the study metallic foams fabricated by means of PM and the SHP technique. The complete Python script was developed using less than 120 code lines, when comment lines are not counted, and only the Python libraries numpy and numba, along with the three build-in modules math, random, and sys have been used.

The proposed implementation showed a signifcant gain in performance time for the task with respect to the currently used technique, which requires the execution of the complete sequence of CAD extrusion and cut operations by the FEA preprocessor. The time consumption to compute the percentage porosity showed a reduction between 84 and 99% when geometries containing between 100 and 4000 spherical pores were analyzed.

When compared to other similar software, the proposed implementation has shown to be able to achieve consistently smaller errors in approximating the percentage porosity of foam geometries than McVol. These better results are believed to be related to a better suited defnition of the domain restrictions of the set. While McVol relies on

defning a bounding box that contains the whole union of spheres, the proposed implementation gives a more precise restriction of the outer boundary of the domain, whether it is a box (i.e., a cube) or a cylinder. Although, when cylinders are considered for analysis, a trade-off must be done, and sometimes, performances are loss in exchange for precision. In addition, the proposed implementation relies on a statistical criterion to stop computations, rather than a unique measure or an arbitrary samples size.

The DMCS has been showed to provide a simple yet powerful tool in estimating the porosity percentage in 3D non-convex analytical geometries, as the accuracy in the computed results provides an estimation error below the prescribed uncertainty, with respect to the percentage porosity obtained by the generation of the geometries by the FEA preprocessor, in seven of the eight subsets of the tested data set. This estimation error represents a neglectable diference for the primary purpose for which the algorithm has been developed.

In addition to this implementation, future work related to this algorithm includes further performance improvements by means of parallel computing and its extension to more general geometries such as polyhedrons and other volumes.

### <span id="page-15-0"></span>**Appendix A**

In this appendix, the complete Python code script is presented. This code runs in single core configuration allowing to estimate the porosity of cylindrical- or cubic-shaped foam geometries containing randomly distributed spherical pores when a complete analytical description of it (i.e. overall dimensions and pores location and dimension) is given in a JavaScript script fle. The code uses Eq. [15](#page-5-4) to average the results obtained by a series of Monte Carlo simulations, based on an LHS strategy, according Eqs. [5](#page-4-5), [6,](#page-4-2) [7,](#page-4-4) and [8.](#page-4-3) The code will iterate until two established criteria are met, which are a minimum number of iteration and a maximum standard error, according to Eq. [18](#page-5-5). This code requires the user to provide the flename with its extension as system argument (e.g., Cube\_100\_1.js). The path to the fle is assumed to be the current work directory



```
35. 1_H, 1_D = 0, 0
36. else: # Acts if doe
   s not detect neither a cube nor a cylinder
37. l_A, l_D, l_H = 0, 0, 0<br>38. print('Error retrieving38. print('Error retrieving file info. Not a Cylinder nor a Cube was found.')
39. sys.exit(2)
40. return l_H,l_D,l_A,np.asarray(x),np.asarray(y),np.asarray(z),np.asarray(radii),n_
  void
41. #
42. def monte_carlo_exp(H, D, A, x_vec, y_vec, z_vec, r_vec, seeds):
43. # Allocates list for sampling results
44. sampling_list = np.zeros([nx, ny, nz])<br>45. for k in range(nz):
        45. for k in range(nz):
46. for j in range(ny):
47. for i in range(nx):
                    # Takes a smple from the [i,j,k] stratum
49. if A!=0: # CUBE
50. dart = [random.uniform(dx*(i),dx*(i+1)), random.uniform(dy*(j),dy*(j+1)), random.uniform(dz*(k),dz*(k+1))]51. if (D != 0 and H != 0): # CYLINDER<br>52. tmp = [random.uniform(dr*math.sqrt
                        time = [random.uniform(dr*math.set(i),dr*math.set(i+1)), random.
  uniform(dt*(j),dt*(j+1)), random.uniform(dz*(k),dz*(k+1))]53. dart = [tmp[0]*np.cos(tmp[1]), tmp[0]*np.sin(tmp[1]), tmp[2]]
54. # Evaluates the sampling point<br>55. Sampling list[i,j,k] = member
                    sampling_list[i,j,k] = member_oracle(np.asarray(dart), head, list, np
   .asarray([x_vec, y_vec, z_vec]), r_vec, vecDelta, vecMin)
56. # # Computes expected value E[x]<br>57. hits num = sum(sum(sum(sampling
57. hits_num = sum(sum(sum(sampling_list)))<br>58. # Returns the computed probability p(x,
        # Returns the computed probability p(x,n) = E[x]/n59. return float(hits_num/seeds)
60. #
61. @numba.jit
62. def member_oracle(vecP, head, list, vecPores, vecRadii, vecDelta, vecMin):
63. # Initiates variable assuming no hit
64. Cell = np.zeros(3)<br>65. # Defines Central
        # Defines Central Cell
66. Cell = [math.floor((vecP[i]-vecMin[i])/vecDelta[i]) for i in range(3)]
        67. # Evaluates around the central cell
68. for cell_x in [int(Cell[0]), int(Cell[0]-1), int(Cell[0]+1)]:
69. for cell_y in [int(Cell[1]), int(Cell[1]-1), int(Cell[1]+1)]:
70. for cell_z in [int(Cell[2]), int(Cell[2]-1), int(Cell[2]+1)]:
71. test = int(head[cell_x, cell_y, cell_z])
72. while test != 0:
73. #Evaluate Distance to the Pore
74. if (vecP[0]-vecPores[0,test-1])**2 + (vecP[1]-vecPores[1,test-
  1])**2 + (vecP[2]-vecPores[2,test-1])**2 <= vecRadii[test-1]**2:
75. return 1
76. #Update Test Pore
77. test = int(list[test-1])<br>78. teturn 0
        return 0
79. #
80. def build_lists(x_vec, y_vec, z_vec, r_vec):
81. # Temp Lists<br>82. x ext = [np.
        x<sub>ext</sub> = [np . amin(x<sub>vec</sub> - r<sub>vec</sub>), np . amax(x<sub>vec</sub> + r<sub>vec</sub>)]
83.  y_{ext} = [np_{\text{.}} \text{min}(y_{\text{.}} \text{vec} - r_{\text{.}} \text{vec}), np_{\text{.}} \text{max}(y_{\text{.}} \text{vec} + r_{\text{.}} \text{vec})]84. z_ext = [np.amin(y_vec - r_vec), np.amax(z_vec + r_vec)]<br>85.        r max = max(r vec)
        r_max = max(r_vec)
86. \# Cells definition<br>87. C = \{math, \text{math.ceil}(x \in \mathbb{R}\})C = [math, c = [math.cei]((x_ext[1] - x_ext[0])/(2*rmax)), math.ceil((y_ext[1] - y_ext[0]))(2*r_{max}), math.ceil((z_ext[1] - z_ext[0])/(2*r_max))]
```

```
88. diff = [(x_ext[1] - x_ext[0])/C[0], (y_ext[1] - y_ext[0])/C[1], (z_ext[1] - z_ext
   [0])/C[2]]
89. # Arrays Allocation
90. list = np{\text{.}zeros(len(r\_vec))}<br>91. head = np{\text{.}zeros(lc[0].c[1].}head = np{\text .}zeros((C[0], C[1], C[2]))92. # Lists Filling
93. for i in range(len(r_vec)):
94. CELL = [math,float] (x \text{ vec}[i] - x \text{ ext}[0])/diff[0]), math.floor((y vec[i]-
    y_ext[0])/diff[1]), math.floor((z_vec[i]-z_ext[0])/diff[2])]
95. list[i] = int(head[CELL[0],CELL[1],CELL[2]])
96. head[CELL[0],CELL[1],CELL[2]] = int(i+1)97. return head, list, diff, [x_ext[0], y_ext[0], z_ext[0]]
98. #
99. ####################
100. # MAIN PROGRAM
101. ######################
102. # Input Parameters
103. n_grid = 50    # LHS Grid size per dimension<br>104. c int = 5.0*10**(-4)    # Confidence Interval Size
                                           # Confidence Interval Size
105. min_iter = 5 # Minimum MC iterations
            # Variables
107. inppath = sys.path[0]+"/" # Work Path<br>108. citer = 1 # Iteration
                                           # Iterations Counter
109. usum = 0    # Sum of computed porosities
110.  sum2 = 0   # Sum of the square of computed porosities<br>111.  stderr = 1   # Std Error of computed porosities
                                           # Std Error of computed porosities
112. [nx, ny, nz] = [n_grid]*3 # Latin Hypercube Sampling Grid Size
113. ni=n_grid**3 # Monte Carlo Sampling Size (1x each stratum)
114. try:
115. file = sys.argv[1]<br>116. print("File: "+fil
116. print("File: "+file)
117. # Retrieve File Information<br>118. H.D.A.Xvec.Yvec.Zvec.Rvec.n
                118. H,D,A,Xvec,Yvec,Zvec,Rvec,n_sph = retrieve_file_info(inppath + file)
119. # Definition of Strata Sizes<br>120. if A != 0: [dx, dy, dz] = [A
                if A != 0: [dx, dy, dz] = [A/n\_grid]*3# Strata sizes for CUBE
121. if D !=0 and H != 0: [dr, dt, dz] = [D/(2*math.sqrt(n_grid)), (2*np.pi)/n
_grid, H/n_grid] # Strata sizes for CYLINDER
122. # Linked Lists Building<br>123. print('Building Neighboy
                 123. print('Building Neighbour Lists')
124. head, list, vecDelta, vecMin = build_lists(Xvec, Yvec, Zvec, Rvec)<br>125.             # MC Independent Simulations
                 # MC Independent Simulations
126. print("Computing Volume Using "+str(ni)+" MC Points.")<br>127. while stderr >= c int/3 or c iter <= min iter:
                while stderr >= c int/3 or c iter <= min iter:
128. inst_porosity = monte_carlo_exp(H,D,A,Xvec,Yvec,Zvec,Rvec,ni)
129.                    [usum, sum2] = [usum+inst_porosity, sum2+inst_porosity**2]<br>130.                  [mean_val, std_dev] = [usum/c_iter, math.sqrt((sum2-
                     [mean_val, std\_dev] = [usum/c\_iter, math.sqrt( (sum2-(usum**2/c_iter))/c_iter)] \qquad # Mean Value and Std Deviation of Porosity<br>131 stdern = std dev/math sort(c iter)
                     stderr = std_dev/math.sqrt(c_iter)
                         # Standard Error for current iteration
132. c_iter += 1
133. print("Porosity Computation: DONE")
134. # Prints results in screen<br>135. print ('Iterations needed:
                 print('Iterations needed: '+str(c_iter-1))
136. print("Computed percentage porosity: "+str(round(mean_val*100,2))+"%")
            except:
138. print("Error: Unexpected Exit")
139. sys.exit(2)
```
### <span id="page-18-0"></span>**Appendix B**

In this appendix, more detailed information regarding the computed percentage porosity for the data set obtained by means of the proposed algorithm and McVol after 20 independent runs are presented in Table [2](#page-19-0).

<span id="page-19-0"></span>



## <span id="page-20-0"></span>**Appendix C**

algorithm and McVol after 20 independent runs are presented in Table [3.](#page-20-1)

In this appendix, more detailed information regarding the computed runtimes for the data set for the proposed

<span id="page-20-1"></span>**Table 3** Minimum, maximum, and average runtime after 20 independent runs using the proposed implementation and McVol



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