Morse Theory-based Segmentation and Fabric Quantification of Granular Materials

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Abstract This article presents a robust Morse theory-1 based framework for segmenting 3D x-ray computed 2 tomography image (CT) and computing the fabric, rel-3 ative arrangement of particles, of granular ensembles. 4 The framework includes an algorithm for computing the 5 segmentation, a data structure for storing the segmenta-6 tion and representing both individual particles and the 7 connectivity network, and visualizations of topological 8 descriptors of the CT image that enable interactive ex-9 ploration. The Morse theory-based framework produces 10 superior quality segmentation of a granular ensemble as 11 compared to prior approaches based on the watershed 12 transform. The accuracy of the connectivity network 13 also improves. Further, the framework supports the effi-14 cient computation of various distribution statistics on 15 the segmentation and the connectivity network. Such 16 a comprehensive characterization and quantification of 17 the fabric of granular ensembles is the first step towards 18 a multiple length scale understanding of the behavior. 19

- 20 Keywords Granular ensemble · fabric · segmentation ·
- ²¹ particle connectivity network · Morse theory ·
- ²² Morse-Smale complex · topological persistence.

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1 Introduction

Understanding of the mechanical behavior of geomateri-24 als from the micro-scale (*i.e.*, inter-particle interactions) 25 to the macro-scale (or, continuum scale) has significantly 26 benefited engineering of physical infrastructure. Such 27 an understanding straddling multiple length scales has 28 established beyond reasonable doubt that the ensemble 29 mechanical response is an integration of the arrange-30 ment and interaction of individual particles in geomate-31 rials such as sand, clay, structured sand, and structured 32 clay [46, 34, 7]. Numerical techniques such as discrete ele-33 ment methods and imaging experiments using x-ray com-34 puted tomography (CT) have enormously contributed 35 to this understanding of geomaterial behavior. Prior to 36 use of x-ray CT, fabric and microstructure studies were 37 limited to only two-dimensional packing [33, 34]. Access 38 to CT has enabled extraction and quantification of the 39 three-dimensional fabric and its evolution under different 40 stress paths [11, 49, 20, 21, 3, 16, 1]. Within the overall 41 workflow of quantification of fabric of soils, the method 42 used for image segmentation crucially determines the 43 quality of the results. In this paper, we present a novel 44 approach for segmentation based on Morse theory, which 45 is a branch of topology that studies the relationship be-46 tween the connectivity of spaces and scalar functions 47 defined on them. Crucial to the goal of quantification 48 of fabric, this Morse theory-based approach simultane-49 ously supports a data structure that efficiently stores 50 both the segmentation and the connectivity network 51 that represents the inter-particle arrangement. 52 53 1.1 Previous work

Fabric Quantification. Initial quantification of the 54 fabric of sands using CT techniques has been reported 55 by multiple authors [2, 3, 16, 24, 26]. More recently, the 56 fabric of weakly cohered glass-ballotini has also been in-57 vestigated [44, 43]. The basic premise of this CT-based 58 analysis of fabric is the collection of images through 59 the cross-section of a laboratory or natural sample, and 60 concomitant image segmentation and analysis. Segmen-61 tation refers to the process of labeling or identification 62 of different phases or objects present in the image. In 63 this study, images of weakly cemented granular materi-64 als captured through x-ray CT are analyzed. The small 65 amount of epoxy present mediates the contact between 66 particles. Air voids present in this ensemble, are also 67 segmented. 68

Once the particles and contacts are identified, the inter-particle arrangement is studied by computing the coordination number distribution, or more commonly by computing a fabric tensor. The *fabric tensor* is a measure of the orientation distribution of the particles [32]. Characteristics of the fabric tensor and its formulation have been enumerated in recent work [43].

Watershed-based segmentation. In previous work, 76 including the above-mentioned studies, variants of the 77 watershed algorithm [30, 37] have been typically used 78 to segment the particles and to identify the contacts 79 (or epoxy bonds). These watershed-based algorithms re-80 quire as input a scalar field that contains, in its topology, 81 exactly one catchment for each particle *i.e.*, the presence 82 of exactly one field minimum / maximum corresponding 83 to each particle. Further, [48] show the contact orienta-84 tion obtained through conventional and random walker 85 based watershed implementation is either error-prone 86 or computationally expensive. 87

The watershed algorithm identifies and labels each 88 catchment (particle) and its ridge line (epoxy bonds). 89 The segmentation method begins by first binarizing the 90 volume image corresponding to a particle-void gray level 91 cutoff. Following this, the binarized particles are tested 92 to ensure that they do not contain inherent flaws such 93 as regions of low attenuation intensity. Next, a distance 94 transform is applied on the complementary image to 95 obtain a scalar field such that each particle contains 96 exactly one maximum. Due to the irregular shape of the 97 particles, the distance transform generates one dominant 98 peak and several small peaks. These small peaks are 99 removed by applying the h-maxima transform, which 100 suppresses all the maxima (peaks) that are lower than 101 the specified tolerance [45]. An appropriate threshold for 102 103 the h-maxima transform is chosen manually by an expert user. This transform provides exactly one maximum for 104

each particle. Next, the field is inverted such that each 105 particle contains exactly one minimum. The labeled par-106 ticles are computed by applying the watershed algorithm 107 to the inverted modified distance field. This algorithm 108 progressively fills the basins defined by each minimum 109 from below by uniformly raising the "water level". Wa-110 ter from two adjacent basins meet along their common 111 boundary, the separating ridge, that passes through the 112 contact region. Figure 1 illustrates the different steps in 113 the watershed-based approach and contrasts it with our 114 proposed Morse theory-based approach thereby high-115 lighting the benefits of the latter. 116

An important limitation of the watershed technique 117 is that it fails for non-convex shaped particles, partic-118 ularly those that contain a neck, and results in over-119 segmentation. A carefully chosen threshold for the h-120 maxima transform addresses this problem to some ex-121 tent. However, a single threshold uniformly applied to 122 the entire scalar field fails to capture all possible cases 123 and the method often results in over-segmentation. Also, 124 the quality of the result naturally depends on the manual 125 choice of threshold for the h-maxima. Other extensions 126 and variants have been proposed [50, 26] to handle the 127 over-segmentation problem. These methods either rely 128 on a single absolute user-specified threshold or adaptive 129 local application of watershed, which is computationally 130 inefficient and expensive. 131

Topology-based segmentation. Topological frame-132 works like discrete Morse theory [17] and persistent 133 homology [13] enable effective tools for the study and 134 analysis of scientific data due to their ability to robustly 135 extract shape and structure even in the presence of 136 noise [19, 18, 6]. The physical properties of porous and 137 granular materials are closely related to the geometry 138 and topology of the material structure. So, topological 139 methods have been found to be useful in the study of 140 diverse phenomena within this domain, ranging from 141 the crystallization process in bead packings [38] to per-142 meability in sandstone [22]. Persistent homology-based 143 pipelines have been used for summarizing essential struc-144 tural properties [38, 25] and Morse theory has been used 145 to extract robust geometric representations of the ma-146 terial structure [19, 36, 10]. Robins et al. [36] and Frei-147 drichs et al. [10] first described the benefits of discrete 148 Morse theory for the segmentation and skeletonization of 149 3D images, including an application to the computation 150 of the pore network of a silica sphere pack from 3D CT 151 scan data. The focus of their work is on computing the 152 Morse-Smale (MS) complex and hence the other steps 153 of the CT scan analysis pipeline such as the boundary 154 surface computation and geometry extraction are not 155 discussed. Further, the use of a manually selected sim-156 plification threshold results in the lack of an intuitive 157



Fig. 1: Visual comparison of the watershed-based and the proposed Morse theory-based approaches to process and analyze 3D CT data. The comparison helps identify key differences between the two method pipelines in terms of computation, output, and supported analysis. The unified data structure retrieved from the MS complex supports various analysis tasks.

choice or guide for selecting noise simplification thresh-158 olds. Gyulassy et al. [19] also adapted Morse theory in 159 their pipeline to extract the core filament structure of 160 porous materials. Their curved skeleton-based represen-161 tation is computed by carefully simplifying a distance 162 field in order to identify and extract features of inter-163 est from the MS complex. They are able to identify 164 consistent and stable core structures of porous solids, 165 thus extracting meaningful results about the material 166 properties. Their method design exemplifies the benefits 167 of adapting Morse theoretic ideas to fit the context of 168 an application focused pipeline. 169

Our method falls into the category of Morse theory-170 based approaches that aim to study material packing 171 structures, but it is distinct from previous work on three 172 essential points. First, our proposed method packages 173 the benefits of discrete Morse theory as a part of a 174 comprehensive and coherent pipeline for analysis of CT 175 images with the objective of computing a robust geo-176 metric representation of the granular material structure. 177 Second, the pipeline is designed to support a smooth and 178 convenient end-user experience with intuitively tunable 179 parameters that generalizes well to diverse use-cases. 180 Third, our focus is on the study of the material phase of 181

the packing structure of granular materials as opposed to the study of pore spaces in earlier work.

1.2 Contributions

This paper presents a new Morse theory-based frame-185 work for computing, storing, and exploring the fabric 186 in geomaterials. We apply this framework on x-ray CT 187 images of a mono-dispersed steel sphere packing for 188 benchmarking and on an epoxy bonded sand packing 189 to demonstrate its utility and advantages over existing 190 approaches. The framework incorporates a new method 191 for segmentation, introduces a new connectivity network 192 that supports further statistical analysis, and a visual-193 ization tool that supports interactive exploration of the 194 granular ensemble, see Figure 1. Key contributions of 195 this paper include 196

- A robust algorithm for computing a segmentation
 of a 3D x-ray CT image. The segmentation is represented as the Morse complex of the scalar field
 corresponding to the CT image. Further, the algorithm is amenable to parallel computation.
- An effective method for simplification and noise removal. A topological persistence [15, 13] based 203

method removes both geometric and topological noise resulting in the desired segmentation. The method enables both a flexibility in the choice of integrated geometric and topological criterion for simplification and an intuitive choice of threshold for the criterion.

An improved boundary surface computation method
that constructs the necessary input for the segmentation algorithm. An active-contour [8] based method
computes an optimal boundary surface of the particles in the packing. The surface is locally adapted
to the best separating scalar value and is thus free
of the assumption of a single global threshold.

A unified data structure for storing both the segmentation and contacts. The data structure representing
the Morse complex supports fast queries. Further,
it enables efficient noise removal while guaranteeing
consistency in the topological relationship between
the segments.

Construction of a geometrically meaningful connec-223 tivity network between contacting particles in the 224 packing. The arcs of the network adapt to the ge-225 ometry of the particles and are guaranteed to lie 226 within the packing as they connect particle centers 227 and contacts, thus accurately representing the pack-228 ing structure even for complex non-convex particle 229 geometries. 230

 An effective method for visualizing the segmentation and the connectivity network. The segmentation, contacts, and connectivity network are visualized via high quality rendering of the cells of the Morse complex.

Experimental results demonstrate that the segmentation 236 results are superior to watershed-based methods, both 237 in terms of segmentation quality and identification of 238 contacts. The proposed method is suitable for both 239 simple and complex CT granular datasets with non-240 convex morphology, and has the potential to generalize 241 to broad range of particle sizes including those with 242 particle breakage. 243

244 2 Challenges

In this section, we describe the main challenges faced in
the process of analyzing 3D CT images using existing
techniques. These challenges serve as the underlying
motivation for the design of our approach.

Characteristics of reconstructed CT data. Reconstructed 3D CT images of granular ensembles can be
very challenging to process because of the reconstruction artifacts, noise, and irregular geometry of the particles.
The x-ray CT scans suffer from noise and artifacts, the

incompatibility between the reconstructed values and 254 their attenuation densities, due to the inconsistencies in 255 x-ray source (polychromatic source), object (cylindrical 256 shape of the object), detector, and reconstruction algo-257 rithm [23]. Figure 2 shows the reconstructed volume and 258 a representative slice of the epoxy-bonded sand speci-259 men. The background intensity and the reconstructed 260 values of sand particles vary across the specimen due 261 to artifacts and noise. The reconstructed image is af-262 fected by beam hardening — the average reconstructed 263 values of air, epoxy, and sand are significantly higher 264 near the periphery than the central region. The partial 265 volume effect results in dark streaks near the periphery. 266 We also observe occasional low reconstructed valued 267 regions in sand due to flaws within the particles that 268 manifest into intra-particle voids after thresholding. Fur-269 ther, the epoxy and pore air have similar attenuation 270 coefficients that, together with the background noise, 271 makes the segmentation of the epoxy phase challenging. 272 In addition to reconstruction-based artifacts, the actual 273 geometry of the particles in the packing too can often 274 be highly non-convex, irregular, and show great vari-275 ation in shape, size, and volume. This combination of 276 artifacts and diversity in particle geometry necessitates 277 specifically designed segmentation techniques that are 278 not only inherently robust but make no assumptions 279 about the particle geometry and distribution of noise. 280

Manual selection of boundary surface. A watershed-281 based segmentation method begins by computing a sur-282 face that bounds the particles. Otsu's method is the 283 popular choice of boundary surface extraction method. 284 It often reports a surface that fails to accurately capture 285 the geometry of the packing. One approach to address 286 this shortcoming is to manually explore the histogram 287 of values in the CT scan and fine tune the choice of 288 isosurface that represents the boundary surface. This 289 manual approach, in addition to being tedious, may 290 still fail to correctly capture the packing geometry. For 291 instance, we highlight one such situation in Figure 3, 292 where a volume rendering of the CT scan with a well 293 tuned color and opacity map is used to illustrate the dif-294 ference between the actual particle geometry and what 295 is identified by a carefully selected isosurface. Further, 296 we also observe that isosurfaces corresponding to dif-297 ferent isovalues capture the correct packing geometry 298 within different regions of the CT data. This indicates 299 that a single uniform choice of isosurface is not sufficient 300 to model the boundary surface geometry. An incorrect 301 boundary surface has the potential to grossly affect 302 downstream analysis, making it difficult to compute 303 accurate segmentation and contact information. 304

Simplification and noise removal. The tendency 305 of watershed-based segmentation to over-segment non- 306



Fig. 2: Left: Direct volume rendering of CT scan of a sand sample using a grayscale color map and opacity which increases linearly with the CT intensity values. Note how the particle boundaries are not clear and there is noise in the data. Moreover, the sand particles have varying intensity, and some imaging artifacts appear as horizontal bands of varying intensity values. Right: A cross section of the CT volume highlights similar issues. Additionally, imaging artifacts at the periphery of the sample is apparent.



Fig. 3: Left: Direct volume rendering of the CT data using a fine-tuned color map. Right: Manually selected isosurface guided by a histogram. Notice how the isosurface does not agree well with the CT data, specially in the region enclosed by the red box.

convex geometry necessitates the employment of regionmerging heuristics such as the h-maxima transform. The
h-maxima transform requires the user to select the desired depth value up to which local maxima are to be
suppressed. A good manual choice is essential for achiev-

ing good results. However, this choice of parameter is tedious and unintuitive. It is often difficult to choose a parameter value with a high level of confidence because evaluating the effect of this choice requires the computation of the entire segmentation followed by statistics 316



(c) h-maxima threshold=2.0

(d) h-maxima threshold=3.0

Fig. 4: In the watershed-based approach to segmentation, it is challenging to identify a "correct" h-maxima threshold. Note how the chosen threshold affects the final segmentation. Often, this threshold is chosen using a trial-and-error process. In some cases, the threshold identification is guided by prior knowledge of the sample, say the expected particle size or expected number of particles in the sample.

on the segmented particles. Further, small changes in 317 h-maxima value may result in large variations in the 318 segmentation, which implies that approximate solutions 319 may not be satisfactory. In Figure 4, we illustrate the 320 watershed segmentation of a granular ensemble at dif-321 ferent h-maxima thresholds. Note how relatively similar 322 h-maxima parameter values result in different segmen-323 tation results. 324

Network extraction. The natural step after computing the segmentation and a set of contacts between particles is to construct a connectivity network. To achieve this, the particles associated with each contact are identified by searching within the neighborhood (26 pixel connectivity) of each contact. A simple network, where connectivity of particles is represented by straight lines between particle centres that share a contact, is used to represent the packing. This is an abstract representation and does not represent the geometry of the contact or connectivity.

3 Background

This section reviews the necessary mathematical background for the topological methods described in this paper. We introduce terminology and key concepts on Morse functions, MS complex, and topological simplifica-340



Fig. 5: Morse complex and its cells. (a) A synthetic scalar field shown using pseudo colors. Points identified as *maxima* are shown as red spheres. (b) Morse complex partitions the domain of the scalar field based on *descending manifolds* of maxima. (c) A pair of adjacent maxima are highlighted, their descending manifolds are shown in yellow and green. The 2-saddle at the interface of these two maxima is shown as a blue sphere. The descending manifold of this 2-saddle is the grey surface at the interface between the descending manifolds of the two maxima. The bold black line is the *ascending manifold* of the 2-saddle which also connects the two maxima. (d) The integral lines within the descending manifolds of the two maxima. Notice how these lines converge towards the two maxima and form a separation surface around the 2-saddle. This structure and partition is robustly extracted using the MS complex.

tion, which are required for understanding the definition
of our proposed connectivity network, the segmentation,
and particle extraction methods described in the next
section.

345 3.1 Morse function and Morse complex

Morse theory studies the relationship between the topological properties of a space and the critical points of a smooth real valued function defined on the space. We introduce the necessary terms from Morse theory and refer the reader to books on this topic for further details [28, 13].

Consider a smooth (twice differentiable) scalar function $f : \mathbb{R}^3 \to \mathbb{R}$. A point $p_c \in \mathbb{R}^3$ is called a critical point of f if the gradient of f at p_c is zero,

$$\nabla f_{p_c} = \left(\frac{\partial f}{\partial x}, \frac{\partial f}{\partial y}, \frac{\partial f}{\partial z}\right)_{p_c} = \mathbf{0}.$$
 (1)

The critical point p_c is called non-degenerate if its Hessian (matrix of second partial derivatives at p_c) is nonsingular. The function f is called a *Morse function* if all its critical points are non-degenerate and have distinct function values. Any smooth function f can be infinitesimally perturbed into a Morse function. The number of negative eigenvalues of the Hessian corresponds to the *Morse index* of the critical point. Critical points of a three-dimensional function f can be of four types: minima (index-0), 1-saddle (index-1), 2-saddle (index-2), and maxima (index-3). An *integral line* is a curve in \mathbb{R}^3 whose tangent vector at a point on the curve is parallel to the gradient of f at that point. The integral line passing through a point p is the solution to

$$\frac{\partial}{\partial t}L(t) = \nabla f(L(t)), \forall t \in \mathbb{R}$$
(2)

with initial value L(0) = p. The function f monotoni-352 cally increases or decreases along an integral line. The 353 end points of an integral line corresponding to the limits 354 as t approaches $-\infty$ and ∞ , its origin and destination 355 respectively, are critical points of f. The set of all in-356 tegral lines originating at a critical point p_c together 357 with p_c is called the ascending manifold of p_c . Simi-358 larly, p_c together with the set of all integral lines whose 359 destination is p_c is called the *descending manifold* of 360 p_c . Figure 5 shows a simple 3D scalar field, its critical 361 points, and descending manifolds. 362

The descending manifolds (similarly, ascending man-363 ifolds) of all critical points of f partition the domain 364 of f. This partition is called the *Morse complex* and 365 consists of cells of dimensions 0, 1, 2, and 3. The de-366 scending manifold of a critical point with index i has 367 dimension i. Therefore, the descending manifold of a 368 maximum is a three dimensional volume (3-manifold), 369 the descending manifold of a 2-saddle is a two dimen-370 sional sheet (2-manifold), the descending manifold of a 371 1-saddle is a one dimensional arc (1-manifold) and the 372 descending manifold of a minimum is the minimum itself. 373 Conversely, the ascending manifold of a critical point 374 with index *i* has dimension 3 - i. We propose the use of 375 the Morse complex to represent the individual particles 376 and the connectivity network between the particles. 377

The overlay of the ascending and descending mani-378 folds results in a finer partition of the domain of f called 379 the Morse-Smale complex (MS complex). Each cell of 380 the MS complex is a set of integral lines that share a 381 common origin and destination. The MS complex is a 382 well studied topological structure and there are several 383 efficient algorithms developed for computing hierarchical 384 representations of the complex [40, 42]. More important 385 in the context of this paper, the Morse complex can be 386 extracted directly as a subset of the MS complex. 387

388 3.2 MS complex computation

CT scan data is available as a 3D density image, which 389 can be modeled as a continuous scalar function defined 390 on the vertices of a 3D cube grid and interpolated within 391 each cell. The cube grid represents the domain of the 392 scalar function. Ideas from Morse theory can be trans-393 ferred to piecewise continuous functions to define critical 394 points [15] via local comparison of function values. Fur-395 ther, a simulated perturbation ensures that neighboring 396

vertices do not have equal function values and hence the 397 gradient is zero only at critical points. This perturba-398 tion helps classify the critical points. Earlier algorithms 399 employed numerical approaches to compute the MS 400 complex and were affected by the errors in the computa-401 tion of interpolants and derivatives. In contrast, we use 402 combinatorial methods for MS complex computation. 403 We use a parallel algorithm based on *discrete Morse* 404 theory, which introduces a discrete analog of gradients 405 as a directed edge between grid cells and computes inte-406 gral lines as paths in a directed graph [40]. By avoiding 407 derivative computations, this method ensures robust-408 ness of the results. In addition, the method provides 409 theoretical guarantees for the correctness of adjacency 410 relationship between cells of the output complex. 411

The 3D cube grid represents the domain as a col-412 lection of cells, a *cubical complex*. Voxels of the input 413 are vertices (0-cells) of the grid; edges, faces, and cubes 414 are the 1-, 2-, and 3-cells. The parallel algorithm of 415 Shivashankar and Natarajan [40] computes discrete rep-416 resentations of the ascending and descending manifolds 417 of the critical points of the function and supports oper-418 ations for topological simplification as described in the 419 following section. The descending manifold of a max-420 imum is a collection of connected 3-cells in the grid, 421 a volumetric region. The descending manifold of a 2-422 saddle is a collection of connected 2-cells resulting in 423 a surface. The descending manifold of a 1-saddle is an 424 arc consisting of a sequence of edges (1-cells) and the 425 descending manifold of a minimum is a solitary 0-cell, 426 namely the minimum. The ascending manifolds are com-427 puted similarly and represented as a collection of 3-cells 428 (minimum), 2-cells (1-saddle), 1-cells (2-saddle), and 429 0-cell (maximum). 430

3.3 Topological simplification

A pair of critical points of f, that is connected via a 432 single arc in the MS complex, may be removed via a 433 critical point pair cancellation, resulting in a simpler 434 MS complex [15, 14]. Morse theory guarantees the ex-435 istence of a simpler function f' corresponding to the 436 simpler MS complex. Pairs of critical points represent 437 topological features. For example, the 2-saddle-maximum 438 pairs in Figure 5c represents the two volumetric regions 439 (yellow and green). We employ the critical point pair 440 cancellation for topological simplification, iteratively re-441 moving small features that result from high frequency 442 noise or sampling artifacts in the CT scan. Two crucial 443 advantages of this approach are that (a) the algorithm 444 is combinatorial and does not suffer from errors due 445 to numerical computation, and (b) it supports con-446

trolled simplification where the data outside of a localneighborhood is not affected.

The cancellation changes both the combinatorial 449 structure of the MS complex as well as the geome-450 try of the ascending and descending manifolds of the 451 surviving critical points. In this paper, we focus on 2-452 saddle-maximum cancellation because we are interested 453 in computing the individual particle segments. Let p_2q_3 454 represent a pair of critical points where the subscript 455 indicates the index of the critical point. First, p_2, q_3 , 456 and all arcs incident on them are deleted. Next, arcs 457 incident on q_3 are routed to the surviving maximum 458 that is connected to p_2 in the MS complex. Finally, the 459 geometry of the descending manifold of q_3 is merged 460 into the descending manifold of the surviving maximum 461 that was connected to p_2 . 462

The order of the cancellations plays an important 463 role in determining the structure and geometry of the 464 resulting simplified MS complex. Intuitively, given the 465 aim of the simplification, the order should depend on 466 the importance or size of a feature represented by a 467 pair of critical points. While this importance can be 468 defined in different ways, a commonly used and effective 469 definition is the difference in function value between the 470 pair of critical points. We iteratively apply critical point 471 pair cancellation ordered by the difference in function 472 value between the pair of adjacent critical points in the 473 MS complex. 474

⁴⁷⁵ 3.4 Persistence diagrams and persistence curves

An important practical consideration is the choice of the 476 amount of simplification to be performed. The simplifi-477 cation threshold is defined as the maximum difference 478 in function value between the canceled critical point 479 pairs. This difference in function value is also referred 480 to as the *persistence* of a critical point pair. Visual rep-481 resentations called persistence diagrams and persistence 482 curves are often used to aid the choice of a threshold 483 that separates noise from features of interest [13, 9]. A 484 *persistence diagram* is a 2D scatter plot of the function 485 values of the canceled critical point pairs. A *persistence* 486 curve is a graph plot of the number of surviving critical 487 points on the y-axis against an increasing simplification 488 threshold on the x-axis. A dense collection of points 489 very close to the diagonal in the persistence diagram 490 is visually representative of a set of noisy features. A 491 simplification threshold to remove such a collection of 492 critical point pairs can be computed by locating the 493 494 knee of the persistence curve, which corresponds to a sharp change in the slope of the curve. 495

4 Methodology

In this section, we describe our method for the analysis 497 of 3D CT scans of material packing. The method first 498 constructs a geometric and a topological structure – a 499 segmentation of the packing into its constituent particles, 500 and a topologically accurate contact-based connectivity 501 network between the particles. The structures are con-502 structed in two steps. First, we extract the material's 503 boundary surface and compute its associated distance 504 field. Second, we use the shape and connectivity in-505 formation as captured by the gradient of the distance 506 field to inform the segmentation and network extrac-507 tion. Figure 1 presents an overview of the method while 508 comparing it with the watershed-based approach and 509 Figure 6 illustrates the individual steps of the method 510 using a synthetic 3D dataset. 511

We begin with a description of the dataset followed ⁵¹² by the individual steps of the method. ⁵¹³

4.1 Dataset and preprocessing

We analyze a 3D CT scan of a packing of sand particles 515 coated with epoxy at the contacts. The packing has 516 a contact bound structure *i.e.*, the cementation exists 517 only at the contacts in a skeleton of sand particles [44]. 518 The scan has a dimension of $888 \times 912 \times 1360$ voxels 519 with a resolution of 12.5 µm per voxel. For the data to 520 fit in memory, we first downsample the scan by a factor 521 of 4 across each dimension. The downsampled image is 522 computed using a simple mean operation across 4×4 523 \times 4 sized cubes. We then run our analysis algorithms 524 on the downsampled data of size $222 \times 228 \times 340$. 525 Further, the epoxy bonds are not segmented separately 526 due to the weak contrast in the reconstructed values 527 of epoxy and pore air. However, this does not affect 528 the segmentation accuracy because the packing has a 529 contact bound structure and the epoxy bonds form at 530 the particle contact. 531

4.2 Boundary surface extraction

The material boundary surface is extracted using an automatic bi-modal threshold computation followed by an iterative local refinement of the corresponding isosurface. 536

We use Otsu's method [35] to compute the threshold for the initial boundary isosurface. Otsu's method takes as input the 3D image and essentially searches for an optimal threshold that divides the image into two classes - foreground and background. Optimality, 540

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496

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Fig. 6: Method pipeline. First, compute a bounding surface based on an automatically determined bi-modal threshold and a local active contour-based refinement. Next, compute a signed distance field from this surface and the MS complex for this scalar field. Simplify the MS complex via iterative cancellation of pairs of adjacent critical points in the MS complex. The difference in function value between the pair of critical points, called its persistence, is used to determine the order of cancellation. The threshold for simplification is determined by analyzing the persistence diagram. The segmentation is computed as the descending manifolds of the maxima in the simplified complex. Apply geometry driven pruning to compute the connectivity network.

in this search, is defined as a minimum intra-class vari-542 ance. While the Otsu threshold-informed isosurface is a 543 good starting point, attempting to pick a single scalar 544 threshold that minimizes intra-class variance globally 545 for the two classes may result in a surface that is locally 546 non-optimal. Specifically, a global threshold restricts 547 the search to isosurfaces of the function. The boundary 548 surface is not necessarily modeled well by an isosurface, 549

especially in regions where a locally better surface geom-550 etry is ignored due to the restriction and assumption of 551 a single globally optimal isovalue. We therefore employ 552 a local refinement procedure using an active contour 553 model to construct a better boundary surface. Active 554 contour models apply local updates on an input surface 555 to optimize a given objective function. In particular, 556 we adapt the Chan-Vese method [8], also known as the 557

Active Contours Without Edges (ACWE) method. The 558 ACWE method, as opposed to using a gradient based 559 objective function to extract a boundary surface, tries 560 to minimize intra-class variance both inside and outside 561 the surface within an iteration. The primary advantage 562 of the ACWE method is its ability to robustly handle 563 images with noisy boundaries without relying on nu-564 merical and potentially unstable gradient computations. 565 We adapt the ACWE model to our setting by ignor-566 ing the smoothness and rigidity terms in the standard 567 ACWE objective function to allow for sharp features in 568 the boundary surface. As a result, our model effectively 569 minimizes the following objective: 570

$$F(C, c_1, c_2) = \lambda_1 \int_{Inside(C)} |f(x, y, z) - c_1|^2 dx dy dz$$
$$+ \lambda_2 \int_{Outside(C)} |f(x, y, z) - c_2|^2 dx dy dz,$$

where C is the surface in the current iteration, f is 571 the image intensity scalar function defined at each voxel, 572 c_1 is the average intensity of voxels inside C, c_2 is the 573 average intensity of voxels outside C, λ_1 is the normal-574 izing constant for initial number of points and range of 575 intensity inside C, and λ_2 is the normalizing constant for 576 initial number of points and range of intensity outside 577 C. 578

⁵⁷⁹ Our active contour framework successfully gives us ⁵⁸⁰ high quality particle boundary surfaces across packings ⁵⁸¹ of a diverse set of materials and geometries.

4.3 Segmentation and connectivity networkcomputation

Next, we segment and compute the connectivity network 584 of the individual particles in the ensemble based on the 585 geometry of the extracted boundary surface. A popu-586 lar two-step approach for segmenting connected regions 587 that touch each other first computes a distance trans-588 form from a bounding surface followed by a watershed 589 transform [37], described in section 1.1. In watershed 590 transform, the central point of the contact region be-591 tween two catchment basins is a 2-saddle of the distance 592 field. The 2-saddles are therefore natural representatives 593 of the regions of contact between two particles in the 594 packing. Watershed-based methods are however prone 595 to over-segmentation of non-convex regions. Such re-596 gions may contain multiple local maxima of the distance 597 field. So, it is necessary to design region merging proto-598 cols that fix the over-segmentation errors. In our work, 599 we achieve robustness and computational efficiency in 600

the two-step pipeline by using a fast Chamfer distance transform [27] and the topological framework of the MS complex. We use the MS complex for (a) characterizing and computing the catchment basins and regions of contact, and (b) to fix over-segmentation using an elegant combinatorial topological simplification procedure.

Initial catchment basin computation. The descend-607 ing 3-manifolds of maxima of the distance field corre-608 spond to the catchment basins and aid in computing the 609 segmentation. We retrieve the set of all descending mani-610 folds of maxima from the Morse complex of the distance 611 field. The set of these descending manifolds, further 612 restricted to lie within the material boundary surface, 613 is a segmentation of the packing into its individual par-614 ticles. This step may produce over-segmented regions 615 due to imaging noise or small variations in the distance 616 field may introduce multiple local maxima within some 617 non-convex regions. 618

Topological simplification. Next, we apply topolog-619 ical simplification to merge such regions and fix the 620 over-segmentation. As described earlier, the topological 621 simplification proceeds through an iterative cancellation 622 procedure, canceling critical point pairs with persistence 623 below a given simplification threshold. In our method, 624 we restrict such cancellations to 2-saddle-maxima pairs. 625 Intuitively, the cancellation operation essentially merges 626 regions represented by low persistent maxima connected 627 with the higher persistence regions. Persistence of a 628 2-saddle-maxima pair, can be loosely defined as the 629 difference in function values, thus representing the dif-630 ference between the radius of the contact (represented 631 by the distance value of the 2-saddle) and the radius of 632 the particle (represented by the distance value of the 633 maxima). High persistent pairs are therefore particles 634 where the contacts are much closer to the boundary sur-635 face than the central points in the particles. Conversely, 636 low persistent pairs are particles where the width of 637 the contact is comparable to the width of the entire 638 particle thus implying that the particle may possibly be 639 over-segmented. 640

We select a persistence threshold for simplification 641 by identifying a knee in the persistence curve, a plot of 642 decreasing persistence values over all critical point pairs. 643 The presence of a clear knee in the curve indicates a 644 clear separation of features from noise. Following noise 645 removal via topological simplification, the quality of 646 the segmentation can be further improved based on 647 additional geometric criteria. For instance, we observed 648 over-segmentation in particles with high-width contacts 649 relative to particle size. We fix it by using the ratio of 650 the distance field values at the 2-saddle and maximum. 651 In particular, particles with the aforementioned ratio 652 greater than 0.75 were often found to be over-segmented 653 and were thus merged with their neighbor particle, as defined by the 2-saddle.

Contact identification. A 2-saddle of the distance 656 field with a positive value represents a connection be-657 tween two contacting particles because the descending 658 manifolds of the maxima associated with the particles 659 meet at the 2-saddle. The ascending 1-manifold of the 660 2-saddle is therefore a steepest ascent arc connecting the 661 2-saddle to the maxima associated with the contacting 662 particles. The set of all ascending manifolds of 2-saddles 663 is thus a geometrically meaningful connectivity network 664 between the contacting particles in the packing. How-665 ever, retrieving this entire set from the MS complex 666 results in a number of spurious and clustered 2-saddles 667 and multiple arcs connecting a pair of maxima. Since 668 multiple contacts may exist between a pair of parti-669 cles, it is important to distinguish between the primary 670 2-saddles that represent the central point within a con-671 tact region and spurious 2-saddles that are alternate 672 673 representations of a given contact region. We achieve this by computing all explicit contact regions between 674 two connected particles and picking the saddle with the 675 highest function value from each such extracted region. 676 For each pair of particles, the contact regions are 677 computed by clipping the descending 2-manifold of the 678 highest 2-saddle. The descending 2-manifold is clipped 679 by the boundary surface *i.e.*, all 2-cells containing ver-680 tices with distance value less than 0 are removed. The 681 connected components of the remaining 2-cells repre-682 sent the contact regions. While extracting the connected 683 components, we mark the highest 2-saddle within each 684 component and retain its arc in the connectivity net-685 work. We remove all other spurious arcs and therefore 686 extract a robust contact-based connectivity network for 687 the granular ensemble packing. 688

5 5 Implementation

In this section, we describe implementation details of 690 the proposed method and the practical experience of 691 executing the pipeline shown in Figure 6. We highlight 692 the unified data structure that supports efficient and 693 robust processing, and the principled simplification that 694 results in good quality segmentation. From a user's 695 perspective, executing the pipeline consists of three 696 main steps: 697

Boundary extraction. The first step processes the
input CT image, computes and stores the boundary
surface that separates material from the surrounding
volume. The surface is stored as a distance field, sampled
at the same resolution as the input CT image. The user
provides the CT scan as input and the software runs the

active contour optimization, automatically computing 704 the required parameters, and returns the distance field. 705

MS complex computation and simplification. The 706 second step processes the distance field to construct the 707 MS complex, computes the persistence curve, and dis-708 plays it to the user. Guided by the curve, the user selects 709 an appropriate persistence threshold. An appropriate 710 choice is the knee in the curve as indicated in Figure 6. 711 The iterative cancellation based simplification directed 712 by a measure of importance such as difference in scalar 713 function values simplifies all gradient based manifolds 714 consistently. The persistence curve is a succinct and 715 abstract visual representations of the distribution of 716 extracted particle segments corresponding to a selected 717 simplification measure. It provides a data-driven process 718 to the user for selecting a simplification threshold. The 719 MS complex is simplified based on the selected thresh-720 old to separate scan noise and concave protrusions from 721 potentially important features. 722

The resulting simplified MS complex is stored. All 723 ascending and descending manifolds are stored in a 724 topologically consistent fashion. Specifically, three types 725 of manifolds are computed - the 3D descending mani-726 folds of maxima that represent the segmentation into 727 individual particles, the 2D descending manifolds of 728 2-saddles that are processed to compute the contact re-729 gions between particles, and the 1D ascending manifolds 730 of 2-saddles that represent the connectivity network. 731 The computed manifolds are stored in a unified data 732 structure that supports fast queries by separating the 733 storage of the connectivity of the MS complex from its 734 geometry. The nodes and arcs (0-cells and 1-cells) of the 735 MS complex are stored as a graph and the geometry is 736 extracted on demand. The data structure also enables 737 flexible and efficient processing of the manifolds while 738 guaranteeing that the topological relationships between 739 the manifolds are maintained consistently. Specifically, 740 it supports global simplification of the MS complex to 741 remove noise and the local clustering of 2-saddles in the 742 connectivity network using the contact regions. 743

The algorithms for computing the MS complex are 744 amenable to parallel computation and execute on multi-745 core CPUs and GPUs resulting in small run times [40] 746 Note that seemingly time-intensive tasks such as com-747 puting the persistence curve are executed efficiently in 748 our pipeline due to fast combinatorial algorithms that 749 require only the connectivity of the MS complex for 750 these steps. 751

Geometry extraction. The third step focuses on extracting the desired geometric and topological structures from the simplified MS complex. The user can optionally extract and store the segmentation, contact information, and the connectivity network. The segmentation consists 756 of the segmented volume together with associated particle centres. The contact information consists of contact
regions between a pair of particles and the associated
contact points. The connectivity network connects the
particle centers and contact points. This comprehensive
collection of structures can then conveniently be used
to facilitate user-desired downstream analysis.

In addition to the intuitive parameter-free pipeline, the code is easy to install and execute. The core algorithms used in the pipeline, including the boundary surface and discrete MS complex computation are available from existing software packages [47, 39, 29, 41]. We further plan to release our source code in the public domain for use by the community.

771 6 Benchmarking and Experimental Results

In this section, we present results and experiments to 772 validate our approach and illustrate the key benefits 773 of our method compared to standard watershed-based 774 pipelines for the analysis of granular materials. The sec-775 tion is divided into three main parts. We first benchmark 776 our method by computing physically expected statis-777 tics for a packing of mono-dispersed steel beads. After 778 validating our approach on a well-studied dataset, we 779 demonstrate the utility of the method for the analysis 780 of complex packing structures by processing a cemented 781 sand packing. We compute and visualize the segmenta-782 tion and connectivity network for the packing and high-783 light observed statistical trends. Finally, we compare 784 the quality of computed results and ease of computation 785 with a recent watershed-based pipeline for the analysis 786 of cemented granular materials [44]. All computational 787 experiments were performed on a workstation with an 788 Intel Xeon E5-1660 v4 @ 3.8 GHz processor with 8 789 cores, 128 GB main memory, and an Nvidia Quadro 790 P2000 graphics card with 6 GB RAM. As described ear-791 lier, the MS complex is computed using a GPU parallel 792 algorithm [41] 793

⁷⁹⁴ 6.1 Benchmarking: mono-dispersed steel sphere packing

We use the tomography scan of a packing of mono-795 dispersed steel beads to benchmark our algorithm. We 796 compare physically expected statistics like the average 797 coordination number and diameter (in mm) in the pack-798 ing to perform the benchmarking. The original scan has 799 a dimension of $681 \times 681 \times 1004.$ In order to segment 800 this large volume, the analysis is carried out by splitting 801 the volume into five blocks (1-201, 202-402, 403-603, 802 604-804, and 805-1004 along the height) and four in-803 termediate blocks (131-271, 332-472, 533-673, 734-874 804

along the height). These blocks were chosen such that 805 each particle is completely captured in at least one of 806 the blocks. The center of each particle was remapped 807 to the original volume and all duplicate entries and 808 partially split particles were removed. Table 1 presents 809 the average diameter and coordination number for both 810 methods. The average diameters (0.98 mm and 1.0 mm) 811 are consistent within the resolution $(16\mu m)$ of X-ray CT. 812 Theoretically, the average coordination number of a ran-813 dom packing of mono-dispersed spherical particles with 814 infinite friction is known to lie in the interval [4,6], as 815 calculated using Maxwell counting [4]. A deviation from 816 the perfect spherical shape increases the average coordi-817 nation number [12]. The average coordination number 818 in our computation (6.04 and 6.08) is close to the the-819 oretical range and within the experimentally obtained 820 range [5]. 821

Table 1: Comparing statistics for mono-dispersed steel beads.

	Diameter (mm)	Coordination number
MS Complex - High res. ($681 \times 681 \times 1004$)	0.98	6.04
Watershed - High res. $(681 \times 681 \times 1004)$	1.00	6.08

We also study the effect of data resolution by com-822 puting the statistics for downsampled versions of the 823 data. We create a medium resolution (341 \times 341 \times 824 502) and a low resolution $(171 \times 171 \times 251)$ version 825 by downsampling by a factor of 2 and 4, respectively. 826 Figure 7 shows the segmentation and connectivity net-827 work for medium resolution. The average diameters were 828 found to be 0.96 mm and 0.91 mm for medium and low 829 resolution, respectively. The difference from the actual 830 diameter increases with decreasing resolution (error in 831 the measurement), namely 32 μm (medium) and 64 μm 832 (low). The average coordination numbers were 6.03 and 833 6.06 for medium and low resolution volumes. These re-834 sults are also consistent with the watershed results at 835 the original resolution. This indicates that our method is 836 robust to downsampling for the computation of average 837 statistics. 838

6.2 Visualizations of extracted geometric and topological structures 840

In this section, we demonstrate our proposed method's potential for analyzing complex packing structures by 842



Fig. 7: The segmentation (left) and connectivity network (right) for the mono-dispersed steel bead packing.

Processing a cemented sand packing with varied particle geometries. From the computed MS complex, we first extract the following geometric and topological structures:

- Segmentation of the packing structure into its con stituent individual particles.
- Topological network of contact-based connections
 between particles, embedded in the domain using
 geometrically meaningful arcs.
- Geometric regions of contact between connected par ticles.

Access to these structures facilitates both visual exploration and statistical analysis of the material packing
structure. In particular, the computed segmentation and
connectivity network is used to generate high-quality
visualizations of the individual particles in the packing
(Figure 8) and their contact structure (Figure 9).

The MS complex data structure and persistence-860 based visualizations of the extracted segments (see Fig-861 ure 11) further allow for the convenient selection and 862 visualization of specific segments and contacts of in-863 terest. The selection may be directed by a variety of 864 geometric and topological criteria such as the coordina-865 tion number, volume, and relative width of the particle 866 compared to the contact region. Figure 8 (right) and 867 Figure 9 (right) highlight visualizations of such selected 868 particles and Figure 10 illustrates the visualization of a 869 selected contact and its extracted properties. 870

⁸⁷¹ 6.3 Statistical Analysis

In addition to localized exploration and analysis, our
framework supports robust and efficient computation
of detailed statistical metrics to study global trends in
the packing structure. Figure 12 presents a comparison

between the statistics. We highlight the similarities and 876 differences between the histograms of the coordination 877 number (with and without multiplicity), volume, and 878 the orientation of contacts for individual particles in the 879 packing. We report statistics computed using watershed-880 based segmentation computed using both manual and ac-881 tive contour based boundary surfaces. The histogram of 882 coordination number with the inclusion of multiple con-883 tacts is significantly different from the watershed-based 884 method. The histogram computed by excluding multiple 885 contacts show similar trends, for example the most fre-886 quent coordination number is observed in the range of 887 4 to 6. The distribution of volume of particles is similar 888 but there is a small difference in the actual counts. Both 889 methods predict that most contacts are oriented along 890 the z-direction (direction of gravity); this observation 891 is consistent with previous studies on fabric of granu-892 lar materials and cemented granular materials [31, 44]. 893 The differences in counts of histograms is due to over-894 segmentation of the particles by the watershed-based 895 method (discussed in section 6.4). 896

We also present a spherical histogram representation of the sand packing obtained from this analysis, see Figure 13. Details on creating these spherical histograms are provided elsewhere [43]. Further details on the quantification of the fabric of this cemented sand ensemble is presented in the supplementary material.

6.4 Watershed comparisons

In this section, we present a detailed comparison between our method and a recently proposed watershed-based pipeline for the analysis of cemented granular materials [43]. Using the cemented sand packing as a case study, the key benefits of our method are highlighted below in a comparison divided across the fundamental steps 909



Fig. 8: Global (left) and local (right) visualization of the segmentation of the packing into its constituent individual particles. Particles are colored by their segmentation identifiers.



Fig. 9: Global (left) and local (right) visualization of the geometric embedding of the extracted connectivity network of the packing. In the local view (right), bead centres are represented as red spheres and contacts as blue spheres.

of boundary extraction, segmentation, and connectivitynetwork extraction.

Boundary surface extraction. The histogram of scalar 912 values is often used to select a global scalar threshold 913 that separates the scan into regions containing material 914 and the surrounding medium. A good separating surface 915 consists of a collection of surface pieces that are defined 916 by different scalar thresholds within different regions 917 of the scan. As a result, a boundary surface that is de-918 fined by a single threshold contains noticeable artifacts 919 that range from incorrectly merged particles to highly 920 distorted particle geometries. 921

In comparison, our method's active contour-based optimization computes an optimal boundary surface (Figure 14) that locally adapts to the best separating value, and is thus free of the assumption of a single global threshold. Figure 14 highlights artifacts in a boundary surface that is computed using a global threshold and compares it against the improved surface computed using our active contour-based approach. The higher quality of the boundary surface results in an improvement in the quality of downstream analysis.

Segmentation. The difference in quality of the ex-932 tracted boundary, coupled with the watershed trans-933 form's tendency to over-segment in the presence of noise 934 and non-convex geometry, causes a tangible quality dif-935 ference between the segmentation results of our Morse 936 theory-based approach and watershed-based methods. 937 As discussed in Section 2, post-processing heuristics 938 such as the h-maxima transform merge over-segmented 939 regions based on a given threshold. However, a precise 940 and optimal choice of threshold for the transform is 941



Fig. 10: Extracted region and orientation of contact between two sand particles colored in blue and brown. The translucent gray surface represents a clip of the extracted boundary surface of the particles near the contact region. This surface is the descending 2-manifold of the 2-saddle that defines the contact. Constituent points of the contact region are represented as white spheres, with the arrow displaying a fitted normal vector to the contact.



Fig. 11: Persistence diagram (left) and curve (right) for the cemented sand packing. The knee in the persistence curve (0.7) informs the choice of a noise threshold that separates features from noise. We can see how the selected threshold removes the cluster of low-persistence noise (white region) from the remaining features (grey region) in the persistence diagram.

elusive, and thus this merging process can potentially
both struggle to correct over-segmentation or result in
under-segmentation.

Our persistence-based framework alleviates this in-945 convenience by providing an intuitive and statistically 946 optimal choice of threshold for simplification. In par-947 ticular, we use the knee of the persistence curve (see 948 Figure 11) to identify a precise and well-motivated choice 949 of threshold. The persistence curve is a plot of the num-950 ber of segments for increasing values of simplification 951 threshold. It can be computed efficiently due to the uni-952 fied data structure, which supports simple and efficient 953 updates. Computing a similar curve for an h-maxima 954 transform requires the repeated application of the trans-955 form for a large number of values while counting the 956 segments, and still arriving at an approximation of the 957 persistence curve. Figure 15 shows the watershed seg-958 mentation using a manually selected h-maxima thresh-959

old, which is identified via a visual inspection process960followed by an expert, and highlights the improvement961in segmentation achieved using the automatically computed persistence curve-based threshold. The segmentation962tion quality is better both in terms of the geometry of964individual particles and in terms of fewer cases of under-965and over-segmentation.966

The quality of segmentation also implicitly influences 967 the accurate identification of contacts in the packing. 968 While geometric inaccuracy in a segmentation due to 969 a non-optimal boundary surface can affect the identi-970 fied geometric position of the contacts, issues such as 971 under-segmentation and over-segmentation can result 972 in missing or spurious additional contacts. For instance, 973 true contacts may be missed due to under-segmentation, 974 and over-segmentation of a single particle leads to spu-975 rious contacts between incorrectly identified segments. 976 Figure 16 compares the set of contacts identified by the 977



Fig. 12: Statistical analysis of sand granular ensemble computed using the Morse theory-based and watershed-based methods. Histogram of coordination numbers is similar when counting contacts with multiplicity, else the histograms are similar. The distribution of volume of particles is similar but the actual counts are different.

two methods by plotting them on a manually tuned 978 volume rendering of the CT scan. The volume rendering 979 helps us visualize key differences in the contacts identi-980 fied by the two methods. The highlighted circles bring 981 to attention particularly problematic regions, where 982 either essential contacts are missed or are incorrectly 983 placed or spurious additional contacts are identified by 984 the watershed-based approach. Further, the large dis-985 parity in the number of identified contacts highlights 986

the magnitude of difference in results between the two approaches.

987

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An additional benefit of our method is that the crite-989 rion used for simplification can be suitably modified to 990 accommodate different packing structures with variation 991 in particle size and geometry. The topological simplifi-992 cation step naturally supports alternative criteria for 993 iterative region-merging. The persistence diagram can 994 be used as an intuitive visual tool to explore the dis-995 tribution of the packing based on the selected criteria, 996



Fig. 13: (left) Normalized particle stereographic histogram and (right) normalized contact spherical histogram for sand packing.



Fig. 14: High quality boundary surface extraction. A side-by-side visualization of a volume rendering of the CT scan (left) along with the boundary surfaces extracted using the isosurface method (middle) and active contour optimization (right). The inset highlights a region containing multiple artifacts in the isosurface boundary. The geometry of an elongated particle is not captured correctly, it merges with other particles in the packing. The volume rendering is generated using a manually tuned color and opacity map that delineates the individual particles.

⁹⁹⁷ thus providing a unified framework for the intuitive exploration of diverse choices of criterion and associated
⁹⁹⁸ thresholds.

Connectivity network extraction. In order to rep-1000 resent the contact structure, existing methods use a 1001 network of straight lines connecting particle centers and 1002 contacts. However, in the case of non-convex particle 1003 geometries, such a network fails to account for the ge-1004 ometry of the packing [44]. For example, arcs of the 1005 network that are represented as line segments may lie 1006 outside the individual particles. 1007

Our method instead uses the gradient of the distance 1008 field to compute a geometry-aware connectivity network 1009 that resides within the packing structure, see Figure 9. 1010 The combination of the discrete Morse theory-based 1011 gradient and the unified persistence-based simplification 1012 allows us to efficiently compute a numerically robust 1013 connectivity network that accurately captures the global 1014 packing structure. 1015



(a) Direct volume rendering

(b) Watershed-based approach with h- (c) Morse theory-based approach maxima threshold = 2.0

Fig. 15: Improved segmentation. Comparison of particle segmentation obtained using the watershed and our Morse theory-based approach. Red box: Geometry of the segments computed using the Morse complex agree better with the original CT data thanks to the improved quality of the boundary surface. Green and Blue boxes: Multiple cases of under segmentation in the watershed approach. In both cases, two particles are incorrectly identified as a single particle whereas the Morse complex identifies them correctly. The volume rendering uses a manually tuned color and opacity map to delineate individual particles.

1016 7 Conclusions

In summary, we outline a robust and efficient approach 1017 to compute the segmentation and connectivity network 1018 of granular material packings from x-ray CT images. 1019 Through the combination of active-contour optimization 1020 and Morse theory, our approach functions as a unified 1021 and fully-automated framework to compute, query, and 1022 visualize the geometric and topological properties of 1023 the packing structure. The automated nature of the 1024 algorithm allows for convenient large-scale computation 1025 and the persistence-based simplification approach en-1026 ables the intuitive exploration of variation in particle 1027 geometries. Salient features of our method, such as the 1028 locally-optimal boundary extraction, an efficient and 1029 flexible framework for noise removal and geometry-aware 1030 connectivity network, together ensure that the approach 1031 generalizes to diverse packing structures with variation 1032 in particle size, geometry, and material density. 1033

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Conflict of interest

The authors declare that they have no conflict of interest 1042

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(a) Direct volume rendering

(b) Watershed-based approach with h- (c) Morse theory-based approach maxima threshold = 2.0

Fig. 16: Improved identification of contacts. Comparison of particle contacts obtained using the watershed and our Morse theory-based approach. A total of 2746 contact points are identified using watershed approach while 1842 points are identified by our method. Blue circle: The watershed-based method fails to identify the contact point. Red and green circles: The watershed-based approach incorrectly identifies a large number of contact points. The contacts identified by our approach appear to be reasonable when compared to the volume rendering.

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