Semantic Segmentation and Model Fitting for Triangulated Surfaces using Random Jumps and Graph Cuts

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Abstract: Recent advances in 3D reconstruction allows to acquire highly detailed geometry from a set of images. The outcome of vision-based reconstruction methods is often oversampled, noisy, and without any higher-level information. Further processing such as object recognition, physical measurement, urban modeling, or rendering requires more advanced representations such as computer-aided design (CAD) models. In this paper, we present a global approach that simultaneously decomposes a triangulated surface into meaningful segments and fits a set of bounded geometric primitives. Using the theory of Markov chain Monte Carlo methods (MCMC), a random process is derived to find the solution that most likely explains the measured data. A data-driven approach based on the random sample consensus (RANSAC) paradigm is designed to guide the optimization process with respect to efficiency and robustness. It is shown, that graph cuts allow to incorporate model complexity and spatial regularization into the MCMC process. The algorithm has successfully been implemented and tested on various examples.

1 INTRODUCTION

The reverse engineering of CAD models based on triangulated surfaces definitely belongs to the most important fields of 3D computer vision. The main challenge is to find a valid decomposition of the surface which allows to fit a set of geometric models. This is a typical chicken-and-egg problem. Many related algorithms, however, try to tackle this problem by applying a set of sequential processing steps. In this paper, we argue that the high presence of noise on the one hand, and the bidirectional dependencies of these steps on the other hand, require a global approach to find an optimal solution. The proposed processing flow is motivated in Figure 1. Especially mechanical objects can be described as a composition of bounded geometric primitives (a). Towards CAD-based applications, we aim at reconstructing a geometric representation from a noisy triangulated surface (b). Simultaneously, the surface is decomposed into meaningful segments (c), where each segment corresponds to a component of the object. The object as well as the labeling of the vertices define a "world state". From a Bayesian perspective, we want to find the state which most likely explains the observation. The resulting optimization problem is defined in a highdimensional and non-continuous domain. The jumpdiffusion framework seems to be particularly suited for these complex problems. It allows to optimize energy functions which are defined in a complex space that includes several sub domains of different dimensions. Secondly, it can escape from local minima.

1.1 Contributions

This paper contains several new contributions. An MCMC process is designed for the purpose of mesh processing. The semantic segmentation is formulated as an inference problem based on a triangulated surface. Compared to related algorithms, we incorporate model complexity and spatial regularization into the random-jumps process. A crucial aspect of MCMC methods is the computation of the proposal probabilities. We show, that RANSAC allows to compute a non-uniform distribution that balances efficiency and robustness against local minima.

1.2 Overview

The remaining part of the paper is structured as follows. A brief overview of related approaches is given in Section 2. In Section 3, we formulate the task as an inference problem. An energy function is derived to evaluate the likelihood of the current state. In Sec-

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Figure 1: Schematic illustration of the reverse-engineering problem. An object is given in (a). The object is measured by a vision- or laser-based reconstruction tool chain which results in a noisy triangulated surface (b). The problem is now to recover the original object based on the measurement and some general prior knowledge. Especially man-made geometry can be described by a set of bounded geometric primitives. The proposed algorithm robustly decomposes the triangulated surface into meaningful segments (c) and recovers a set of components accordingly (d).

tion 4, we briefly introduce geometric primitives and define the corresponding cost functions. Finally, we design a MCMC process in Section 5 in order to robustly minimize the energy function. The resulting algorithm has been successfully implemented and an evaluation can be found in Section 6. The paper concludes with Section 7.

2 RELATED WORK

The presented approach is related to a broad field of algorithms which can roughly be subdivided into multi-model fitting problems, mesh segmentation and classification. In the following, an overview of related work is given.

2.1 Mesh Segmentation and Classification

There has been a lot of attention to the problem of mesh segmentation. Many segmentation algorithms are inspired by classical machine learning methods, such as hierarchical clustering, k means, or mean shift. Based on the fitting of three different geometric primitives, Attene et al. propose a fast greedy method that segments a given mesh by an iterative clustering scheme (Attene et al., 2006). Another important group of algorithms follows the k-means strategy. Cohen-Steiner et al. present an algorithm referred to as variational shape approximation that is based on k means (Cohen-Steiner et al., 2004). The partition step is realized by a modified region growing scheme that assures connected segments. Several other authors propose modified or extended versions of the variational approach (Wu Leif Kobbelt, 2005)(Yan et al., 2012). The most important problems of variational methods are a suitable initialization and the adequate choice of the number of clusters.

Especially suited for man-made geometry, feature-based algorithms achieve an implicit segmentation by detecting the blending regions. These algorithms do not explicitly fit geometric models (Várady et al., 2007)(Mangan and Whitaker, 1999). A (semantic) segmentation can also be achieved by designing a classification problem, where each vertex is assigned with a label which can have a semantic meaning. This motivates to use methods like Markov random fields or graph cuts. The spatial regularization allow to achieve smoother boundaries than variational or hierarchical methods. However, the set of classes is fixed and need to be known a priori. Related algorithms are often designed to learn and apply a natural segmentation based on large datasets of models such as animals, humans or furniture(Longjiang et al., 2013)(Lavoué and Wolf, 2008)(Lai et al., 2008). They decompose objects with respect to topology or the minima rule which is mainly suited for organic shapes such as humans or animals. Some approaches require a set of similar training models which is often not available in reverse-engineering scenarios (Kalogerakis et al., 2010)(Benhabiles et al., 2011)(Lv et al., 2012).

2.2 Multi-model Fitting Problems

The challenge described in this paper is clearly related to the class of multi-model fitting problems. Recently there has been a lot of attention to this field. Roughly summarized, there are three different approaches to the fitting of multiple models.

2.2.1 Random Sample Consensus

First, RANSAC is a well-known strategy to estimate model parameters in presence of a high number of outliers. In each iteration, a model candidate is parametrized, based on a minimal set that is randomly drawn from the input data. Each candidate is scored by the number of inliers that are sufficiently explained by the model. Schnabel et al. propose an efficient approach where RANSAC is used to extract primitive shapes from a noisy point cloud in an iterative manner. The result significantly depends on a set of parameters that are required to detect the inliers in the sampling procedure (Schnabel et al., 2007). Papazov et al. use a RANSAC-like strategy for the recognition of high-level objects in scattered environments (Papazov and Burschka, 2011). The greedy nature of multi-RANSAC algorithms does not guarantee a global optimum. Li et al. uses RANSAC to globally fit a set of primitives with respect to alignment constraints (Li et al., 2011). There approach makes strong assumptions on the geometry of the object which is not always given.

2.2.2 Multi Labeling

Secondly, multi-model fitting can be solved by graphical models such as MRFs or graph cuts. Delong et al. have extended the α -expansion algorithm for incorporating label costs. The additional term in the optimization function allows to minimize the number of used classes or models explaining the data (Delong et al., 2010). Based on multi-label optimization, Isack and Boykov introduce an algorithm referred to as PEaRL which iterates a two-step procedure (Isack and Boykov, 2012). Given a set of geometric models, the algorithm of Delong et al. is used to compute a labeling which incorporates label costs and spatial regularization. In the second step, the set of all models is reconfigured by inserting new models, deleting obsolete models, or updating their parameters. The work of Woodford et al. introduces a new type of contraction move for the optimization of multi-model fitting problems (Woodford et al., 2012). The drawback of these methods is that the update of the model configuration is deterministic and thus they cannot escape from local minima.

2.2.3 Random Jumps

A third powerful and very general framework for multi-model fitting is given by random jumps. Tu and Zhu proposed an image segmentation approach based on random jumps (Tu and Zhu, 2002). The image is assumed to be a composition of a set of heterogeneous models that allow to describe typical regions such as clutter, texture, or shading. The global energy function takes data costs and boundary smoothness into account. As one of the key contributions, they use data-driven techniques such as expectation maximization or mean shift to compute new proposals and their corresponding probabilities. Han et al. use a very similar framework to segment range images based on geometric models (Han et al., 2004). Image regions which cannot be fitted by any of these models are described by a cluttered type which is a 3D histogram. Larfarge et al. use geometric primitives and the MCMC framework for the reconstruction of socalled hybrid models from a set of images and known camera poses (Lafarge et al., 2013). Similar to Hang et al., a hybrid model can either be a primitive or an arbitrary shape which is represented by a triangular mesh. The data term in the energy function enforces photo consistency which measures the error between the image projections of two cameras on a geometric primitive. Model complexity as well as spatial regularization in the labeling of the mesh is not considered.

3 STOCHASTIC PROBLEM FORMULATION

Consider a polyhedral surface $S = \{V, E, T\}$, which consists of vertices V, edges E, and triangles T. S is assumed to be a measurement (observation) of an object *O*. The hidden world state is defined as:

$$\mathbf{P} = \{O, \mathcal{X}\}, \ |\mathcal{X}| = |\mathbf{V}| \tag{1}$$

where Φ is the model *O* itself and the labeling *X* which explicitly assigns a component of *O* to each vertex on **S**. *O* is decomposed into *K* components:

$$\mathcal{O} = \left\{ K, \left\{ \mathcal{M}_k, \mathcal{A} \right\}, \ k = [0, \dots, K] \right\}, \qquad (2)$$

where a component $\mathcal{M}_k = \{r_k, \partial_k, \theta_k\}$ is given by the type of the underlying geometric primitive r_k , a parameter vector θ_k , and a boundary ∂_k . Geometric primitives are represented by implicit equations that allow to efficiently compute distances or projections (Schnabel et al., 2007). \mathcal{A} is introduced as a default model. From a classification perspective, \mathcal{A} can also be seen as an "unassigned"-class that represents all vertices which cannot be explained by any other class in O. All components of O are associated with a label $l \in \mathcal{L}$:

$$\mathcal{L} := \{ l_{\mathcal{M}}^{k} \mid k \in [0, \dots, K], l_{\mathcal{A}} \}, \qquad (3)$$

where *l* is the *k*-th unit vector. The labeling of the vertices $l \rightarrow x$ implies a semantic segmentation on **S**, $\mathbf{S} = \{S_0, \ldots, S_K, S_A\}$, where S_k corresponds to the set of all vertices which are being labeled with l^k . ∂_k is the boundary of the corresponding segment on \mathbf{S} , $\partial_k = \partial S_k$. Finally, the overall problem is to find a world state Φ^* that most likely explains a given observation, **S**:

$$\Phi^* = \arg\max_{\Phi \in \Omega} p(\Phi \mid \mathbf{S}) . \tag{4}$$



Figure 2: Effect of the prior term for the Grabber model. The result in (a) is based on data cost only and 200 initial shape candidates. The result of the MCMC process with a low, non-zero prior can be seen in (b). If the regularization parameters are set to higher values, the algorithm produces a smaller number of segments and smoother boundaries, as it can be seen in (c).

 $p(\Phi|\mathbf{S})$ can be decomposed into prior and likelihood:

$$p(\Phi|\mathbf{S}) \propto p(\mathbf{S}|\Phi)p(\Phi)$$
. (5)

For the sake of simplicity, the optimization problem in Equation 5 is transferred to the log-probability domain:

$$\Phi^* = \arg\min_{\Phi \in \Omega} U(\Phi, \mathbf{S}) = U_D(\Phi, \mathbf{S}) + U_M(\Phi) . \quad (6)$$

 $U_D(\Phi, \mathbf{S})$ corresponds to the likelihood in Equation 5 and is referred to as data term in the optimization community. $U_M(\Phi)$ represents prior knowledge about the spatial regularity and the model complexity of Φ .

3.1 **Prior Distribution**

As soon as assumptions or knowledge about the demanded model O is available, the prior distribution is non uniform and allows to enforce specific properties on Φ . The complexity of O is given by the complexity of the models \mathcal{M}_k :

$$U_{\mathcal{M}}(\mathcal{O}) = \sum_{k=1}^{K} |\mathcal{M}_{k}|, \mathcal{M}_{k} \in \mathcal{O}, \qquad (7)$$

where $|\mathcal{M}_k|$ represents the complexity cost for each type of model. As specified in Table 4, we prefer simple over more complex models. $U_{\mathcal{M}}(O)$ is also referred to as label cost term. Each vertex of **S** is assigned to a class and the set of all vertices implies the boundary of the corresponding surface patch ∂_k . The label-based representation allows to minimize ∂_k based on α -expansion (Delong et al., 2010):

$$U_{\partial}(\mathcal{X}) = \sum_{(i,j)\in\mathcal{N}_{\mathbf{T}}} \phi(x_i, x_j) , x_i, x_j \in \mathcal{X} , \qquad (8)$$

where $\phi(x_i, x_j)$ penalizes adjacent sites with different labels. The overall term is then:

$$U_M(\Phi) = \alpha U_{\mathcal{M}}(O) + \beta U_{\partial}(X) .$$
(9)

The strength of the regularization as well as of the model-complexity term is controlled by α and β respectively. If both values are set to zero, each vertex is simply assigned to the closest component of O, as it can be seen in Figure 2(a). The label costs eliminate redundant components, while the smoothness term enforces straight boundaries. The effect is shown in (b). If the values for α and β are increased further, the segmentation captures only the most important features, as it is seen in (c).

3.2 Likelihood

The likelihood term U_D quantifies how well the state Φ fits the observation S:

$$U_D(\Phi, \mathbf{S}) = \sum_{x_i \in \mathcal{X}, v_i \in \mathbf{S}} \langle u(v_i, \mathcal{O}), x_i \rangle.$$
(10)

 $u(v_i, O)$ is a vectorial function, where the *k*-th entry of *u* contains the cost for assigning $l^k \to x_i$, given the observation $v_i \in \mathbf{S}$. According to Equation 3, x_i is represented as a unit vector. The energy contribution of each site can then be written as a scalar product of $u(v_i, O)$ and x_i .

4 GEOMETRIC MODELS

A component \mathcal{M}_k is given by the underlying geometric primitive which is fully defined by its type *r* and its parameter vector θ_k . For all considered geometric models an orthogonal projection on \mathcal{M} can be computed for a given point: $\Pi_{\theta}(\cdot)$. The distance of a vertex v_i and a geometric primitive θ is defined as:

$$L^{e}(v_{i}, \boldsymbol{\theta}) = \|p_{i} - \Pi_{\boldsymbol{\theta}}(p_{i})\|, \qquad (11)$$

where a vertex is represented by its point p_i . $\Pi_{\theta}(q_i)$ is the projection of p_i on the surface of θ . A component \mathcal{M}_k has a likelihood which is assumed to be:

$$\phi(v_i, \mathcal{M}_k) = \begin{cases} L^e(v_i, \theta_k), & \text{if } p_i \in \mathcal{N}_{\varepsilon}(\mathcal{M}_k) \\ \infty, & \text{otherwise.} \end{cases}$$
(12)

 $\phi(v_i, \mathcal{M}_k)$ has infinite cost if v_i is not in the environment of the component $\mathcal{N}_{\varepsilon}(\mathcal{M}_k)$, which is defined later. The overall cost vector for assigning v to each model in O is:

$$u(v, \mathcal{O}) = \begin{pmatrix} (\phi(v, \mathcal{M}_1), \dots, \phi(v, \mathcal{M}_K))^T \\ c^{\mathcal{A}} \end{pmatrix} .$$
(13)

For a given vertex v, \mathcal{A} is the preferred label if no other component yields cost values lower than $c^{\mathcal{A}}$. In practice, $c^{\mathcal{A}}$ only needs to be set to a sufficiently large number. This encourages the MCMC process to insert models.

4.1 **Bounded Geometric Primitives**

Assume that a geometric primitive $\theta_k \in \mathcal{M}_k$ is given which fits a subset of the triangulated surface **S** according to Equation 12. Intuitively speaking, the likelihood for assigning a vertex $v \in \mathbf{S}$ to \mathcal{M}_k should depend on the distance of v to θ_k according to Equation 11. This is illustrated by the blue shading in Figure 3. However, geometric primitives are models with infinite extend and a boundary is not explicitly represented in L^e . This leads to many small clusters of vertices which are assigned to θ_k , as shown in Figure 3. We use the topology of **S** to recover the boundary $\partial \mathcal{M}_k$ on **S**. A function is defined, that indicates all vertices which are spatially close to the geometric primitive:

$$b(v, \theta_k) = \begin{cases} 1, & \text{if } L^e(v, \theta_k) \le e_{\varepsilon} \\ 0, & \text{otherwise.} \end{cases}$$
(14)

 $b(v, \theta_k)$ implies a partition of the surface into connected components:

$$\mathbf{S} = \{\mathbf{S}_{\mathcal{M}_k}^{b=1}, \mathbf{S}_{\mathcal{M}_k}^{b=0}\}.$$
 (15)

Moreover, $\hat{S}_{\mathcal{M}}^{b=1}$ is defined to be the largest connected component in $\mathbf{S}_{\mathcal{M}_k}^{b=1}$. The local environment of a model is then defined as the subset of segments in $\mathbf{S}_{\mathcal{M}_k}^{b=1}$ which are bigger than a predefined threshold $c_s |\hat{S}_{\mathcal{M}_k}^{b=1}|$:

$$\mathcal{N}_{\varepsilon}(\mathcal{M}_k) = \{ \forall S \in \mathbf{S}_{\mathcal{M}_k}^{b=1} : |S| \ge c_s |\hat{S}_{\mathcal{M}_k}^{b=1}| \}, \quad (16)$$

where c_s is the ratio that controls the minimum size of a connected component. $\{S \in \mathbf{S}_{\mathcal{M}_k}^{b=1}\}$ is the set of all connected subsets in $\mathbf{S}_{\mathcal{M}_k}^{b=1}$. $c_s = 0.1$ turns out to be a suitable value for all experiments.

5 OPTIMIZATION

In the following, an MCMC process is designed in order to approximate Φ^* in Equation 6. Consider the processing flow presented in Figure 4. RANSAC is used to fit a set of k initial shape candidates. The MCMC process samples the energy function by randomly selecting a move. The computation of the proposal probabilities is based on RANSAC. The new state is accepted according to a non-uniform distribution. The acceptance probability depends on the energy gain which is achieved by the current move. This simulates an annealing behavior which allows to escape from local minima. The approximation process is modeled by a Markov chain with the stationary probability of $p(\Phi|\mathbf{S})$. Assuming the current state of



Figure 3: Problem of finding the boundary of a model \mathcal{M}_k . Objects are described by a set of bounded surface segments. Geometric primitives such as planes, however, are of infinite extend. In this approach all vertices which are in the local environment of the model θ_k are decomposed into connected segments on **S**. The size of each segment relative to the largest connected component $\hat{S}_{\mathcal{M}_k}^{b=1}$ is used to robustly extract the boundary $\partial \mathcal{M}_k$.



Figure 4: Overview of the presented algorithm.

the Markov chain to be Φ_o , the probability of a move $\Phi_o \rightarrow \Phi_n$ is:

$$P(\Phi_o \rightarrow \Phi_n) = G(\Phi_o \rightarrow \Phi_n) A(\Phi_o \rightarrow \Phi_n) , \qquad (17)$$

where the move is proposed with the probability $G(\Phi_o \rightarrow \Phi_n)$ and accepted with probability $A(\Phi_o \rightarrow \Phi_n)$. Given a random proposal, the well-known Metropolis choice defines the acceptance probability to be (Suzuki, 1993):

$$A(\Phi_o \rightarrow \Phi_n) = \min\left(1, \frac{G(\Phi_n \rightarrow \Phi_o)}{G(\Phi_o \rightarrow \Phi_n)} \frac{p(\Phi_n | \mathbf{S})}{p(\Phi_o | \mathbf{S})}\right).$$
(18)

The first term is the ratio of the proposal probabilities G and describes the dynamic for selecting the move $\Phi_o \rightarrow \Phi_n$. If probabilities are expressed with respect to the energy function $U(\Phi, \mathbf{S})$, the second fraction in Equation 19 results in:

$$\frac{p(\Phi_n|\mathbf{S})}{p(\Phi_o|\mathbf{S})} = \exp\left(-\frac{U(\Phi_n, \mathbf{S}) - U(\Phi_o, \mathbf{S})}{\lambda T}\right), \quad (19)$$

where T is the temperature and λT simulates an annealing behavior. T is decreased over time to allow

random jumps in the beginning of the optimization procedure while the decisions become more deterministic with increasing number of iterations (Lafarge et al., 2013).



Figure 5: RANSAC-based computation of proposal probabilities. Consider the parameter space for a geometric model $(\theta^{(1)}, \theta^{(2)})$. The probability distribution $p(\mathcal{M}_j|S)$ for a geometric model given a triangulated surface segment (blue line) is difficult to evaluate. However, RANSAC is used to compute model candidates that can be seen as samples from the unknown distribution. Based on these samples, $p(\mathcal{M}_j|S)$ is approximated by a kernel-based density estimator.

Table 1: Overview of random moves and their corresponding proposal probabilities. 1-2 are diffusion moves and 3-7 are implementing discrete jumps.

	move	proposals $G(\Phi_o \rightarrow \Phi_n)$
1	opt. \mathcal{M}_k	$p(l^k)$
2	opt. X	1
3	split	$p(l^k)p(\mathcal{M}_{i,j} S_k)$
4	merge	$p(l^i)p(l^j l^i)p(\mathcal{M}_k S_{i,j})$
5	insert	$p(\mathcal{M}_n S_u)$
6	delete	$p(l^k)$
7	swap	$p(l^k)p(\mathcal{M}_n S_k)$

5.1 **Proposal Probabilities**

One of the most crucial aspects in the design of an MCMC process is the computation of the proposal probabilities. In the original paper by Metropolis et al. a uniform distribution is assumed. This means that, given a current state Φ_o , every state Φ_n within a bounding box of Φ_i is equally likely. However, this leads to a slow convergence since the ratio $p(\Phi_n)p(\Phi_o)^{-1}$ is close to zero for most of the proposed states (only a small subset yields high stationary probability). On the other hand, if randomized choices are replaced by deterministic decisions, the algorithm degenerates to a greedy method. A set of seven moves are defined for the stated problem (1-2 diffusion, 3-7 jumps). A move is selected randomly according to a uniform distribution. Each move requires a sequence of randomized decisions. The proposal probabilities for each move are given in Table 1.

5.1.1 Optimize Model (Diffusion)

A model \mathcal{M}_k is randomly chosen according to $p(l^k)$. The computation of $p(l^k)$ is explained in Section 5.1.6. Given a fixed partition $S_k \in \mathbf{S}$, the parameters θ_k are optimized based on the gradient $\frac{\partial U_D(\Phi, \mathbf{S})}{\partial \theta_k}$. In this work, a standard Levenberg-Marquardt algorithm is used to optimize the model parameters.

5.1.2 Optimize Labeling (Diffusion)

The second diffusion move is used to minimize the energy of the prior term in Equation 9. If the model O is assumed to be fixed, the energy function can be minimized by graph cuts. We perform one α -expansion iteration to compute the labeling X (Delong et al., 2010). The method allows to optimize spatially regularized classification problems while minimizing the number of used labels. A class l^k yields label costs as soon as at least one site in X is assigned with this label. This nicely allows to incorporate model complexity into the random-jumps framework.

5.1.3 Split and Merge Model (Jump)

Given a randomly selected class l^k , two new models $\mathcal{M}_{i,j}$ are proposed according to $p(\mathcal{M}_{i,j}|S_k)$. $p(\mathcal{M}_{i,j}|S_k)$ is computed based on a data-driven approach which is described in Section 5.2. The complementary move is to merge two regions which are randomly chosen according to $p(l^i)p(l^j|l^i)$. In order to estimate $p(l^j|l^i)$, we consider all segments which are adjacent to S_i . on **S**. Finally, RANSAC is used to compute $p(\mathcal{M}_k|S_{i,j})$.

5.1.4 Insert New and Delete Old Models (Jump)

All vertices which could not been assigned to existing models with a data cost below $c^{\mathcal{A}}$ are assigned to the default class \mathcal{A} . The set of all vertices $v \in S_{\mathcal{A}}$ is used to propose new models according to $p(\mathcal{M}_n|S_{\mathcal{A}})$. The delete move selects an existing model by $p(l^k)$ for removal.

5.1.5 Swap Model (Jump)

The swap move allows to switch between different types of models. This includes to select an existing model $p(l^k)$ and propose a new model $p(\mathcal{M}_n|S_k)$ using RANSAC. Note that the choice of the target model \mathcal{M}_n is independent of the former model \mathcal{M}_k .

5.1.6 Selecting a Candidate

A component k which is considered in a move is selected according to $p(l^k)$. Instead of assuming a uniform distribution, $p(l^k)$ could also be estimated with respect to the size of the underlying segment $|S_k|$. A larger segment is more likely to be described by two instead of only one model. Two smaller segments, however, might yield a lower energy if they are represented by one common model. However, this is only a heurisitic which does not turn out to be significantly better than a uniform distribution in our experiments.

5.2 Data-driven Proposals

Even if a very limited set of moves is considered, the likelihood of decreasing Equation 6 significantly depends on $p(\mathcal{M}|S)$. The Data-driven method proposed by Tu and Zhu uses bottom-up algorithms to estimate $p(\mathcal{M}|S)$ such that this is close to the actual probability density (Tu and Zhu, 2002). We want to adapt that method for our purpose. Methods such as RANSAC might not always compute a certain optimum but "proposals" of such methods are likely to decrease $p(\Phi|\mathbf{S})$. In this work, the algorithm of Schnabel et al. is modified for the domain of triangulated surfaces. The algorithm is referred to as Γ . Suppose that S is a subset of the triangulated surface $S \in \mathbf{S}$ and $\theta_{\Gamma} = \{e_{\varepsilon}, n_{\varepsilon}\}$ is a set of parameters which significantly influence the result. The e_{ε} -environment and the maximally allowed normal deviation n_{ε} are the inlier thresholds for the RANSAC procedure. The remaining parameters are set to the proposed standard values. Please refer to the corresponding paper for further details (Schnabel et al., 2007). The optimal set for θ_{Γ} is not known and a random sample $\theta_{\Gamma} \sim p(\theta_{\Gamma})$ is drawn to generate a set of shape candidates:

$$\{\mathcal{M}_k \in \mathbf{G}\} = \Gamma(S, \theta_{\Gamma}) . \tag{20}$$

Each candidate is associated with a weight ω_k which is a normalized sum of over all vertices belonging to the considered segment *S*:

$$\omega_k = \frac{1}{|S|} \sum_{\nu \in S} p(\nu | \mathcal{M}_k) , \qquad (21)$$

where $p(v|\mathcal{M}_k)$ is the likelihood of v belonging to \mathcal{M}_k . $p(v|\mathcal{M}_k)$ can also be interpreted as a vote of v for \mathcal{M}_k . In order to fulfill the requirements of a MCMC process, all possible configurations of \mathcal{M}_k should have $p(\mathcal{M}_k|S) > 0$:

$$p(\mathcal{M}_k|S) = \frac{1}{\sum_{i=0}^{|\mathbf{G}|} \omega_i} \sum_{|\mathbf{G}|} \omega_i K(\boldsymbol{\theta}_k - \boldsymbol{\theta}_i) , \qquad (22)$$

where K() is a kernel function such as Parzen window. $p(\mathcal{M}) > 0$ is required for the ergodicity of the Markov chain. The computation of the proposal probabilities is visualized in Figure 5.

Table 2: Segmentation score for the three considered algorithms, random jumps (b), variational shape approximation (c), and hierachical clustering (d). The number in each cell represents the normalized score of a segmentation. A vertex contributes a positive value to the overall score if correctly assigned to the most overlapping segment. The overall score is given as percent per total number of vertices.

Mdl.	Planar			Mixed		
Alg.	N1	N2	N3	N1	N2	N3
(b)	85.4	71.8	70.4	87.1	82.0	80.7
(c)	80.7	63.1	62.9	45.0	40.2	38.2
(d)	81.8	61.2	57.5	73.0	57.9	54.9



Figure 7: Visual comparison of the segmentation for two synthetic data sets. Row (1) shows the result for the planar model (N1) and row (2) for the mixed model (N2). Each row starts with the original, synthetic CAD model (a), followed by the resulting segmentation of the three considered algorithms, random jumps (b), variational shape approximation (c), and hierarchical clustering (d). In the table below, the number in each cell represents the normalized score of a segmentation. A vertex contributes a positive value to the overall score if correctly assigned to the most overlapping segment. The overall score is given as percent per total number of vertices.

6 EVALUATION

The presented algorithm is qualitatively and quantitatively compared to two related approaches. The greedy algorithm presented by Attene et al. performs segmentation by fitting spheres, cylinders and planes in a hierarchical clustering scheme (Attene et al., 2006). Driven by Lloyd's algorithm, the variational framework introduced by Cohen-Steiner et al. decomposes a mesh in approximately planar regions based on the normal vectors (Cohen-Steiner et al., 2004).

6.1 Qualitative Comparison

Both competitive methods require the number of clusters to be specified manually. In this experiment, it is



Figure 6: Result of the qualitative comparison for three different models, Clamp (1), Carburetor (2), and Train (3). The result of our method is shown in (a) and (b). (a) visualizes the synthesized model while the segmentation is shown in (b). The results for the segmentation algorithm based on hierarchical clustering can be found in (c), while the outcome of the variational shape approximation is shown in (d).

set according to the number of components extracted by our algorithm. For the entire qualitative evaluation, $\alpha = 1000, \beta = 20$ are used as prior parameters. The result of the qualitative evaluation can be seen in Figure 6. We apply the considered methods to three different models that have been acquired by the 3D reconstruction tool chain proposed by Jancosek et al. : Clamp (1), Carburetor (2), and Train (3) (Jancosek and Pajdla, 2011). The result of the presented method is shown in the first two rows. The synthesized object *O* is visualized in (a) and the semantic segmentation *X* is presented in (b). It can clearly be seen, that the O captures the most important geometric features of the model that allow to decompose **S** into smooth segments. The boundary curves (black) are free of perturbations and run along the principle curvature directions. Due to the prior term, O balances fitting quality and model complexity. The variational approach decomposes the surface according to the normal vectors which leads to approximately planar segments (c). If the model consists of planar segments which are separated by sharp feature lines, the variational approach delivers satisfying results, as it is shown in (1,c). However, the partition step is driven by region

Table 3: Overview of the label costs for all considered model types.

	type	$ \mathcal{M}_k $
1	plane	2
2	cylinder	6
3	sphere	8
4	torus	10
5	cone	10
7	default	1

Table 4: The intensity of the noise is given by the ratio of the maximal deviation and the global scale of the data. In this work, we apply fractal Brownian motion (fBm) and ridged multifractal noise (rmf).

	noise	r _o	
N1	fBm	.02	
N2	rmf	.04	
N3	fBm+rmf	.06	

growing which leads to scattered segment boundaries on less curved regions of the surface (bottom region in (1,c)). The result of the Lloyd algorithm also depends on the initialization of the clusters. Another problem are small, degenerated segments (black dots in (2,c)and (3,c)). This problem remains even when seeds are "teleported" as suggested in the paper of Cohen-Steiner et al. The hierarchical clustering algorithm is very sensitive to noise and produces very scattered boundary curves, especially on approximately flat regions of the surface. Due to its greedy nature, it fails to reasonably decompose the surface into spherical and cylindrical regions as it can be seen in (2,d).

The convergence behavior of the MCMC process for the Carburetor model is plotted in Figure 8 for ten optimizations. The graph shows the mean and the variance for the total energy (Equation 6) as well as for the three components, data costs (Equation 10), smoothness (Equation 8), and model complexity (Equation 7). For each run, the algorithm is initialized with four randomly detected primitives. The energy clearly converges to a minimum.

6.2 Quantitative Comparison

In a second experiment, we use two CAD models as ground truth and simulate the measurement process by surface reconstruction and random deformation. Using Boolean operations, the planar model is a union of boxes while the mixed model is composed of boxes, cylinders, and spheres. This yields a decomposition into faces that can be described by bounded geometric primitives. The CAD models are sampled at a very high resolution to obtain a set of reference vertices (> 1m). A standard surface reconstruction algo-



Figure 8: Convergence behavior for the clamp model. The plot shows mean and covariance of the energies for ten optimization processes. The total energy (black) is composed of data (green), label (blue), and smoothing costs (red). The algorithm was initialized with random settings for each run.

rithm is then applied to compute a test mesh which is used as input data for the algorithms. We apply fractal Brownian motion (fBm) and ridged multifractal noise (rmf) to randomly deform the test mesh at different intensity levels (Ebert, 2003). Please refer to Table 4. The intensity is given as the ratio r_{σ} of the maximal deviation and the global scale of the test mesh. After applying the algorithm, a score is computed by comparing the estimated labels with the ground truth for each vertex. Since we do not have explicit classes, we compute a unique mapping between estimated labels and ground truth data such that the overlap is maximized. A vertex contributes one to the overall score if correctly classified, otherwise zero. The normalized score is given in Table 2. 100 means all vertices in the test mesh are correctly classified, zero means no vertex is classified correctly. The score is the average of ten runs per algorithm. The number of clusters is manually specified to be the true number of clusters for both competing algorithms. The results are also presented for visual inspection in Figure 7. In (1), the segmentation is applied to the planar model (N1). The result for the mixed model that is deformed by rmf noise (N2) is shown in (2). The presented algorithm outperforms the other methods in all categories. However, both competitive algorithms also show a feasible performance at the lowest noise level. The performance of the hierarchical clustering algorithm drastically decreases with higher noise levels. The score for the variational algorithm mainly suffers from the sensitivity to bad cluster initializations.

7 CONCLUSION

A new and robust approach for the simultaneous classification and model fitting for triangulated surfaces is presented. The problem is formulated as a minimization of a global energy function. A MCMC framework allows to robustly compute a solution that explains the measured data. RANSAC turns out to be very efficient for computing new proposals that are likely to minimize the energy function. In future research, the algorithm will be extended by more advanced models, such as b-spline patches or quadrics surfaces.

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