# SIMPLIFICATION OF INDOOR SPACE FOOTPRINTS

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#### ABSTRACT

Simplification is one of the fundamental operations used in geoinformation science (GIS) to reduce size or representation complexity of geometric objects. Although different simplification methods can be applied depending on one's purpose, a simplification that many applications employ is designed to preserve their spatial properties after simplification. This article addresses one of the 2D simplification methods, especially working well on human-made structures such as 2D footprints of buildings and indoor spaces. The method simplifies polygons in an iterative manner. The simplification is segmentwise and takes account of intrusion, extrusion, offset, and corner portions of 2D structures preserving its dominant frame.

Keywords simplification · building · indoor space · footprint

#### **1** Introduction

Simplification is one of the methods used for generating data at different levels of detail (LoDs) from precise data. The more compact size of simplified data is desirable in a variety of data processing tasks including data transmission. Let P be a polygon, P' be a simplified polygon, and D(P, P') be the distance between P and P'. The computational geometry community distinguishes between two variants of curve simplification: While the min-# problem is to find P' with the minimum number of vertices such that  $D(P, P') \leq \varepsilon$ , the min- $\varepsilon$  problem is to find P' of at most k vertices such that D(P, P') is minimized. Depending on the constraints on the location of vertices of P', the problem can be categorized into (1) vertex-restricted, (2) curve-restricted, and (3) non-restricted simplification, see [1]. The Ramer-Douglas-Peucker (RDP) algorithm [2, 3] is widely used in practice and provides a vertex-restricted approximation to simplify 2D polylines and polygons. However, it does not reflect a specific form (see Figure 1b). Although Figures 1b and 1c are similar in terms of the number of segments, the figures demonstrate the RDP cannot preserve spatial features such as intrusion, extrusion, offset, and corners. In geographic information science (GIS), simplification is considered a type of generalization process [4]. These processes or operations consider not only metric constraints but also topological, semantic, and Gestalt constraints. In particular, Gestalt constraints are used to preserve the characteristics of spatial features such as a room.

This article presents a method that progressively simplifies 2D polygons by preserving their spatial properties in a iterative manner. The method was originally introduced by Kim and Li [5] to simplify 3D indoor spaces (e.g., rooms and hallways) that can be represented with the prism model [6, 7], which is an alternative 3D data model. As shown in Figure 1c, the simplification can be applied to footprints of complex buildings such as shopping malls or subway

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Figure 1: Example of simplification of a complex indoor space: (b) and (c) have 103 and 102 segments, respectively

stations. This article elaborates on the simplification of 2D footprints of [5] so that anyone can implement and apply it to their applications. In the next section, the detailed method is described.

# 2 Simplification of Polygons

Let P be a simple planar polygon without holes. P can be represented by the circular sequence of n line segments  $\langle s_0, \ldots, s_{n-1} \rangle$  describing the boundary of P in counterclockwise order, where  $p_i \in \mathbb{R}^2$  and each pair of vertices  $s_i = (p_i, p_{i+1})$  represents a line segment, for all  $0 \le i < n$ . The sequence of vertices of P is circular such that  $p_i = p_{(i \mod n)}$  for  $i \in \mathbb{Z}$ . Let  $\overline{s_i}$  be the length of  $s_i$ , and let  $\hat{s_i}$  be the internal angle between two consecutive segments  $s_i$  and  $s_{i+1}$ , i.e.,  $\hat{s_i} = \angle p_i p_{i+1} p_{i+2}$ . In the following, Q denotes a priority queue containing line segments  $s_i$  sorted by their length  $\overline{s_i}$ . Given a distance threshold  $\tau$ , an angle threshold  $\varepsilon$ , a collinearity threshold  $\delta$ , and a joining distance threshold  $\gamma$ , the simplification is designed to preserve the overall shape of a 2D polygon according to the following rules:

- *Rule-1*: The shorter the line segment, the smaller the effect on the overall shape of the geometry.
- *Rule-2*: Only segments longer than the tolerance  $\tau$  are considered to reflect the overall shape of the polygon.
- *Rule-3*: Consistent simplifications should be performed for feature types such as intrusion/extrusion, offset, and corner.



Figure 2: Flow chart of the algorithm of simplification of 2D polygon

Figure 2 depicts the flow of the simplification algorithm<sup>2</sup> that considers the rules, and Algorithm 1 shows the corresponding pseudo-code. In order to simplify shorter segments first in accordance with Rule-1, all segments of P are

<sup>&</sup>lt;sup>2</sup>The flow chart of [5] was modified according to the notation and context.

Algorithm 1: Simplification input :P// Polygon to simplify input : $\tau$ // Tolerance distance input : $\varepsilon$ // Tolerance angle input : $\delta$ // Angle threshold used to determine if consecutive segments are collinear input : $\gamma$ // Distance threshold used to determine whether to join neighboring segments 1  $Q \leftarrow \emptyset$ ; // Initialize a priority queue 2 foreach  $s \in P$  do  $| Q \leftarrow Q \cup s;$ // Insert all segments of polygon P into the queue Q3 4 while  $|Q| \ge 0$  and  $|P| \ge 3$  do  $s_i \leftarrow Q.dequeue();$ // Dequeue next segment 5  $\alpha \leftarrow |\widehat{s_{i-1}} - \widehat{s_i}|;$ // Angle difference between the angles entering and leaving  $s_i$ 6 7 if  $\pi - \delta < \widehat{s_i} < \pi + \delta$  then  $RemoveMiddlePoint(s_i)$ ; 8 // Merge two approximately collinear consecutive segments 9 else if  $\overline{s_i} \leq \tau$  then if  $0 \le \alpha \le \varepsilon$  then 10  $SegmentRegression(s_i);$ 11 else if  $\pi - \alpha \leq \varepsilon$  then 12  $TranslateSegment(s_i);$ 13 14 else 15  $q \leftarrow Intersect(s_{i-1}, s_{i+1});$  // Intersection of two lines obtained by extending  $s_{i-1}$  and  $s_{i+1}$ if  $dist(s_i, q) \leq \gamma$  then 16  $JoinSegment(s_i, q);$ 17 else if  $\overline{s_{i-1}} < \overline{s_{i+1}}$  then 18  $RemoveMiddlePoint(s_{i-1});$ 19 20 else  $RemoveMiddlePoint(s_i);$ 21 22 return P; // Return the simplified polygon

sorted by length and inserted into a priority queue Q (lines 1-3). Simplification is repeatedly carried out on the next segment in the queue until the queue is empty (line 4). The shortest line segment is dequeued from the queue as the current segment  $s_i$  (line 5). To merge collinear segments, it is checked whether  $\pi - \delta < \hat{s}_i < \pi + \delta$ , where  $\delta > 0$  is a small angle threshold that is used to evaluate collinearity (line 7). If so, the middle point between  $s_i$  and  $s_{i+1}$  is removed so that a new line segment between the two end points is created (line 8; see also Algorithm 2). If the length of current segment is less than or equal to  $\tau$ , simplification is performed (lines 9-21). In consideration of Rule-3, one of four methods is chosen depending on the angle difference  $\alpha = |\hat{s}_{i-1} - \hat{s}_i| \leq \pi$  (lines 10-21). Figures 3-5 illustrate this process.

- If 0 ≤ α ≤ ε, regress two segments s<sub>i-1</sub> and s<sub>i+1</sub>, considering their lengths and tangents as shown in Figures 3(c),
  (d) and (e). The detailed process is outlined in Algorithm 3.
- If π − α ≤ ε, this is considered an intrusion/extrusion, and the current segment is translated in order to remove the intrusion/extrusion as shown in Figure 4 (see Algorithm 4).
- Otherwise, it is checked whether the two segments can be joined by extending  $s_{i-1}$  and  $s_{i+1}$  until  $s_{i-1}$  and  $s_{i+1}$  intersect (see Figure 6). Let q be the intersection point, dist a distance function, and  $\gamma$  the distance threshold.
  - If  $dist(s_i, q) \leq \gamma$  as shown in Figures 6(a)-(d), then remove  $s_i$  and join  $s_{i-1}$  and  $s_{i+1}$  (see Algorithm 5 and Figure 6(e)-(h)).
  - Otherwise, do not join the segments because the extended part of the segments is too long as shown in Figures 7(a)-(e). Instead, remove the middle point between  $s_i$  and  $s_{i-1}$  (or  $s_{i+1}$ ) (see Figure 7(f)-(k)).

Pseudocode for the functions invoked in Algorithm 1 is described as Algorithms 2, 3, 4, and 5. Assume that P and Q defined in Algorithm 1 can be accessed from Algorithms 2, 3, 4, and 5. Python implementation and examples can be found at the git repository (https://github.com/joonseok-kim/simplification).

Algorithm 2: RemoveMiddlePoint

#### input : $s_k$

1  $Q \leftarrow Q \setminus s_{k+1}$ ;

- 2  $s' \leftarrow (p_k, p_{k+2});$
- $P \leftarrow P \setminus (s_k \cup s_{k+1});$
- 4  $Q \leftarrow Q \cup s'$ ;

# Algorithm 3: SegmentRegression input : $s_k$

- $\mathbf{1} \ Q \leftarrow Q \setminus (s_{k-1} \cup s_{k+1});$ 2  $r \leftarrow \overline{s_{k-1}}/(\overline{s_{k-1}} + \overline{s_{k+1}});$  $p \leftarrow s_k.PointAlong(r);$ 4  $\theta \leftarrow \tan(\widehat{s_{k-1}} \cdot r + \widehat{s_{k+1}} \cdot (1-r));$ **5**  $q_1 \leftarrow Projection(s_{k-2}, p, \theta)$ ; 6  $q_2 \leftarrow Projection(s_{k+2}, p, \theta)$ ; 7  $s_k \leftarrow (q_1, q_2);$ **8**  $s_{k-1} \leftarrow (p_{k-1}, q_1)$ ; 9  $s_{k+1} \leftarrow (q_2, p_{k+2});$
- 10  $Q \leftarrow Q \cup s_{k-1} \cup s_k \cup s_{k+1}$ ;

Algorithm 4: TranslateSegment

#### // Segment

// Remove  $s_{k+1}$  from the queue Q// Create a new segment  $(p_k, p_{k+2})$  by merging  $s_k$  and  $s_{k+1}$ // Remove existing  $s_k$  and  $s_{k+1}$  from P // Insert the new segment into the queue Q

#### // Segment

// Ratio to be used as a weight. // A point p on  $s_k$  such that p is at distance  $\overline{s_k} \cdot r$  from  $p_k$ . // The slope of a regression line for  $s_{k-1}$  and  $s_{k+1}$ . // Intersection of  $s_{k-2}$  with the line through p with slope  $\theta$ . // Intersection of  $s_{k+2}$  with the line through p with slope  $\theta$ . // A regression line segment for  $s_{k-1}$  and  $s_{k+1}$ . // Update  $s_{k-1}$ // Update  $s_{k+1}$ // Add the new three segments into the queue

	input : $s_k$	// Segment
1	$Q \leftarrow Q \setminus (s_{k-1} \cup s_{k+1});$	// Remove $s_{k-1}$ and $s_{k+1}$ from the queue $Q$
2	if $\overline{s_{k-1}} < \overline{s_{k+1}}$ then	
3	$p' \leftarrow \overrightarrow{p_{k+1}} - \overrightarrow{s_{k-1}};$	// Translate vertex $p_{k+1}$ by vector $s_{k-1}$
4	$s_k \leftarrow (p_{k-1}, p');$	// Update $s_k$
5	$s_{k+1} \leftarrow (p', p_{k+2});$	// Update $s_{k+1}$
6	$P \leftarrow P \setminus s_{k-1};$	// Remove existing $s_{k-1}$
7	$Q \leftarrow Q \cup (s_k \cup s_{k+1});$	// Add $s_k$ and $s_{k+1}$ into $Q$
8	else if $\overline{s_{k-1}} > \overline{s_{k+1}}$ then	
9	$p' \leftarrow \overrightarrow{p_k} - \overrightarrow{s_{k+1}};$	// Translate vertex $p_k$ by vector $s_{k+1}$
10	$s_{k-1} \leftarrow (p_{k-1}, p');$	// Update $s_{k-1}$
11	$s_k \leftarrow (p', p_{k+2});$	// Update $s_k$
12	$P \leftarrow P \setminus s_{k+1};$	// Remove existing $s_{k+1}$
13	$Q \leftarrow Q \cup (s_{k-1} \cup s_k);$	// Add $s_{k-1}$ and $s_k$ into $Q$
14	else	
15	$s_k \leftarrow (p_{k-1}, p_{k+2});$	// Update $s_k$
16	$P \leftarrow P \setminus (s_{k-1} \cup s_{k+1});$	// Remove existing $s_{k-1}$ and $s_{k+1}$
17	$\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ $	// Add $s_k$ into $Q$

Algorithm 5: JoinSegment	
input : $s_k$	// Segment
input :q	// Intersection point
$ Q \leftarrow Q \setminus (s_{k-1} \cup s_{k+1}); $	// Remove $s_{k-1}$ and $s_{k+1}$ from the queue $Q$
$2 \ s_{k-1} \leftarrow (p_{k-1}, q) ;$	// Update $s_{k-1}$
<b>3</b> $s_{k+1} \leftarrow (q, p_{k+2});$	// Update $s_{k+1}$
4 $P \leftarrow P \setminus s_k$ ;	// Remove existing $s_k$
$ 5 \ Q \leftarrow Q \cup (s_{k-1} \cup s_{k+1}) ; $	// Add $s_{k+1}$ and $s_{k+1}$ into $Q$



Figure 3: Removing points and merging into regression segment



Figure 4: Translating current segment



Figure 5: Removing of points and joining segments



Figure 6: Joining segments  $(\gamma = \overline{s_i})$ 



Figure 7: Removing middle points ( $\gamma = \overline{s_i}$ )

# **3** Experiments

IndoorGML data for the Lotte World Mall<sup>3</sup> (LWM), see Figure 8, one of the most complex shopping malls in Seoul, is used to conduct a comparative experiment on the performance of the introduced simplification. IndoorGML is one of OGC standards to provide a standard framework of semantic, topological, and geometric models for indoor spatial information [8]. The dataset for the experiment is compatible with CityGML LoD4 [9] so that data can be visualzed via any CityGML viewer. Figures 9 and 10 show the visual and quantitative results of simplification of one large and complex corridor for varying  $\tau$ , given  $\varepsilon = \pi/36$ ,  $\delta = \pi/180$ ,  $\gamma = \overline{s_i}$ . As  $\tau$  increases, gradual simplification of shorter segments is observed, in particular at extremities, dominant features such as intrusions, extrusions, offsets, and corners are preserved.

While Figure 10 focuses on simplification results for one polygon, Figure 11 shows the comparison between the original LWM data, the indoor simplification, and RDP for all spaces on a floor. Note that a needle-shaped polygon will vanish after simplification if the length of its width is less than  $\tau$ .



Figure 8: Lotte World Mall dataset

Figure 9: The number of segments left after simplification of the corridor shown in Figure 10

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<sup>&</sup>lt;sup>3</sup>IndoorGML data (core module) for Lotte World Mall (IndoorGML 1.0.3), http://www.indoorgml.net/resources/



Figure 10: Results of simplification for varying  $\tau$ 



Figure 11: Comparison between the original data, the indoor simplification ( $\tau$ =2), and RDP (tolerance=1.2)

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