

SOLVING THE T-JOINT PROBLEM IN RECONSTRUCTING 2-D OBJECTS

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Abstract: This paper describes a solution to the T-joint problem in matching 2D fragments of an object. Matching fragments of an object is useful for solving puzzles or reassembling archaeological fragments. Many factors, such as the number of pieces and the complex shapes of pieces make this a difficult problem. Various approaches to this problem exist. This paper presents an approach to solving the T-joint problem, which comes up in assembling fragments. The work described in this paper starts with a 2D object that should be easy to extend to 3D problems.

1 INTRODUCTION

The assembly of fragments of an object using computer software has significant usages in the real world. Applications include reconstructing torn documents or fragmented pottery and artefacts while excavating ruins of ancient civilizations. It can be a challenging and formidable task reconstructing an entire object. A manual approach is time consuming, especially for a large number of fragments and requires direct contact with objects, which increases the chance to damage those artefacts. In contrast, an automated approach could efficiently reconstruct a large number of pieces. In addition, the trace of the reassembly can be kept in an electronic medium for study.

An additional problem is that in naturally occurring fragmentation of objects, fragments do not commonly match at pairs of corners. Most objects fragment in a way that forms triple junctions, such as T-joints. McBride and Kimia (2003) reported that in samples of broken ceramic tiles, T-joints ranged from 70-89% of the junctions, while other triple junctions ranged from 6-9%, and non-triple junctions ranged from 5-20%. Figure 1 shows a T-joint.

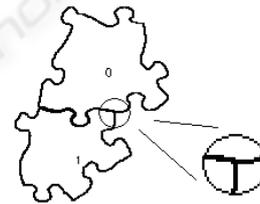


Figure 1: Two puzzle pieces showing T-joint match.

The assembly of the fragments of 3D objects is complicated. This work focuses on 2D objects, rather than 3D. In this way, all the drawbacks in 2D approaches are resolved before leaping to the 3D world to simplify the computational complexity. In section 2, previous work on matching objects is discussed. Section 3 introduces a method to perform 2D curve matching for pieces *with T-joints*. Section 4 describes the results of processing puzzle pieces with the method introduced in section 3. Section 5 presents conclusions and future issues to explore.

2 HISTORY REVIEW

Numerous previous works attack matching issues in reconstructing fragmented objects. Most of them compare the boundaries of the objects, represented

by some sort of spline curve (Lee, Clark, & Araman, 2003; Krebs, Korn & Wahl, 1997) or polygonal approximation. The algorithms split curves at boundary points and then match sub curves (Freeman & Garder, 1964; Kong & Kimia, 2001; McBride & Kimia, 2003; Radack & Badler, 1982; Stringfellow, Simpson, Bui, Peng, & Hood, 2008; Weiss-Cohen, Halevi, 2005). Many approaches perform local shape analysis first, which results in ambiguous matches and require backtracking to resolve the mismatches (Freeman & Garder, 1964; Kong & Kimia, 2001; Radack & Badler, 1982; Stringfellow, Simpson, Bui, Peng, & Hood, 2008).

Finding accurate matches is more often than not very time consuming. Curve fitting methods often require a least squares fit or similar process to determine the associated error of each fitted curve. As the number of curves that require processing increases, the complexity of each fit becomes more of an issue. This is the situation that is encountered in the reconstruction of 2D fragmented objects. Each collection of N fragments, where each fragment has on the average M distinct edges, requires $O((MN)^2)$ curve fits. Of course, problem specific information can be used to reduce this number.

The approaches used by researchers vary based on the characteristics of the fragmented objects. These characteristics concern orientation of pieces, whether there are missing pieces, whether the exterior boundary is known beforehand (such as a rectangular grid), whether there is a unique solution, and what types of junctions between pieces are allowed (Kleber, 2009). Some approaches consider image features of the pieces, such as color and texture (Nielsen, Drewsen, & Hansen, 2008) and others consider shape of pieces (Da Gama Leitao & Stolfi, 2002; Freeman & Garder, 1964; Goldberg, Malon, & Bern, 2002; Horst & Beichl, 1996; Krebs et al., 1997; Kong & Kimia, 2001; Lee et al., 2003; McBride & Kimia, 2003; Radack & Badler, 1982; Stringfellow et al., 2008; Zhu, Zhou & Hu, 2008), while (Weiss-Cohen & Halevi, 2005; Yao & Shao, 2003) consider both.

Many of the approaches have problems. Freeman and Garder (1964) repeatedly search all pieces and the best matches are merged to form new pieces until only one piece is left. Insufficient constraints result in mismatches and require backtracking. The algorithm by Goldberg et al. (2002) is efficient, but not fully automated and only works on pieces with four corners. Yao and Shao (2003) present no method to solve for triple junctions.

Kong and Kimia (2001) McBride and Kimia (2003) and Zhu et al. (2008) present two-step approaches to solving reconstruction of fragmented objects. In both, the first step finds likely candidate pairs of pieces by computing affinity measures using polygonal approximation of pieces. In Kong and Kimia (2001) McBride and Kimia (2003) triples that arise from generic junctions (Y- and T-joints) are formed from this rank-ordered list of the top-ten pairings. Their second step compares these candidate pieces at a finer level. If a match, the pieces are merged and removed from the piece list and the merged piece is inserted. The results show some mismatches, probably due to issues in the second matching step. In Zhu et al. (2008) the second step uses a confidence number assigned to each pair and then maximizes the consistency until the confidence reaches one. This creates the advantage of overall checking of joins to decide on matches or false positives. However, it only results 20-30 percent matches.

Stringfellow et al. (2008) present a method that works on pieces with discrete closed boundaries, represented by points (not curves or polygonal approximations). It does not require pieces to fit a grid. It does not match pieces with T-joints, but pieces may "join" a puzzle, if an adjacent edge without a T-joint is matched to another different piece. It does require a very small amount of user interactions to verify automated results.

3 METHOD

This paper builds on the work of Stringfellow et al. (2008) to solve the T-joint problem. The approach, which is now a 2-step matching process is briefly described.

First, boundary points representing outlines are extracted from scanned images of puzzle pieces. Then the convex corner points are detected using a non-parameterized algorithm (Staples & Hood, 2008). In order to match curve segments of pieces, a fitness function is introduced to determine whether two curve segments are candidates for matching. Figure 2 shows the curve segment is divided into four small segments. It also shows the start points and end points for the four sub-segments. The middle point of a segment is chosen to divide the segment into sub-segments. The distance lengths of the sub-segments are denoted $d_1 - d_4$, respectively. The four sub-segments form three angles $\Theta_1, - \Theta_3$. The distance of a segment is computed in two ways: the number of pixels between *start* and *end* points

denoted as PD and the straight line distance between *start* and *end* points and is denoted as D. The fitness value, $F(S_i, S_j)$, is calculated using absolute or relative differences between the nine criteria. The smaller the fitness value, the better chance the two segments are actually a match. If the fitness is too large, then the pair is rejected as a candidate pair.

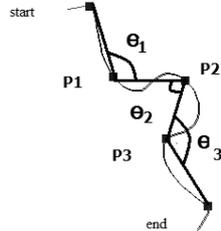


Figure 2: Break down of a curve segment in order to calculate its fitness value.

Other tests are applied to determine if the mid-points and quarter points of segment S_i and segment S_j are close to each other after a rotation and a translation are performed. If they are too far away from each other, then the fitness $F(S_i, S_j)$ is set to a large value (not a candidate for matching pair.) Figure 2 shows two original segments and the segments after a rotation and a translation.

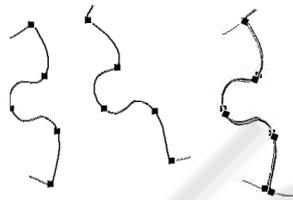


Figure 3: S_i and S_j and their sub-segments before and after a rotation and translation.

Next, the algorithm (shown in Figure 4) uses a local approach to match curves. The algorithm selects a puzzle piece with four or more corners. If there is no piece with four or more corners, the algorithm selects just any piece. However, the more corners a piece has, the better the matching pool generated. Next, pieces with the matches for the piece's curve segments are found and the adjacent pieces are pushed onto a stack. The next piece out of the stack is popped and the best match for its curve segments is found. The process is repeated until the stack is empty.

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push piece [i] in the stack
while the stack is not empty
    pop (current_piece)
    for every segment of current_piece
        find the best fit segment
        if there is a match and the adjacent piece
            with the best fit is not in the stack already
                move adjacent piece to new position
                push the adjacent piece onto the stack
    
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Figure 4: Regular curve matching algorithm (Stringfellow et al., 2008).

After regular curve matching, T-joint matching is performed. The unmatched segments are sorted (by length) into a list. The T-joint algorithm (shown in Figure 5) starts with a unmatched pair of segments selected from the sorted segment list. It calculates a sub-segment in the longer segment equal in length to the shorter segment starting at the *start* point of the longer segment and applies the matching criteria to see if they are a match. If not, it calculates a sub-segment in the long segment starting at the *finish* point and determines if they are a match. If a match is found (from either end point), it subdivides the long segment into a matched (sub) segment and a leftover (sub) segment. The leftover segment is inserted into the sorted segment list for future consideration.

- 1: Get distance of short segment.
- 2: Find sub-segment in long segment that is equal length to short segment starting at the *start* point of long segment.
- 3: The new sub-segment is separated into 4 (equal) segments: calculate the midpoint and quarter points and other matching criteria.
- 4: Compare the sub-segment to the short segment to see if they match using the criteria.
- 5: If they match by a certain fitness value, then compute the left over part of the long segment.
- 6: If segments do not match, create a sub-segment from the long segment with same distance as short segment starting at *finish* point of long segment.
- 7: Perform Steps 3-5 for this sub-segment.
- 8: If either of the sub-segments of the long segment match the short segment, compute the left over segment of the long segment.
9. The matched sub-segment is removed from the segment list and the leftover segment is inserted in the sorted segment list.

Figure 5: T-joint matching algorithm.

The criteria for determining if segments are matches in regular matching and T-joint matching are essentially the same, except segment length criteria is disregarded in the overall fitness value for T-joint matching (since the segments compared are made to be the same length). The tolerance values for the criteria can be made different for T-joint matching, but with these puzzles they were left the same. Only the overall fitness value was decreased in the T-joint matching (to compensate for the segment length criteria).

Overall, regular and T-joint matching is $O(S^2)$, where S is the number of segments in all the pieces. Some efficiency is gained in regular matching, by only comparing segments to segments of other pieces, as long as those segments are within its neighbourhood in the sorted list. (A neighbourhood is size 10.) In addition, if the differences between two segments in one of the nine criteria are too large, then no other comparison calculations are performed. When a pair of segments is matched, a piece gets attached to the puzzle, and *that* piece becomes the current piece for further matching. If a piece has none of its segments match, the algorithm goes back to a previous piece and checks its remaining unmatched segments. It is possible to join an N -piece puzzle with $O(N)$ segment matches. The worst case scenario is that no pieces match, and $O(S^2)$ segments are compared. After regular matching is performed, the unmatched segments are stored in a sorted unmatched segment list – this takes $O(S)$. The T-joint algorithm, worst case, would need to compare every unmatched segment to every other unmatched segment twice – taking $O(S^2)$, but if regular matching found most of the matches, there would be fewer unmatched segments in the sorted list to compare.

4 RESULTS

The application was developed using C# within the Microsoft Visual Studio.Net 2005 platform. Puzzle pieces were scanned in using a HP ScanJet 5200C scanner, several pieces at a time. RGB format is transformed to greyscale and then the images are converted to a binary format, so all pixels are either black or white. Boundaries are extracted and corner points are detected, then regular matching is performed. Figure 6 shows a puzzle put together after regular matching is performed. The two pieces not joined only have segments with T-joints. There are a few pieces joined in the puzzle that have

segments with T-joints, but they are matched to other pieces using segments without T-joints.

Figure 7 shows the results after the T-joint matching is performed. The bottom segment of the shaded piece is matched to the joined puzzle. (There is a bit of round-off error in the translation and rotation of pieces.)

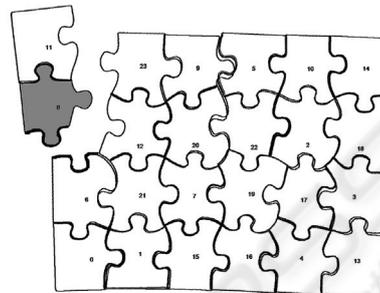


Figure 6: Puzzle after regular matching performed (Stringfellow et al., 2008).

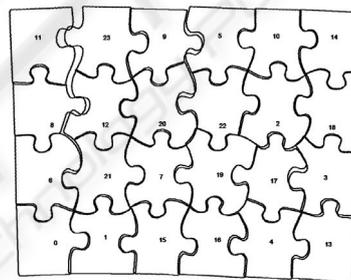


Figure 7: Final result after T-joint matching.

Figures 8 – 10 show a puzzle with *many* T-joints. As can be seen in Figure 11, only 2 pairs of pieces are matched using regular matching. After the T-joint matching algorithm is applied, every piece is joined to the puzzle.

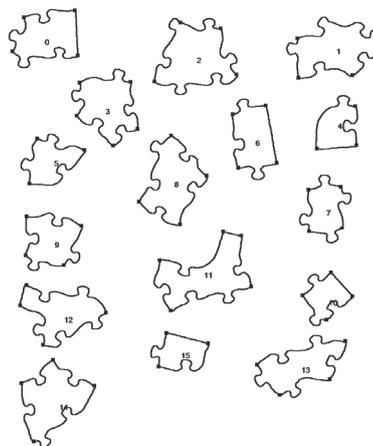


Figure 8: Puzzle 2 with *many* T-joints.

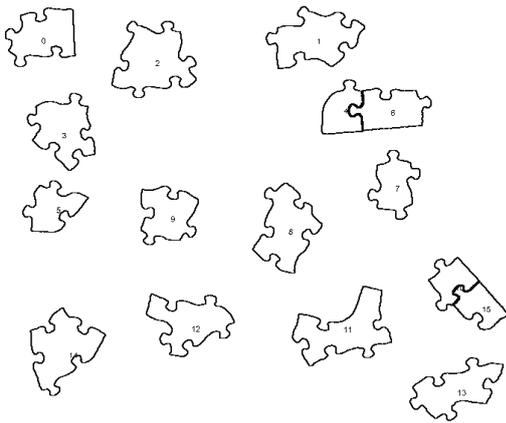


Figure 9: Puzzle 2 after regular matching.

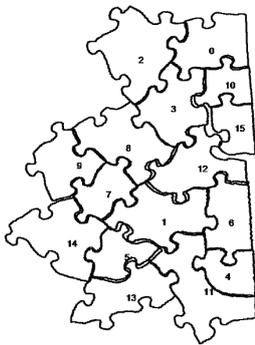


Figure 10: Puzzle 2 after T-joint matching performed.

Figures 11 and 12 show a puzzle that has pieces with smoother, longer curves and many T-joints. All pieces are matched by the T-joint matching algorithm. Note that this puzzle has joints where two pieces match, but do not have either a start or end point in common (designated as a Type 3 junction in [8]).

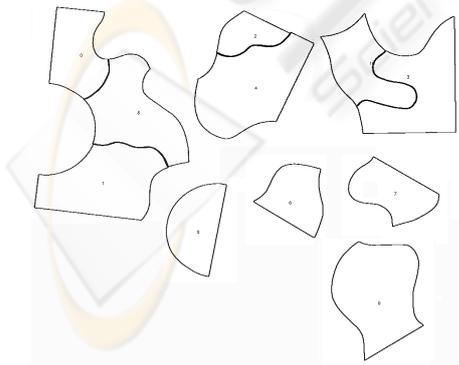


Figure 11: Puzzle 3 after regular matching is performed.

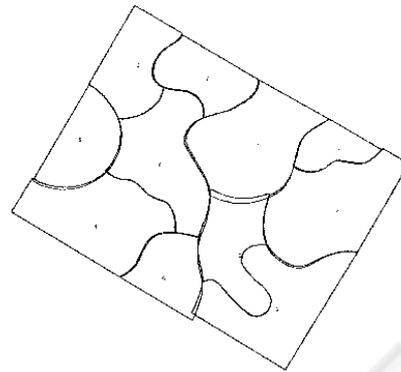


Figure 12: Puzzle 3 after T-joint matching performed.

Figures 7, 10 and 12 show that the algorithm in Figure 5 is successful in putting the remaining puzzle pieces with T-joints together. This algorithm also solves the problem of type 3 junctions. These joints get resolved when the piece's segment is divided into a matched part (to another piece) and a leftover part that is then inserted into the segment list to be matched with a future piece. The type 3 joints reduce to T-joints. In this way, the semi-circular piece in puzzle 3 is matched to three pieces.

The threshold values for some of the fitness criteria for a match were made stricter (that is, the difference between two segments' criteria had to be smaller to be considered a match). In puzzles 1 and 2, the criteria had to be relaxed in order to get them completed. There is a trade-off between using stricter criteria and the number of correct matches. When the criteria was made more strict, fewer matches were made (although not much fewer), but it did not give fewer false matches, except in puzzle 2. In puzzle 3, a large number of the false T-joint matches were due to the semi-circular piece, which tended to match a lot of other arcs in other pieces' segments.

False T-joint matches are more likely than false regular matches, due to the fact that two criteria are eliminated. (The T-joint matching algorithm makes pixel segment length and actual segment lengths the same.) Making the remaining criteria tighter, may result in fewer false T-joint matches, but it may also reduce true T-joint matches.

5 CONCLUSIONS AND FUTURE WORK

This paper proposes a method and its implementation for semi-automatic reconstruction of 2D jigsaw puzzles that have pieces with T-joints.

Applied to a 24 piece puzzle, the implemented application is able to put all 24 pieces together. Puzzles 2 and 3 are also entirely put together.

Efficiency in this approach is achieved in several ways. First only pairs of segments nearby in the sorted list of segments are compared during regular matching, and only if they meet the other distance and angle criteria do they undergo more calculations. T-joint matching is only performed on unmatched segments after all regular matching is done. Type 3 junctions reduce to T-joints after matched segments are subdivided into matched and leftover segments.

The suggested approach is considered a successful and reliable one. This approach reconstructed all the pieces of a given set of objects. Restricting the fitness criteria may reduce the number of matches and increase the number of false matches, but it still results in successful matching. Future work needs to be done to base the fitness criteria on properties of the fragments, such as their size.

Future research will also consider matching combined pieces. By applying this approach, there exists a potential to resolve any few remaining unmatched pieces. Applying this approach to the reconstruction of ancient artefacts would result in considerably less pieces having to be touched over and over again.

Finally, for the jump to 3D, the work will likely follow a similar technique to Krebs et al. (1997) using 2D slices of 3D objects, although rather than using splines the approach will consider the points in a cloud to represent the edges or boundaries of the object.

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