

Fig. 2.

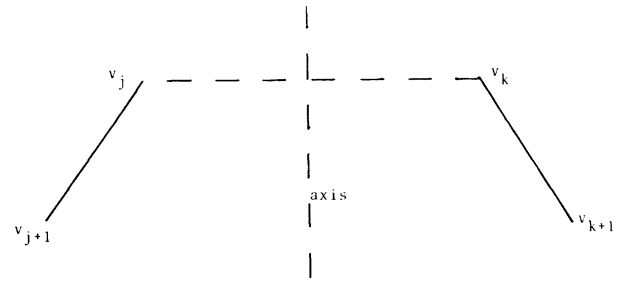


Fig. 4.

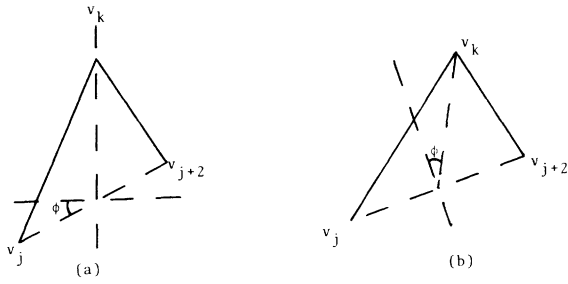


Fig. 3.

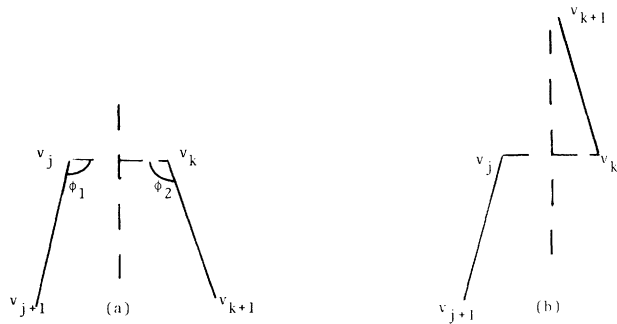


Fig. 5.

*Case 1* (all numbers modulo  $n$ ): Sides  $\mathcal{S}_j$  and  $\mathcal{S}_k$  meet at a vertex. Suppose  $k = j + 1$ . Then the two sides meet at vertex  $v_{j+1}$ . A likely candidate for the axis of symmetry  $a$  is either

- 1) the angle bisector of vertex  $v_{j+1}$ , or
- 2) the perpendicular bisector of the line joining vertices  $v_j$  and  $v_{j+2}$  (see Fig. 2(a), (b)).

A good measure of the cost of the symmetry  $c(a)$  in this case is just the difference in the lengths of sides  $\mathcal{S}_j$  and  $\mathcal{S}_k$ . However, it would be better for the measure to be scale invariant; the cost is related to angle  $\phi$  in Fig. 3(a), (b). If  $\phi$  is  $0^\circ$ , then the symmetry is perfect. If  $\phi > 0^\circ$ , then the symmetry deviates from the ideal.

*Case 2.* Sides  $\mathcal{S}_j$  and  $\mathcal{S}_k$  do not meet. In this second case, there are four possible axes, one for each pairing of an end of  $\mathcal{S}_j$  with an end of  $\mathcal{S}_k$ . If the pairing is  $(v_j, v_k)$  then the hypothesized axis  $a$  is the perpendicular bisector of the line joining  $v_j$  and  $v_k$  (see Fig. 4). The measure of the cost of the symmetry is (see Fig. 5(a), (b)):

- a)  $|\phi_1 - \phi_2|$  if  $v_{j+1}$  and  $v_{k+1}$  are in the same half plane with respect to the line  $v_j v_k$ ;
- b) otherwise some high number.

In practice, a threshold is set on these cost measures, and any proposed symmetry with higher cost is discarded.

The next stage of the process is to cluster the local axes of symmetry into aggregates. If each axis,  $a_i$  is represented by an equation of the form

$$\rho_i = -x \sin \theta_i + y \cos \theta_i$$

then clusters of local axes correspond to clusters of points in  $(\rho, \theta)$  space [7]. The clustering algorithm designed here for grouping axes works in the following way:

Define the span of an axis,  $\text{span}(a_i)$ , as the set of sides that are symmetric with respect to that axis. Define the span fraction  $s(a_i)$  as the ratio of the length of  $\text{span}(a_i)$  to the perimeter of the polygon. Choose a local axis having minimal cost and maximal span as a cluster center. Call this cluster  $A_i$ . Let

$$P_i = \sum_{a_i \in A_i} \rho_i / |A_i|$$

and

$$\theta_i = \sum_{a_i \in A_i} \theta_i / |A_i|.$$

Now find a local axis  $a_j$  having minimal cost and maximal span, satisfying

- a)  $|\theta_i - \theta_j| < t_\theta$
- b)  $|\rho_i - \rho_j| < t_\rho$

where  $t_\theta$  and  $t_\rho$  are dynamically defined thresholds that depend on  $A_i$  (e.g.,  $t_\rho$  may be the standard deviation of the  $\rho_i$ 's). If no such  $a_j$  exists, then the cluster is complete, and a new cluster can be constructed from the remaining local axes. The process continues until all local axes belong to some cluster.

Two examples of the local axes determination and clustering procedures are shown in Fig. 6. Fig. 6(a), (b) shows the polygons, with the sides and vertices labeled. Fig. 6(c), (e) are tables of the local axes detected in the polygons. (Note in Fig. 6(e) that many of the local axes do not correspond to any axis that would ordinarily be perceived by a human observer.) Fig. 6(d), (f) list the clusters constructed from the local axes.

These clusters form the basic units with which subsequent processes will work. These processes will construct hierarchical representations of the symmetries of the polygon. An example of the types of representation that will be constructed should clarify the later discussion of how the processes work. Consider Fig. 6(b), and the list of clusters for that polygon in Fig. 6(f). Fig. 7 shows one possible description of the symmetries of Fig. 6(b). The level 1 description indicates that the vertical axis comprises the highest level (level 1) of the hierarchy. It splits the figure in "half" and enables us to concentrate on finding representations of the symmetries in only half the figure. Next, level 2 contains a horizontal axis that spans all of the sides in half of the level 1 representation. Similarly, the level 3 description contains just the diagonal axis that spans all of the level 2 sides.

This example indicates that the first step in building a hierarchical symmetry representation is to choose an axis for level 1. A reasonable choice is one that "accounts for" the largest

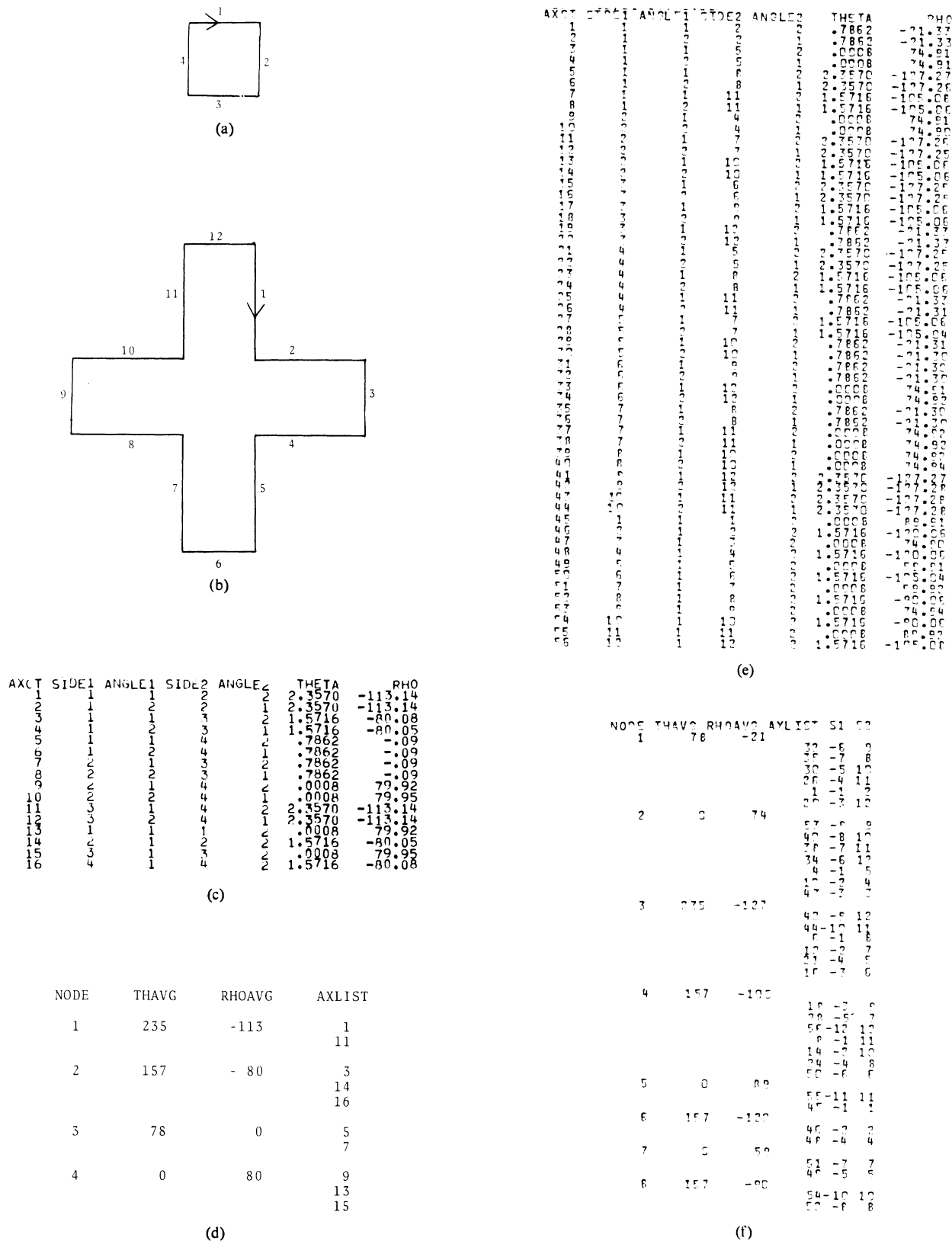


Fig. 6. (c) List of axes for (a). (d) Clusters for (a). (e) Axes for (b). (f) Clusters for (b). Note: For each side, angle 1 is tail of arrow and angle 2 is head (for (c)-(f)).

amount of the polygon in the best way. The particular figure of merit that was adopted here was to find the cluster  $A_i$  such that  $\sum_{a_i \in A_i} s(a_i)$  is maximal, while  $\sum_{a_i \in A_i} c(a_i)$  is minimal; i.e., the cluster that accounts for the greatest part of the perimeter at minimal cost. To avoid building descriptions of figures that are

simply not symmetric, a threshold is placed on the  $\sum_{a_i \in A_i} s(a_i)$  term. If no cluster accounts for an appreciable percentage of the polygon, then no description is generated. On the other hand, there may be several equally good clusters (as is the case for both polygons in Fig. 6). In this case, many hierarchies of symmetry

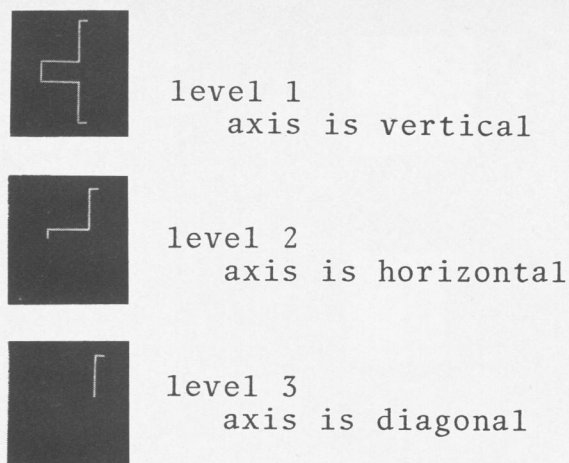


Fig. 7.

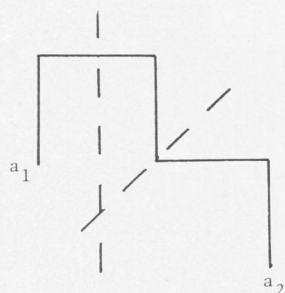


Fig. 8.

are possible. At times it may be necessary to create them all; or, there may be *a priori* biases concerning the "best" level 1 axis (e.g., the longest, the one closest to vertical). These biases will tend to affect the control of the description building mechanism, but not their underlying computations (see the recent papers by Marr [4], [5] on the role of context in low level vision).

The level 1, or major, axis splits the polygon into two sets of vertices—those in the positive half plane determined by that axis, and those in the negative half plane. The next phase of the process involves finding a set of clusters that account for the symmetries between the sides of the polygon that have at least one vertex in the positive half plane of the major axis. In order to do this we made use of two definitions:

**Definition 1:** Two clusters of axes  $A_1$  and  $A_2$  will be called *incompatible* if there exist  $a_1 \in A_1$  and  $a_2 \in A_2$  such that  $\text{span}(a_1) \cap \text{span}(a_2) \neq \emptyset$ .

The notion of incompatibility tries to capture the fact that there are pairs of axes of a figure that cannot be attended to simultaneously. For example, in Fig. 8, the axes  $a_1$  and  $a_2$  are incompatible. It should be pointed out that incompatibility is a relation that is only meaningful when comparing two axes at the same level in the hierarchy—i.e., if two clusters are incompatible this means that they cannot both occur at the same level of the hierarchy. However, it is quite possible for them to occur at different levels. A relation that extends across levels in the hierarchy is:

**Definition 2:**  $A_2$  is *inconsistent* with respect to  $A_1$  if there exists an  $a_2 \in A_2$  such that elements of  $\text{span}(a_2)$  occur entirely in both the positive and negative half planes determined by  $A_1$ . For example, in Fig. 9,  $A_2$  is inconsistent with respect to  $A_1$  because sides  $s_1$  and  $s_2$  are contained one in the positive, and the other in the negative, half plane defined by  $A_1$ .

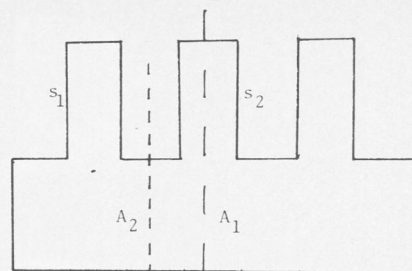


Fig. 9.

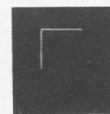
level 1  
vertical  
{1,3,4}



level 1  
diagonal  
{1,2}



level 2  
horizontal  
{1,4}



level 2  
diagonal  
{2}



level 3  
diagonal  
{1}



Fig. 10.

Given these two definitions, we are now in a position to describe the construction of the lower levels of the hierarchy. Call a node  $N$  in the hierarchy a tip node if the operation described below has not been applied to it. Initially, the root of the hierarchy, which corresponds to the major axis, is the only tip node. Each node has an axis  $A_N$  associated with it. Let  $S_N$  be the set of sides of the polygon associated with the tip node. At the root initially,  $S_N$  is the set of sides of the polygon that

- are symmetric with respect to the major axis,
- have at least one vertex in the positive half plane defined by the major axis.

The expansion operation is as follows.

1) Choose a tip node  $N$  from the hierarchy—if there are none, then the hierarchy is complete.

2) Find all clusters  $A_i$  such that a)  $A_i$  is not already in the hierarchy, and b)  $A_i$  is not inconsistent with respect to  $A_N$ .

Let  $PD$  be the set of these  $A_i$ ;  $PD$  represents the set of possible descendants of node  $N$  in the hierarchy. Let  $D_N$  be the set of descendants of node  $N$ . Initialize  $D_N = \emptyset$ .

3) Choose from  $PD$  the axis  $A_j$  such that  $s(\text{span}(A_j \cap S_N)) > 0$  is maximal, and  $A_j$  is not incompatible with any element of  $D_N$ . If no such  $A_j$  exists, then all of the descendants of node  $N$  have been generated, and we go to step (1).

4) Set  $S_N = S_N - (\text{span}(A_j) \cap S_N)$ . Set  $PD = PD - A_j$ , and  $D_N = D_N \cup A_j$ . Insert node  $N'$  into the hierarchy with  $A_{N'} = A_j$ , and  $S_{N'} = \{\mathcal{S}_i \mid \mathcal{S}_i \in \text{span}(A_j) \cap S_N \text{ and } \mathcal{S}_i \text{ is in the positive half plane determined by axis } A_j\}$ . Go back to step (4).

Given different major axes, different hierarchies will be generated. Fig. 10 shows two such hierarchies for the square in Fig. 6(a). At each node in the hierarchy we display  $S_N$  and

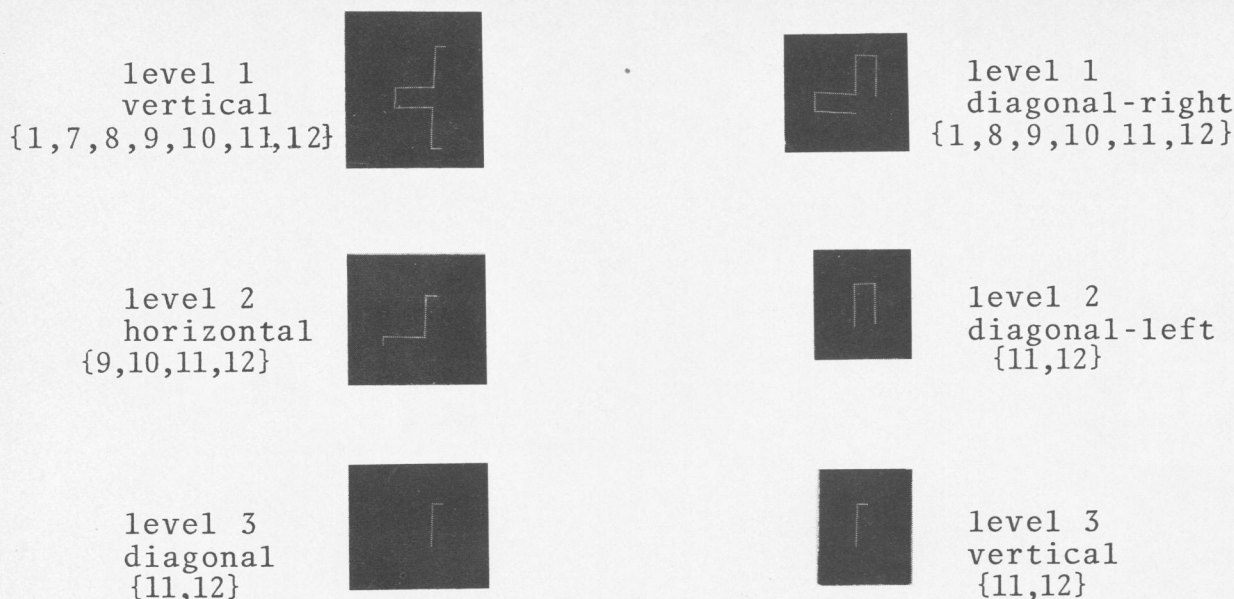


Fig. 11.



Fig. 12. (a) Sharp angle. (b) Blunt angle.

list the axis associated with that node. Fig. 11 contains two hierarchies for the cross in Fig. 6(b).

III. EXTENSION TO SMOOTHLY CURVED SIDES

This section will consider how the approach described in Section II is modified to deal with sides that may curve gently. A smooth curve segment is taken to be one with no significant curvature maxima. Significance is a function of both the resolution and magnitude of the curvature maxima that do exist along a curve segment. In the context of hierarchical descriptions, a smooth side is a side with no substructure. Thus the process that constructs the hierarchical description implicitly defines the significance of curvature maxima.

In order to augment the symmetry understanding mechanisms to deal with smooth curves, a notion of shape similarity is incorporated into the microsymmetry detection procedures. It is important to emphasize that we are interested in coarse measures of similarity; therefore, curves that are exactly similar according to the criteria we will define will not necessarily be identical. The problem of matching two curves will be discussed elsewhere. This entails modifying the microsymmetry cost function  $c$  to reflect the degree of similarity between two curves. A second, more subtle, difference between contours with smooth sides and these with only straight sides is that angles may no longer be corners; an angle may be sharp or blunt, depending on the curves that meet at the angle (see Fig. 12). Here again, the microsymmetry cost function is where this new information enters the system. This means that neither the cluster building nor description building procedures have to be changed.

The remainder of this section is divided into three parts:

A) A discussion of similarity measures between smooth curves.

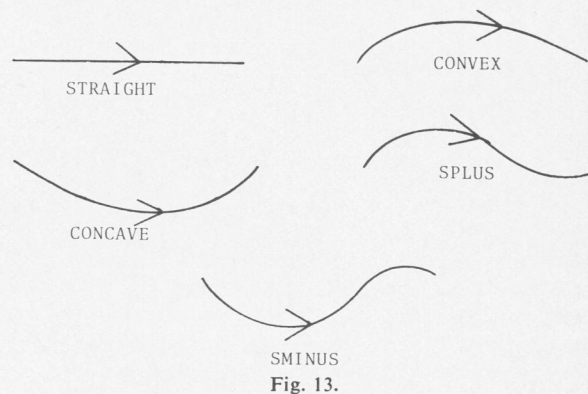


Fig. 13.

B) A discussion of functions that compute the "sharpness" of an angle.

C) Analysis results for some smoothly curved contours.

A. Similarity Measures Between Smooth Curves

Our similarity measure is based on assigning fuzzy memberships in a class of sets to a curve. The sets correspond to the allowable description types for the class of contours being considered—e.g., straight, convex, concave. The names of these sets constitute a vocabulary for describing smooth sides. We will consider only the following five descriptions—STRAIGHT, CONVEX, CONCAVE, SPLUS, SMINUS (see Fig. 13).

Let the sequence of coordinate pairs of a side  $s_i$  be

$$v_i = (x_{i_0}, y_{i_0}), (x_{i_1}, y_{i_1}), \dots, (x_{i_n}, y_{i_n}) = v_{i+1}.$$

Let  $b_{s_i} = (\theta_{s_i}, \rho_{s_i})$  be the parametric description of the straight side in the hierarchy of sides corresponding to  $s_i$ . For example,  $b_{s_i}$  might be  $\vec{v}_i v_{i+1}$ . A particularly bad choice is the least squares line defined by the points  $(x_{i_0}, x_{i_1}), \dots, (x_{i_n}, y_{i_n})$ , since it must pass through one of the points in the sequence. Let  $d(x, y, b_s) = \rho_s - x \sin \theta_s + y \cos \theta_s$  be the distance of the point  $(x, y)$  from the line  $b_s$ .

The first step in determining the membership values for  $s_i$  is to construct the set

$$\mathcal{D}_{s_i} = \{d(x_i, y_i, b_{s_j}); 1 \leq j \leq n - 1\}$$

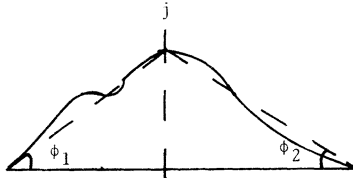


Fig. 14.

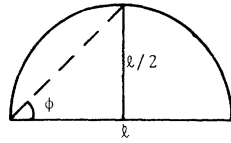


Fig. 15.

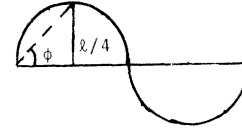


Fig. 16.

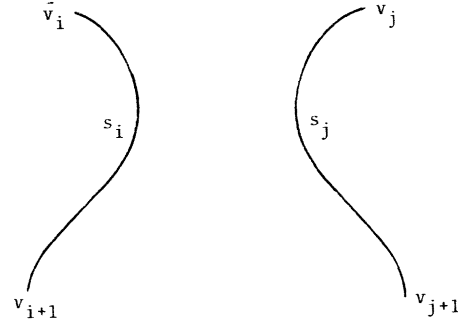


Fig. 17.

which is the set of distances of all points of  $s_i$  from  $b_{s_i}$ . Define

$$\mathcal{D}_{s_i,r}^+ = \{j: d(x_i, y_i, b_{s_i}) > 0 \wedge d(x_{i_k}, y_{i_k}, b_{s_i}) < d(x_i, y_i, b_{s_i}), \quad |j - k| \leq r\}$$

$$\mathcal{D}_{s_i,r}^- = \{j: d(x_i, y_i, b_{s_i}) < 0 \wedge d(x_{i_k}, y_{i_k}, b_{s_i}) > d(x_i, y_i, b_{s_i}), \quad |j - k| \leq r\}.$$

$\mathcal{D}_{s_i,r}^+$  is the set of points that are local maxima of distance from  $b_{s_i}$  over a range (smoothing factor  $r$ ), and  $\mathcal{D}_{s_i,r}^-$  similarly is the set of local minima. We should choose  $r$  large enough so that the local distance maxima corresponding to curvature maxima of fine resolution will not be detected.  $\mathcal{D}_{s_i,r}^+$  can be computed by a parallel-like algorithm (see Davis [1]).

We sort  $\mathcal{D}_{s_i,r}^+$  by  $d$  so that we can refer to  $\mathcal{D}_{s_i,r}^+(1)$ , the element of the set with greatest  $d$  value, etc. Similarly, we sort  $\mathcal{D}_{s_i,r}^-$  so that  $\mathcal{D}_{s_i,r}^-(1)$  is the element of the set with minimal  $d$  value. We use the ordered sets  $\mathcal{D}^+$  and  $\mathcal{D}^-$  to construct the fuzzy membership values for side  $s_i$ . Denote  $\mathcal{D}_{s_i,r}^+(1)$  by  $i^+$  and  $\mathcal{D}_{s_i,r}^-(1)$  by  $i^-$ . If  $\mathcal{D}^+$  is empty, let  $i^+ = -1$ , and similarly for  $\mathcal{D}^-$ . We can now present the fuzzy membership functions for the various sets.

1) STRAIGHT: We take the ideal straight side to be one whose  $\mathcal{D}^+$  and  $\mathcal{D}^-$  are empty. The straightness of a side is related to the angular stickout of the elements of  $\mathcal{D}^+$  and  $\mathcal{D}^-$  from the line  $b_{s_i}$ . The angular stickout of point  $(x_i, y_i)$  from  $b_{s_i}$  is (see Fig. 14)

$$ASO(j) = \frac{\sin^{-1}(|d(x_i, y_i, b_{s_i})|)}{\max[\sqrt{(x_{i_0} - x_{i_j})^2 + (y_{i_0} - y_{i_j})^2}, \sqrt{(x_{i_n} - x_{i_j})^2 + (y_{i_n} - y_{i_j})^2}]}, \quad \text{if } 1 \leq j \leq n$$

$$= 0, \quad \text{otherwise.}$$

If  $i^+ = -1$ , then  $ASO(i^+) = 0$ . To measure STRAIGHT ( $s_i$ ), then, we compute

$$\max(0, (\max(ASO(i^+), ASO(i^-)) - t_{ST}) \cdot (1/t_{ST}))$$

That is, the straightness of side  $s_i$  is related to the maximum angular stickout of any point of side  $s_i$  from  $b_{s_i}$ . If this angular stickout is greater than some threshold  $t_{ST}$ , we set the membership of  $s_i$  in the set STRAIGHT to 0.  $t_{ST}$  will correspond to the angular stickout of an ideal convex side. Thus we are saying that a side whose membership in CONVEX is 1.0 has membership 0 in STRAIGHT.

2) CONVEX: Our model for a convex side is a semicircle. For side  $s_i$  of length  $l = |\vec{v}_i \vec{v}_{i+1}|$ , a semicircle with  $s_i$  as diameter will have maximum angular stickout from  $b_{s_i}$  of  $\pi/4$  (see Fig. 15). So, we define CONVEX ( $s_i$ ) =  $\min(ASO(i^+) - ASO(i^-), 0) \cdot 4/\pi$ . If  $ASO(i^-) > ASO(i^+)$ , then the side is more concave than convex.

3) CONCAVE ( $s_i$ ) =  $\min(ASO(i^-) - ASO(i^+), 0) \cdot 4/\pi$ .

4) SPLUS: This case represents smooth sides that have a convex segment followed by a concave segment. Clearly SPLUS blends into STRAIGHT, CONVEX, and CONCAVE as the  $ASO$  values go to 0, or the lengths of the convex or concave segments go to 0. Our model for an SPLUS side is a convex semicircular arc joined with its reflection, a concave semicircular arc (see Fig. 16), and we take

$$\text{SPLUS}(i, s_i) = \text{if } i^+ < 0 \vee i^- < 0, \text{ then } 0,$$

$$\text{else if } i^- < i^+, \text{ then } 0,$$

$$\text{else } \min(ASO(i^+), ASO(i^-)) \cdot 1/t_{SP},$$

where  $t_{SP} = \pi/8$  (see Fig. 16).

5) SMINUS is defined analogously to SPLUS.

Using these functions, a side is assigned a membership in each set. As a measure of dissimilarity between two sides we have chosen the maximum absolute difference between corresponding membership values. The cost function for microsymmetries is changed in the following way. Any pair of sides not symmetric according to the old criterion is rejected, but now any pair of sides whose dissimilarity measure  $> 0.5$  is also rejected. The value 0.5 was chosen because it indicates a reasonable certainty that one side is contained in one set (e.g., CONVEX), while the other side is not.<sup>1</sup> The new value for the cost is the old value plus a scale factor times the dissimilarity:

$$c'(a) = c(a) + k \cdot \text{dissimilarity of } (s_i, s_j).$$

## B. "Cornerity" Measurement

In [3], Freeman and Davis proposed a "cornerity" measure that was based on the curvature at an angle and the straightness of the sides of that angle. This section presents an alternative method based on the local curvature measurements that underlie the angle detection procedures described in Davis [1].

Briefly, this curvature measure is defined as follows: Let  $\{(x_i, y_i)\}_{i=0}^n$  be a sequence of points that describe a simple curve. Let  $\vec{p}_{i,j}$  be the vector from  $(x_i, y_i)$  to  $(x_j, y_j)$ . Then  $c_{i,r}$ , the curvature at point  $i$  and resolution  $r$ , is defined to be  $\cos^{-1}(\vec{p}_{i,i-r} \cdot \vec{p}_{i,i+r})$ . This is the size of the smaller included angle determined by the two vectors.

<sup>1</sup> Note that when computing the dissimilarity between  $s_i$  and  $s_j$  for determining the degree of microsymmetry between the sides, if we are using vertex pairs  $v_i$  and  $v_{j+1}$  (or  $v_{i+1}$  and  $v_j$ ), then we compare SPLUS ( $s_i$ ) with SMINUS ( $s_j$ ) and SMINUS ( $s_i$ ) with SPLUS ( $s_j$ )—see Fig. 17.



Fig. 18.

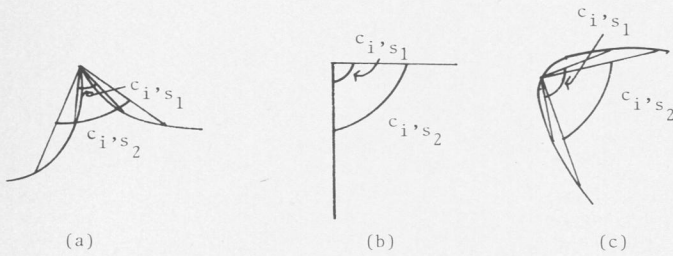


Fig. 19.

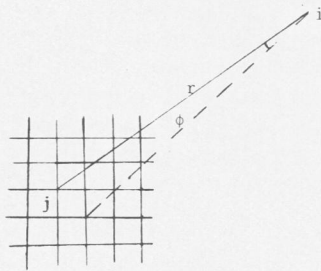


Fig. 20.

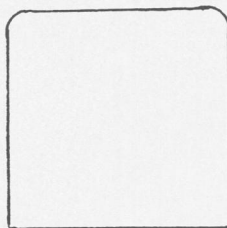


Fig. 21.

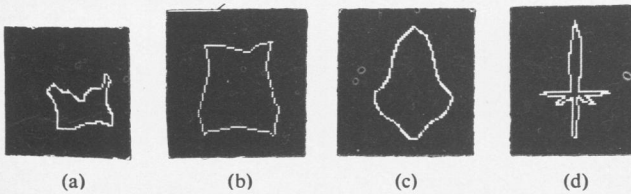


Fig. 22. Test curves.

Now, suppose that the angles in Fig. 18 are to be distinguished on the basis of their sharpness. The angle in Fig. 18(a) can be described as cusp-like, that in Fig. 18(b) as corner-like, and that in Fig. 18(c) as rounded.

The method is based on the rate of change of  $c_{i,r}$  with respect to  $r$ —i.e.,  $dc_{i,r}/dr$ . Fig. 19(a)–(c) shows that we would expect  $dc_{i,r}/dr$  to be positive for a cusp-like angle, zero for a corner-like angle, and negative for a rounded angle.

In order to estimate  $dc_{i,r}/dr$ , we fit a least squares line to a set of  $c_{i,r}$ 's and regard the slope as  $dc_{i,r}/dr$ . Since different resolutions are attributed to different angles, the set of  $c_{i,r}$ 's used to calculate  $dc_{i,r}/dr$  for a particular angle should be determined by the resolution of that angle. The largest size used

TABLE I  
FOURIER COEFFICIENTS FOR FIG. 22(c)–(d)

(a)	$n$	$\alpha_n$	$\gamma_n$
	0	20	0
	1	20	0
	2	20	0
	3	20	0
	4	20	0

$$r(\theta) = \sum_{n=0}^4 \alpha_n \exp[\cos(\theta + \gamma_n)]$$

(b)	$n$	$\alpha_n$	$\gamma_n$
	0	4.	0
	1	4.5	0
	2	5.0	0
	3	5.5	0
	4	6.0	0

$$r(\theta) = \sum_{n=0}^4 \exp[\alpha_n \cos(\theta + \gamma_n)]$$

TABLE II  
FUZZY MEMBERSHIP VECTORS FOR THE SIDES OF FIG. 23(a)–(d)

(a)	SIDE	DEGREES OF MEMBERSHIP	(STRAIGHT, CONVEX, CONCAVE, SPLUS, MINUS)
	1	(.87, 0, .12, 0, 0)	
	2	(.74, .25, 0, 0, 0)	
	3	(.71, 0, .28, 0, 0)	
	4	(.85, 0, .08, 0, .11)	
	5	(.82, 0, .17, 0, .11)	

(b)	1	(.74, .25, 0, 0)
	2	(.88, 0, .02, .17, 0)
	3	(.79, .20, 0, 0, 0)
	4	(.88, 0, 0, 0, .22)

(c)	1	(.88, .05, 0, .16, 0)
	2	(.84, .15, 0, 0, 0)
	3	(.85, .14, 0, 0, 0)
	4	(.86, .08, 0, 0, .10)

(d)	1	(.92, .07, 0, 0, 0)
	2	(1.00, 0, 0, 0, 0)
	3	(.70, .24, 0, 0, .10)
	4	(.88, .11, 0, 0, 0)
	5	(1.00, 0, 0, 0, 0)
	6	(.40, .54, 0, .09, 0)
	7	(.89, .10, 0, 0, 0)
	8	(.86, .13, 0, 0, 0)

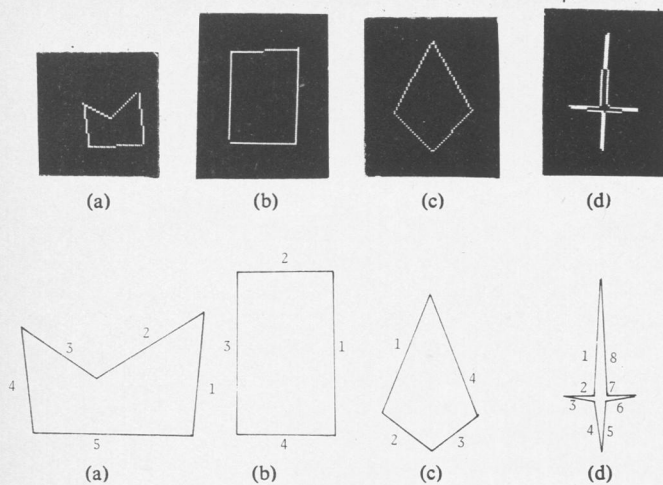


Fig. 23. Top-level approximations.

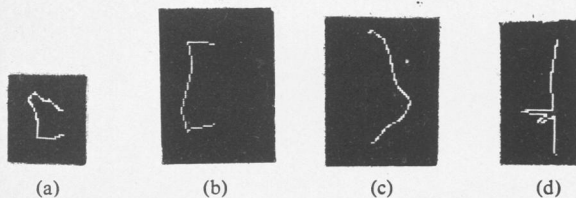


Fig. 24. "Halved" figures using best axis.

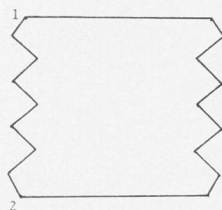


Fig. 25.

should reflect the localness of the perception of sharpness, and so will probably not usually be comparable to the resolution of the angle. The smallest size, on the other hand, should be large enough so that curve digitization error cannot appreciably influence the computations (if  $l$  is the length of  $\vec{p}_{ij}$ , then the angular uncertainty in  $\vec{p}_{ij}$  is proportional to  $\arctan 1/l$ —see Fig. 20). Alternatively, one can weight the curvature values according to the expected error due to digitization. However, we will assume that our curves are well digitized (Freeman [2]), so that digitization effects can be ignored.

It is not obvious how to compare the importance of similar angle descriptions and side descriptions in evaluating the quality of an axis of symmetry, and so the cornerity measure has not yet been integrated into the microsymmetry cost function. Fig. 21, though, seems to indicate that the description of angles does play a role in human perception of symmetry; intuitively, its role is less important than that of side descriptions.

### C. Experimental Results

Fig. 22(a)–(d) shows four smooth curves. Fig. 22(a) is a jigsaw puzzle piece taken from Freeman [2]; Fig. 22(b) is a hand-drawn square-like object; while Fig. 22(c), (d) were generated using Fourier methods due to Shepard [8] and described in Davis [1]. Table I lists the coefficients used to generate Fig. 22(c), (d).

TABLE III  
CORNERITY MEASURES FOR THE ANGLES IN FIG. 23(a)–(d)

angle	cornerity (r=8)
(a)	
1	.05
2	.16
3	.13
4	.28
5	-.05
(b)	
1	-.02
2	0.
3	.03
4	-.01
(c)	
1	-.01
2	0.
3	.10
4	.01
(d)	
1	0.
2	.02
3	0.
4	.01
5	.04
6	.02
7	0.
8	.02

The top level of the angle/side hierarchy for each of the contours is shown in Fig. 23(a)–(d). Table II(a)–(d) lists the fuzzy membership values of each of the sides of the hierarchies in Fig. 23(a)–(d). Table III(a)–(d) displays the cornerity measures for the angles in the hierarchies, and Fig. 24(a)–(d) shows the best symmetry descriptions of each of the contours. Note that especially for Fig. 22(b), if the descriptions of the sides were not taken into account, then the vertical axis would not have been preferred over, say, the horizontal.

### V. SYMMETRY UNDERSTANDING FOR HIERARCHICAL DESCRIPTIONS OF SHAPE

Suppose that instead of restricting the sides of a contour to be smooth, we allow them to have substructure in the form of subsides. (Methods of obtaining hierarchical representations of contours in terms of sides and subsides were discussed in an earlier paper [1].) For example, in Fig. 25, it does not seem sufficient to describe the side between angles 1 and 2 as simply being straight; some notice should be taken of the many small sharp angles and short sides that underlie the gross straight nature of the side. The presence of this substructure should also be taken into account when describing the symmetries of Fig. 25.

The discussion that follows will be mostly speculative; instead of presenting extensions or modifications to the algorithms of Sections II and III, we will examine certain issues that would be central to any such modifications. Some of these issues should be relevant to a computer model of human shape understanding. The first question to be dealt with is: Are the smooth sides discussed in Section III a subset of the set of



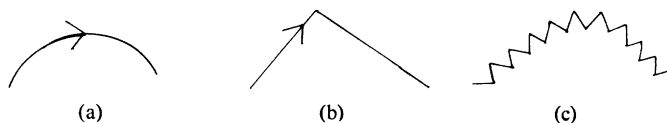


Fig. 26.

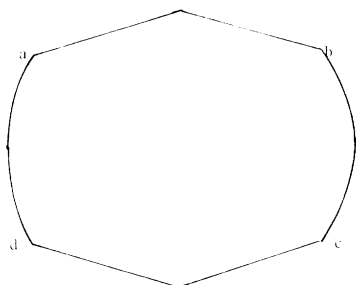


Fig. 27.



Fig. 28.

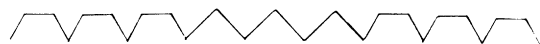


Fig. 29.

sides that have substructure, or are the sets disjoint? On the one hand, we have regarded a side as being smooth just in case it is assigned no substructure. Yet, on the other hand, a piecewise-linear approximation to a smooth side may be represented as a side with substructure, and this can happen in many ways.

For example, Fig. 26(b), (c) shows piecewise-linear approximations to the smooth curve in Fig. 26(a). If we regarded all three curve segments in Fig. 26 as smooth, and generated the sets  $\mathcal{D}^+$  and  $\mathcal{D}^-$ , then the description generated based on these sets would probably be the same for all three segments: primarily convex curves. The substructural components of Fig. 26(b), (c) correspond to yet another level of description. Fig. 26(b) is a convex side with one obtuse angle located at the point of maximal displacement from the baseline, while Fig. 26(c) is a convex side that is jagged. This suggests that in detecting symmetries between two sides, we might compare the descriptions based on the assumption that the sides are smooth, and then refine this comparison using substructure information. So, for example, in Fig. 27, all four sides would be characterized as convex, but sides  $ab$  and  $cd$  would have a common substructural description—namely, the obtuse angle.

Fig. 26(b), (c) also illustrates an important aspect of the representation of substructure. Fig. 26(b) has as substructure one obtuse angle, while we described the substructure of Fig. 26(c) as "jagged." The distinction between substructural description that are statistical (Fig. 26(c)) and enumerative (Fig. 26(b)) is a fuzzy one. In the former case, we are abstracting a common feature of many substructural components, and assigning that quality to the side as a whole; while in the latter case, no such global description of the side is possible. The situation can become more complex, as in Fig. 28, where there is a combination of statistical and enumerative features—i.e., we can describe the curve segment as a straight jagged segment, followed by a smooth, straight segment, followed by a straight wavy segment.

Even more perplexing is the need to notice repetitive features.

Figs. 27 and 28 can probably be adequately dealt with on the basis of a simple taxonomy of angles as sharp or rounded. But in curve segments such as in Fig. 29, it is important to be able to isolate the elements  $\frown$  and  $\smile$ ; otherwise, we cannot obtain a satisfactory description of the curve, i.e., a description that is in accordance with our own perception.

## VI. DISCUSSION

Symmetry certainly plays a prominent role in human perception of shape. In computer understanding of shape, symmetry can serve as a basis for compact representation schemes, as well as providing clues for the completion of shapes with gaps. A subsequent report will deal with the latter problem.

The previous sections of this report have presented algorithms that can understand symmetry for polygon-like shapes, and have described modifications to these algorithms that allow the analysis of shapes with smooth sides. Several examples were presented to illustrate the capabilities of the algorithms. Finally, some of the difficulties that would be encountered in attempting to extend the symmetry analysis to shapes with a hierarchy of side and angle representations were discussed.

## ACKNOWLEDGMENT

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## Signature Verification Experiment Based on Nonlinear Time Alignment: A Feasibility Study

MAKOTO YASUHARA, MEMBER, IEEE, AND MASATOMO OKA

**Abstract**—Automatic signature verification experiments are described, in which we select one of the most promising techniques described in the literature of automatic speech recognition of multisyllabic utterances: nonlinear time alignment by dynamic programming, and apply it to our problem to check its feasibility. Our experiments show that it is very useful under certain conditions.

## INTRODUCTION

This correspondence discusses one of the techniques applied to the automatic verification of real-time signatures.

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M. Yasuhara is with the Research Laboratory of Communication Sciences, University of Electro-Communications, Tokyo 182, Japan.  
M. Oka is with Nippon Electric Co., Tokyo, Japan.