

A Continuous Model for
Salient Shape Selection and Representation

by

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The good life is one inspired by love and guided by knowledge.

- Bertrand Russell /1872 – 1970/, “What I Believe”

To My Family

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Abstract

We propose a new framework for shape representation and scenery shape selection. Various topics including figure/ground separation, shape axis construction, junction detection and illusory figure finding will be discussed.

The model construction is inspired by the Gestalt studies. They suggest proximity, convexity, symmetry, etc, as cues for figure/ground separation and visual organization. By our distributed systems, we quantify those attributes for complete/partial shapes and use them for shape evaluations and representations. In particular, the shape convexity instead of other well-studied shape attributes such as the symmetry axis or size, will be emphasized.

Two models are proposed. The decay diffusion process is applied in predicting figure/ground phenomenon, based on a convexity measure for figure/ground sharing the same area. The orientation diffusion process, adopting orientation information on shape boundaries/edges, will discuss the figure/ground separation or shape convexity comparison for regions not owning the same size. A Kullback-Leibler convexity measure is proposed, with a flexible scenario. Through a parameter, we are allowed to choose between a size-invariant convexity measure or one with small-size prefer-

ence. For convexity comparison of perfectly convex shapes, a preference of circles over triangles will be given, as well as the preference of squares over rectangles. These two models are also used in generating the symmetry information. In particular, the symmetry information suggested by the orientation process is computed by only local operations. The junction information will be derived similarly, where junctions are considered no more than “boundary axis points”.

Our framework, based on variational formulations will produce the static-state results. The simulation is continuous, rely on no artificial binary thresholds. For convexity measurement, other than the mathematical 0-1 definition, we distinguish between “more” or “less” convex shapes. For axis construction, we provide the information which continuously describes strength of the axes for natural axis pruning. For junction detection, the transition from low-curvature or high-curvature curves to curves with a discontinuous curvature will be seen.

The decay diffusion process, with help of the convexity/entropy measure will also be applied in shape selection. Hence, our framework integrally combining many different functions is useful as a universal low- to middle-level vision simulation.

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Chapter 1

Introduction

A goal in computer vision is to simulate human visual system with computer routines. There are variations in the visual system among different individuals or even among different occasions for the same individual. Thus, the general simulation must be defined in a statistical sense. In general, the success of computer vision model is based on some percentage of agreements between experiments in human visual system and prediction of the computational simulation.

Statistics and Bayes Rule The performance of a visual system on a particular experiment can be modeled as the outcome of a certain random variable. The randomness implies that, no outcome from any particular experiment will be essential as testing data for our simulation. However, we pursue the general behavior for a group

of random variables. Two cases can be studied. When we try to build a theory across different individuals, our principle is to search the agreement between the prediction of our simulation and majority of outcomes from human tests. In this case, one “diverse” human visual system can be treated as an “imperfect machine”. More likely, we accept a weaker, the computational view. The simulation is the goal to find a model with the help of so-called *method of undetermined parameters*, waiting for the hidden parameters owned by each individual to be filled in.

On the other hand, we can also pursue the theory built for a single individual among different occasions. When the outcomes are sorted by time, *learning* is involved.

One basic rule applied in the decision making process is the Bayes Rule. Based on an observation recorded by the vector* \mathbf{x}_0 , an object A can be categorized as one of k categories $\{\omega_i; i = 1, \dots, k\}$ by investigating value of the conditional probability† of ω_i on \mathbf{x}_0 ,

$$P(\omega_i|\mathbf{x}_0) = \frac{p(\mathbf{x}_0|\omega_i)P(\omega_i)}{p(\mathbf{x}_0)}, \quad (1.1)$$

where

$$p(\mathbf{x}_0) = \sum_{i=1}^k p(\mathbf{x}_0|\omega_i)P(\omega_i). \quad (1.2)$$

*A **bold** face will be used for vectors to distinguish from scalars.

†We use an upper-case P to denote the probability mass function and a lower-case p to denote the probability density function.

For a fixed observation \mathbf{x}_0 , the result is decided by the conditional probability $p(\mathbf{x}_0|\omega_i)$ and the *a priori* probability $P(\omega_i)$. Moreover, in the case where $P(\omega_i)$ is a constant, independent of the categories ω_i , we can adopt a more useful form,

$$P(\omega_i|\mathbf{x}_0) \propto p(\mathbf{x}_0|\omega_i), \quad (1.3)$$

known as the maximum likelihood method. A larger conditional probability $P(\omega_i|\mathbf{x}_0)$, called the *a posteriori* probability suggests the decision “ A belongs to the category ω_i ”. In the case where Eq. 1.3 can be assumed, we have a universal theory, independent from the “time” and different individuals. When Eq. 1.3 is not applicable, for a fixed individual, we look for an asymptotic result within a period, if there is any.

Let us discuss $P(\omega_i|\mathbf{x}_0)$ for different applications. In high-level vision problems, a decision may be made for guessing whether or not the object is a cat or a dog based on some features in a given image. In low-level vision problems, the decision may be the detection or not of a feature in a particular location of an image. Middle-level vision makes the transition from a distributed (local) low-level information to a more abstract, object oriented, perhaps symbolic representation. It may address the problem of distinguishing between the figures and the background in images.

High-level x low-level vision In general, the concept of different level tasks in human intelligence can be illustrated in Tab 1.1. The tasks in levels n , $n+1$ and $n+2$

Level	Task
⋮	
$n + 8$	Tasting baptism water sweeter than normal water
$n + 7$	Knowing “Path Finder” may not find the path
$n + 6$	Realizing he is your grandfather if your father calls him father
$n + 5$	Ignoring the ads on the web
$n + 4$	Reading text through a mirror
$n + 3$	Object recognition
.....	
$n + 2$	Object detection
$n + 1$	Texture detection
n	Detection of intensity difference
⋮	

Table 1.1. The level of human tasks. The tasks above the dot line may not be accomplished by using only the visual system. The level $n + 5$ and the levels below it are called the *literal levels*. ■

may be accomplished by introducing only the visual system. The tasks in levels n , $n + 1$ may be accomplished by applying only local considerations in images.

There are differences between low-level and high-level vision and we want to expand on this topic.

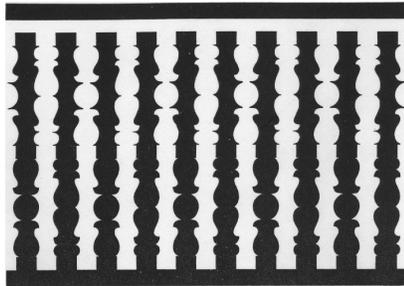
- i In low-level vision many of the random variables, each associated with an image feature, are defined everywhere in the image. In higher level vision, the variables are more global (not defined everywhere in the image) and possibly there are fewer variables than for the low-level vision case. Therefore, a variable in the high-level vision is expected to have higher complexity than the one in the low-level vision. It can lead to tough challenge for the high-level simulation.
- ii The distributions associated with low-level tasks are more ambiguous than for high-level ones. One is expected to have, in general, more difficulties in deciding whether there is a corner or not in an image location than to decide if there is a face of a person in an image location. The distribution associated with face recognition consists of so many different random variables (distributions) that it is expected to be a much sharper/more peaked distribution (see Fig. 1.1).
- iii There are more variations across human visual systems over the task-distributions in the high-level vision than in the low-level vision. Someone that has never

seen birds will have a very different high-level description/distribution of a bird when seeing one, than most of us who have seen birds. High-level descriptions/distributions are learned through experience and therefore have more varieties according to individual experiences. In this way, it is easier to collect consistent data for a low-level human task than for a high-level one.

- iv The outcome from the high-level task usually overrides the outcome from the low-level task when there is a conflict. If an edge-boundary is needed to complete a face figure the high-level system will “see” the edge, even if no intensity gradient exists.
- v The high-level task will interact more strongly with other brain activities (other types of intelligence), such as in speech or upon playing chess.

From low-level to high-level, figure-ground separation and visual organization

The middle-level vision is where the transition from low-level vision to high-level vision occurs. It transforms a distributed (local) set of informations into a coherent, object oriented, abstract, description of the world, but yet, without naming objects or recognizing objects from past experience (memory). In our view, it is where local properties are integrated into surfaces or objects. The choice of intergration is called *visual organization*.



(a)



(b)

Figure 1.1. The figure/ground problem (adapted from [77]) and how it interacts with low-level x high-level vision processes. (a) Wrangling rungs. Typically, humans perceive white or black as figure with equal probability. (b) Sara Nader. A high-level knowledge is involved for making the decision to see a woman face or a man playing saxophone. Each figure, (a) and (b), admit two solutions as figure/background, but in (b) the solutions are more stable than in (a), i.e., one does not keep switching from woman face to saxophonist as often as in (a) from black to white. ■

Fig. 1.1 illustrates the task of *figure/ground separation*. The visual system chooses one side of the edges as figure and as background for the other side. One can also formulate it as a *border ownership* problem [71] [72] [59]: given the complete description of the boundary contours, how does a system decide on which side of the

boundary is the surface that gives rise to that border? It is the simplest visual organization problem where the search of organization is equivalent to the binary selection of F/G. As formulated by the border ownership problem, usually, we discuss the F/G separation through pairs of experiments of inverse intensities, as in the convexity-versus-symmetry images in Fig. 1.6(a1) & (a2). Without the contrast polarity, the result of F/G is decided by geometry of the (partial) shapes. To describe the F/G problem by Bayes Rule, we can assume a constant *a priori* probability $P(\omega_i)$ with $\omega_i = \text{figure, ground}$ in Eq. 1.1.

There is a case which the intensity edge is absent. The illusory curve in Fig. 1.2(a), introduced by line-endings, will not be detected by any naive edge detector. For the detection of the “Kanizsa square”[‡] in Fig. 1.3(a), people believe that a global consideration is necessary [39] [48] [19].

Vision as an inverse problem with the “simplicity” criteria The real world is described and reasoned by our visual system and intelligence. The 3-D real world, the scene \mathfrak{M} , reflects lights and is projected to the image space \mathfrak{J} . Our description of

[‡]In principle, people can be trained to perceive up to 5 different visual organizations. Besides those two demonstrated in Fig. 1.3(b) & (c), we have (3) a white square in front, occluding white plane with 4 holes; (4) white plane with 4 holes occluding a white square in the back and (5) plane with 4 holes of pecman shape.

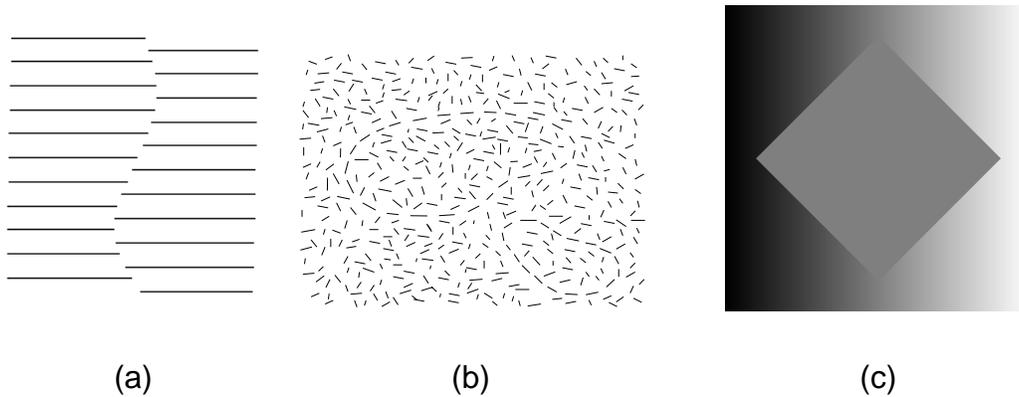


Figure 1.2. (a) The illusory curve (adapted from Schumann [79]). (b) Random oriented dots with the suggestion of closure. (c) Constant intensity square with 1-D linear gray from 100% black to 100% white in the background from left to right, inspired by Shapley and Gordon [76]. The illusory gradient is perceived inside the square without any support of intensity changes. In particular, the gradient is extended in the direction perpendicular to the square boundaries. ■

the scene is an inverse process from the image to the scene.

$$\mathfrak{W} \stackrel{f}{\rightleftharpoons} \mathfrak{I}.$$

We consider searching of the inverse of f , the mapping $\mathfrak{I} \mapsto \mathfrak{W}$, to be the computer vision research. Given a single image, the mapping $\mathfrak{W} \xrightarrow{f} \mathfrak{I}$ is considered not 1-1. It is neither an onto mapping in the sense that the majority part of \mathfrak{I} is with low probability within the range of f . Most images in the image space are just white

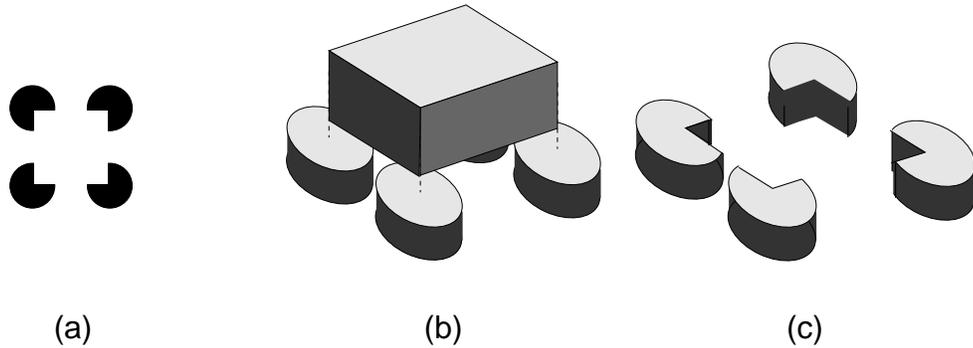


Figure 1.3. The Kanizsa square and two of its 3-D constructions. The human visual system prefers the perceptual organization (b) which gives the white square, 4 black circles and the background as three successive layers in the scene rather than (c) which consists of only two layers, four pecmen and the background. Each layer may contain one or more than one object. ■

noise. Real images, coming from real scenes, are clustered in a small subset of the space \mathcal{I} . The Kanizsa square in Fig. 1.3 shows us an example where a 3-D scene, W_1 , with four (4) black pecmen and another one, W_2 , with a salient white square occluding four (4) black circles in the back, both in \mathfrak{W} with $f(W_1) = f(W_2) = I \in \mathcal{I}$, but $W_1 \neq W_2$. Other examples involving depth and lighting can be referred to the discussion of visual organization and depth perceiving in [8] or Fig. 1.4.

We consider computer vision as searching for some mapping \tilde{f} out from f whose inverse will bring a given image $I \in \mathcal{I}$ to a scene $W \in \mathfrak{W}$ with $W = \tilde{f}^{-1}(I)$. We need to mention that, first, various elements in \mathcal{I} should be rejected from our

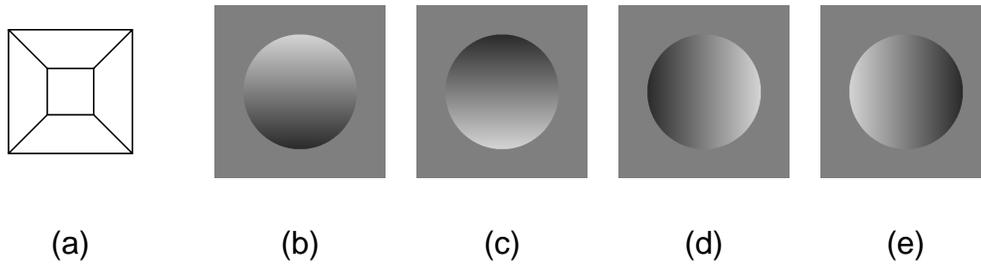


Figure 1.4. The various shapes with different depth interpretations. (a) is usually seen as cuboid popped up. (b) & (c) suggest the assumption of nature lighting effect which (b) looks like a bump and (c) looks like a dent. Without the nature lighting effect, (d) & (e) suggest the preference of perceiving objects as convex ones rather than concave ones. (b)-(e) are generated by changing gray level linearly from 4 different directions. ■

discussion if they are out of the range of f/\tilde{f} . Secondly, the selected scene by the (computer) vision system may not be the “true” scene W_{true} that gave rise to the image I via the true projection f_{true} ; i.e., $I = f_{\text{true}}(W_{\text{true}}) = \tilde{f}(W)$.

The preference of one interpretation over others usually indicates an order between the representative and other elements. For Bayesian models, the representatives can be chosen by the Maximum A Posteriori (MAP) estimation [21]. Another semantic description selects the “simplest” version among them as the representative. It is known as the Law of Prägnanz, introduced by Wertheimer [90]. Koffka [45] described the law as:

Of several geometrically possible organizations that one will actually occur which possesses the best, simplest and most stable shape.

For the Kanizsa square, the law can be expressed by having less corners to account for the square/4-circle interpretation than the 4-pecman one. In noisy images, a more regular geometry is easier to describe as it costs less information (in a compression sense) and so one produces methods of image restoration to identify the “simplest” image representative of the noisy image.

For the search of the function \tilde{f} , it is possible that the search is exponentially hard on the size of the images which one wants to explain. The so-called “visual organization research” is introduced to describe the physical or semantic structure from the 2-D input. Operationally, this information dramatically reduces the number of combinatorial alignments needed for searching possible figures in the scene, from exponential to polynomial [25]. The evidences of experiments on human visual systems imply that the “depth”[§] information is appropriate for such mission [56].

Occlusions The real world has occlusions occurring very frequently. Solving the visual organization with occlusions requires selection and reduction between combinatorial choices (see Fig. 1.5). The occlusion suggests the depth change based on

[§]The depth here can accept a weaker sense which indicates the relative depth between pixels, rather than the absolute distance from the viewers.

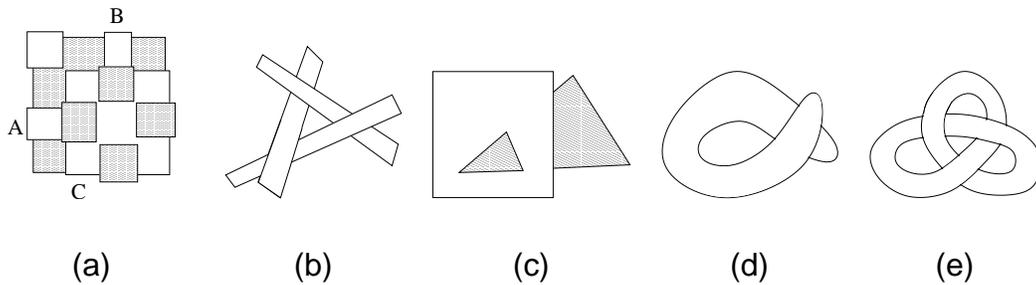


Figure 1.5. Various samples may introduce combinatorial explosion in the identification process(adapted from Cooper [10]). (a) square identification, and various objects violating the assumption of “flat objects” in $2\frac{1}{2}$ -D reconstruction, from simple to difficult: (b) nonrigid, (c) broken, (d) self-occluded and (e) self-occluded with cycles. ■

local interpretations. In the Kanizsa square image, and various examples in Fig. 1.6, the line endings, T- (or L-) junctions, convexity, size or parallelism suggest information beyond 2-D [39] [48] [63] [32] [8] [56] [92] [79]. These are local cues in images, an additional step from local to global is necessary [48] [86] [27].

Some past works are stated here. Kumaran et al. [48] approached the organization problem from considering the local (corner) configurations. To conquer the input with discretization, a perfect corner detector was assumed, same as the assumption made in Mumford [58] and Williams and Jacobs [91] for the occluded edge completion by Elastica, with the assumption of “good continuation” for edges. Also, all of their works showed no intention of handling different scales of images. The work of

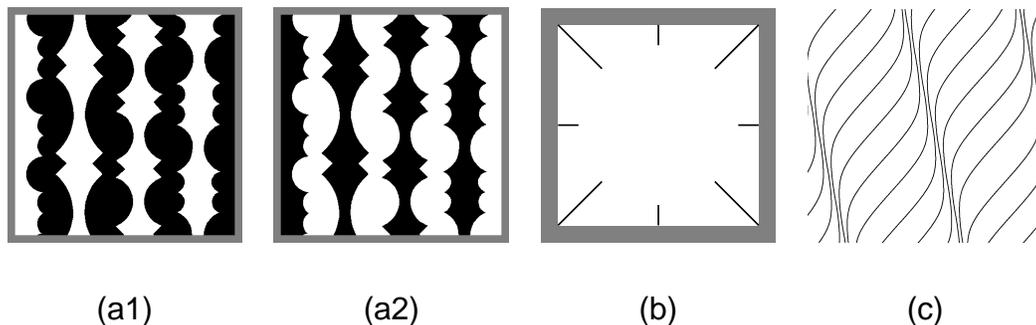


Figure 1.6. The images contain more than 2-D information. (a1) (a2) the figures Kanizsa illustrated the convexity as the dominant influence over the symmetry for the F/G separation (adapted from Kanizsa [39]). Part (a2) is provided to remove the polarity effect. (b) a pop-up circle shows the depth change provided by the line endings. (c) the parallelism evokes the impression of contours lying on a 3-D surface (adapted from Witkin et al [92]). ■

Leung & Malik [50] combined the *elastica* [58] [16] and the *normalized cuts* [78] to solve the image partition problem for various images including real images, paintings and synthesized images with subjective contours. But the concept of occlusion is missing, hence no depth information can be derived from it. From a simulation viewpoint, the works by Kumaran et al. [48] are closer to our visual system in the sense of being able to reason more than one interesting organizations (see Geiger et al. [19]). For example, the Kanizsa square has 5 different organizations where either of them can be visualized actively.

Visual Organization, Figure/Ground, Shapes and Gestalt Laws We will consult the Gestalt Laws [45] [46] [90] [8] for 2-D shapes and visual organization to sketch the plan for salient figure selection. The Gestalt psychology is usually known by the assertion “the whole is greater than the sum of its parts”. As we argued, the F/G separation is the simplest example for the search of visual organization. It is the case where the search of organization is equivalent to the binary selection from several regions, which is also equivalent to investigation of the shape geometry. It is the place where the Gestalt laws will be invoked.

The Gestalt laws says that proximity, convexity, similarity, good continuation, closure, relative size, surroundedness, orientation, symmetry are useful for F/G separation or visual organization construction [45] [46] [90] [8]. Our plan is first, construct a model to quantify the Gestalt Laws for 2-D shapes and the quantified result will be used in shape representations and evaluations. The salient surface is chosen based on such information. In other words, we will design the shape representation with those Gestalt laws included explicitly or implicitly.

1.1 Representation of Shapes

Our ultimate goal is to account for the Gestalt Laws of proximity (Ch. 2, 3), convexity (Ch. 2, 3), good continuation (Ch. 5), closure (Ch. 5), relative size (Ch. 3), surround-

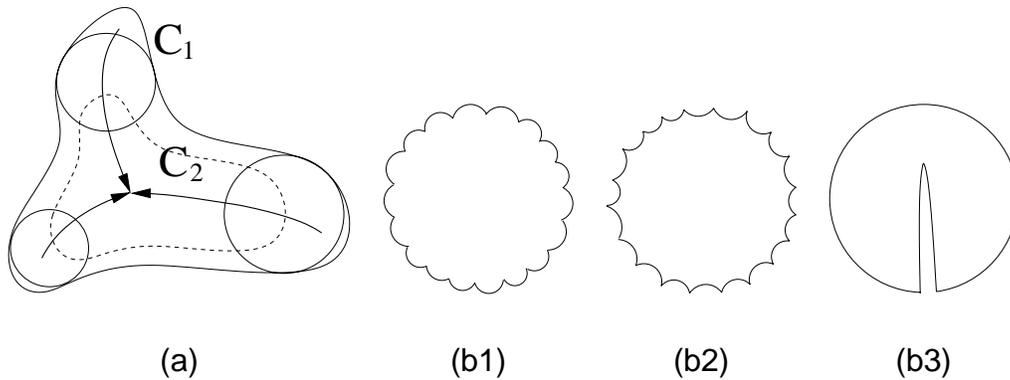


Figure 1.7. Shape representation and some Gestalt properties of shapes. (a) a shape with its representation: the internal skeletons show its symmetry axes. (b1) (b2) two shapes are convex if considered in coarse scales. (b3) the shape considered to own high convexity by the definition of Rosin [70], measuring difference of the shape area and the area of its convex hull. ■

edness (Ch. 2, 3), orientation (Ch. 3), symmetry (Ch. 4) to construct a model of shapes. We are here studying the 2D shapes, with either closed or non-closed boundaries. The non-closed shapes are considered in the F/G simulation. That is, image frames will not be treated as shape boundaries, inspired by the fact that the image frames are usually perceived in front of F/G phenomenon with figure/background extended in behind.

Convexity Kanizsa et al. [40] [39] have suggested that convexity is essential in F/G judgment. Measuring the shape convexity is thus necessary. The mathematical

definition of convexity says:

Definition 1.1.1 *A region $\Omega \in \mathbf{R}^2$ is called a convex region if the line segment joining any pair of points $A, B \in \Omega$ lies entirely in Ω . A region is called concave if it is not convex.*

This definition is binary: either a shape is or is not convex. For non-convex shape, i.e., for concave shapes, there is no concept of being more or less convex. In the experiments by Kanizsa and Gerbino [40] [39] (see Fig. 1.6(a1) & (a2)), all the shapes are not convex, but perceptually they appear to be more or less convex. Thus, a perceptual and continuous measure of convexity/concavity is requested. Voss [87], Sklansky [81], Gaafar [17], and Kim et al. [43] have suggested various definitions of convexity for discrete regions. But their proposed definitions are binary instead of continuous ones, therefore not appropriate for our purpose.

An important issue on analysis of shapes is scale. In the case of convexity, fine scale structures should not contribute too much in the final measure of shape convexity (see Fig. 1.7(b1) & (b2)). Realizing this scale property will ensure the robustness under micro perturbations of the boundary.

There are other works related to the issue of convexity. In Jacobs [36], Huttenlocher et al. [35], convexity has been used as criterion for feature grouping. But their works only select salient objects that are convex, and do not differentiate among con-

cave or convex shapes in a continuous manner. In Weiss [89], convexity was used as the criterion for F/G separation without considering the small-size preference for F/G (see next paragraph). Besides, the regional consideration was left out. All of their works can not account, nor aim to do so, for the Kanizsa and Gerbino experiments. Also, the concept of scale was missing in those approaches. Results can be altered by fine-scale perturbations.

In the research of shape analysis, various definitions of convexity was proposed for shape decomposition. One approach is to examine the difference between a shape and its convex hull. For instance, Rosin [70] used the area difference between the shape and its convex hull to define a continuous convexity. In Held et al. [29], a continuous measure of convexity, called *approximate convexity* was defined based on the fraction of region boundary coinciding with convex hull of the region. In many cases, shape attributes like area and perimeter are not enough to specify the shape properties. For the shape in Fig. 1.7(b3), the area difference between the shape and its convex hull is small, therefore, own a high convexity by the definition of Rosin [70]. We claim that missing of the internal axis information is the key for the failure. The shape in Fig. 1.7(b2) will obtain a bad convexity measure in the sense of approximate convexity by Held et al. [29].

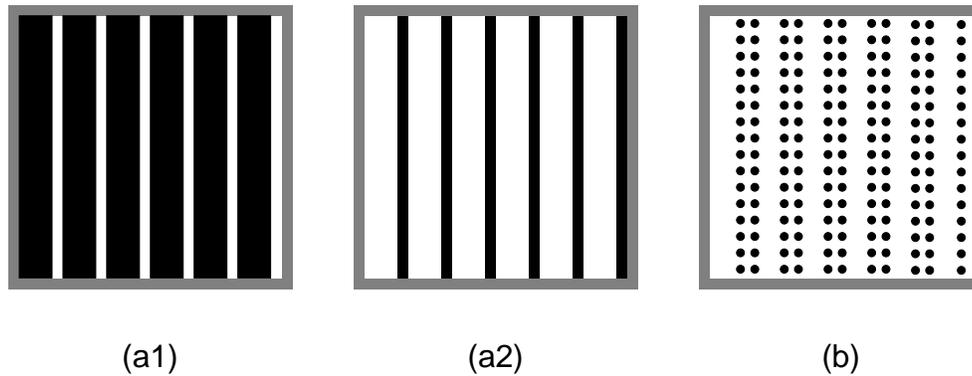


Figure 1.8. F/G separation that uses “size” as the factor to decide figures from background (adapted from Koffka [45]). (b) shows their edge sampling. By the law of proximity, we can build the same prediction which says white/black regions in (a1)/(a2) respectively tend to be perceived as figures rather than background. ■

Size and Proximity A Gestalt law discussed by Koffka [45] suggests that smaller objects are more easily perceived as figures. It gives the interpretation that size of a figure has a dominant effect on the result of F/G, as shown in Fig. 1.8(a1) & (a2). This law may be driven from the fact that usually in images, the objects of interest/focus are smaller than the rest of the image (the background). This law is a very effective rule and should be present on studying F/G separation.

Another simple law, but a very important one, is the Gestalt law of proximity. It says that features in an image that are closer should be grouped together. A typical

image example is given with set of black dots in white background and perceptually the closer ones are grouped together.

We can relate these two laws by the following argument. Suppose we have smaller objects in an image, once features such as edges are extracted, they will be grouped by the proximity law. The border ownership problem will then be resolved. I.e., edges that belong to larger objects will give the ownership to smaller objects since the latter ones will be grouped together. Then smaller objects become salient/figures. One can see this effect in Fig. 1.8(b). Once the edges are extracted and grouped by proximity, they will belong to the smaller width strips, and so smaller width strips will be figures and the larger width strips will be the background.

Symmetry Axis and Symmetry Measure The symmetry axis was first introduced by Blum [4] as the *grass fire* description. For a given curve $\mathcal{C}_0(s)$ parametrized by s , Kimia et al. [44] [80] extended the idea of Blum by giving a nice description of shape and medial axis via the evolution equation

$$\begin{aligned}\partial_t \mathcal{C} &= (\beta_0 - \beta_1 \kappa) \vec{N}, \\ \mathcal{C}(s, 0) &= \mathcal{C}_0(s).\end{aligned}$$

with \vec{N} denotes the normal vectors in boundaries. The coefficients β_0 and β_1 were used to produce the constant flow and the curvature flow respectively. The medial



Figure 1.9. Similarity test suggests that similarity is not a metric (adapted from Mumford [57]). It depends on the context which we are concerned. To ask the most similar shape to the shape (b), we obtain the shape (a) in the left group and end up with the shape (d) in the right group. ■

axis was created by various types of shocks. Sethian [73] also discussed the 2-D and 3-D flow propagations for shape description. Both of their works are based on the geometry fact provided by Gage et al. [18] and Grayson [22]. In addition, Liu et al. [52] found the symmetry axis and SA-tree by using double parameterization on boundaries via a variational energy. Both of the boundary and the regional considerations were made in their work. The approach with only the boundary consideration can lead to inappropriate result. For instance, tiny perturbations on boundaries can make a “matched neck” in a shape “mismatched” [57]. Other than the dynamic approach used in Kimia et al. [44] [80], we pursue a framework in which a static state will be considered and a map (a field) inside the shape will be computed. All measures will be derived from it.

For shape comparison, we emphasize that *shape similarity is not a metric*, which

was discussed in Mumford [57]. First, the matching process between shapes is not symmetric. If we say A is similar to B , what we mean is B is some kind of prototype in a category which includes A . Also, the similarity between A and B depends strongly on contexts. E.g., in Fig. 1.9, by given a shape, we are asked to choose the most similar one among others. In the experiment I(the left group), we are asked the most similar one to b among others. In the experiment II(the right group), e is removed and the same question is asked again. Many people reported that a is the most similar one to b in the experiments I while d is the most similar one to b in the experiment II. It gives ¶

$$d(b, a) < d(b, c), d(b, d), d(b, e) \quad \text{and} \quad d(b, d) < d(b, a), d(b, c).$$

A contradiction occurs. Therefore, to search for a universal concept of shape similarity is not appropriate.

1.1.1 A Continuous Simulation

We consider the shape representation from a brand-new approach. We call it the *continuous simulation*.

For the convexity measure, instead of using the binary definition adopted in mathe-

¶Suppose there is a non-symmetric metric “ d ” and $d(x, y)$ denotes the difference for “ x compared to y ” (not vice versa).

matics, we would like to search for a continuous version definition. By this definition, we can discuss “more convex” shapes and “less convex” shapes, for two non-convex ones. Our comparison will be meaningful even between two perfectly convex shapes.

For study of the symmetry axis, we would like to derive a new axis representation of shapes. On this representation, more than giving the information telling a point is on or off an axis, we would like to provide the information which can describe “how likely” a point is on the axis or “how strong” this axis is.

We will also provide a way to detect junction features. For the junction detection, we look for a representation which instead of giving a brutally 1 or 0 declaration, will provide the transition from low-curvature curves, high-curvature curves to curves with discontinuous curvature.

1.1.2 A Global Simulation

In our simulation, we consider the shape information as a whole. For convexity measurement, we would like to have the ability to discuss the coarse scale convexity.

For the symmetry axis, the information which we look for is based on a global consideration. So a small axis coming from noise in fine scales shall not be confused with main axes of the shape.

For the junction detection, we will not only concentrate on the finest scale fea-

tures. We are also able to pick up junction features in coarse scales, with a stronger declaration.

1.2 Problem Proposed

For a given single and still image, my proposed problems are

- i How can we characterize 2D shapes via local computations and diffusion processes where this characterization is particular useful in perceptual simulation?
- ii How can the Gestalt studies help us in defining such shape characterization ?
- iii How can we find one (or more than one) plausible organization(s) from an image based on the study of 2-D shapes ?

For the first two topics, in Chapter 2 and Chapter 3, two models will be proposed, called the *decay diffusion process* and the *orientation diffusion process* so as two measures called the *decay diffusion measure* and the *orientation diffusion measure*. The F/G separation and shape convexity will be discussed through these two measures. In Chapter 4, we use two ways, from the results of the decay process and the orientation diffusion process to generate the shape axis. Also, junction information can be derived from the result of orientation process. For the third topic, in Chapter 2 and 3, we use the result from shape analysis to predict the F/G. In Chapter 5, we

discuss salient shape selection and visual organization. We conclude our result and sketch the future work in Chapter 6.

Chapter 2

Convexity and Size in Figure/Ground

Separation

2.1 From Binary to Continuous Definitions

For a (compact) shape Ω and its boundary set $\partial\Omega$, we discuss the idea of convexity.

In mathematics, convexity is described by

$$tP_1 + (1 - t)P_2 \in \Omega . \quad \forall t \in [0, 1], P_1 \& P_2 \in \Omega \quad (2.1)$$

The condition of the definition can be changed to an equivalent one as $\forall t \in [0, 1]$, $\forall P_1 \& P_2 \in \partial\Omega$, checking only boundary points. Sometimes, only part of Ω or $\partial\Omega$ is visible due to restriction of image frame. E.g., in Fig. 2.2, only Ω_a is visible out of

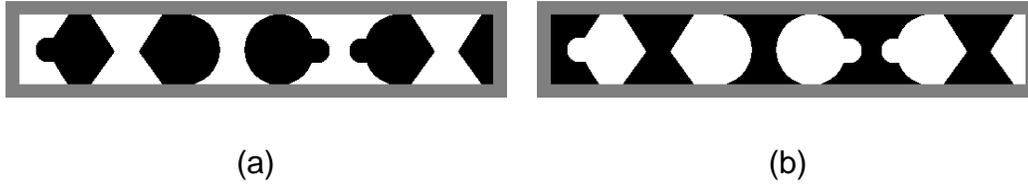
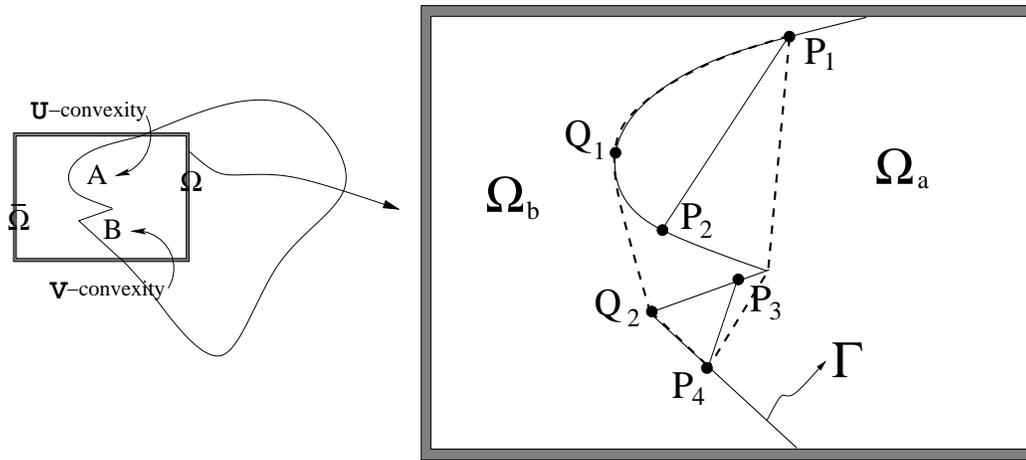


Figure 2.1. The figures adapted from Kanizsa et al. [39] [40] to illustrate the important role of convexity in F/G separation. Human results show the black/white area in (a)/(b) respectively is more likely to be considered as figures. ■

a complete region Ω with its partially visible boundary curves $\Gamma \subset \partial\Omega$. A definition similar to Eq. 2.1 can be formulated for the partial cases such as the region Ω_a or its partial boundary Γ . The equivalence argument between checking boundary points or interior points can also be made.

The idea of convexity for F/G separation was proposed by Kanizsa et al. [39] [40] in their convexity versus symmetry experiments. In Fig. 1.6(a1) & (a2) and Fig. 2.1, the idea of “convexity” (other than symmetry) is proposed as the dominant criteria for F/G decision. However, the convexity of these regions does not follow the mathematical definition of convexity in the sense that most regions are considered concave. Therefore we need a continuous version definition for convexity which allows us to distinguish between “more” and “less” convex shapes. Various definitions will be discussed.



(a)

Figure 2.2. The convexity examination of the region Ω and its complement $\bar{\Omega}$ sitting on two sides of the boundary $\partial\Omega$. The regions Ω , $\bar{\Omega}$ and boundary $\partial\Omega$ may be only partially explored and end up with Ω_a , Ω_b and Γ respectively. We measure convexity by shortest paths connecting two points through two (partial) regions Ω_a and Ω_b separately. ■

Measuring convexity via shortest path

Let us analyze Fig. 2.2 and try to propose a continuous measure of convexity. We have a rectangular frame, an image \mathbf{I} , divided into two regions Ω_a and $\Omega_b \equiv \mathbf{I} - \Omega_a$, both with the same area (within the frame) and a common edge boundary Γ . Before we can decide the border ownership problem, we assume $\Gamma \subset \Omega_a$ and $\Gamma \subset \Omega_b$. Both regions, if interpreted as shapes, are concave and yet, we perceive one region as more

convex than the other.

We observe that in Fig. 2.2(a), length of the segment $\overline{P_1P_2} \subset \Omega_a$ is shorter than length of the curve $\widetilde{P_1P_2} \subset \Omega_b$. For points P_3 and P_4 , the straight segment $\overline{P_3P_4} \subset \Omega_a$ connects two points in the convex side, but a longer path, the segments $\overline{P_3Q_2} \cup \overline{Q_2P_4} \subset \Omega_b$ is chosen in the concave side. The smaller the angle $\angle Q_2$ is, the bigger the difference between the length of the two shortest paths (s.p.) in both regions. The same is true for the case where (the integral of) curvature in the curve $\widetilde{P_1P_2}$ gets larger. In Fig. 2.2(a), based on the evidence points P_1 and P_4 , the region Ω_a is more convex than the region Ω_b by owning the shorter path $P_1P_3P_4 \subset \Omega_a$ compared to the path $P_1Q_1Q_2P_4 \subset \Omega_b$.

We then propose a definition of convexity as follow. The version which we will give is for a complete shape Ω . A similar version for partial explored Ω_a , Ω_b or Γ can also be formulated similarly.

Definition 2.1.1 *Given a shape Ω and its boundary $\partial\Omega$, we estimate the shortest path $\text{sp}(P_i, P_j)$ between pair of points P_i, P_j where $\text{sp}(P_i, P_j) \subset \Omega$ & $P_i, P_j \in \partial\Omega$. Its length is denoted by $E_{\text{sp}}(P_i, P_j)$. Suppose $\partial\Omega$ is parameterized by its arc length, we define s.p. convexity as the integral of such distance over all pairs of points on $\partial\Omega$,*

$$\text{CV}_{\text{sp1}}(\partial\Omega) = \int_0^L \int_0^L E_{\text{sp}}(P(s_1), P(s_2)) ds_1 ds_2, \quad (2.2)$$

where L is length of $\partial\Omega$. It can be normalized by the Euclidean distance $\|P(s_1) -$

$P(s_2)$ ||, which gives

$$\text{CV}_{\text{sp2}}(\partial\Omega) = \frac{\int_0^L \int_0^L E_{\text{sp}}(P(s_1), P(s_2)) ds_1 ds_2}{\int_0^L \int_0^L \|P(s_1) - P(s_2)\| ds_1 ds_2}. \quad (2.3)$$

A rough estimation gives

$$\text{CV}_{\text{sp1}}(\partial\Omega) \leq ML^2,$$

where M denotes the maximum length E_{sp} for pair of points in $\partial\Omega$. For a perfectly convex shape Ω , M is its diameter. For the normalized version, we have $\text{CV}_{\text{sp2}}(\partial\Omega) = 1$ for convex shapes, and $\text{CV}_{\text{sp2}}(\partial\Omega) > 1$ for concave shapes. It provides a continuous definition of convexity. On the other hand, we can write a variant of the definition which checks all points in Ω instead of $\partial\Omega$. We can also consider the case when image frame exists. We compare convexity for two (partially explored) regions Ω_a and Ω_b with a shared edge Γ by computing convexity of them via either the boundary or the regional version of the definition.

Let us take a look at these measures. Suppose we sample $\partial\Omega$ by N points, the computation of these measures requires $O(N^2)$ operations and it can not be parallelized below $O(N)$ computations. One needs to verify, for each point $P(s(i))$, all the other points $P(s(j))$'s. Moreover, it assumes that the shape is given. I.e., it does not provide an automatic way to extend this approach when shapes need to be detected, e.g., the case of noisy images or illusory figures where the shape is not clear

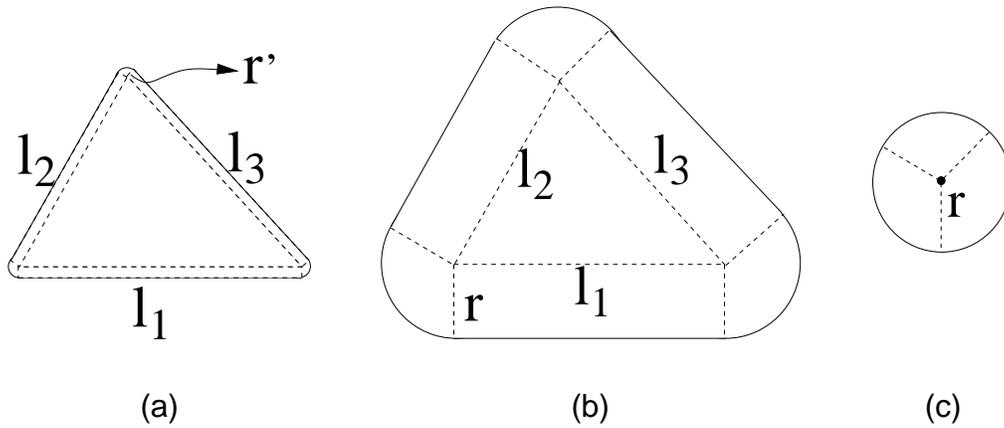


Figure 2.3. Shapes with different convexity, examined by $CV_{\kappa 2}$. We have $CV_{\kappa 2}(\text{pseudo triangle}) > CV_{\kappa 2}(\text{rounded triangle}) > CV_{\kappa 2}(\text{circle}) = 1$ and $CV_{\kappa 2}(\text{pseudo triangle}) \rightarrow +\infty$ as $r' \rightarrow 0$. ■

or is partially broken.

Alternatively, we would like to develop a measure of convexity that is based on a distributed system, based on local computations that can be added to yield global measures. For that we invoke the machinery of diffusions processes and Markov random fields.

Measuring convexity via curvature on boundary

We propose another version of the definition based on curvature information on boundaries.

Definition 2.1.2 Given a shape Ω and its boundary $\partial\Omega$, parameterized by the arc

length s . The curvature convexity is given by

$$CV_{\kappa 1}(\partial\Omega) = \frac{1}{2\pi} \int_0^L |\kappa(s)| ds, \quad (2.4)$$

where κ denotes curve curvature on $\partial\Omega$. Or we can try a quadratic form*

$$CV_{\kappa 2}(\partial\Omega) = \frac{L}{4\pi^2} \int_0^L |\kappa(s)|^2 ds. \quad (2.5)$$

For shapes with smooth boundary, we have $CV_{\kappa 1}(\partial\Omega) = 1$ for any convex shapes and $CV_{\kappa 1}(\partial\Omega) > 1$ for concave shapes. For non-smooth boundaries where some undefined curvature points were occurred, we can make them “curved” and compute the convexity on the perturbed ones. It will be well-defined. It gives us

$$CV_{\kappa 1}(\text{circle}) = CV_{\kappa 1}(\text{triangle}) = 1.$$

The second definition $CV_{\kappa 2}$ can only be discussed on smooth curves. For a circle of arbitrary radius, we have $CV_{\kappa 2}(\text{circle}) = 1$. In fact, $CV_{\kappa 2}$ is invariant under scaling.

For figures in Fig. 2.3, we have

$$CV_{\kappa 2}(\text{pseudo triangle}) > CV_{\kappa 2}(\text{rounded triangle}) > CV_{\kappa 2}(\text{circle}) = 1.$$

*A similar one called *average bending energy*, normalized by the curve length instead, was given by $E(\partial\Omega) = \frac{1}{L} \int_0^L |\kappa(s)|^2 ds$ in Young et al. [93]. Our definition is made to be size-invariant.

The computation gives $CV_{\kappa^2}(\text{rounded triangle}) = 1 + (l_1 + l_2 + l_3)/2\pi r$. So we have

$$\begin{aligned} CV_{\kappa^2}(\text{rounded triangle}) &\rightarrow +\infty && \text{if } r \rightarrow 0 \\ CV_{\kappa^2}(\text{rounded triangle}) &\rightarrow 1 && \text{if } l_1 + l_2 + l_3 \rightarrow 0. \end{aligned}$$

In fact, Young et al. [93] proved that circle shape will minimize the criteria, either the curvature convexity CV_{κ^2} or their average bending energy, if the curve length is fixed. It is possible to have a convex shape owning bigger measure of CV_{κ^2} than a concave shape.

Let us check how effective these measures are. First of all, both definitions are not appropriate in the F/G problem, either for measuring convexity of complete regions Ω & $\bar{\Omega}$ or incomplete regions Ω_a & Ω_b with shared edges on an image (Fig. 1.6(a1) & (a2) and Fig. 2.1). Those regions will have the exact same value on both sides no matter what we consider, the smoothed or non-smoothed versions. Again, we need to know the shape before we can execute the computation involved in both definitions.

Compactness

We discuss a related criteria called *shape compactness*[†], which has been discussed in Bacus et al. [2], Green [23], Rosenfeld [69] and Bribiesca [7].

[†]In Young et al. [93], the average bending energy, similar to the convexity definition CV_{κ^2} was used to substitute the conventional definition of shape compactness.

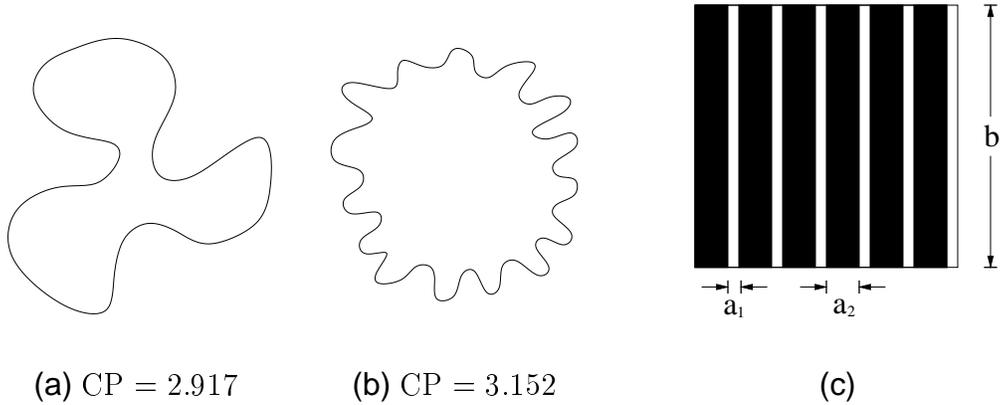


Figure 2.4. The Shape (a) is considered more compact than the shape (b), suggested by the conventional shape compactness measure CP. (c) the measure CP will not select the white regions over the black regions as figures. ■

Definition 2.1.3 Given a shape Ω and its boundary $\partial\Omega$. Its shape compactness is defined by

$$CP(\partial\Omega) = \frac{|\partial\Omega|^2}{4\pi \cdot |\Omega|}, \quad (2.6)$$

with $|\partial\Omega|$ and $|\Omega|$ denoting perimeter and area of Ω respectively.

A shape with smaller value is considered to be a more compact shape. It is a continuous definition with the similar behavior to $CV_{\kappa 2}$ by the fact that the circle also minimize the criteria with $CP(\text{circle}) = 1$ for any circle shapes. However, it is not appropriate in vision applications as two significantly different shapes can own the same measure [93] [7]. In Fig. 2.4, the shape (a) is considered to be more compact

than the shape (b), if measured by CP. To apply this measure to F/G simulation, it will not give a correct prediction. For example, we can design a F/G experiment such as in Fig. 2.5(a) where both regions own the same area. It gives the same compactness for both regions if only the edge boundaries are considered as shape boundaries. If we do consider the image frame as part of the shape boundaries, we will not select the white regions in Fig. 2.4(c) as the favored regions. Their measures are $CP_{\text{white}} = 4(a_1 + b)^2 / (4\pi a_1 b)$ and $CP_{\text{black}} = 4(a_2 + b)^2 / (4\pi a_2 b)$. It gives $CP_{\text{white}} > CP_{\text{black}}$ if $b^2 > a_1 a_2$ or $b > a_1, a_2$, which is the case in Fig. 2.4(c). Namely, for this size/proximity experiment, we will collect a result against the perceptual phenomenon, predicting black regions as figures. In fact, the area term is assigned as the denominator. Therefore, larger regions will be favored given the same perimeter. To accomplish the task of F/G separation, a measure with the small-size preference is preferred, if a measure with size preference is offered. On the other hand, the convexity definition CV_{sp1} , given by examining shortest paths within shapes, is more appropriate than the shape compactness CP, for deciding F/G. Smaller shapes will be favored, with shorter shortest-paths.

U-convexity and V-convexity

Let us discuss two different types of convexity. In Fig. 2.2, the region A and B are considered different as on boundaries of the region A (an arc or a **U**), curvature

$\kappa = \dot{\theta}(s)$ (suppose orientation is parameterized by the arc length s) shares the same sign while boundaries of B (an angle or a \mathbf{V}) has curvature equal to 0 except at the angle peak. The first kind is called arc convexity or \mathbf{U} -convexity and the second kind is called angle convexity or \mathbf{V} -convexity. We say that the convexity is concentrated on a single point on the boundaries with \mathbf{V} -convexity (region B) while it is more uniformly distributed over the boundaries with \mathbf{U} -convexity (region A). It tells one important difference between them. When a small aperture is randomly put on these two kinds of boundaries, we have a better chance of knowing where the convex side is by putting it on the boundaries with \mathbf{U} -convexity than the boundaries with \mathbf{V} -convexity. We argue that this local suggestion makes the \mathbf{U} -convexity more favorable than the \mathbf{V} -convexity. Another evidence of our argument comes from the experiments provided by Stevens et al. [82] or Fig. 3.2. The concave cusp feature (more like owning \mathbf{V} -convexity) suggests that it is the background side rather than the figures. Also, in the Kanizsa image in Fig. 1.6(a1), we do have similar number of \mathbf{U} 's (arcs) or \mathbf{V} 's (angles) for the black and white regions. Although, the perception provides the preference of black regions (with more \mathbf{U} -convexity) as figures.

In the next section and in Chapter 3, we proceed to construct two kinds of convexity measure. The *decay diffusion measure* is derived by a process called the *decay diffusion process* and the *orientation diffusion measure* is from a process working on the domain of $\mathbf{I} \times [0, 2\pi)$, called the *orientation diffusion process*. They are the 1st

and 2nd order of Markov processes respectively. Those images and perceptual experiments that we have discussed will be studied again under these measures. As we have discussed, the measures that we will propose is based on a distributed system, a highly-parallelizable algorithm. Besides, the frameworks should not be recognized as having only the goal to derive the (convexity) measures for F/G prediction. Other features of shapes, such as symmetry or junctions, can also be obtained as side results. It will be discussed in Chapter 4.

2.2 Decay Diffusion Process

We seek a computational model to measure convexity of a complete or partial shape Ω or Ω_a in a continuous manner. Based on boundary input, the first step of our framework is designed to generate a field inside[‡] the shape while the value of each point can be viewed as convexity on that point. The field called *decay diffusion field* is derived by a leaking energy diffusion process where the discrete formulation can link this process to a 2-D decay walk, starting from the boundaries (Appendix B). It means that the computation during this part can be done locally through interactions between adjacent pixels. A judgment of convexity, called convexity measure is given

[‡]The inside (or outside) means just one side of an closed contour, depending on where the interesting part is. For a partial shape, the interesting region is restricted within the image.

by computing (point-wise) entropy of the field belonging to Ω . A shape with a smaller measure is called a more convex shape. More discussions will show that not only convexity, but also size preference, will be caught by this measure. As we know, both are important to F/G. From now on, we will not clearly distinguish a complete shape Ω from a partial one Ω_a or $\partial\Omega$ from Γ unless it is necessary to do so. We use Ω and $\partial\Omega$ throughout our formulation.

2.2.1 Decay Diffusion Process as Energy Minimization

Notations and Assumptions

Given a binary synthesized image $\mathbf{I} \equiv [0, A] \times [0, B] \subset \mathbf{R}^2$ with the characteristic function $I(\mathbf{x})$ of the region Ω as intensity function. The feature set of the image, is given by the edge/boundary set $\partial\Omega = \{\mathbf{x} \in \mathbf{I} : \exists \mathbf{N}_{\mathbf{x}} \text{ s.t. } I(\mathbf{x}) \neq I(\mathbf{N}_{\mathbf{x}})\}$ where $\mathbf{N}_{\mathbf{x}}$ is drawn from the neighborhood set of the point \mathbf{x} . A discretization in the Euclidean grid gives four neighbors as $\mathbf{N}_{\mathbf{x}} = \{(x_1+1, x_2), (x_1-1, x_2), (x_1, x_2-1), (x_1, x_2+1)\}$ for the point $\mathbf{x} = (x_1, x_2)$. We record this edge set by its characteristic function $e(\mathbf{x})$, which is,

$$e(\mathbf{x}) = \begin{cases} 1, & \text{if } \mathbf{x} \in \partial\Omega \\ 0. & \text{if } \mathbf{x} \in \Omega - \partial\Omega \end{cases} \quad (2.7)$$

Rest of the formulation is set related to this function (as well as the intensity function

I). The input σ_0 , also called the inducer or hypothesis function, is given by

$$\sigma_0(\mathbf{x}) = \begin{cases} 1, & \text{if } \mathbf{x} \in \partial\Omega \\ 0. & \text{if } \mathbf{x} \in \Omega - \partial\Omega \end{cases} \quad (2.8)$$

The area of Ω is recorded by $|\Omega|$, and $|\partial\Omega|$ denotes length of the edge set.

Variational Formulation

We adopt a variational formulation in our framework. Solving of the decay diffusion field is the minimization process on a given energy. The energy functional which we are interested in is

$$E_{\text{decay}}(\sigma|\sigma_0, \Omega, \lambda) = \int_{\Omega} \Lambda(\mathbf{x}) (\sigma(\mathbf{x}) - \sigma_0(\mathbf{x}))^2 + M(\mathbf{x}) \|\nabla\sigma(\mathbf{x})\|^2 d\mathbf{x}, \quad (2.9)$$

for a given input σ_0 . The function $\sigma : \Omega \rightarrow \mathbf{R}$ is the field we need to evaluate. The function $\Lambda(\mathbf{x})$ in the first term (data fitting) is given by

$$\Lambda(\mathbf{x}) = \begin{cases} \delta_2(\mathbf{x}), & \text{if } \mathbf{x} \in \partial\Omega \\ \lambda, & \text{if } \mathbf{x} \in \Omega - \partial\Omega \end{cases} \quad (2.10)$$

where $\delta_2(\mathbf{x})$ stands for the 2-D Dirac delta function s.t. $\int_{\partial\Omega} \delta_2(\mathbf{x}) d\mathbf{x} = 1$. A complete definition can be found in Appendix A. In the discrete case, it is provided by $\Lambda(\mathbf{x}) = 1/\epsilon, \mathbf{x} \in \partial\Omega$ for a small constant ϵ , called the delta function coefficient (Appendix B). On the inside part of Ω , we choose $\Lambda(\mathbf{x}) = \lambda$, a small positive constant called the decay coefficient. The function $M(\mathbf{x})$ in the second term (smoothness),

called smoothness function is simply assigned as $M(\mathbf{x}) = 1$ for the homogeneous case.

The first part stands for fidelity to the input. The second part stands for the smoothness assumption, minimizing square of the gradient value $\|\nabla\sigma(\mathbf{x})\|$. Let us reorganize the energy functional into two parts $\Omega - \partial\Omega$ and $\partial\Omega$ and write down its Euler-Lagrange equation

$$\begin{aligned}\Delta\sigma(\mathbf{x}) &= \lambda\sigma(\mathbf{x}), & \mathbf{x} \in \Omega - \partial\Omega \\ \sigma(\mathbf{x}) &= \sigma_0(\mathbf{x}), & \mathbf{x} \in \partial\Omega\end{aligned}\tag{2.11}$$

where $\Delta = \partial_{xx} + \partial_{yy}$ is the Laplacian operator. When Ω touches image frame, an *absorbing barrier* is assumed for the boundary condition.

We solve the equation by the finite difference method which will be demonstrated in Appendix B. The solution, written as $\sigma^*(\mathbf{x})$ is the decay diffusion field we need. Practically, for the convexity comparison of two shared-edge regions within an image, we solve the field for both regions Ω and $\mathbf{I} - \Omega$ simultaneously by discretely assigning $\sigma_0 = 1$ and -1 to both sides of the edges. As for the original image Fig. 2.5(a), we assign black/white color as $-1/1$ to pixels on both sides of the edges, and assign 50% gray or 0 to pixels in $\mathbf{I} - \partial\Omega$ representing $\sigma_0(\mathbf{x})$, so called the inducers or the hypothesis function. A simple example of F/G image is shown in Fig. 2.5.

2.2.2 Entropy Criteria

In order to obtain a convexity measure, first, we convert[§] the field $-1 \leq \sigma^*(\mathbf{x}) \leq 1$ into a probability distribution at each pixel, via a linear mapping

$$p^*(\mathbf{x}) = \frac{1}{2}(1 + \sigma^*(\mathbf{x})). \quad (2.12)$$

For a F/G problem, $p^*(\mathbf{x})$ can be treated as the probability of being in front (salient). In Appendix B, we will give a random walk formulation to describe the solution σ^* as well as the probability field p^* . As we will show in Appendix B, $p^*(\mathbf{x})$ is the probability to reach any “figure inducers” before reach any “background inducers”, for a walk starting at \mathbf{x} . Either with or without broken edges, the probability to reach “background inducers” can be greater than zero. The reason is that the walks stopped in interior points will be considered having half of the probability contributing to the “background part”, as it lacks of knowledge to commit. Therefore, it is natural to characterize the region Ω as the set $\{\mathbf{x} : p^*(\mathbf{x}) \geq 0.5\}$ and $\mathbf{I} - \Omega$ as the set $\{\mathbf{x} : p^*(\mathbf{x}) < 0.5\}$. When two regions on both sides of edges are solved simultaneously, the (discrete) pixels on the edges, where $p^*(\mathbf{x}) = 1/0$ (or $\sigma^*(\mathbf{x}) = 1/-1$) is considered as the strongest commitment to be salient/background respectively. The place with

[§]It is possible to have $\sigma^*(\mathbf{x}) \leq 0$ as we are solving the field simultaneously on both sides of the edges. Later when we introduce the mechanism of organization selection for illusory figures, we have another case of $\sigma^*(\mathbf{x}) \leq 0$. Please see Chapter 5 for more discussion.

$p^*(\mathbf{x}) = 0.5$ (or $\sigma^*(\mathbf{x}) = 0$) stands for the place being neutral or non-committant.

The criteria is given by calculating (point-wise) entropy of the shape Ω ,

$$\mathcal{S}(p^*) = -\frac{1}{|\Omega|} \int_{\Omega} p^*(\mathbf{x}) \log p^*(\mathbf{x}) + (1 - p^*(\mathbf{x})) \log(1 - p^*(\mathbf{x})) d\mathbf{x}. \quad (2.13)$$

A shape with *smaller* measure \mathcal{S} is considered more convex. As we said, we can substitute the sub-index of the summation by an equivalent condition $p(\mathbf{x}) \geq 0.5$. We compute this measure for both regions Ω and $\mathbf{I} - \Omega$, indicated by \mathcal{S} and $\overline{\mathcal{S}}$ respectively. Conventionally, $p^*(\mathbf{x}) = 1/0$ is reserved for convex/concave side respectively and we use \mathcal{S} to denote the measure for convex side and $\overline{\mathcal{S}}$ for concave side if the convex or concave side can be easily distinguished. The sharper the diffusion, the closer to $1/0$ $p^*(\mathbf{x})$ for $\Omega / \mathbf{I} - \Omega$ is, the smaller the entropy we can obtain. The entropy is a point-wise entropy or entropy of the region normalized by the area of Ω in the discrete case.

2.2.3 Convexity and Decay Coefficient

We analyze our model by the random walk formulation. In Appendix B, a discrete form energy functional will be given and the process becomes a decay walk in 2-D. From the random walk viewpoint, the solution $\sigma^*(\mathbf{x})$ is *the summation of probability carried by all the walks starting from inducers on boundaries (may be with different weights) to the pixel \mathbf{x}* . The decay coefficient λ controls how strong the decay effect

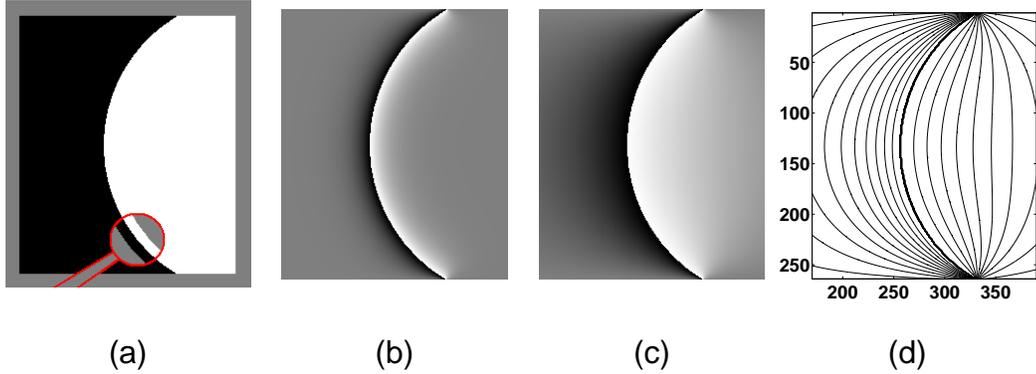


Figure 2.5. (a) F/G image with two regions, one is convex and concave is another. Both of them own the same area and the number of inducers. (b) (c) the maps according to the typical hypothesis shown within the magnifier in (a), with white/black/gray representing the salient/background/neutral assumption respectively. We set $\lambda = 5 \times 10^{-3}$, 1×10^{-4} in (b), (c) respectively. The maps show the decay when it is away from the arc (stronger decay in (b)). The entropy (convex/concave or S/\bar{S}) for them are (b) 0.972/0.974 and (c) 0.880/0.916. (d) the iso-contour (iso- $\sigma^*(\mathbf{x})$) of (c). It shows how the diffusion expands larger distance on the convex side. ■

is where particles are with the chance of $\Lambda(\mathbf{x})/(\Lambda(\mathbf{x}) + \mu(\mathbf{x}, \mathbf{N}_{\mathbf{x}}))$ to vanish ¶ after each jump (Eq. B.15) or simply $\lambda/(\lambda + 4)$ for pixels not on the boundaries. A smaller λ creates less particle vanishing and more diffusion due to the fact that each walk can

¶The function $\mu(\mathbf{x}, \mathbf{N}_{\mathbf{x}})$ records number of neighbors for the point \mathbf{x} , when no edge blocking is assumed, we got a constant 4, see Appendix B.

remain for a longer period. The effect is illustrated in Fig. 2.5(b) and (c). The idea why convexity is caught by our model can be presented in the following way. First of all, random walks tend not to cross over boundaries. The chance of particles vanishing on boundaries is given by $1/(1 + \epsilon\mu(\mathbf{x}, \mathbf{N}_x))$ with a small ϵ . So we can assume that the walks are more likely to visit the boundaries only once by choosing a small enough ϵ . Therefore, we can discuss walks in the convex side and the concave side separately. For walks starting from boundary inducers to points in Ω or $\mathbf{I} - \Omega$, shorter ones in the convex side will be accumulated which mean less decay while longer ones will be chosen in the concave side (see Fig. B.1). In Fig. 2.5(d), it can be seen that the convex side has wider iso-contour than the concave side, which means a sharper diffusion and a smaller convexity measure.

Continuous Convexity Measure

To understand why our criterion gives a continuous convexity measure, we test our measure on parallel pentagon images with various angles from $\pi/6$ to π . In Fig. 2.6(d), the difference of the convexity measure gets bigger when the angle becomes smaller. The diffusion results for angles equal to $\pi/6$ and $5\pi/6$ are shown in Fig. 2.6(a2) & (b2) respectively. Smaller angles provide stronger intensities of diffusion.

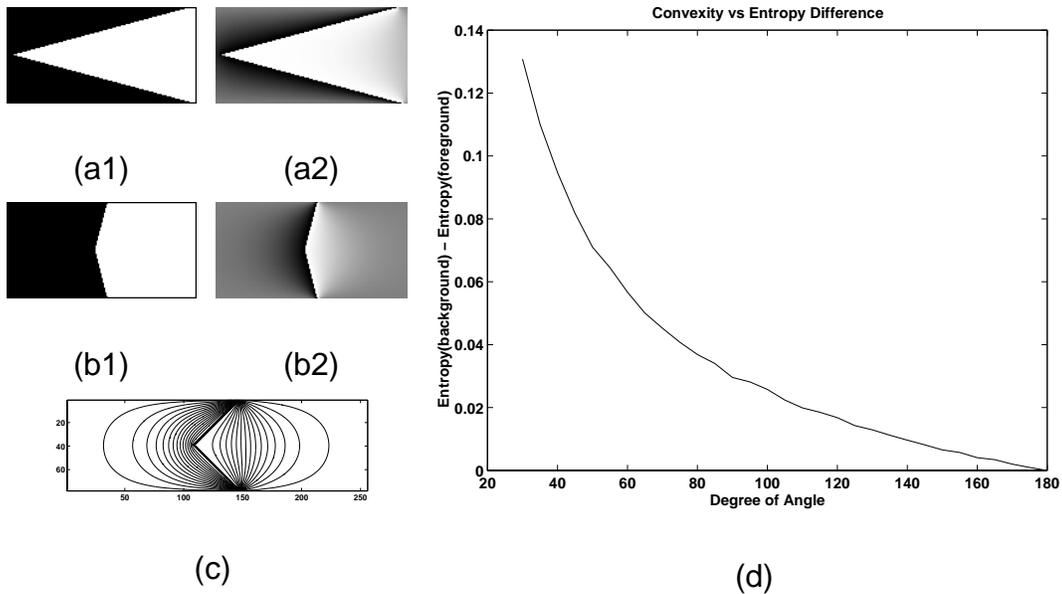


Figure 2.6. (a1) (b1) source images with angles of $\pi/6$ and $5\pi/6$ and (a2) (b2) their diffusion result (only middle parts are shown). The entropy values for (a2) and (b2) are 0.721 / 0.870 (convex/concave or S / \bar{S}) and 0.944 / 0.950 respectively, with $\lambda = 1 \times 10^{-4}$. (c) The iso-contour with angle equal to $\pi/2$ (complete map). (d) The difference between entropy for the convex and concave regions as a function of the angle (“inverse of convexity”). ■

Figure/Ground Separation

Let us check the image of “convexity versus symmetry”, adopted by Kanizsa [39] in Fig. 2.7(a), The result supports the idea proposed by Kanizsa, which tells us that the black regions rather than white regions tend to be perceived as figures, and convexity

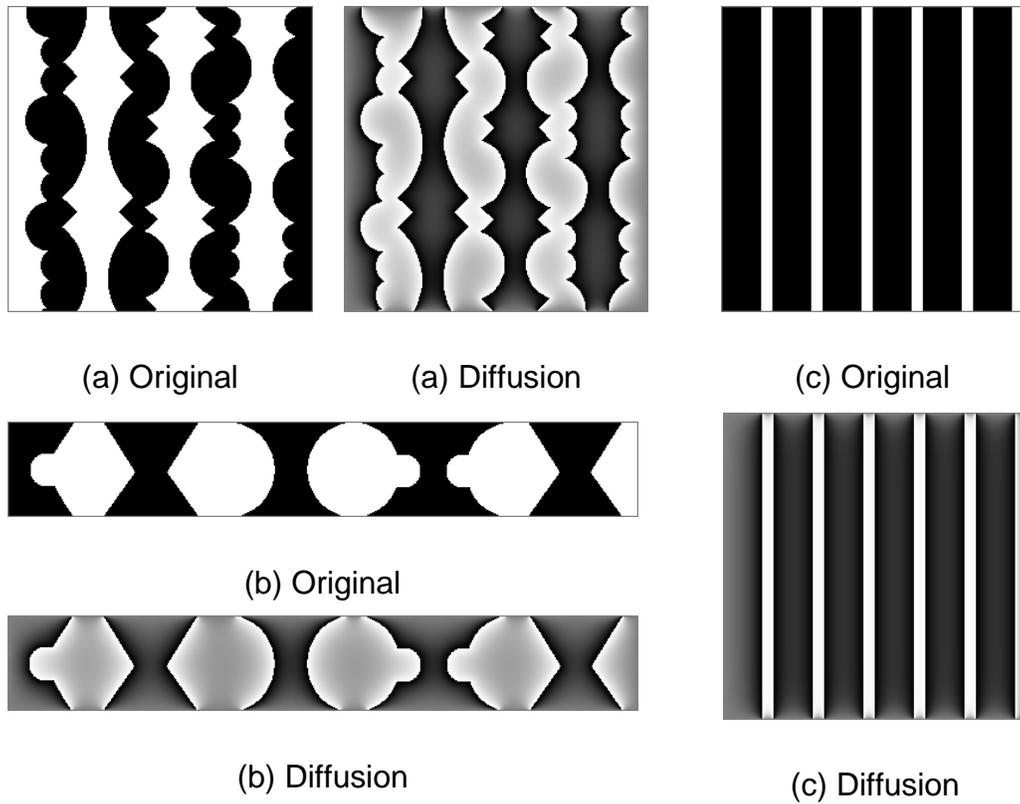


Figure 2.7. F/G separation by the decay process, with the convexity and size preferences. (a) the Kanizsa figure for the convexity-versus-symmetry experiments. The entropy values are $S/\bar{S} = 0.546/0.560$ for the black/white regions respectively. Beware that we have more diffusion near the neck area. (b) another F/G dominated by the figural convexity. The decay measure gives the measure as $S/\bar{S} = 0.723/0.747$, favoring the “shell” shapes, the white regions. (c) The size preference is given by the measure of $S/\bar{S} = 0.220/0.602$, favoring the white/small strips. ■

is the key for this phenomenon. Another example given in Fig. 2.7(b), prefers the “shell” shapes. We discuss other properties of our model in the following sections.

2.2.4 Size and Proximity

As we mentioned, the idea of using size as a criteria for F/G separation is highly related to the Gestalt law of proximity. In our model, as we can realize in the random walk argument that proximity between the inducers of a shape and any points in the shape is the key for large intensities of diffusion. So for the size preference test in Fig. 2.7(c), without any effect of convexity, our model will prefer the white regions which are the regions with smaller size. This is because the white regions have inducers closer to each other, where inducers will support the diffusion in the nearby regions before the particles vanish in an exponential decay. An extreme case can be given by neglecting the smoothing in Eq. 2.9. We have $\sigma^*(\mathbf{x}) = \sigma_0(\mathbf{x})$. The convexity measure is given by

$$\mathcal{S}(p^*) = \frac{|\Omega| - |\partial\Omega|}{|\Omega|} = 1 - \frac{|\partial\Omega|}{|\Omega|}. \quad (2.14)$$

A smaller area (or more number of inducers) will yield a smaller measure. It matches our visual perception, a known fact proposed by Koffka [45].

Size Invariance and Tuning of λ

We would also like to discuss the relation between *similar* shapes with different sizes.

It can be seen from the Euler-Lagrange equation in Eq. 2.11. In Eq. 2.11, for a scaling of s from the shape Ω to Ω' , we substitute \mathbf{x} by $\mathbf{x}' = s\mathbf{x}$ and let $\sigma'(\mathbf{x}') = \sigma(\mathbf{x})$, also $\sigma'_0(\mathbf{x}') = \sigma_0(\mathbf{x})$. A new equation is obtained as

$$\begin{aligned} s^2 \Delta \sigma'(\mathbf{x}') &= \lambda \sigma'(\mathbf{x}'), & \mathbf{x}' \in \Omega' - \partial\Omega' \\ \sigma'(\mathbf{x}') &= \sigma'_0(\mathbf{x}'). & \mathbf{x}' \in \partial\Omega' \end{aligned} \quad (2.15)$$

It means that if we choose a new decay coefficient $\lambda' = \lambda/s^2$, we can collect a similar field σ'^* to the original field σ^* with $\sigma'^*(\mathbf{x}') = \sigma^*(\mathbf{x})$.

To speak of the convexity measure, we consider Eq. 2.13. For two probability maps p^* and q^* with $p^*(\mathbf{x}) = q^*(\mathbf{x}')$ derived from σ^* and σ'^* on Ω and Ω' respectively, we have

$$\begin{aligned} \mathcal{S}(q^*) &= -\frac{1}{|\Omega'|} \int_{\Omega'} q^* \log q^* + (1 - q^*) \log(1 - q^*) d\mathbf{x}' \\ &= -\frac{1}{s^2 |\Omega|} \cdot s^2 \int_{\Omega} p^* \log p^* + (1 - p^*) \log(1 - p^*) d\mathbf{x} = \mathcal{S}(p^*). \end{aligned} \quad (2.16)$$

Therefore, the same convexity measure can be obtained for both shapes if the decay coefficient is carefully adjusted.

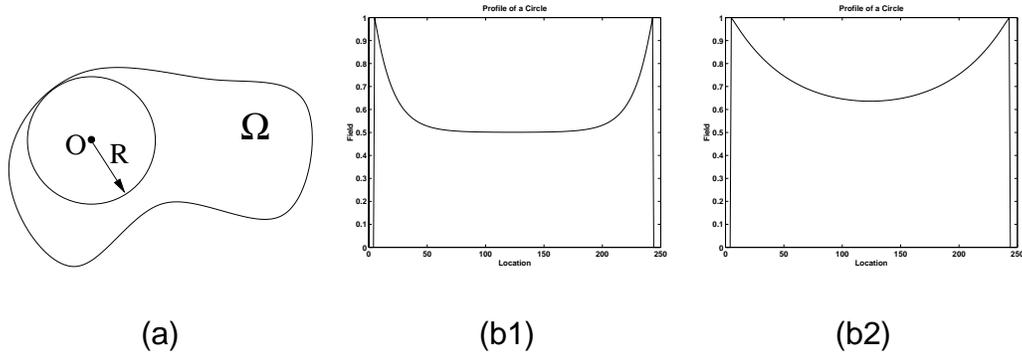


Figure 2.8. (a) The point O , center of an inscribing circle of radius R has an exponential decay as R getting bigger. (b1) & (b2) Profiles of the diffusion field $\sigma^*(\mathbf{x})$ at diameter of a circle shape when the decay coefficient is assigned by $\lambda = 5 \times 10^{-3}$ and $\lambda = 5 \times 10^{-4}$ respectively. Their convexity measures are $\mathcal{S}_{(b1)} = 0.890$ and $\mathcal{S}_{(b2)} = 0.667$. ■

2.2.5 The Speed of Decay

We would like to estimate speed of the decay in the decay process. For a shape Ω , a point $O \in \Omega$ with $d(O, \partial\Omega) = R$, shown in Fig. 2.8(a), the decay in O can be bounded by

$$\sigma^*(O) \leq e^{-\frac{\sqrt{\lambda}R}{2}}. \quad (2.17)$$

For a circle shape, the decay in the circle center O can be approximated by

$$e^{-\frac{\lambda R^2}{4}} \leq \sigma^*(O) \leq e^{-\frac{\sqrt{\lambda}R}{2}}. \quad (2.18)$$

Profiles of the fields in the diameter are shown in Fig. 2.8(b1) & (b2). For a shape, this

exponential decay weakens the possible description of its coarse scale structures. For “highly convex” shapes with minor perturbations such as the shapes in Fig. 1.7(b1) & (b2), an appropriate convexity measure should depend on not only fine scale convexity, but more importantly, coarse scale convexity. To improve this, in next chapter, the orientation measure will be introduced based on the orientation diffusion process.

2.2.6 Implementations

We choose the parameter set as the decay coefficient $\lambda = 5 \times 10^{-3}$ and the delta function coefficient $\epsilon = 10^{-1}$. This set is fixed throughout all our experiments unless different notification is given.

For the F/G problem, we design the input images with the same size in both of the black and white regions. By doing this, we remove the size bias which may be created in Eq. 2.14. Also, we would like to discuss the boundary condition with more details.

Boundary Condition

As we mentioned, all experiments are tested under the assumption of absorbing barriers for image frames, which has $\sigma^*(\mathbf{x}) = 0$ if $\mathbf{x} \in \partial\mathbf{I}$. To be more careful, we can choose the frame used for computation to be a little bigger than the image frame. The principle is to choose a frame for computation big enough s.t. we can assume

all walks going beyond the (computation) frame are likely to be stopped. So those walks can be ignored in computation without losing too much precision for the final measurement. When the image frame is extended, in Eq. 2.13, we use the criteria of $\{\mathbf{x}:p(\mathbf{x}) \geq 0.5\}/\{\mathbf{x}:p(\mathbf{x}) < 0.5\}$ to characterize figure/ground respectively.

In perception, this assignment is interpreted as the F/G phenomenon prevailing below image frames with certain range. For an evidence, we can refer to Fig. 2.7(c) where we can perceive a fatter strip near the boundaries in the right end.

Chapter 3

Orientation Diffusion Process

The purpose of quantifying convexity of shapes was inspired by figure/ground separation. Therefore, a convexity measure was pursued and applied to F/G problems. We consider this measure a definition of shape convexity, with the small-size preference. With the perceptual evidences provided by Kanizsa [39], this definition is useful for the F/G problems, by providing F/G predictions statistically similar to the results from our visual systems. Computationally, our model was designed in a distributed system; hence, a highly parallelizable scheme can easily be obtained. Let us discuss how well this decay diffusion model can fit our goals from different viewpoints. Again, we do not try to distinguish between partial or complete shapes. Although as a measure to suggest F/G it is more likely to discuss partial shapes, while as an abstract definition for shape convexity, we are interested more in the complete ones.

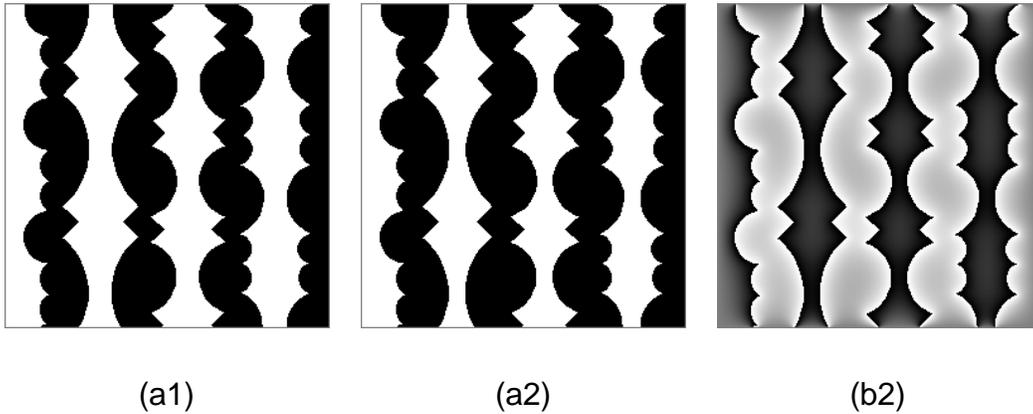


Figure 3.1. When we horizontally translate the boundaries in (a1), a copy of Fig. 2.7(a1) to (a2), we have a different convexity judgment from the decay model. The diffusion result of (a2) is in (b2) with the convexity measures $S/\bar{S} = 0.588/0.515$, predicting the white regions as figures, opposite to the result for (a1) which gives $S/\bar{S} = 0.546/0.560$, predicting the black regions as figures. Nevertheless we did not perceive much difference between (a1) & (a2) which has just pixels difference in location. ■

Figure/Ground Separation

From the F/G point of view, presumably, we can use the output from the decay model to predict the answer of F/G. More precisely, based on the studies of the convexity and size preferences of the decay measure in the last chapter, we should be able to use the measure in F/G prediction if the F/G phenomenon in input images are dominantly decided by these two factors. The remaining question is that, in the

decay model, we need to check if these two are “well-balanced/normalized” as in our visual systems. We design a simple experiment by horizontally translating the boundaries of the convexity-versus-symmetry image in Fig. 3.1. Under the convexity shape preference which yields the preference of black regions in (a1) as figures, the knowledge of small shape preference also tells us the following phenomenon:

When we translate the boundaries via the direction of expanding/shrinking the black/white regions respectively, we can expect the white regions will eventually become the prevailing parts as figures rather than background.

This phenomenon can be seen either through our model or the experiments in our visual systems (e.g. Stevens et al. [82]). For each of them, a “no difference” point exists where both regions share the same preference there. So the succeeding question is: Will they happen to be identical to each other? From Fig. 3.1(a1) to (a2), we horizontally translate the boundaries for a few pixel distance. The result from the decay model shows the result of $\mathcal{S}/\overline{\mathcal{S}} = 0.546/0.560$ for (a1) and $\mathcal{S}/\overline{\mathcal{S}} = 0.588/0.515$ for (a2). It suggests a “black region preference” in (a1) and a “white region preference” in (a2), opposite to our perceptual observation where we can hardly describe the difference between them if one of them is hidden from the other. Namely, our model is too sensitive to the size preference. It indicates a failure and we would like to improve it by accounting for this observation.

In the new framework as well as the new convexity measure that will be proposed,

a free parameter is provided and two no-difference points from the simulation and our visual systems can be synchronized by tuning this parameter.

Definition of Convexity

In the decay model, a convexity measure favoring small size regions has been proposed. If this measure is considered as a quantitative definition for shape convexity, it is natural to ask for a size-invariant one. By adjusting the decay coefficient λ (Eq. 2.9, 2.10), it is possible to give the same measure from the decay process for similar shapes with different sizes, according to Eq. 2.15 and Eq. 2.16. But the choice of λ needs region area as an input. In general, this knowledge is not available before the whole shape outline is detected. So this adjustment is not preferable for images containing illusory shapes, or noisy images. We would like to put it forward, with more flexibility. We would like to construct a convexity measure with or without the preference of small size regions.

In the next two sections, we will discuss some technical issues and see why the decay diffusion model is not appropriate under such considerations.

U-convexity, V-convexity, Boundary Curvature and 2nd Order Transition

On boundaries, we examine two typical types of convexity, the **V**-convexity and the **U**-convexity, defined in Sec.2.1 or Fig. 2.2. For the curves with **V**-convexity, the

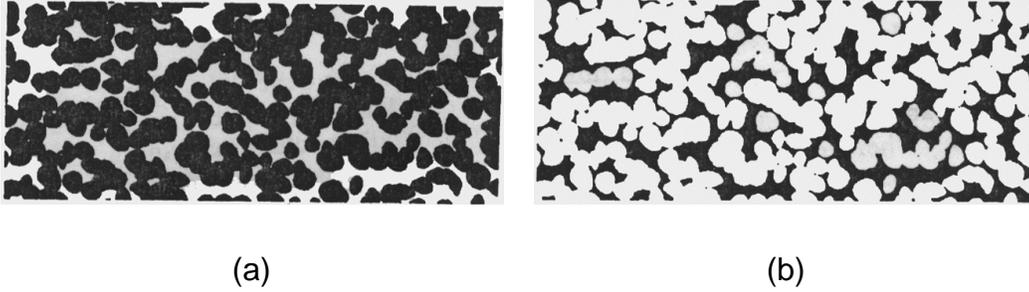


Figure 3.2. The concave cusp feature serves as the cue for F/G separation (adapted from Stevens et al. [82]). The result suggests that black/white regions in (a)/(b) respectively are more easily to be perceived as figures. ■

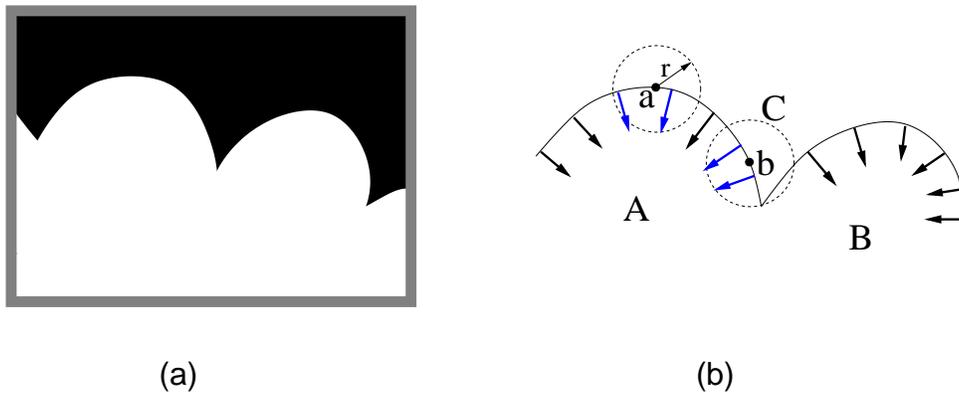


Figure 3.3. concave cusp feature as the dominator for F/G, suggests the white region as the most salient part. The orientation diffusion process, as opposed to the decay diffusion process, will give a more stable result (see Fig. 3.8). ■

convexity is concentrated on one peak of the boundaries, while for the ones with **U**-convexity, the convexity is more uniformly distributed along the boundaries.

Perceptually, we say that the **U**-convexity is more preferred than the **V**-convexity. As we argued in Sec. 2.1, if boundaries with both types are sampled through a *small* aperture, more likely, the F/G can be picked up from boundaries with **U**-convexity than with **V**-convexity, for the reason that most areas (except the peak) in **V**-convexity boundaries provide no clue for the F/G. For the **U**-convexity boundaries, F/G is more likely to be decided in a small scope, and in a separated way. Namely, for the **U**-convexity ones, we can decide the F/G by a small scope observation and by only one of the numerous observations. It is important, for the reason that profiles projected from objects in the 3-D world are mostly partial, broken or occluded and more collections of such local suggestions give a better chance to decide F/G efficiently. Furthermore, the angle peaks/junctions in **V**-convexity regions sometime suggest a potential visual organization movement. Different organization constructions lead to different F/G choices and figural hypotheses can be assigned on either the convex or concave side [48] [19] [32] [33] [34]. In various situations, junctions, the place with the **V**-convexity, are produced by object overlapping seen through a particular perspective, and hence, are of no physical significance. All these reasons show more importance of the **U**-convexity than the **V**-convexity, and it may lead to the preference of the **U**-convexity for F/G separation. It is true that information collected by

a small aperture is local, sometime incompatible with other local decisions and can be altered by a global consideration in visual organization construction. However, we discuss the boundaries with only the “balanced” presence of **U**-convexity and **V**-convexity. In this situation, the F/G decision will be chosen favoring the **U**-convexity rather than the **V**-convexity. It is exact the case in the convexity-versus-symmetry image provided by Kanizsa [39] (Fig. 1.6(a1), (a2) or Fig. 3.1) where number of **U**-convexity and **V**-convexity curves are similar.

Another similar evidence is provided by Stevens et al. [82]. In Fig. 3.2, they argue that the concave cusp feature is the dominator for F/G. We tend to choose black area as figures in (a) or white area as figures in (b). For a concave cusp (more like owning the **V**-convexity) in Fig. 3.3, the white region (owning the **U**-convexity) will be the favored part.

To speak of the decay diffusion model, a discussion can be made in the concave cusp in Fig. 3.3(b). We draw a small circle at either the point a or b with the same radius r . For the diffusion result, we say that the point b gets better support on the black side than the point a on the white side because the circle centered in b covers more inducers than the circle centered in a . This fact may lead a better result in the black region than in the white region. A numerical evidence is shown in Fig. 3.8.

We can consider curves with the **U**-convexity and **V**-convexity from a different view. Suppose the diffusion from boundaries are set to be oriented, i.e., the bound-

ary inducers, σ_0 in Eq. 2.8 or Eq. 2.9 are oriented ones, depending on the boundary normals and delivered by an oriented transition (a tendency to keep the previous orientations). In the curves with **U**-convexity, we can see diffusion comes from neighboring points and “converging” with similar orientations while for the curves with **V**-convexity, we either have no such “convergence” or have “convergence” from non-adjacent neighboring inducers. A 2nd-order Markov chain will be used as the transition mechanism to carry those normal information to form a regional description of shapes.

We are interested in another experiment which may be related to the discussion in this section. In Fig. 1.2(c), an image generated by a square with constant intensity and 1-D linear 0 – 100% gray on the background (inspired by Shapley et al. [76] where a circle figure was used). The intensity difference on boundaries suggests the illusory gradient inside the square, even no intensity change is provided. In particular, the gradient is extended in the direction perpendicular to the square boundaries. We are interested in what process can simulate this effect.

The Speed of Decay and Coarse Scale Structures

In Sec. 2.2.5, we estimated the speed of decay in the decay process. The decay is exponential, therefore the field lose structures when it is far from inducers. For the convexity measure, few structure in coarse scales will be considered. For convex

shapes with fine-scale perturbations, such as Fig. 1.7(b1) & (b2), an appropriate convexity measure should depend on not only fine scale convexity, but more importantly, coarse scale convexity.

3.1 Orientation Diffusion in $\mathbb{R}^2 \times \mathbb{S}^1$

The new convexity measure based on an orientation diffusion process will be proposed to improve the decay model in the following sense:

- i. The new measure allows the flexibility of switching between measuring convexity with or without size preference.
- ii. The 2nd order Markov chain is used as the transition mechanism to simulate the preference of **U**-convexity over **V**-convexity.
- iii. Fine scale protrusion, either convex or concave can only have minor contribution to the final measure.
- iv. The measure can distinguish two perfectly convex shapes. The result shows a simulation of the Law of Prägnanz for 2-D shapes (described in Chapter 1 or by Wertheimer [90] and Koffka [45]).
- v. Many shape attributes, such as symmetry and junction feature can be naturally obtained through computation of the measure.

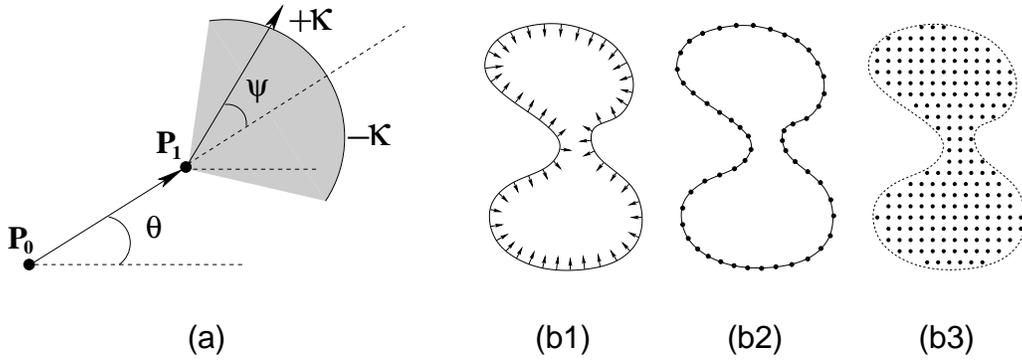


Figure 3.4. (a) an oriented particle moves from P_0 to P_1 with the orientation changed slightly. We say that the particle has its configuration changed from $(x, y; \theta)$ to $(x + \cos \theta, y + \sin \theta; \theta + \psi)$ as it goes from the point P_0 to P_1 . The shadow indicates possible trails the particle may choose, with different weights according to the deviation angle ψ . For a bell shape Ω , we show its hypothesis set, σ_0 in Eq. 3.2, by a vector representation s.t. $\sigma_0(\mathbf{x}; \theta)$ is indicated by the vector in \mathbf{x} with length $\sigma_0 \geq 0$, pointing to the direction $\mathbf{u}_\theta = (\cos \theta, \sin \theta)$. The 0-length vectors are indicated by dots. (b1) represents $\sigma_0(\mathbf{x}; \theta)$ for $\mathbf{x} \in \partial\Omega$, $\mathbf{u}_\theta \cdot \mathbf{n}_x \geq 0$, where \mathbf{n}_x is the normal on \mathbf{x} . Only the normal directions are shown. Other parts of the hypothesis include: (b2) $\sigma_0(\mathbf{x}; \theta) = 0$ for $\mathbf{x} \in \partial\Omega$, $\mathbf{u}_\theta \cdot \mathbf{n}_x < 0$, represented by vectors of length 0 (again, only those with the orientations in the outward normal directions are shown); and (b3) $\sigma_0(\mathbf{x}; \theta) = 0$ for $\mathbf{x} \in \Omega - \partial\Omega$, $\theta \in [0, 2\pi)$. The whole hypothesis set is given by $\sigma_0 = \{(b1)\} \cup \{(b2)\} \cup \{(b3)\}$. ■

We will discuss the first four topics in this chapter and postpone the fifth topic to the next chapter.

The plan is designed similarly to the decay model. After a field is generated by a process, we derive a measure by a point-wise computation. The new diffusion process, called the orientation diffusion, augment the working space for diffusion from the image space \mathbf{R}^2 to $\mathbf{R}^2 \times \mathbf{S}^1$ with an additional dimension $\Theta \equiv [0, 2\pi)$. The idea of orientation diffusion can be illustrated in Fig. 3.4(a). Apart from the decay diffusion which is considered as a random walk or a Brownian motion $W(t)$ with the property

$$\text{The increment } W(t) - W(s) \text{ is independent of } W(s). \quad \forall t \geq s$$

The walk carries no memory other than its location as it proceeds. The orientation diffusion takes additional memory in the orientation dimension. With a known orientation θ at time s , it is more likely to keep this orientation to the next status of time t , up to some noise. In Fig. 3.4(a), a particle with the configuration (x, y, θ) at the time s could end up with the next step of time t with a new configuration $(x + \cos \theta, y + \sin \theta, \theta + \psi)$ where ψ is created by noise. While the previous model caught convexity by the support of two near points, we now catch convexity in a different sense. *The support of orientation diffusion is always from neighboring inducers*, unlike the support of decay diffusion which may come from two non-adjacent

ones. E.g., in the diffusion map in Fig. 2.7(a) & (b) or Fig. 3.1(b2), results generated by the decay process, we can find more diffusion near the neck area of the white regions.

Some works have been done with models dealing with oriented data or working on an oriented space. Tang et al. [85] collected the unary oriented information as input and a process, called the *direction diffusion* was applied to the data regularization. Some other works related to image regularization and segmentation by isotropic or anisotropic diffusion can be referred to Weickert [88]. Mumford [58], William and Jacobs [91] used an oriented process and applied the Euler *Elastica* of two oriented sources (junctions) to complete occluded-curves. Unlike those works, our works are derived from boundary input, and the result is recorded by a vector-bundle field instead of a scalar or a vector field. Besides, our interests are concentrated on describing 2-D shapes, including deriving convexity measure, computing shape axis and junction information.

3.1.1 Variational Formulation and Energy Functional

Notations

For a given region Ω (represented by its boundaries $\partial\Omega$) in an input image \mathbf{I} , either partial or complete, we discuss its convexity. A set of useful information is collected

as follows. For an intensity function $I(\mathbf{x})$, the oriented edge map

$$e(\mathbf{x}; \theta) = \frac{1}{M_e} \left| \frac{\partial I(\mathbf{x})}{\partial \mathbf{u}_\theta} \right| \in [0, 1], \quad \mathbf{x} \in \partial\Omega \quad (3.1)$$

records the normalized magnitude of the intensity change at the point \mathbf{x} in the direction of $\mathbf{u}_\theta = (\cos \theta, \sin \theta)$, namely, the (unsigned) directional derivative in \mathbf{u}_θ , normalized by $M_e = \max\{|\partial I(\mathbf{x})/\partial \mathbf{u}_\theta| : \mathbf{x} \in \mathbf{I}, \theta \in \Theta\}$, the largest directional intensity edge in the image \mathbf{I} . Also, on $\partial\Omega$, a hypothesis set is defined by

$$\sigma_0(\mathbf{x}; \theta) = \begin{cases} e(\mathbf{x}; \theta), & \text{if } \mathbf{u}_\theta \cdot \mathbf{n} \geq 0 \wedge \mathbf{x} \in \partial\Omega \\ 0, & \text{if } \mathbf{u}_\theta \cdot \mathbf{n} < 0 \vee \mathbf{x} \in \Omega - \partial\Omega \end{cases} \quad (3.2)$$

as inducers. On the boundary, the vector \mathbf{n} is the unit normal vector chosen with the inward direction. Intuitively, we say a direction closer to the inward normal is in the *source mode*, or otherwise, in the *sink mode*. That is, on boundaries, we prefer a value of greater than/equal to 0 for the orientations going inward/outward respectively. It is a different design from the decay model. The positive inducers only occur in the inward orientations. Finally, a field

$$\sigma(\mathbf{x}; \theta) : \Omega \times \Theta \rightarrow \mathbf{R}$$

will be evaluated by our orientation diffusion process.

We will use vector notation to “visualize” σ (as well as σ_0), where $\sigma(\mathbf{x}; \theta)$ will be denoted by a vector at \mathbf{x} with length $|\sigma(\mathbf{x}; \theta)|$, pointing to the direction θ . All vectors located at \mathbf{x} form a vector bundle of \mathbf{x} .

Local Hypotheses and Data Fitting

The field $\sigma(\mathbf{x}; \theta)$ should take the local hypothesis value $\sigma_0(\mathbf{x}; \theta)$, where they are available. In this case, they are available at all intensity edge points (points on $\partial\Omega$) according to their strength $e(\mathbf{x}; \theta)$. I.e., we seek the $\sigma(\mathbf{x}; \theta)$ which minimizes

$$E_{\text{data}}(\sigma|\sigma_0) = \int_{\Omega \times \Theta} \Lambda(\mathbf{x}; \theta) (\sigma(\mathbf{x}; \theta) - \sigma_0(\mathbf{x}; \theta))^2 d\mathbf{x} d\theta, \quad (3.3)$$

where the function $\Lambda(\mathbf{x})$ is given by

$$\Lambda(\mathbf{x}; \theta) = \begin{cases} e(\mathbf{x}; \theta) \cdot \delta_2(\mathbf{x}), & \text{if } \mathbf{u}_\theta \cdot \mathbf{n} \geq 0 \wedge \mathbf{x} \in \partial\Omega \\ \lambda, & \text{if } \mathbf{u}_\theta \cdot \mathbf{n} < 0 \vee \mathbf{x} \in \Omega - \partial\Omega \end{cases} \quad (3.4)$$

with the 2-D Dirac delta function $\delta_2(\mathbf{x})$ s.t. $\int_{\partial\Omega} \delta_2(\mathbf{x}) d\mathbf{x} = 1$ (Appendix A). For a non-smooth, a binary intensity function $I(\mathbf{x})$ valued in $\{0, 1\}$ characterizing Ω , we can assign the edge map by

$$e(\mathbf{x}; \theta) = |\mathbf{u}_\theta \cdot \mathbf{n}|.$$

The directions perpendicular to boundaries are the one with the maximum value in $e(\mathbf{x}; \theta)$, with either an inward ($\sigma_0 = 1$) or outward ($\sigma_0 = 0$) hypothesis.

For the homogeneous regions (where no intensity edge is present), the functions $\sigma_0(\mathbf{x}; \theta)$ and $\Lambda(\mathbf{x}; \theta)$ are given by respectively assigning a local hypothesis of value 0 and a small constant strength $0 < \lambda \ll 1$, known as the decay coefficient *. As before, the coefficient λ controls the decay effect of the energy when it is away from

*When this functional is analyzed as a Markov chain, the parameter λ plays the role of

the inducers (hypothesis set). A larger λ has a stronger effect in bringing points away from the intensity edges to take value 0 (see Fig. 3.5(d) & (e)).

Smoothness

Propagation of the local hypothesis is done by adding a smoothness term that prefers neighboring points with similar orientations to share similar values. For instance, in Fig. 3.4(a), the vector in position P_0 with the orientation θ and its neighbor P_1 with the orientation $\theta + \psi$ are encouraged to share similar values in σ . A simple quadratic form is applied here by minimizing

$$\frac{1}{\kappa} \sum_{\psi=-\kappa}^{\kappa} M(\mathbf{x}; \theta, \psi) (\sigma(x + \cos \theta, y + \sin \theta, \theta + \psi) - \sigma(\mathbf{x}, \theta))^2,$$

for each point \mathbf{x} . A continuous form gives the smoothness functional as

$$E_{\text{smooth}}(\sigma) = \int_{\Omega \times \Theta} \left[\frac{1}{\kappa} \int_{-\kappa}^{\kappa} M(\mathbf{x}; \theta, \psi) (\cos \theta \sigma_x + \sin \theta \sigma_y + \psi \sigma_\theta)^2 d\psi \right] d\mathbf{x} d\theta. \quad (3.5)$$

The function $M(\mathbf{x}; \theta, \psi)$ is simply provided by $\cos(\psi \frac{\pi}{2k})$, giving the cosine weighting for the smoothness that concentrates particularly on small deviations. A more complicated version of the function M may also depend on \mathbf{x} (non-homogeneous) or θ (anisotropic).

the decay parameter, or the probability of “vanishing”, of the random walk associated with this functional.

A new parameter is introduced, $\kappa \in (0, \pi)$, called the deviation factor[†], controlling how likely moving particles will keep their orientations. Later in the discussion of normalization between convexity and size, we can tune this factor to give a convexity measure between a size-invariant one and a measure with small shape preference.

Energy Functional

The total energy functional is summation of the functionals Eq. 3.3 and Eq. 3.5,

$$E(\sigma|\sigma_0) = E_{\text{data}}(\sigma|\sigma_0) + E_{\text{smooth}}(\sigma). \quad (3.6)$$

For Eq. 3.5, the functional within the bracket, called $\mathcal{H}(\sigma|\mathbf{x}; \theta)$ can be rewritten by

$$\begin{aligned} & \mathcal{H}(\sigma|\mathbf{x}; \theta) \\ &= (\cos \theta \sigma_x + \sin \theta \sigma_y)^2 \cdot \frac{1}{\kappa} \int_{-\kappa}^{\kappa} M(\mathbf{x}; \theta, \psi) d\psi \\ & \quad + 2\sigma_\theta (\cos \theta \sigma_x + \sin \theta \sigma_y) \cdot \frac{1}{\kappa} \int_{-\kappa}^{\kappa} \psi M(\mathbf{x}; \theta, \psi) d\psi \\ & \quad + \sigma_\theta^2 \cdot \frac{1}{\kappa} \int_{-\kappa}^{\kappa} \psi^2 M(\mathbf{x}; \theta, \psi) d\psi \\ &= A (\cos \theta \sigma_x + \sin \theta \sigma_y)^2 + 2B \sigma_\theta (\cos \theta \sigma_x + \sin \theta \sigma_y) + C \sigma_\theta^2. \end{aligned} \quad (3.7)$$

For the simplest case where $M(\mathbf{x}; \theta, \psi) = \cos(\psi \frac{\pi}{2k}) \forall \mathbf{x} \in \Omega, \theta \in \Theta$, we have

$A = 4/\pi, B = 0$ and $C = \frac{4\kappa^2}{\pi}(1 - \frac{8}{\pi^2})$. In this case, its Euler-Lagrange equation can

[†]We avoid the degenerate case, $\kappa = 0$, when the smoothness functional becomes $\sum_{\mathbf{x}, \theta} M(\mathbf{x}, \theta, 0)(\sigma(\mathbf{x}, \theta) - \sigma(x + \cos \theta, y + \sin \theta, \theta))^2$, no longer depending on ψ , and the minimization process can be operated separately through different θ 's.

be given by

$$\cos^2\theta \sigma_{xx} + 2 \sin\theta \cos\theta \sigma_{xy} + \sin^2\theta \sigma_{yy} + \kappa^2 \left(1 - \frac{8}{\pi^2}\right) \sigma_{\theta\theta} = \frac{\lambda\pi}{4} \sigma, \quad (3.8)$$

with the boundary condition $\sigma(\mathbf{x}; \theta) = \sigma_0(\mathbf{x}; \theta)$, if $(\mathbf{x}, \theta) \in \partial\Omega \times \Theta$.

Numerically, the energy function is quadratic and the minimizer can be straightforwardly obtained by rewriting Eq. 3.6 in matrix form and solving the matrix inverting problem. The minimizer will be written as $\sigma^*(\mathbf{x}; \theta)$. By the random walk formulation, the result $\sigma^*(\mathbf{x}; \theta)$ can be viewed as the expectation of σ_0 at the first vanishing point of walks starting at \mathbf{x} with orientation θ . The vanishing is done by either hitting the boundaries or decay along the walk. Equivalently, up to a constant, the result can also be viewed as summation of all walks starting at boundary points, carrying the value σ_0 . Some examples of $\sigma^*(\mathbf{x}; \theta)$ are shown in Fig. 3.5(a) & (b). Also, Fig. 3.5(f1) shows the result among all orientations, by using the vector representation.

3.1.2 Kullback-Leibler Measure

Once the field $\sigma^*(\mathbf{x}; \theta)$ is obtained, we can evaluate the convexity. First of all, from the solution space $\Omega \times \Theta$ to the image space Ω , we perform a vector summation considering $\sigma^*(\mathbf{x}; \theta)$ as a vector in the direction of $\mathbf{u}_\theta = (\cos\theta, \sin\theta)$ with length

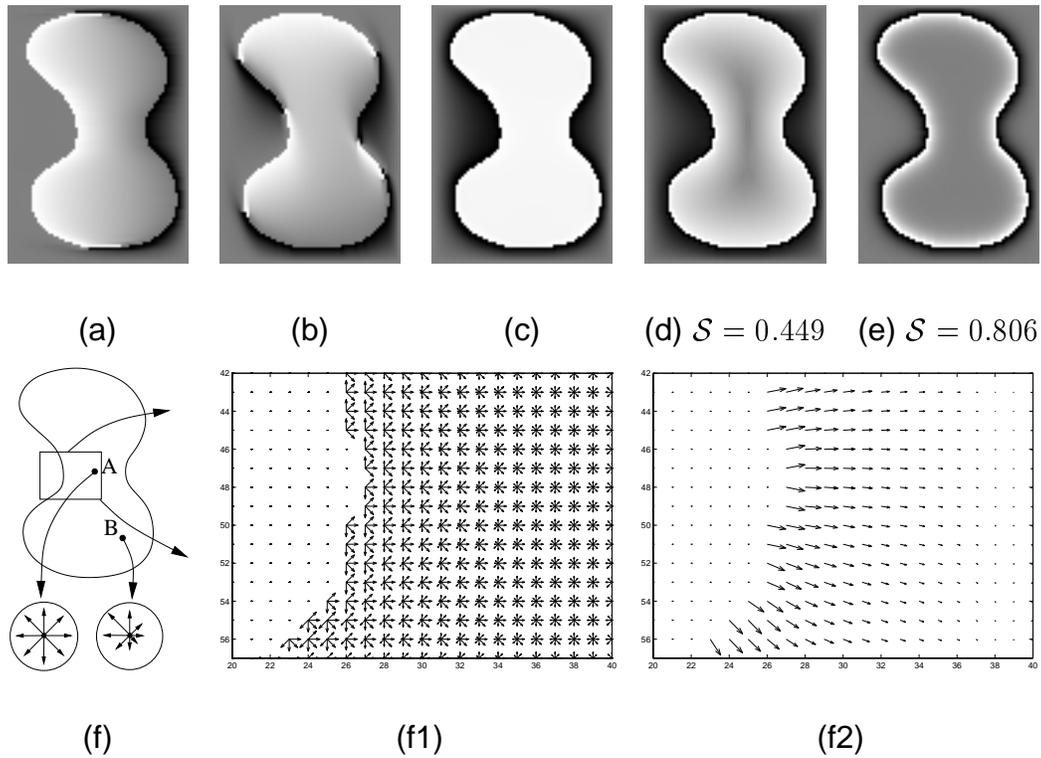


Figure 3.5. Results of orientation diffusion process by two presentations. (a) $\sigma(x, y; 0)$, (b) $\sigma(x, y; \frac{27}{16}\pi)$, (c) maximum magnitude among all orientations, and (d) (e) relative entropy $D(P_x||Q)$ (Eq. 3.12) with $\lambda = 0.1/|\Omega| = 2.876 \times 10^{-5}$ and $\lambda = 500/|\Omega| = 0.144$ respectively. The difference shows effect of the energy decay. Larger λ gives stronger decay and less diffusion. (f1) (f2) Vector presentations. (f1) shows vector bundles and convexity vector $\hat{\sigma}^*$ for part of the bell shape figure in (f). (f1) the vector bundles with choice of 8 orientations. We say the point A has a more balanced (symmetry) result along different orientations than the bundle of the point B . (f2) the convexity vector $\hat{\sigma}^*$. ■

$\sigma^*(\mathbf{x}; \theta)$ (always non-negative),

$$\hat{\sigma}^*(\mathbf{x}) = \hat{\sigma}_{\text{cv}}^*(\mathbf{x}) = \int_0^{2\pi} \sigma^*(\mathbf{x}; \theta) \mathbf{u}_\theta d\theta, \quad (3.9)$$

where direction of the resultant vector $\hat{\sigma}_{\text{cv}}^*$ will be recorded by θ^* , if it has non-zero length. We shall use $\hat{\sigma}^*$ to denote the resultant vectors if no confusion can be made. Examples of the vector field $\hat{\sigma}^*(\mathbf{x})$ can be seen in Fig. 3.5(b2). The idea is that this vector summation will remove the symmetry information of the solution field, and thus what is left is “convexity”. For example, the center of a circle is perfectly symmetric and will result in $\hat{\sigma}^* = \mathbf{0}$ while other internal points will have the symmetry removed and still produce evidence of convexity. In our approach, convexity works on the complementary information as symmetry does. Detecting symmetry will be done by investigating which coordinates have a vector summation yield zero value (cancellations), while convexity examines the non-zero valued information of the vector summation. Indeed, the convexity measure we propose measures how well the “vector bundle” is accumulated around this summation vector. Let us be more precise.

At a fixed location \mathbf{x} , for each pair of opposite orientations θ and $\theta + \pi$, we compute the net effect, called the map $\tau^*(\mathbf{x}; \theta)$ by

$$\begin{aligned} \tau^*(\mathbf{x}; \theta) &= \frac{1}{2} \{ \sigma^*(\mathbf{x}, \theta) - \sigma^*(\mathbf{x}, \theta + \pi) + |\sigma^*(\mathbf{x}, \theta) - \sigma^*(\mathbf{x}, \theta + \pi)| \} \\ \tau^*(\mathbf{x}; \theta + \pi) &= \frac{1}{2} \{ \sigma^*(\mathbf{x}, \theta + \pi) - \sigma^*(\mathbf{x}, \theta) + |\sigma^*(\mathbf{x}, \theta + \pi) - \sigma^*(\mathbf{x}, \theta)| \} \end{aligned}$$



Figure 3.6. A simple experiment used for F/G separation and its relative entropy $D(P_{\mathbf{x}}||Q)$. While right side is convex and left side is concave, we have $S/\bar{S} = 0.605/0.671$ respectively. Both regions are equally sized. ■

$$= \tau^*(\mathbf{x}; \theta) - \sigma(\mathbf{x}; \theta) + \sigma(\mathbf{x}; \theta + \pi), \quad (3.10)$$

where a cyclic boundary condition $\theta = \theta + 2\pi$ is used. Note that equation (3.9) can be rewritten as

$$\hat{\sigma}^*(\mathbf{x}) = \int_0^{2\pi} \tau^*(\mathbf{x}; \theta) \mathbf{u}_\theta d\theta.$$

The second step is to transform $\tau(\mathbf{x}; \theta)$ to $P^*(\mathbf{x}; \theta)$ or simply $P_{\mathbf{x}}(\theta)$, by a linear transformation, similar to what we did for the decay model in Eq. 2.12. It is

$$P^*(\mathbf{x}; \theta) = \frac{1}{2}(1 + \tau^*(\mathbf{x}; \theta)).$$

Unlike any previous work, we propose a Kullback-Leibler distance (or relative entropy) to measure convexity. The map $P_{\mathbf{x}}(\theta)$ is compared to a Gaussian map in \mathbf{S}^1 (not normalized)

$$Q(\theta) = \frac{1}{1 + \epsilon_N} \exp\left(-\frac{(\theta - \theta^*)^2}{2\sigma^2}\right), \quad -\pi \leq \theta \leq \pi \quad (3.11)$$

where the center of Q is located in the direction θ^* , the direction of vector $\hat{\sigma}^*$ (here σ stands for standard deviation, not the diffusion field. It is set to 2.0 throughout all the experiments). A small positive constant ϵ_N is given to avoid the case of $Q(\theta) = 1$. The Kullback-Leibler distance, computed at every point \mathbf{x} , between $Q(\theta)$ and $P_{\mathbf{x}}(\theta)$ is given by

$$D(P_{\mathbf{x}}\|Q) = \int_0^{2\pi} P_{\mathbf{x}}(\theta) \log \frac{P_{\mathbf{x}}(\theta)}{Q(\theta)} + (1 - P_{\mathbf{x}}(\theta)) \log \frac{1 - P_{\mathbf{x}}(\theta)}{1 - Q(\theta)} d\theta . \quad (3.12)$$

Intuitively, it measures the inefficiency of assuming that the distribution is $Q(\theta)$ when the real distribution is $P_{\mathbf{x}}(\theta)$. That is, we assume the map in \mathbf{S}^1 has a sharp peak at θ^* . $D(P_{\mathbf{x}}\|Q)$ is always equal to or greater than 0, and it can be greater than 1. A smaller measure means a better fit.

This measure is given for each point \mathbf{x} . To obtain a global measure of a shape, we simply average this measure over the points on the figure, i.e., the final measure of a shape Ω is

$$\mathcal{S}(P_{\mathbf{x}}) = \frac{1}{|\Omega|} \int_{\Omega} D(P_{\mathbf{x}}\|Q) d\mathbf{x} ,$$

A shape with a smaller measure is considered more convex than others. A simple example can be seen in Fig. 3.6 as a F/G experiment. We continue to use $\bar{\mathcal{S}}$ to denote the measure for background area.

3.2 Convexity Measurement by 2nd Order Process

The process can be considered as a 2nd order Markov chain with memory of location and orientation. Various studies suggest that the 2nd order Markov chain with the help of Kullback-Leibler measure provides better convexity measurement than the one derived from the decay process. We use numerous experiments to show the difference between them. The result strongly depends on the appropriate choice of κ . But in this section, we will only sketch various experiments that we are interested and their results. The results derived by using different κ 's and their relations will be discussed in Sec. 3.3. Other results are separated in different sections in rest of this chapter. The comparison of two perfectly convex shapes are discussed in Sec. 3.4. In Sec. 3.5, we discuss convexity of the shape with minor perturbations in the fine scales. The implementation details are given in Sec. 3.6.

The first series is organized as F/G experiments. It includes the images with the concave cusp feature in Fig. 3.3, inspired by Stevens et al. [82] and the convexity-symmetry images adapted from Kanizsa [39]. Besides the F/G experiments, in the second series of experiments, we compare convexity of different shapes. It is a figure-to-figure rather than figure-to-ground comparison. The reason that we compare “figure” to “figure” by this model instead of the decay model is because many unnecessary attributes in the decay model such as size or symmetry preference have

been removed here. Therefore, it is appropriate to mention “convexity measure” of shapes as it solely owns the name now. The comparison can thus be applied to a wider range. In most experiments, results from the decay model are provided for comparison.

3.2.1 Figure/Ground Separation

We examine our measure using four prototypes, a simple arc in Fig. 3.6, the colonnade in Fig. 3.7, concave-cusp figures in Fig. 3.8 and the Kanizsa figures in Fig. 3.9. Shapes with higher convexity win on all trials.

Arc For a F/G experiment of arc shape in Fig. 3.6, the measure is given by $S/\bar{S} = 0.605/0.671$, favoring the white region.

Colonnade In the second series shown in Fig. 3.7, we test our measure in colonnade-figure images. The result is consistent through images with different ratios of bk/wt area. Also, we do observe the size effect from (a) to (c): the difference between figures and background becomes less as black/white regions get smaller/larger respectively.

Concave Cusp For concave-cusp images in Fig. 3.8(a) & (b), we apply both of the decay diffusion and the orientation diffusion process as methods for F/G separation.

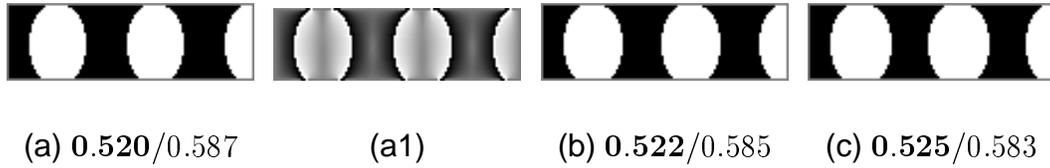


Figure 3.7. Colonnade with boundaries translated through the horizontal direction from (a), a size balanced one to (c), with white regions owning the largest area. The relative entropy of (a) is shown in (a1). The differenced between the measures in figure/ground get smaller from (a) to (c), as black/white regions get smaller/larger. It shows the size/proximity preference of the model. ■

The results show that the orientation diffusion is more capable of providing a correct prediction than the decay diffusion process. For both images, the orientation measure predicts that white regions are favored over black regions which is not always provided by the decay measure.

Convexity vs Symmetry For the Kanizsa images in Fig. 3.9(a3) where white regions own the largest area compared to Fig. 3.9(a1) & (a2), the measure of $\mathcal{S}/\bar{\mathcal{S}} = 0.490/0.497$, predicts that the white regions (convexity) prevail as the figures over the black regions (symmetry). This prediction cannot be derived by the decay model which gives the measure of $\mathcal{S}/\bar{\mathcal{S}} = 0.653/0.576$, favoring the black regions (concave and symmetry), a “wrong” prediction. The reason comes from the fact that the

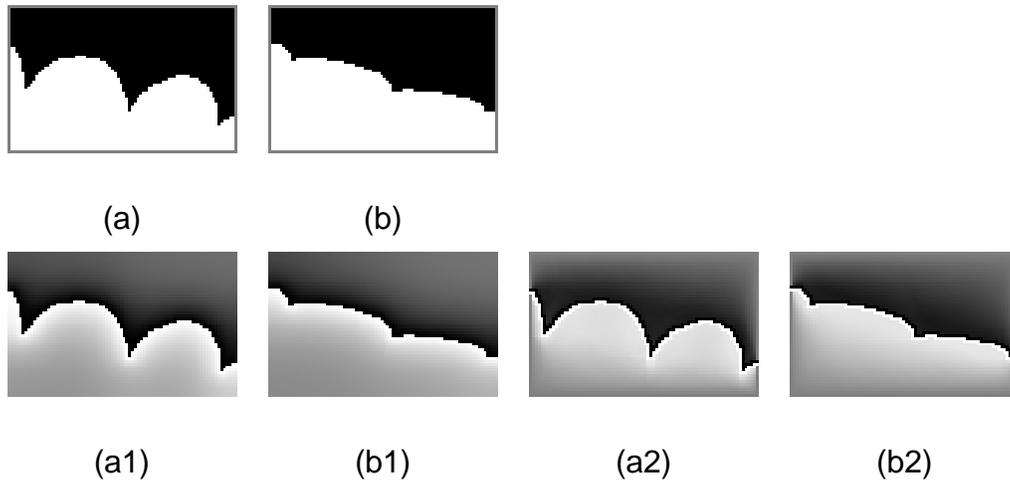


Figure 3.8. Images inspired by Stevens et al. [82] and their F/G predictions. (a) (b) original images, (a1) (b1) results derived from the decay process and (a2) (b2) results from the orientation process. Their convexity measures are (a1) $\mathcal{S}/\overline{\mathcal{S}} = \mathcal{S}_{\text{white}}/\mathcal{S}_{\text{black}} = 0.692/0.716$, (b1) $\mathcal{S}/\overline{\mathcal{S}} = 0.785/0.778$ by the decay model. The improved one from the orientation diffusion gives (a2) $\mathcal{S}/\overline{\mathcal{S}} = 0.507/0.566$ and (b2) $\mathcal{S}/\overline{\mathcal{S}} = 0.556/0.561$. Results provided by the orientation process show more consistency, especially for the case when the feature causing convexity gets weaker. ■

decay model is too sensitive to size and cannot provide a consistent prediction when we move the edge boundaries horizontally, even if the change is small and beyond the detection from our eyes. On the other hand, the oriented measure will give a correct prediction consistently, with a slightly shift of the edge boundaries. More results and

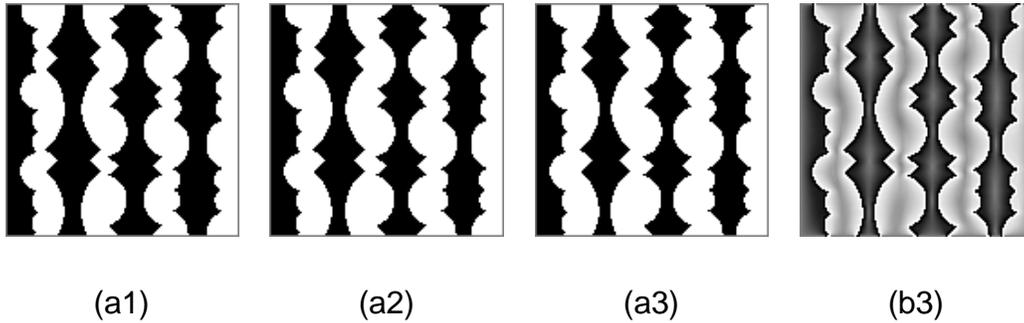


Figure 3.9. Convexity measure for the Kanizsa figures (adapted from Kanizsa [39]). (b3) gives the relative entropy of (a3). For (a3) which gives the largest white space, the convexity measure is $\mathcal{S}/\overline{\mathcal{S}} = 0.490/0.497$ by a test of $\kappa = 0.1$. When the boundaries are translated from (a1) to (a3) with white regions getting bigger, the measure from the decay process shows $\mathcal{S}/\overline{\mathcal{S}} = 0.653/0.576$, a “wrong” prediction. Note that the decay diffusion process will give more diffusion near the neck area while in this case, a bad relative entropy will be given with help of the orientation process. More discussion related to size and the choice of κ can be seen in Tab. 3.1. ■

discussions related to size and the choice of κ will be given in Sec. 3.3.

3.2.2 Convexity Comparison of Shapes

For the figure-to-figure convexity comparison, our attention is turned to Fig. 3.10, a convexity comparison between an imperfect ellipse (convex) and a bell shape (con-

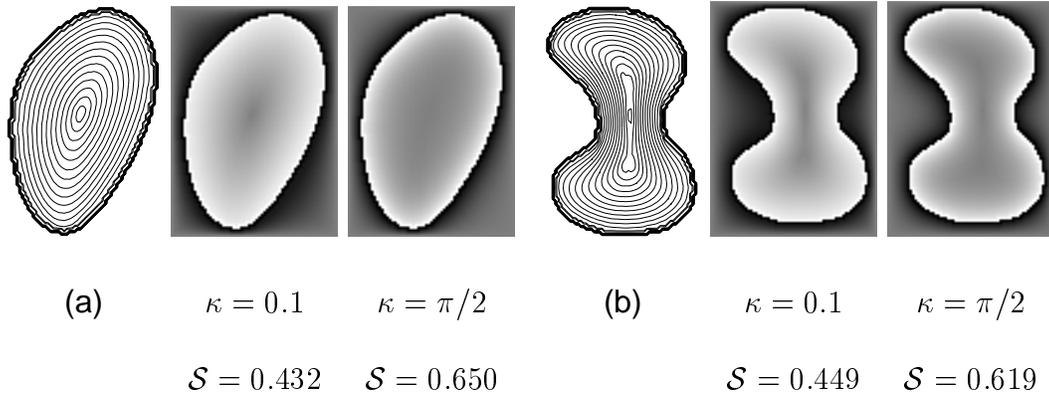


Figure 3.10. The level sets and the relative entropy for the (imperfect) ellipse and the bell shape with the simulations of $\kappa = 0.4$ and $\kappa = \pi/2$. The preference between the ellipse and bell switches as we goes from a small κ to a large κ . A complete comparison of their convexity according to different κ 's will be shown in Fig. 3.12. Results from the decay model give $\mathcal{S}_{\text{ellipse}} = 0.674$ and $\mathcal{S}_{\text{bell}} = 0.607$, by a simulation when $\lambda = 1 \times 10^{-2}$. It gives the same preference as we obtained from the orientation model by a large κ . ■

cave). For a series with κ set to be 0.4, we have $\mathcal{S}_{\text{ellipse}} = 0.432 < 0.449 = \mathcal{S}_{\text{bell}}$, correspondent to our expectation. Results by assigning larger κ as well as the discussion of the choice of κ will be provided in next section (Sec. 3.3.3 and Fig. 3.12).

We can easily find out that the measure derived from the decay process is not appropriate for such arbitrary shape comparison. Besides the necessity of different setting of λ according to different sizes (Sec. 2.2.4), the decay model does not pro-

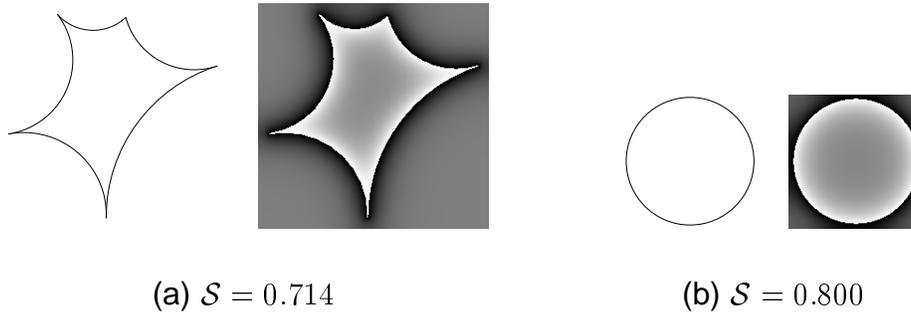


Figure 3.11. A highly concave shape (a) is “more convex” than (b) a circle, according to the measure from the decay process. The measure gives (a): 0.714 and (b): 0.800. Both shapes have the same area. ■

vide a clear meaning for shape convexity, which, for instance, says that a shape in Fig. 3.11(a) is more convex than a circle in (b).

3.3 Convexity versus Size

In the Euler-Lagrange equation in Eq. 3.8, the deviation factor κ serves as the diffusivity in the orientation space Θ . In the random walk formulation, a larger κ provides more noise and less deterministic behavior in orientation when it moves from one location to another neighboring location. A measure with small-size preference is yielded in such a situation. On the other hand, a small κ has more control in orientation. The measure will have less size preference and be closer to ideal convexity measure than the one given in the less-deterministic case. The most extreme case

gives the size invariance measure, provided by letting $\kappa \rightarrow 0$. It will be discussed in Sec. 3.3.1. Results related to different choices of κ will be studied in Sec. 3.3.2 for F/G separation and Sec. 3.3.3 for shape convexity comparison respectively. The choice of κ provides the flexibility of convexity measure between a size-invariant one and the one with small size preference. It cannot be achieved by the decay measure.

3.3.1 Size Invariance by Letting $\kappa \rightarrow 0$

Let us analyze the case when $\kappa \rightarrow 0$, where the walk is totally deterministic in orientation. By taking the limit $\kappa \rightarrow 0$, the functional in Eq. 3.7 becomes

$$\lim_{\kappa \rightarrow 0} \mathcal{H}(\sigma | \mathbf{x}; \theta) = \frac{4}{\pi} (\cos \theta \sigma_x + \sin \theta \sigma_y)^2. \quad (3.13)$$

We want to examine the effect of shape scaling in this limit for the whole functional in Eq. 3.6. After that, we study the scaling effect on the convexity measure \mathcal{S} .

Given two *similar* shapes Ω and Ω' in an image \mathbf{I} , indicated by two characteristic functions χ and χ' s.t. $\chi(\mathbf{x}) = 1$ or $\chi'(\mathbf{x}) = 1$ if $\mathbf{x} \in \Omega$ or Ω' respectively, and $\chi(\mathbf{x}) = 0$ or $\chi'(\mathbf{x}) = 0$ otherwise. Under an appropriately chosen coordinate system, we have $\chi(\mathbf{x}) = \chi'(r\mathbf{x}) = \chi'(\mathbf{x}')$, $\forall \mathbf{x}, \mathbf{x}' \in \mathbf{I}$. Given similar edge maps $e(\mathbf{x}; \theta) = e'(\mathbf{x}'; \theta)$, when $\kappa \rightarrow 0$ and $\lambda \rightarrow 0$, the decay term will vanish and we have

$$E(\sigma') = E(\sigma) = \int_{\Omega \times \Theta} \delta_2 \cdot e \cdot (\sigma - \sigma_0)^2 + \frac{4}{\pi} (\cos \theta \sigma_x + \sin \theta \sigma_y)^2 dx d\theta, \quad (3.14)$$

if we assign $\sigma'(\mathbf{x}'; \theta) = \sigma(\mathbf{x}; \theta)$. So the minimizer for either functional can be obtained easily when the other one has been computed. Two minimizers are related by $\sigma^*(\mathbf{x}; \theta) = \sigma'^*(\mathbf{x}'; \theta)$. Different orientations are worked independently here.

Let us discuss the scaling effect on the convexity measure. For two similar maps σ^*, σ'^* , we can obtain the similarity between $P_{\mathbf{x}}$ and $P_{\mathbf{x}'}$ or $D(P_{\mathbf{x}}||Q)$ and $D(P_{\mathbf{x}'}||Q)$. Therefore, for the convexity measure, we have

$$\begin{aligned} \mathcal{S}(P_{\mathbf{x}'}) &= -\frac{1}{|\Omega'|} \int_{\Omega'} D(P_{\mathbf{x}'}||Q) d\mathbf{x}' \\ &= -\frac{1}{r^2|\Omega|} \cdot r^2 \int_{\Omega} D(P_{\mathbf{x}}||Q) d\mathbf{x} = \mathcal{S}(P_{\mathbf{x}}). \end{aligned} \quad (3.15)$$

It is an ideal case where we can achieve a measure with no sensitivity to the size. Examples in the discrete case can be found in Fig. 3.12 and Sec. 3.3.3. When λ is not zero, we need to set a smaller $\lambda' = \lambda/r^2$ to achieve the similarity $\sigma^*(\mathbf{x}; \theta) = \sigma'^*(\mathbf{x}'; \theta)$.

3.3.2 Figure/Ground Separation in Convexity-Symmetry Image

We will use the Kanizsa figures in Fig. 3.9 to illustrate our points. As we have discussed, the decay model cannot provide a consistent prediction when the area ratio of different regions is changed, even though the change is small and beyond the detection from our eyes. The orientation model will give a consistent prediction, with a small shift of edge boundaries. The trick lies in the free parameter κ which on one

Series (Area)	$\kappa = 0.1$	$\kappa = 0.4$	$\kappa = \pi/2$	Remark
Convexity Measure by Orientation Diffusion (S/\bar{S})				
(a1) (+0/− 0%)	0.487 /0.502	0.490 /0.499	0.560 /0.570	size balanced
(a2) (+6/− 0%)	0.488 /0.500	0.492 /0.497	0.564 /0.565	
(a3) (+28/− 0%)	0.490 /0.497	0.495/ 0.491	0.573/ 0.555	
Convexity Measure by Decay Diffusion (S/\bar{S})				
(a1) (+0/− 0%)	0.621 /0.653			size balanced
(a2) (+6/− 0%)	0.638/ 0.635			
(a3) (+28/− 0%)	0.671/ 0.597			

Table 3.1. Kanizsa figures with various size ratios (Fig. 3.9), measured by the orientation diffusion and the decay diffusion. The series from (a1) to (a3) gives the white region area from small to large. Series (a1) has balanced size. The result can be read from different viewpoints. For the orientation diffusion, a smaller κ gives a purer convexity measure and a larger κ , meaning more uncertainty in orientations in walks, will cause a stronger size preference. The prediction for F/G is likely to be decided by convexity when κ is small and be decided by size/proximity when κ is large. We can get convex predictions for F/G when $\kappa \leq 0.1$. The decay diffusion measure is pretty much a size measure if we do not provide a size-balanced experiment. ■

hand (small κ or the extreme case with $\kappa = 0$) gives a convexity measure without considering small size preference and on the other hand (big κ) gives a measure with the size preference. In Tab. 3.1, we have series (a1), balanced area for black and white regions, series (a2), larger white regions and (a3) white regions with the largest area. Several pairs of comparisons can be studied. When $\kappa = 0.1$, the purest convexity measure among them, gives the expected answer with white in the front, for all three cases. When $\kappa = \pi/2$, a case blended with the most size preference among them, white regions in (a1) & (a2) are preferred, but not for (a3) where the smaller black regions are preferred. The case of $\kappa = 0.4$ gives an intermediate transition. For the perceptual simulation, suppose a case where F/G is decided by both of the convexity and size preferences, we can tune the parameter κ to obtain the appropriate normalization of them, according to the preferences in human visual systems.

3.3.3 Convexity Comparison of Shapes and Tuning of κ

We study our convexity measure for different shapes with the same or different sizes. The source images are provided in Fig. 3.10. The imperfect ellipses $A1$ (large) & $A2$ (small) represent convex shapes, and the bell shapes $B1$ (large) & $B2$ (small) represent non-convex shapes with “strong proximity” in the neck area. For the choice of κ , a similar scenario is used as in the F/G separation experiments. When κ is

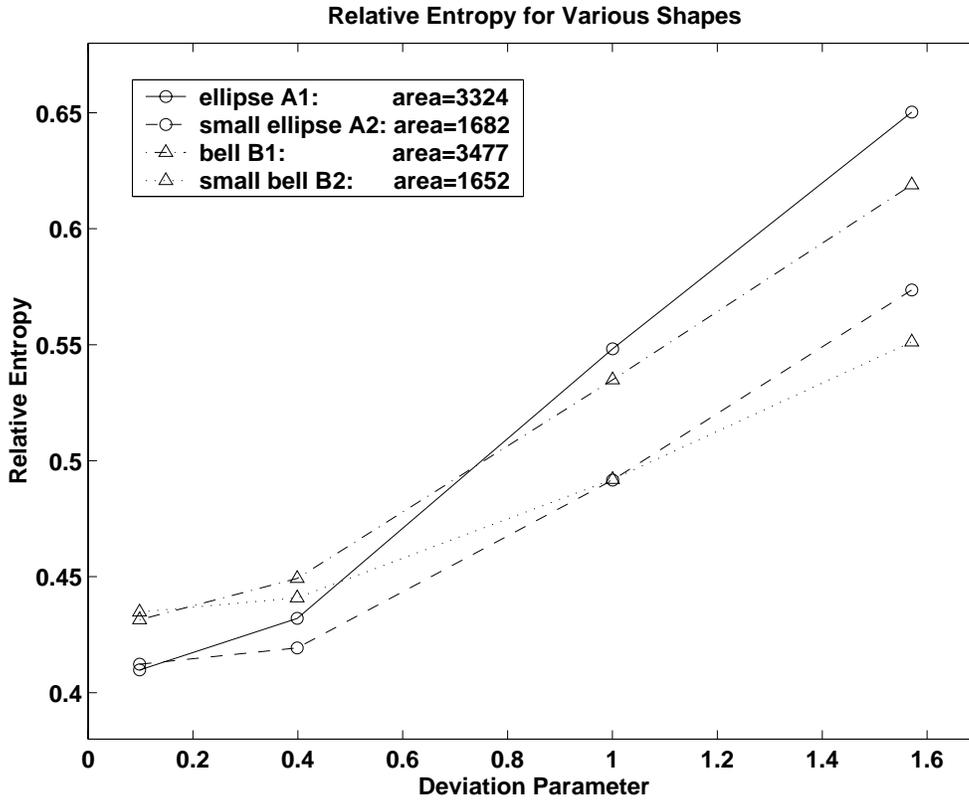


Figure 3.12. A comparison between shapes with different sizes and convexity. Four curves listed in $\kappa = \pi/2$, from top to bottom are for the (imperfect) ellipse $A1$, bell $B1$, small ellipse $A2$ and small bell $B2$, shown in Fig. 3.10. A measure with small κ will pick shapes by convexity. A measure with large κ will pick shapes by size/proximity; therefore, either small sized shapes or the bells which own strong proximity near the neck will be favored. When $\kappa = 0.1$, the case with the least size preference, the measures for all similar shapes with different sizes will coincide with each other. ■

small, we pick shape by convexity and when κ is larger, we either pick the smaller size shapes or pick the bell shape with stronger proximity. The details are shown in Fig. 3.12.

Convexity comparison of shapes with the same size When two shapes share the same size, we can easily pick up the ellipse as the more convex shape by choosing $\kappa \leq 0.6$. A large κ will favor proximity and therefore pick the bell as the more favorable shape. We have $\mathcal{S}_{A1} < \mathcal{S}_{B1}$ and $\mathcal{S}_{A2} < \mathcal{S}_{B2}$ when $\kappa \leq 0.6$.

Convexity comparison of shapes with any sizes To make the measure capable to be applied in a wider range, we compare shapes with different sizes, achieved by using an even smaller $\kappa \leq 0.4$. In this case, no size/proximity issues need to be considered and the measure becomes a “convexity measure”. We have $\mathcal{S}_{A1}, \mathcal{S}_{A2} < \mathcal{S}_{B1}, \mathcal{S}_{B2}$ if $\kappa \leq 0.4$ is provided. The smallest κ or $\kappa = 0.1$ provides a measure with the least sensitivity to the size. As we can see, the values of similar shapes with different sizes are coincided to each other in this situation.

Large κ and size/proximity preference When κ is large, it is the case of diffusion with low certainty in orientation. We will have the result favoring smaller size shapes, as we have $\mathcal{S}_{A2}, \mathcal{S}_{B2} < \mathcal{S}_{A1}, \mathcal{S}_{B1}$. When similar size shapes are compared, a shape with stronger proximity will prevail. Therefore, the bell $B1$ is favored over the ellipse

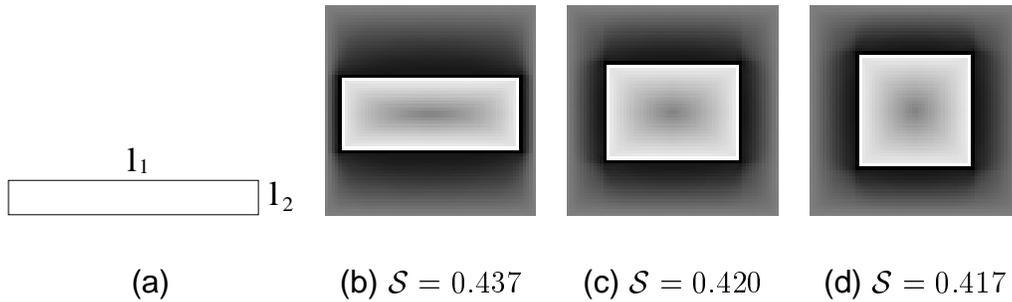


Figure 3.13. (a) rectangle with ratio l_1 to l_2 . (b) (c) (d) samples of different ratios l_2/l_1 and the relative entropy $D(P_x||Q)$. The measures are (b) 0.437, (c) 0.420 and (d) 0.417, given the parameter $\kappa = 0.1$. ■

A_1 and B_2 is favored over A_2 , by having smaller measures.

3.4 Comparison of Convex Shapes and Prägnanz Law

We would like to discuss our convexity measure for convex shapes. That is, for shapes that are categorized into one group, “convex shapes” by the mathematical definition, our measure will give different judgment. We say that our measure has a preference of “simple shapes” which has been discussed by the Law of Prägnanz [90] [45], described in Chapter 1. The result is given by a simulation of $\kappa = 0.1$. When a large κ is applied, we have the preference similar to the one provided by the decay model.

3.4.1 From Rectangle to Square

In Fig. 3.13, we apply the model to a series of rectangles with ratio of side length from 0 to 1. As shown in the experiments, the closer to square the rectangle is, the better the convexity measure. A thin rectangle will get a good score by the decay diffusion measure where more diffusion is promised by good proximity of opposite sides. But this symmetry will be cancelled by the computation of Eq. 3.10.

This same preference can also be given by the shape compactness CP, square of perimeter divided by area, defined in Sec. 2.1. For a square with size length a and a rectangle with size lengths b and c , we have

$$\text{CP}(\text{square}) = \frac{4}{\pi} < \frac{1}{\pi} \left(2 + \frac{b}{c} + \frac{c}{b} \right) = \text{CP}(\text{rectangle}) \text{ if } b \neq c.$$

3.4.2 From Triangle to Circle

In Fig. 3.14, we examine various regular shapes. Unlike the decay model giving a measure with the preference from triangle to circle, this orientation model gives a reverse order, prefers circle most, then, hexagon and triangle. By using the Kullback-Leibler measure, diffusion from neighboring inducers with *similar orientations* will accumulate around θ^* , direction of the resultant vector \hat{o}^* (Eq. 3.9); hence, giving a good score. For a cross check, the square in Fig. 3.13(d) has $\mathcal{S}_{\text{hexagon}} < \mathcal{S}_{\text{square}} < \mathcal{S}_{\text{triangle}}$, falling between the triangle and hexagon. Remember that when a large κ is

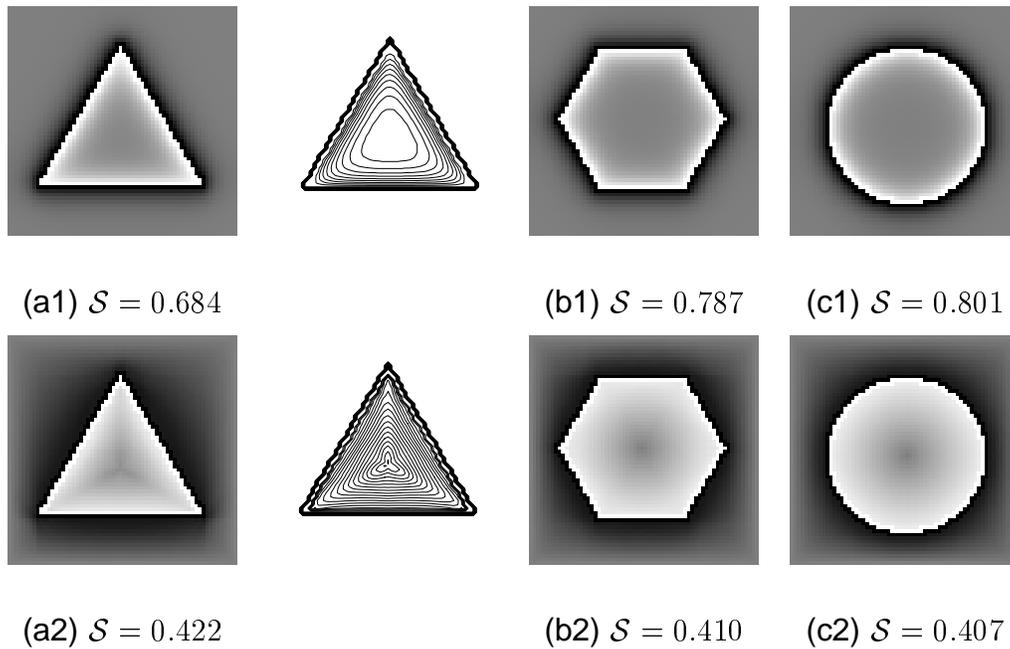


Figure 3.14. The results by both of (1) the decay model and (2) the orientation model for various regular shapes: square, hexagon and circle. (a1) (b1) (c1) the diffusion maps from the decay diffusion process and (a2) (b2) (c2) the relative entropy derived from the orientation process. We also show their isocontours for the triangle shape. Their measures from the decay model are (a1) 0.684, (b1) 0.787, (c1) 0.801 and the orientation model gives the measures as (a2) 0.422, (b2) 0.410 and (c2) 0.407 with $\kappa = 0.1$, a reverse ordering. The result by the orientation diffusion shows a simulation of the Law of Prägnanz (described in Chapter 1). Note that the square in Fig. 3.13(d) falls between triangle and hexagon. Their area is provided by (a) $|\text{triangle}| = 897$, (b) $|\text{hexagon}| = 1361$, and (c) $|\text{circle}| = 1469$ (pixels). ■

applied, we have the reverse preference, the same as the one provided by the decay measure.

The shape compactness CP will also give the same order of preference. As we have discussed in Sec. 2.1, the shape compactness is not appropriate to be applied in F/G problems. For balanced-sized regions, shape compactness will give the same result if the image frame is not considered to be part of the shape boundaries. On the other hand, larger regions will be selected, opposite to our perception, if we do consider the image frame as part of the shape boundaries. The results created by different considerations of the image frame will be discussed in Sec. 3.6.1. Another difference between the shape compactness measure and the orientation diffusion measure is that they have different considerations for structures in different scales. It will be discussed in Sec. 3.5.

3.5 Coarse Scale Structures

The decay diffusion process is not appropriate to measure convexity of shapes with small perturbations in fine scales as in Fig. 3.15 or Fig. 3.16. Due to the exponential drop in the energy, the measure derived from the decay process catches only the properties in fine scales. But for the shapes in (a1) and (a2), we are more interested in the global properties which describe them as circles up to some noise on fine

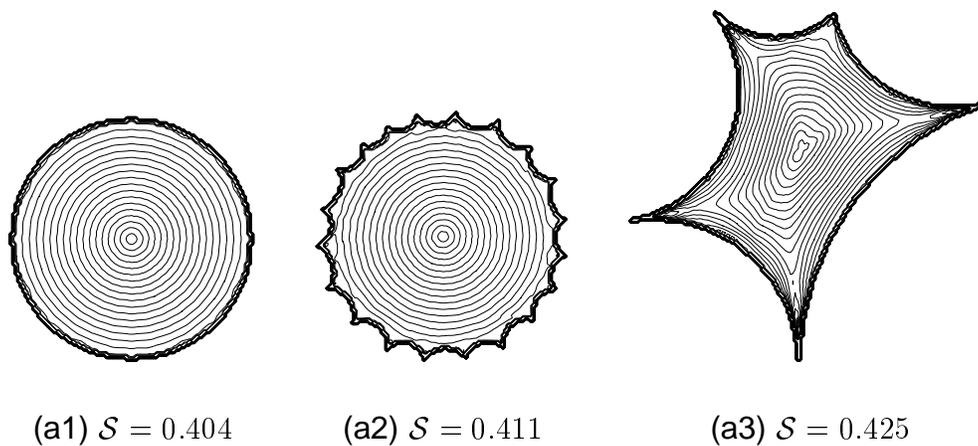


Figure 3.15. Level sets of relative entropy for shapes with different convexity in fine and coarse scales. (a1) circle, (a2) (imperfect) circle with concave perturbations on its boundaries and (a3) concave shape. Their convexity measures are (a1)=0.404 (a2)=0.411 and (a3)=0.425 with $\kappa = 0.1$. ■

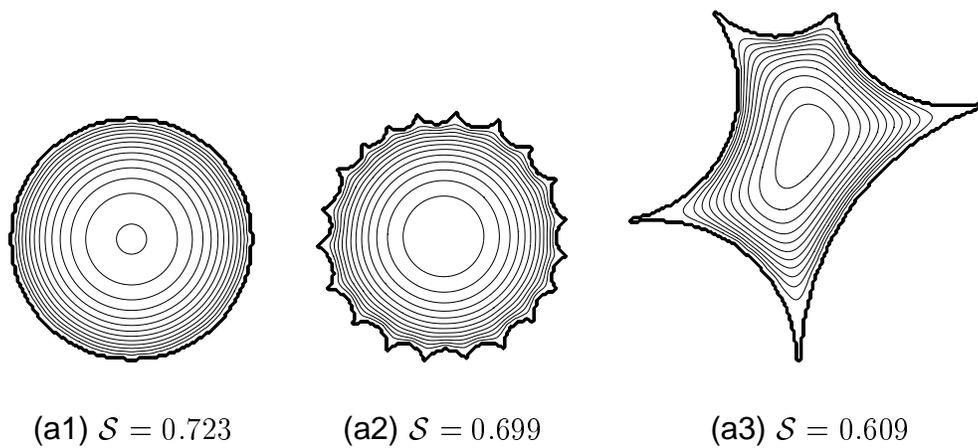


Figure 3.16. Level sets from the decay process for shapes with different convexity in various scales. Their convexity measures are (a1)=0.723 (a2)=0.699 and (a3) 0.609. ■

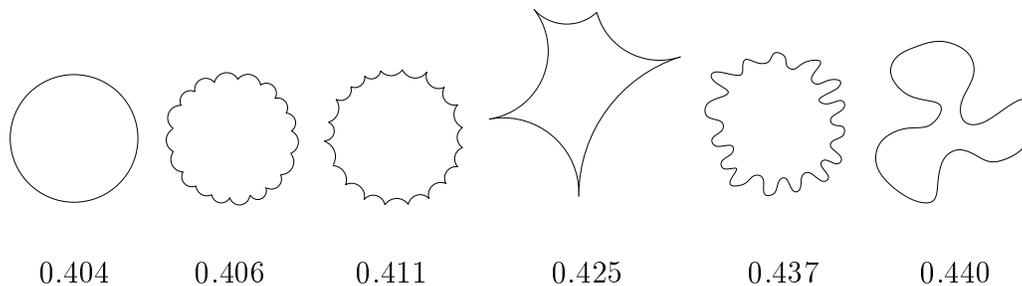


Figure 3.17. Convexity measure for various shapes with different convexity in different scales, from the most convex to the least convex one ($\kappa = 0.1$). ■

scales. In Fig. 3.15, by the orientation process, the result shows (a1) a circle is more convex than (a2) a circle with concave perturbation in fine scales. However, both of them are more convex than the concave shape in (a3). It matches our expectation that both of (a1) and (a2) are more convex than (a3) in coarse scales. Fig. 3.16 shows the result generated by the non-oriented decay model. The result does not give the shape convexity, if convexity is considered in coarse scales. We also study other various shapes, which is shown in Fig. 3.17. The result shows a different preference from preference given by the compactness.

3.6 Implementations

We choose the parameters as $\kappa = 0.4$, $\lambda = 0.1/|\Omega|$, divided by the area of the region Ω and the number of orientations is chosen to be 32 unless different comments are

made. The 2-D delta function is chosen in the same manner as we did in the decay diffusion process and the delta function coefficient is chosen to be $\epsilon = 10^{-5} \cdot |\partial\Omega|$, a constant normalized by the perimeter of Ω . The normalization of the parameters λ and ϵ according to area and perimeter of Ω respectively is not essential for the convexity measurement. As we discussed in Sec. 3.3.1, a small enough λ and κ can bring us to a measurement close to a size-invariance one. The standard deviation used in the Kullback-Leibler measure is set to be 2.0 all the times unless different comment is given.

3.6.1 Boundary Condition and Shape Surroundedness

We adopt the absorbing barriers as boundary/frame condition throughout our simulations. We would like to give an example to illustrate how different choices of boundary condition/shape surroundedness can lead to different judgment of salience.

In the image where size/proximity can dominately decide the F/G phenomenon (e.g., Fig. 2.7 (c)) we can use the orientation measure to predict the “correct” F/G. On the other hand, a different boundary/frame consideration can be made where the image frame is also considered as inducer boundaries. The salience competition becomes the comparison between the (closed-) thinner rectangle shapes and the (closed-) thicker rectangle shapes. The study in Sec. 3.4.1 suggests that the black

thicker rectangles, closer to squares, will have a better convexity measure, therefore provide a different answer. The preference of the white regions in our visual system supports the first choice of boundary condition. It implies that *image frames should not be considered as part of the shape boundaries*.

We would like to relate our models, either the decay model or the orientation model, to the issue of shape surroundedness which is known to be one of the factors affecting the decision of salience. We can easily understand it from our models, with or without the orientation consideration. It predicts that shapes with full surrounded boundaries will have better/smaller measure than the ones without closed boundaries.

Chapter 4

Internal Shape Representation

4.1 Symmetry Information

To illustrate that internal shape representation is included in our diffusion models, we compute shape axis. The idea that shape axis can be captured by our diffusion models is not particularly new or surprising. For example, based on the work of Siddiqi and Kimia [80] or Kimia et al. [44], we know that the diffusion process over shapes will “meet” (yield shocks) at the symmetry axis. However, the results from our models, based on variational approaches are provided in the static state. Therefore, the definition of axis will no longer be decided by pairs of points, but a global property of the boundaries. The term “shape axis” is used instead of “symmetry axis” or “medial axis” to distinguish from the axis generally mentioned in the medial axis transform or

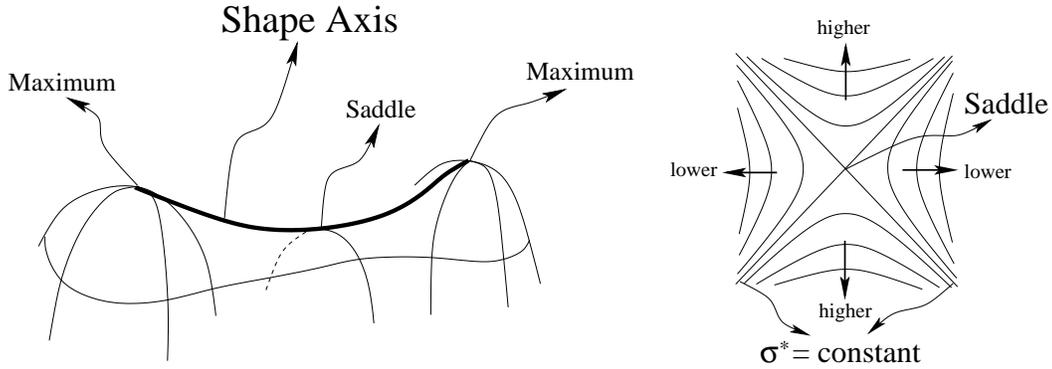


Figure 4.1. Shape axis picked in the surface \mathfrak{S} with its characteristic points. The surface $\mathfrak{S} \equiv (\mathbf{x}, -\sigma^*(\mathbf{x}))$ is derived from the decay process. The right hand side shows its iso-contour near the saddle point. ■

Blum [4] [5]. This shape axis, for being derived by a global consideration, will give more robust result (e.g., Fig. 4.8). We give two types of axis representations, provided by result from the decay process and the orientation diffusion process. To compare to other approaches, our concentration is not aimed at giving a sparse, graph-structured axis or discussing matching between two sparse axes [51] [52]. Instead, we show the axes computed by our models, and give a demonstration level discussion.

4.1.1 Symmetry by Traveling in σ^* -surface

For a 2-D shape Ω , we will define the shape axis by examining the surface $\mathfrak{S} \equiv \{(\mathbf{x}, -\sigma^*(\mathbf{x})) : \mathbf{x} \in \Omega\}$ in 3-D, with the result σ^* derived from the decay diffusion

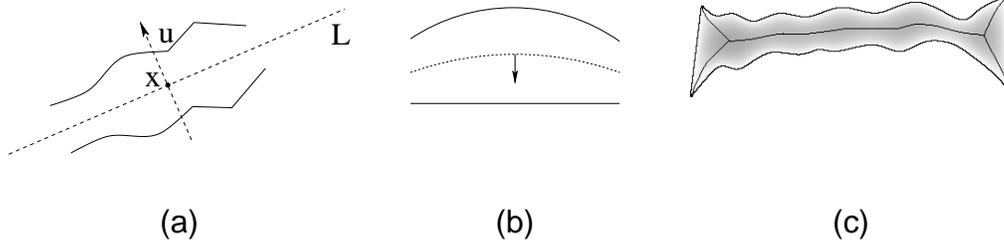


Figure 4.2. (a) Shape mirror symmetric along the line L with $L \perp \mathbf{u}$ at the axis point \mathbf{x} . (b) convexity will “push” axis toward the less-convex side. (c) shape with fuzzy symmetry and its relatively smoother axis, compared to its boundaries and the conventional symmetry axis. ■

process. Let us start from a shape Ω in Fig. 4.2(a), which is mirror symmetric along the line L with $L \perp \mathbf{u}$ at \mathbf{x} for a unit vector \mathbf{u} . Apparently, the line L should be (one of) the axes of Ω . For the axis point $\mathbf{x} \in L$, the decay diffusion result σ^* is expected to have

$$\sigma^*(\mathbf{x} + t\mathbf{u}) = \sigma^*(\mathbf{x} - t\mathbf{u}). \quad (4.1)$$

Divide it by t and take the limit $t \rightarrow 0$, we obtain

$$\lim_{t \rightarrow 0} \frac{\sigma^*(\mathbf{x} + t\mathbf{u}) - \sigma^*(\mathbf{x} - t\mathbf{u})}{t} = 0, \quad \Leftrightarrow \quad \sigma_{\mathbf{u}}^*(\mathbf{x}) = 0. \quad (4.2)$$

It will be used as a necessary condition to judge if a point \mathbf{x} is on any axis of Ω . By the principle of “value symmetry” in Eq. 4.1, we can predict that for a shape with a more convex boundary on one side, as shown in Fig 4.2(b), the shape axis will

appear near the less-convex side, if the axis does exist. It is because points in the convex side will get stronger support from the boundary inducers (Sec. 2.2). This fact makes the shape axis different from the conventional symmetry axis. We define the shape axis set \mathfrak{A} to be the projection (to the plane $z = 0$) of the curve set \mathfrak{C} where each length-parametrized curve $\mathcal{C}(s) \in \mathfrak{C}$ satisfies:

1. The curve $\mathcal{C}(s)$ is a curve going from one characteristic point to another characteristic point on \mathfrak{S} . The set of characteristic points includes saddle points, local maxima and junctions* .
2. At each point $\mathbf{x} \in \mathcal{C}(s)$, in direction of the curve normal \mathbf{n} , we have $\sigma_{\mathbf{n}}^*(\mathbf{x}) = 0$, satisfying the necessary condition Eq. 4.2.

To meet the first condition, e.g., for the surface of a half sphere, we have only the projected circle center as the “axis”. Because there is only one characteristic point on the surface, the maxima in the middle. Before giving the algorithm to compute the axis, let us discuss those characteristic points. The maximum is caught by

$$\nabla\sigma^* = (\sigma_x^*, \sigma_y^*) = \mathbf{0} ,$$

with a positive Hessian matrix \mathbf{H} s.t.

$$\mathbf{t}^T \mathbf{H} \mathbf{t} = \mathbf{t}^T \begin{bmatrix} \sigma_{xx}^* & \sigma_{xy}^* \\ \sigma_{yx}^* & \sigma_{yy}^* \end{bmatrix} \mathbf{t} \geq 0 , \quad \forall \mathbf{t} \neq \mathbf{0}$$

*We mean the place on \mathfrak{S} where its projection on $z = 0$ forms a junction of Ω .

with the row matrix \mathbf{t}^T denoting transpose of $\mathbf{t} = (u, v)$, the small change around \mathbf{x} . Similarly, we can find the minimum if we have $\mathbf{t}^T \mathbf{H} \mathbf{t} \leq 0, \forall \mathbf{t} \neq 0$. When either of them happens for different \mathbf{t} 's, we find the saddle point or we can use the equivalent criteria

$$\det(\mathbf{H}) = \sigma_{xx}^* \sigma_{yy}^* - \sigma_{xy}^{*2} < 0,$$

evaluated at the point \mathbf{x} . In our algorithm, *the junction information will be given as an input.*

We collect all axis points by traveling through the curves from a characteristic point to another characteristic point. We choose the “uphill” direction. By the maximum principle, the field σ^* will decay from the shape boundaries to interior part of the shape (meaning $(\mathbf{x}, -\sigma^*(\mathbf{x}))$ will be rising from the boundaries). Therefore, the only possible trials fitting the condition (1) are trials from a saddle point or a junction to a maximum point or another saddle point. The algorithm to find the axes is therefore given by an initial step and a recursive step:

- i Find the maximum points and saddle points on the surface \mathfrak{S} .
- ii Start from each saddle point and junction, we choose the most steepest uphill direction to visit, until any maximum is reached.

2-D projection of these trails, including all the maxima and saddle points are called the maximum shape axes or shape axes, opposite to those axes going downhill from

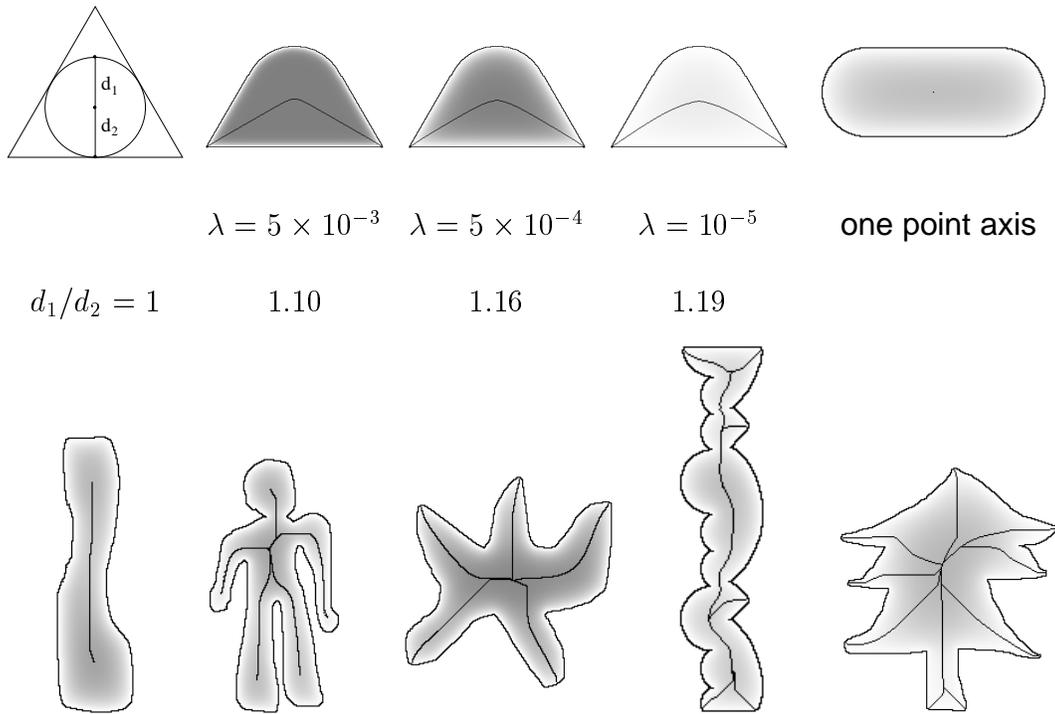


Figure 4.3. Shape axis in various shapes, computed from the result of decay diffusion with the axis superficially added on the decay field σ^* . ■

saddle points, called the minimum shape axes.

4.1.2 Results of σ^* -surface Traveling Method

Some examples of the shape axis are shown in Fig. 4.3. In particular, for shapes like circles, ellipses, rounded rectangles, we have only their center points considered as the axes. In fact, all “rounded-border” convex shapes will have only the center point considered as their axes, because no saddle points will be detected inside those

shapes.

To compare the shape axis and the traditional medial axis, we can refer the shape axes found in Fig. 4.3. In the first row, three shapes are designed by regular triangles with upper angles substituted by their inscribing circles. Their axes found here are lower than their medial axes which should stay in the middle of the shapes. We say that the shape axis is “smoother” than the medial axis defined by Blum [4], especially when the decay coefficient λ is small. In Fig. 4.2(c), we have smoother axes than its boundaries and medial axes.

4.1.3 Symmetry by Local Computation Method

We would like to define another set of shape axes by a *local computation method*, based on the result of orientation diffusion process. We would like to emphasize the differences of this method from previous approaches.

- i The axis information given by this method will not be binary represented. Namely, instead of giving information to tell if a point is on an axis or not, we compute a map to describe “how likely” a point is on an axis or how “strong” the axis is.
- ii We use a local computation technique to catch the shape axis. The axis will be locally decided by the result from orientation process. Once the computation

of the process is finished, the step to collect shape axes is relatively easier and more robust than the approach used in other axis generators.

- iii The junction feature considered as special axis points, will be collected by a similar computation.

In our approach, the axis is shown by a continuous map. The continuous information give us more clue to distinguish between main axis and minor axis. The minor axis will be declared by a weak representation, e.g., the weak axis created by an obtuse angle (first order shock) or a pair of weakly mirror-symmetric boundaries (higher order shock) (Fig. 4.4 - 4.8). It provides a natural way of axis pruning if there is a need. For the second topic, the robustness of our approach comes from two levels. The local computation is robust, compared to, e.g., the traveling method used in the previous section. Because missing the detection of an axis point can prevent any further exploration and miss an axis branch. Secondly, the robustness of our computation comes from the fact that the shape axis computed by us are based on a global property of the shape. We examine the effect of protrusion created on shape boundaries in Fig. 4.8. The result show that we are able to deal with shape boundaries with small protrusion (ignore), boundaries with large protrusion (surrender proportionally) with a transition between them. The third topic as well as the junction detection will be discussed in next section (Sec. 4.2).

Local Computation Method

We take result from the orientation diffusion process to generate a map called symmetry map. In each point of the map, continuous other than binary information will be provided for the declaration of shape axis. This information will be computed by a local computation, opposite to the traveling method used in the previous section.

Let us examine the convexity vector $\hat{\sigma}^*(\mathbf{x})$ used in Eq. 3.9, derived by vector summation of the orientation field $\sigma^*(\mathbf{x}; \theta)$. First, the vectors are normalized to unit vectors, and the divergence of them is computed as the symmetry information. It is

$$\sigma_{\text{sym}}^* = \text{div} \left(\frac{\hat{\sigma}^*}{\|\hat{\sigma}^*\|} \right). \quad (4.3)$$

The idea why we use this quantity to represent the symmetry information can be demonstrated in Fig. 4.4. Recall that $\sigma^*(\mathbf{x}; \theta)$, up to a constant, can be viewed as summation of the random walks starting at boundaries with inward orientations and reaching \mathbf{x} with the orientation θ at certain time. The vector map $\hat{\sigma}^*(\mathbf{x})$ recording vector summation of $\sigma^*(\mathbf{x}; \theta)$ is shown in Fig. 4.4(a). After normalization, only the orientation part is memorized, which is in (b). We can view the shape axis as the place with most sinkage in this unit-vector map. Compare to the shock language in Siddiqi et al. [80] and Kimia et al [44], the axis is the place where diffusion meets each other, after the meeting point, the direction of diffusion can turn with an abrupt angle, as if certain “amount of substance” has disappeared. The result of Eq. 4.3 is

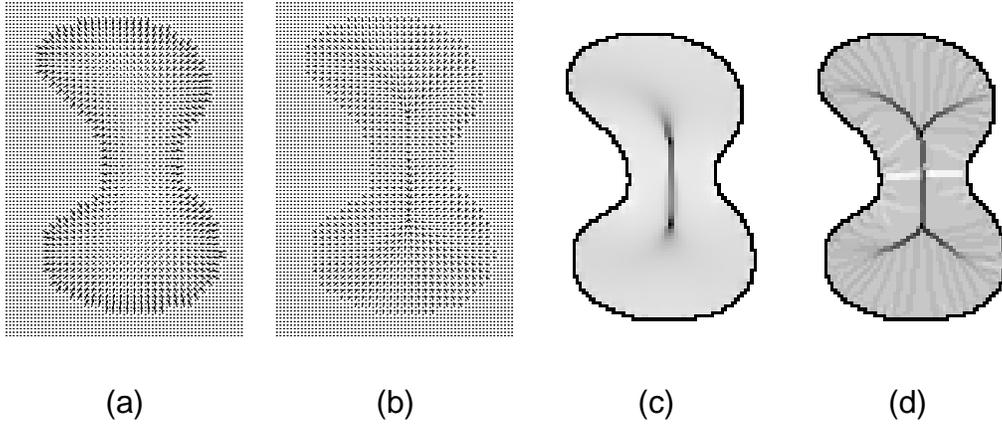


Figure 4.4. Symmetry information of the bell shape, by computing the divergence of unitary $\hat{\sigma}^*$, vector summation of σ^* in (c) or $\hat{\sigma}_M^*$, the maximum vector of σ^* in (d), among all orientations. ■

shown in Fig. 4.4(c).

Let us also provide another similar criteria, by substituting the vector map $\hat{\sigma}^*(\mathbf{x})$ in Eq. 4.3 with the maximum vector $\hat{\sigma}_M^*(\mathbf{x})$ of $\sigma^*(\mathbf{x}; \theta)$, among all orientations. That is,

$$\hat{\sigma}_M^*(\mathbf{x}) = \sigma^*(\mathbf{x}; \theta_M) \cdot \mathbf{u}_M \quad \text{if} \quad \sigma^*(\mathbf{x}; \theta_M) = \max_{\theta \in [0, 2\pi)} \{ \sigma^*(\mathbf{x}; \theta) \},$$

with $\mathbf{u}_M = (\cos \theta_M, \sin \theta_M)$, and similarly

$$\sigma_{\text{symM}}^* = \text{div} \left(\frac{\hat{\sigma}_M^*}{\|\hat{\sigma}_M^*\|} \right). \quad (4.4)$$

The correspondent result of σ_{symM}^* is shown in Fig. 4.4(d). The approach by computing the maximum vector does provide more details. However, in general, it needs

finer mesh (in orientation) or larger κ to ease the quantization error.

4.1.4 Results of Local Computation Method

The results of shape axis, derived by computing the vector summation (Eq. 4.3) and the maximum vector (Eq. 4.4) can be seen in Fig. 4.5 and Fig. 4.6 respectively. The way we recognize the symmetry axis or shape axis is different from the traditional views. Instead of producing the binary symmetry information, we provide a continuous map to tell “how likely” a location stays on any axes or how “strong” this axis is. To produce the common binary symmetry axis, we can do it through a threshold step.

A Continuous Axis

By the continuous representation, we can distinguish the “strong axis” (shown in a strong representation) from the “weak axis” (shown in a weak representation). This representation naturally provides a way for axis-pruning if we want to remove the axis created by noisy boundaries. In the circle shape, many sub-axes are produced, but only the “point axis” in the center are considered as the main axis (e.g., check the symmetry map in Fig. 4.7(b1)). In another example, we test an unbalanced star in Fig. 4.7(a). With the stronger axes indicated by darker intensities, we obtain stronger axes for sharper angles, as we expected. We can better understand this topic by a

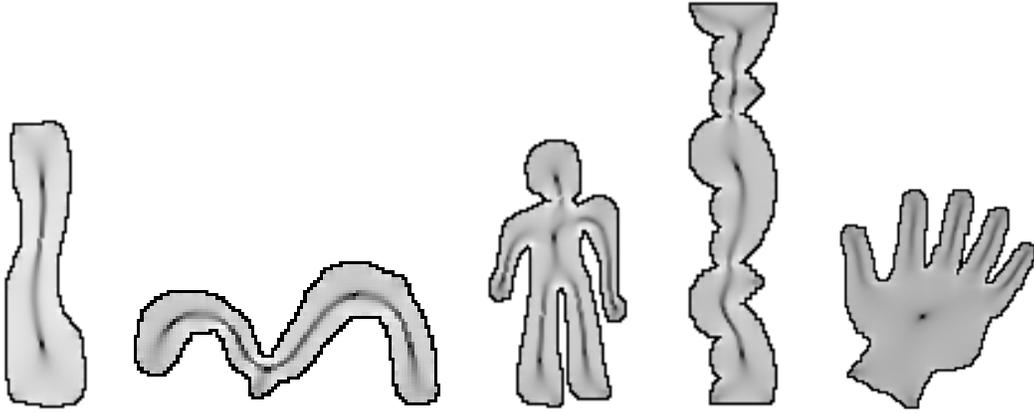


Figure 4.5. Shape axis σ_{sym} in various shapes, computed from result of the orientation process, by using the vector summation result $\hat{\sigma}^*$. ■

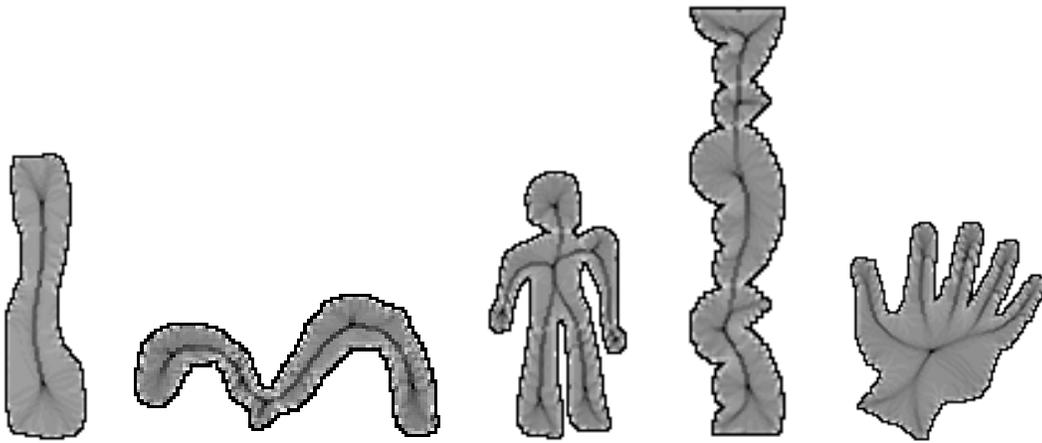


Figure 4.6. Shape axis σ_{symM} , computed from result of the orientation process, by using the maximum vector $\hat{\sigma}_M^*$. A result of $\kappa = \pi/2$ is applied. ■

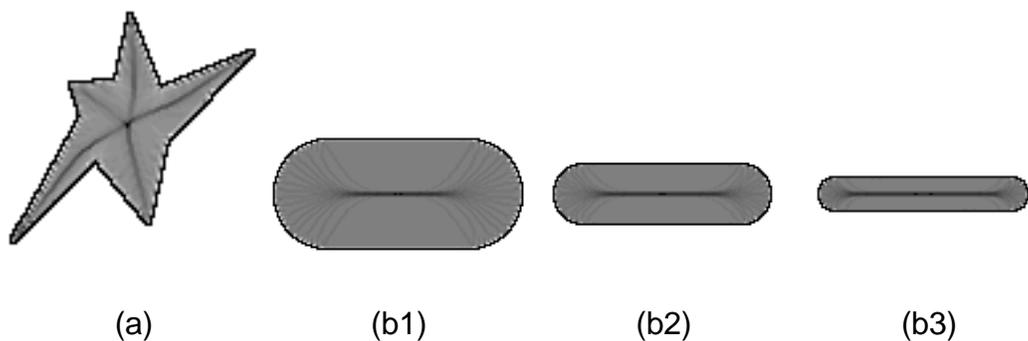


Figure 4.7. Shape axis with a continuous manner. (a) Sharper angles have axes with stronger representations. (b1) - (b3) Circular boundaries have many lighter sub-axes, compared to the axis point in the middle. The axis has similar representation from (b1) to (b3), even the widths are different. We choose the maximum vector $\hat{\sigma}_M^*$, and select the deviation factor $\kappa = \pi/2$. ■

series of experiments from a triangle, square, hexagon to a circle. The triangle will have three strongest axes. Then we have four axes for the square, but with a lighter representation. After all, we end up the lightest (sub-)axes for the circle, except the center.

As another example to illustrate the continuity of our information, we check the hand shape in Fig. 4.6. We obtain stronger axes for the fingers, but weaker axes linking from the finger-axes to center of the hand. Because the boundaries between the fingers and the center are not well symmetric, compared to boundaries of the fingers.

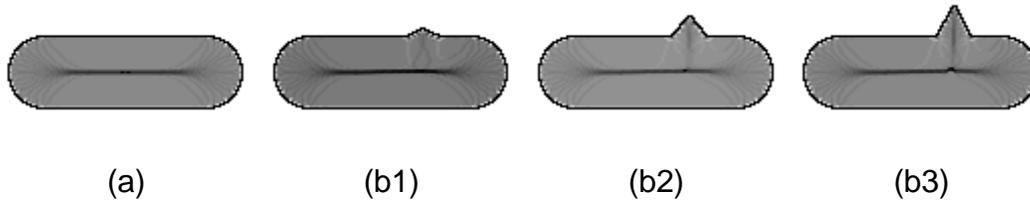


Figure 4.8. Shape axis with protrusion on boundaries. We have the transition from (b1) ignored, (b2) having “sub-axis” to (b3) disturbing the main axis by a faction. The axis created by the protrusion can simply be pruned by a threshold method, if the protrusion is considered as noise. In (b3), a pruning is not appropriate because the main axis is affected. We choose the maximum vector $\hat{\sigma}_M^*$, and select the deviation factor $\kappa = \pi/2$ (Eq. 3.5). ■

In the next section, we study the effect of protrusions on boundaries. The result can only be achieved by considering shape axis as a global feature of shapes.

Robustness and Effect Created by Protrusion

To see how protrusion or noise on boundaries can affect the shape axis, we study the axes in Fig. 4.8. In (b1), the main axis stays the same as we perturb the boundaries from (a) by a small protrusion. But a larger perturbation can (b2) eventually produce a new axis or (b3) even change location of the main axis. As we said in the previous section, the continuous representation provide a natural way to do the axis-pruning. In this case, if only the main axis is needed, the axis created by the protrusion (noise)

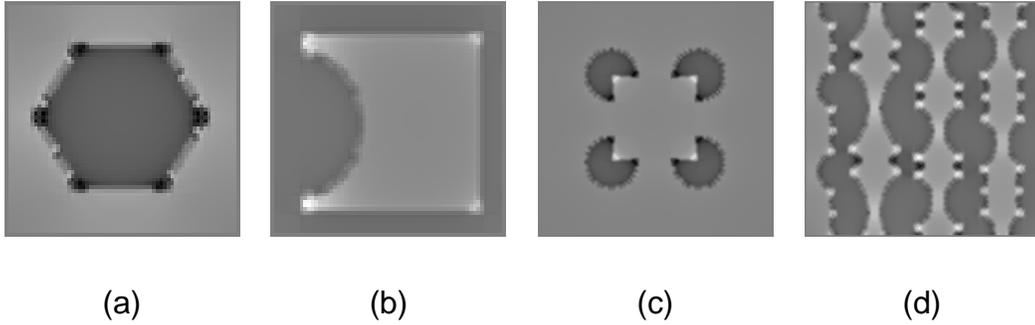


Figure 4.9. Junction information σ_{junc}^* . The black or white color is given due to arbitrary choice of the direction for $\hat{\sigma}^*$ (for specifying inside or outside of the boundaries). ■

can be removed by a simple threshold step.

4.2 Junction Detection

We consider junctions of shapes to be the special axis points. For a given shape Ω , junctions are on the shape axes, and they should be on the boundaries, too. It is described by

$$\mathfrak{J} \equiv \mathfrak{A} \cap \partial\Omega .$$

Follow this idea, it is natural to compute the junction information through the map

$$\sigma_{\text{junc}}^* = \text{div}(\hat{\sigma}^*) . \quad (4.5)$$

Without any normalization in Eq. 4.3, axis points located near the boundaries will



Figure 4.10. (a) Junction information with minor perturbation on boundaries. (b) boundaries with high curvature parts. The result shows the ability of global consideration for junction detection. ■

have strongest intensities of diffusion, possess longest length of $\hat{\sigma}^*$ and largest (or smallest, depends on the choice of sign) divergence; therefore be caught as junctions. We use this map to guide us finding junctions. Some results are given in Fig. 4.9. Either the darkest or brightest area can be considered as the location of junctions.

As we can observe, the sharper the angle is, the stronger the intensity we can derive (e.g., Fig. 4.9(b)). The quantization effect does not have any major contribution to our result.

4.2.1 Junction as a Global Property of Shapes

In our approach, junction feature is considered as a global attribute. We consider two cases where the junctions will not be located on the boundaries.

When the boundaries have minor perturbations, we can still collect correct information even if only the junction feature in the coarsest scale is needed. In Fig. 4.10(a), our approach provides correct detection for a triangle with noisy boundaries. The perturbations, if there is any, can create only “minor” junctions.

In our approach, we do not consider junction feature as a binary information. Instead, we provide the transition from curves with low curvature to high curvature, and eventually become discontinuous. In Fig. 4.10(b), a rounded square is tested, and an eased junction information is obtained.

Chapter 5

Visual Organization

5.1 Introduction

Discovering perceptual organizations is understood as the transition from low-level vision to high-level vision. Without the help from object naming, pixels are formed as surfaces abstractly or symbolically during this process. The illusory figures, such as the Kanizsa square shows an example where the intensity edges are not the only information matters for such intergration, but also geometry of the edges. Fig. 5.1 (a1) gives another example of illusory figures. The open edges around the empty space show cues for occlusions, implying the surface completion with a triangle in the middle. To describe this choice of completion, first of all, the triangle border gives a good (linear) continuation. Secondly, unlike the figure given in (a2), the

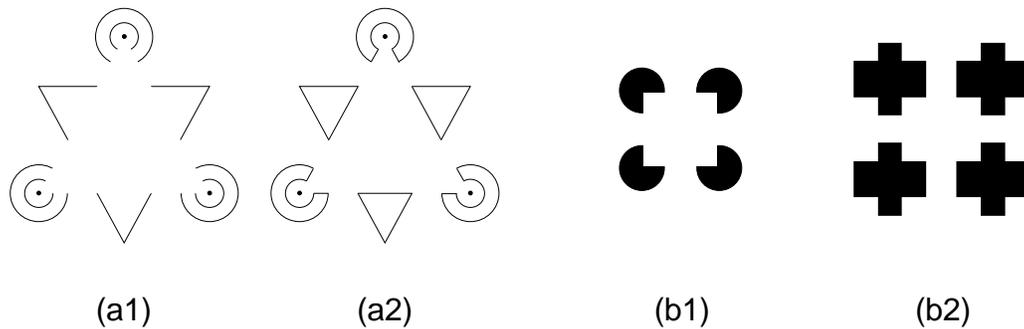


Figure 5.1. (a1) (a2) Illusory triangle, adapted from Kanizsa [39]. (a1) a figure with open edges showing tendency to closure while (a2) a figure with closed edges showing no tendency to amodal completion. (b1) Kanizsa square and a similar figure in (b2), but no salient square is perceived. ■

figure/ground interpretation is consistent with such surface interpretation. Namely, border ownerships of the edges in (a2) implies that those outer shapes should be considered as self-sufficient ones instead of being occluded by the single triangle in the middle. In general, on edges, binary choices, to be on top or on bottom, can be made and the whole selection set is called an organization for surfaces. To speak of junctions, we may have more interpretations, such as in Fig. 5.2 (Kumaran et al. [48], Geiger et al. [19]). The choices or interpretations we made on edges or junctions are called hypotheses.

Our organization reconstruction is based on a diffusion process going from local to global. Sparse information assigned on feature points, called hypothesis is taken

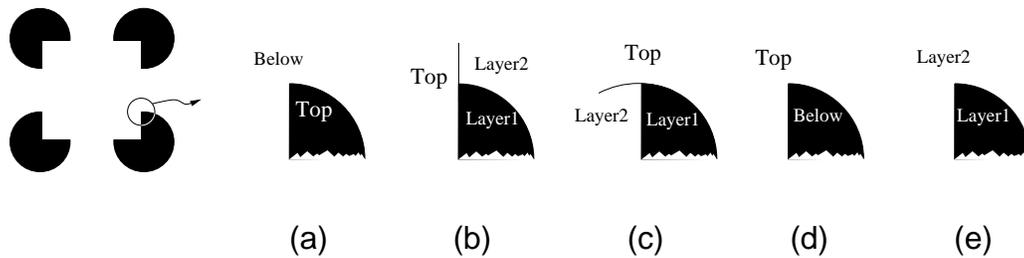


Figure 5.2. Each L-junction (corner) locally suggests five (5) possible salient surface interpretations. Note that (b) and (c) give L- to T-junction transition, suggesting three surface layers, with the hat of the T as the salient one. (e) removes the junction from the salient surface. ■

as the input. This is the place we can freely make assumptions of figure/ground. Afterwards, we decide the visual organization based on evaluation of output of the process. A winner organization is suggested by a winner hypothesis which gives the best evaluation. In the following section, we discuss the method for organization reconstruction. A perfect feature detection scheme is assumed.

Numerous computational models have been proposed to construct the illusory figures. Ullman [86], Heitger & Heydt [28], Guy & Medioni [27], Grossberg & Mingolla [26], Kellman & Shipley [42], Mumford [58], Parent & Zucker [64] and Williams & Jacobs [91] have built models to catch illusory contours, associated with various scenarios to describe the “good continuation” between occluded/broken curves. Brady & Grimson [6] and Nitzberg & Mumford [60] have used the surface

reconstruction formulation to describe the salient/occluded surfaces. Besides the construction of the illusory contours/surfaces, our model provides a reason to describe why the illusory effect happens in one case instead of others by a surface/regional description.

5.2 L-junctions and Prior Distribution of Hypotheses

The idea of picking correct visual organization is inspired by the Kanizsa square. However, we do not perceive the similar phenomenon for another similar figure, the four-crosses in Fig. 5.1 (b2). The key to make this difference comes from the smooth curve preference for surface reconstruction. We consider an organization picking scheme with such preference. In particular, on surface contours, L-junctions, T-junctions and their interpretations will be studied.

For a given image I , we build the bridge between sparse data and global decision by the decay diffusion process used in Sec. 2.2. A discrete version which has been discussed in Appendix B is examined here. We have the hypothesis choice basis

$$H = \{-1, 0, 1\},$$

with 1 and -1 represent foreground and background respectively and 0 is the neutral hypothesis indicating an ignorance. Each edge feature ϕ is assigned by such local

decision as

$$\phi \rightarrow a(\phi) \in H. \quad \forall \phi \in \Phi$$

A hypothesis set for the whole image \mathbf{I} is the set $\{a(\phi) : \phi \in \Phi\}$, for a given (edge) feature set $\Phi \subset \mathbf{I}$. In particular, we separate the feature set to be two parts as $\Phi = \Phi_A \cup \Phi_O$ with the set Φ_A/Φ_O to be the set of assigned/ignored feature respectively. The energy functional is given similar to Eq. B.4, Eq. B.5 and Eq. B.6 as,

$$E_{\text{decay}}(\sigma|\sigma_0) = E_{\text{data}}(\sigma|\sigma_0) + E_{\text{smooth}}(\sigma),$$

trying to minimize the data fitting and the smoothness functionals. However, new setting of the function σ_0 is given by

$$\sigma_0(\mathbf{x}) = \begin{cases} a(\mathbf{x}), & \text{if } \mathbf{x} \in \Phi_A \\ 0. & \text{if } \mathbf{x} \in \mathbf{I} - \Phi_A \end{cases} \quad (5.1)$$

as well as a new function of Λ given by

$$\Lambda(\mathbf{x}) = \begin{cases} \frac{1}{\epsilon}, & \text{if } \mathbf{x} \in \Phi_A \\ 0. & \text{if } \mathbf{x} \in \mathbf{I} - \Phi_A \end{cases} \quad (5.2)$$

The decisions on edges are given by either assigning opposite choices as 1 and -1 on both sides of the edges representing figure and ground or assigning 0 meaning an ignorance. For the Kanizsa square, various hypotheses can be made. Some examples can be found in Fig. 5.3(b1) & (c1), with b/w representing $\sigma_0 = -1/1$ respectively. The hypotheses we particularly interested are those with consistent assignment in

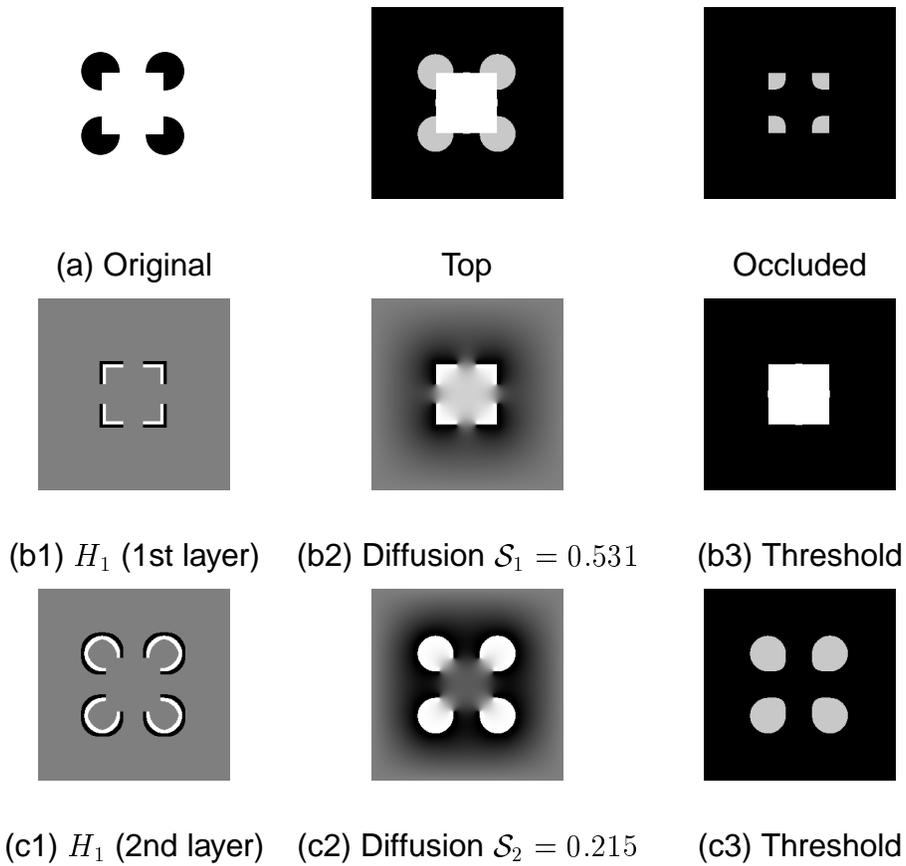


Figure 5.3. Kanizsa Square image with its optimal organization of the top surfaces (white is salient, grey is the 2nd layer and black is the 3rd layer); and occluded layers (amodal completions). (b1) (c1) The winner hypotheses consist of two sets of edge assignments. On each edge, black/white or gray represents $\sigma_0 = -1/1$ or 0 respectively. (b2) (c2) the result of diffusion, showing the map σ^* and (b3) (c3) threshold, given by selecting $\sigma^* > 0$. We have the organization criteria $\mathcal{O} = (0.531 + 0.215)/2 + 4\alpha = 0.653$, with $\alpha = 0.07$. ■

neighboring edges except those neighbors crossing the edges. However, we do allow more freedom in the assignment of junctions, by adopting the interpretations of junctions in Fig. 5.2.

For each hypothesis set, a diffusion result can be derived, called σ^* . By the same linear transform used in Eq. 2.12, we find the probability map p^* .

Zero Energy Decay For the function $\Lambda(\mathbf{x})$ in Eq. 5.2, we mainly follow the choice made in Geiger et al. [19] where the decay coefficient λ is chosen to be 0, no energy decay throughout our discussions and experiments in this chapter.

Modal, Amodal Completion and Multiple Surfaces

After the process, a surface is collected by those points \mathbf{x} with $p^*(\mathbf{x}) > 0.5$, called the salient surface. Usually, scenes contain more than one surface. We will obtain all of them by consecutively applying the above procedure. Namely, after the removal of a surface, the “used/not ignored” features Φ_A where $\sigma_0 \neq 0$ will be removed as well, and another hypothesis set can be assigned for those “not used/ignored” features $\Phi_O = \Phi - \Phi_A$. Multiple surfaces can be found in such manner. We have $\Phi = \bigcup_{s=1}^S \Phi_s$ for a S -surface reconstruction with the assigned feature $\Phi_A = \Phi_s$ in the s -th surface reconstruction. For the Kanizsa square, two surfaces (three layers, including the background) can be formed in Fig. 5.3(b3) & (c3). The top row shows

the typical multiple-surface reconstruction.

Prior Distribution of Hypothesis

The prior distribution for shapes suggests smooth contours. Therefore, a favored hypothesis set should be the one with more cancellation of L-junctions. That is, on L-junctions, we assign non-zero $a(\phi)$ on both sides of the peak in different steps of surface reconstructions. Thus the L-junctions become T-junctions, such as in Fig. 5.2(b) or (c). We evaluate such movement by the cost

$$C_1(\{a(\phi)\}) = -\#(\text{L-junction} \rightarrow \text{T-junction}) .$$

Because number of L-junctions from the input is fixed, the previous criteria can be substituted by another equivalent one as

$$C(\{a(\phi)\}) = \sum_{s=1}^S \#\text{L-junctions}(p_s^*) ,$$

counting number of L-junctions for the result p_s^* on the surface s . The criteria C and C_1 are just a constant away from each other. A smaller one gives a higher preference.

Entropy of Visible Surfaces

To give the decision for the best organization, we also compute the (normalized) entropy for all S surfaces by

$$\mathcal{S}(\{a(\phi)\}) = -\frac{1}{S} \sum_{s=1}^S \frac{1}{N_s} \sum_{\mathbf{x} \in \Omega_s} p_s^*(\mathbf{x}) \log p_s^*(\mathbf{x}) + (1 - p_s^*(\mathbf{x})) \log(1 - p_s^*(\mathbf{x})) , \quad (5.3)$$

where $N_s = |\Omega_s|$ denotes number of pixels in the s -th surface Ω_s .

Organization Criteria

Our final organization criteria is the sum of both criteria

$$\mathcal{O}(\{a(\phi)\}) = \alpha C(\{a(\phi)\}) + \mathcal{S}(\{a(\phi)\}),$$

for a parameter α . For instance, the hypothesis yielding the original Kanizsa image will have $\mathcal{O} = \alpha C(\{a(\phi)\}) + 0 = 12\alpha$ because we have a closed-door diffusion.

The parameter α balances both terms can be decided empirically by the following experiment (Fig. 5.4).

For the Kanizsa square, one set of hypotheses H_0 can be given by the original setting, suggesting the original four pecmen as the (only) salient surfaces. Another set of hypotheses H_1 , the typical hypothesis set suggests (1) a salient square and (2) four occluded circles on the back. The human experiments suggest that for Kanizsa images, salience of the square decreases as we move those four pecmen away from each other as those from Fig. 5.4(a) to (c). Empirically, we can obtain the transition point where the hypothesis H_0 and H_1 yield the same preference. This situation can be described by

$$4\alpha + \mathcal{S}^* = 12\alpha \quad \Leftrightarrow \quad \alpha = \mathcal{S}^*/8,$$

where \mathcal{S}^* is the entropy value of configuration of the transition point (perception

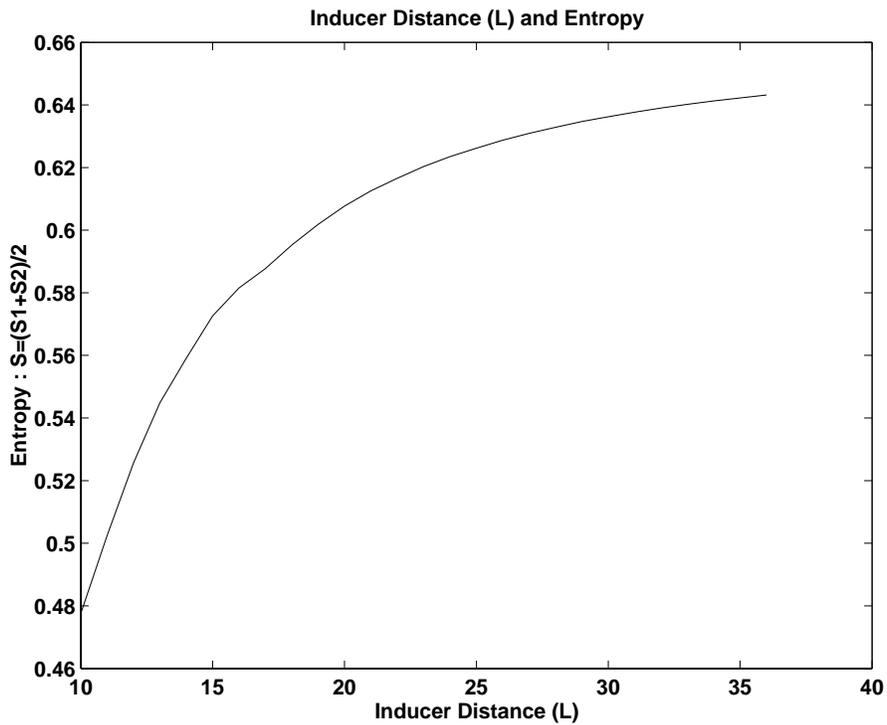
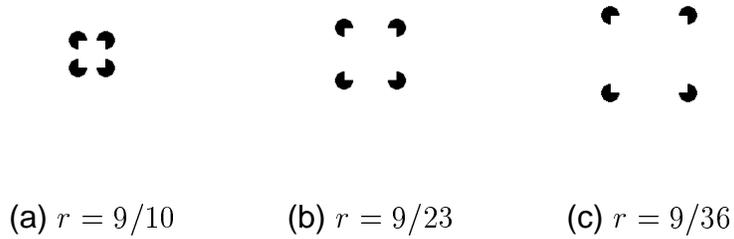
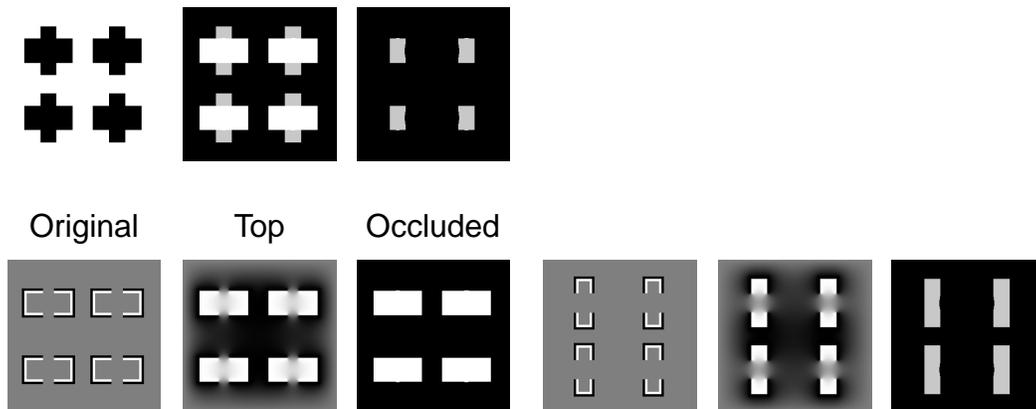


Figure 5.4. A sequence of Kanizsa Square images with different ratios of $r = R/L$, where R is the radius of the circles, and L is the distance between inducers. The graph shows the entropy S versus r for the Kanizsa square. We crudely estimated that the change in perception from the organization where the square is on top to the one where the four pecmen on top occurs for $r = 9/17 = 0.53$. ■



Hypothesis $\mathcal{S}_1 = 0.265$ Threshold Hypothesis $\mathcal{S}_2 = 0.433$ Threshold

Figure 5.5. Four crosses image and its typical surface reconstruction. This reconstruction is suggested by the winning hypothesis set H_D consists of three-layer reconstruction. The 1st layer (left) and the 2nd layer (right) give strips crossing each other. The organization criteria gives $\mathcal{O} = 2.589$. Note that the larger (not the smaller) rectangles are salient because of the lower entropy. ■

wise). We crudely choose $\alpha = 0.07$ and this setting is used throughout all of our experiments. The hypothesis set producing the *smallest* organization criteria will be the winning hypothesis.

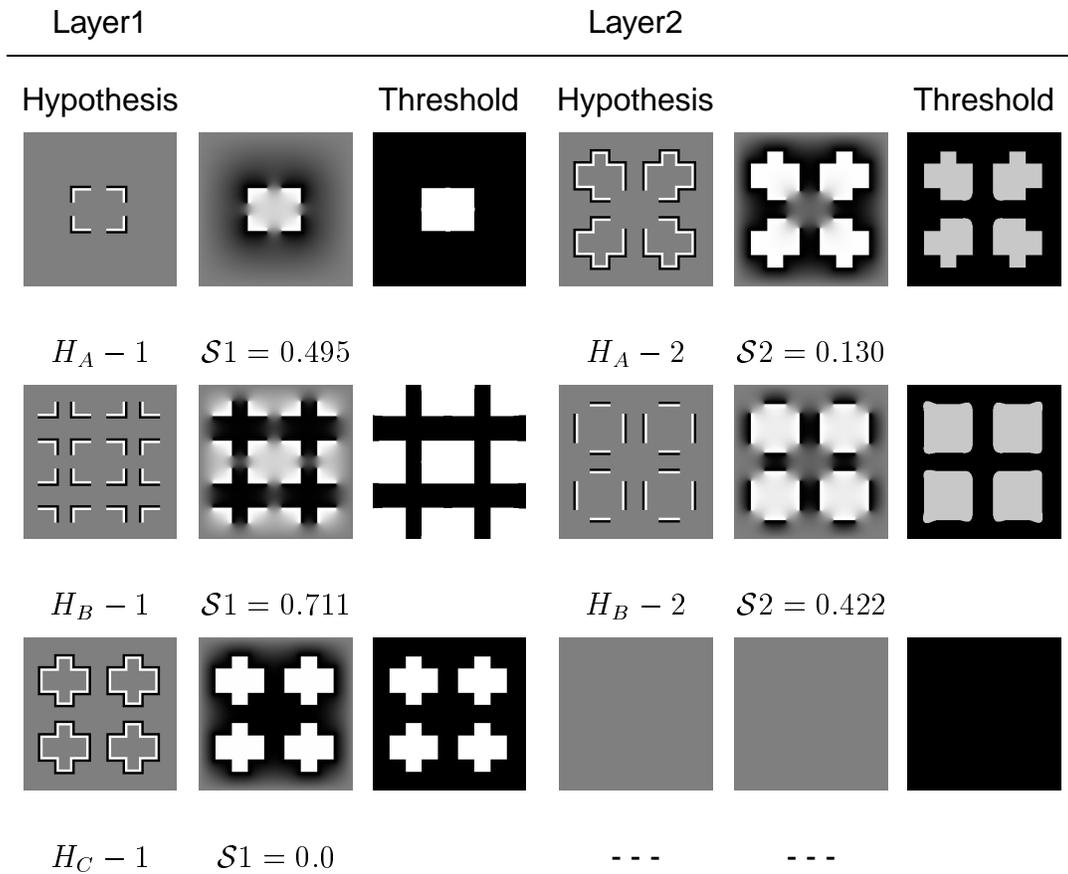


Figure 5.6. Four crosses image with its various organizations based on the hypothesis H_A , H_B , H_C and H_D (in Fig. 5.5). Number of L-junctions for the hypothesis H_A , H_B , H_C and H_D are (1st layer+2nd layer) ($4 + 36 = 40$), ($32 + 0 = 32$), ($48 + 0 = 48$) and ($16 + 16 = 32$). The organization criteria $\mathcal{O}(\{a(\phi)\})$ are 3.113, 2.807, 3.360 and 2.589, making the winner to be the hypothesis H_D . The second voted hypothesis is the hypothesis H_B while the illusory square, hypothesis H_A has a lower score. H_C contains only two layers while others contain three layers. ■

5.2.1 Experiments

We use two images, the Kanizsa square and the four-cross image in Fig. 5.5 to illustrate our ideas. The perceptual difference in these two images leads to two different surface reconstructions. For the Kanizsa image, a salient square in the middle is observed easily, but no such phenomenon exists in the four-cross image. We argue that the preference of smooth contours gives such phenomenon. An organization which is built on contours with less L-junctions or less high curvature points will be chosen. Let us evaluate this by our organization criteria.

In Kanizsa image, the typical hypothesis set gives the salient square, and simultaneously 8 L-junctions are removed in such movement. On the other hand, for the four-cross image, the hypothesis set giving a salient square in the middle is not the most favorable one, compared to another hypothesis set which yields more L-junction cancellations (Fig. 5.5). Consider various hypotheses in Fig. 5.5 and Fig. 5.6, either hypothesis H_A where a single square is in front or H_B where 9 squares are produced will both give bad organization criteria. The best organization is given by the hypothesis H_D , giving crosses as pairs of strips crossing each others. A total of 16 L-junctions are removed by this hypothesis. Its organization criteria is given by $\mathcal{O} = (0.265 + 0.433)/2 + 32\alpha = 2.589$.

By our model, the Kanizsa square image is interpreted as three layers, a salient

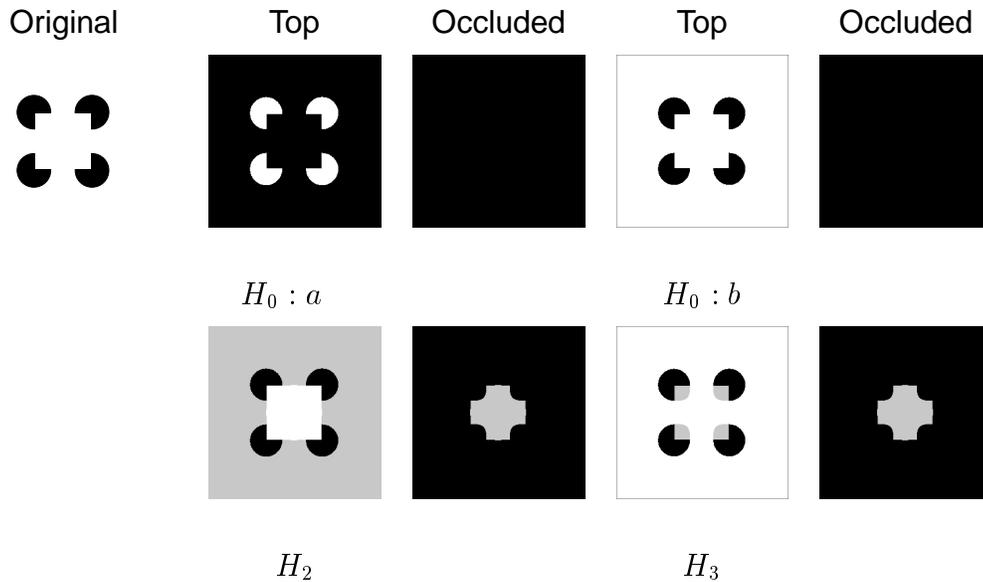


Figure 5.7. Kanizsa square image with its various surface reconstructions.

The hypothesis $H_0 : a$ is the hypothesis keeping the original input. $H_0 : b$ is its dual hypothesis. Remember brightness represents salience. Their organization criteria are $\mathcal{O}(H_0 : a) = 12\alpha = 0.84$, $\mathcal{O}(H_0 : b) = 0.666 + 12\alpha = 1.506$ and $\mathcal{O}(H_2) = \mathcal{O}(H_3) = (0.531 + 0.760)/2 + 4\alpha = 0.926$. The hypothesis H_1 in Fig. 5.3 suggests the best organization by its evaluation $\mathcal{O}(H_1) = 0.653$. ■

square, four (black) circles and background consecutively. The organization criteria is given by $\mathcal{O} = (0.531 + 0.215)/2 + 4\alpha = 0.653$. There are other organizations (Fig. 5.7), but with larger/worse organization costs.

5.2.2 Discussions

First of all, this approach is different from the approach by Kumaran et al. [48]. Each of their hypothesis set might produce 2 dual hypothesis sets, figure/ground or ground/figure in our set of organizations. Namely, no knowledge of F/G were considered in their context. The junction interpretations such as Fig. 5.2(a) and (d) were not distinguished. Accordingly, while their computation of entropy is made throughout all image space, we only compute entropy for the “visible surfaces”, i.e., points for $p^* > 0.5$.

To speak of minors of this approach. The organization criteria contain two parts, entropy of the visible surfaces and the cost of L-junctions. More L- to T-junction transformations can ease the cost of organization criteria and give good organization. Therefore, this approach is highly relies on the input from a good junction detector. In general, we should be able to pick up junctions or high-curvature points on images without making artificial border between them.

Let us discuss complexity of this approach. To pick the best organization, we use a brute force search. So number of the trials is exponential to the number of junctions. Because junctions are the only place the hypothesis can be switched, based on our assumption that only consistent hypothesis (in neighboring points on one side of the edges) is considered.

Chapter 6

Conclusion

We discussed the issues of figure ground separation, shape representation and perceptual organization which are generally considered as low- to middle-level vision problems. Two types of diffusion processes, the decay process and the orientation process were defined, followed by two convexity measures. Our convexity measurement was a support to the traditional 1-0 convexity definition in mathematics which separates shapes into two groups, “convex” and “concave” without any further descriptions. The continuous manner in our convexity measures was established to provide the transition from “highly convex” to “highly concave” shapes. Besides convexity measuring, our processes were able to collect other shape information, including symmetry and junction information.

Decay Diffusion Process The decay process was used in particular for simulating the F/G phenomenon. For a F/G image, when the size of regions and the number of inducers were normalized to be the same, we predicted the “correct” answer corresponding to the outcome from the majority of human tests.

Orientation Diffusion Process For the orientation process, a wider range of images have been tested. First, it could deal with F/G images with sizes of regions not yet normalized to be the same. A brand new design of introducing a free parameter allowed us to surf between the size-invariant convexity measuring and the convexity measuring with the small-size preference. Other than predicting the F/G phenomenon, the model could be considered as a definition for shape convexity. The shapes with either similar or not similar sizes could be compared to each other, by the choice of a small parameter κ . To discuss the perfectly-convex shapes which was categorized as one group by the mathematical definition, our measure showed the ability to distinguish between them. The study showed that the convexity measure based on the orientation process gave the scores of circles better than hexagons, followed by squares and then triangles. Another series of experiment suggested the preference of squares over rectangles. It could be understood as the simulation of the Law of Prägnanz for 2-D shapes. A Kullback-Leibler distance was used as the convexity measure.

Internal Shape Representation Our diffusion processes could also be used to compute the internal shape structure, generally known as the axis study. In our approaches, two so called “shape axes” have been defined, one from the decay diffusion and the other from the orientation diffusion. We defined two operations generating two types of axes. For the decay process, a traveling method on the 3-D surface $\mathfrak{S} \equiv (\mathbf{x}, -\sigma^*(\mathbf{x}))$ was used, based on the result $\sigma^*(\mathbf{x})$ derived from the process. For the orientation process, a simple technique, called the local computation method was used to collect the axes. In this case, we adopted a different point of view for the axis finding. Instead of producing any binary axis information, we derived the symmetry maps σ_{sym} and σ_{symM} to give suggestions for shape axes. Our approach had many benefits. First, once the diffusion process was done, the computation, via some local operations, was easier compared to other algorithms to find the symmetry axis. Secondly, our shape axes were computed by a global consideration. Within a shape, we could distinguish between strong axes and weak axes. When noise, such as protrusion were added to the boundary, our result showed a continuous transition from ignored, having minor axes to disturbing the main structure of the shape. Also, the junction detection was discussed. To us, the junction feature was no more than the *special axis point*, just those happened to be on the borders. Combine with the convexity measuring, we tried to emphasize that our simulations has the goal to cover wide range of the applications.

6.1 Continuous Simulation based on Global Considerations

In our simulations, based on the global considerations, we tried to remove the artificial assumptions generally used in many studies:

- i For the convexity measuring, we did not draw the border between the convex and concave shapes which was used in mathematics. Instead, we defined continuous convexity measures. They are more appropriate than the 1/0 definition in most vision applications.
- ii For axis, we did not assume that the place was on or off the axes, instead, we gave the map telling “how likely” the place was an axis point and how strong the axis is. Therefore, we did not have to apply any additional heuristic or non-heuristic algorithms to deal with the axis pruning or selection. The axes generated by us had different weights themselves to guide the process.
- iii For junctions, we did not distinguish between “discontinuous junctions” and “high curvature junctions”. Instead, a map was derived and the threshold decided later could be used to collect the information whatever fits to our applications.

6.2 Shape Description and Shape Completion

In our approach, the partial and completed shapes were treated similarly. For the perceptual organization problem, the same framework, the decay process and entropy measurement used for the F/G separation, was applied to select the salient shapes.

To improve the decay model used in the perceptual organization finding, we put our future concentrations on:

- i Adopt the orientation diffusion process in the perceptual organization finding.

The assumption of perfect junction detection will no longer be necessary.

- ii Search for an efficient algorithm or approach to achieve the best organization.
- iii Real image studies.

Appendix A

2-D Dirac Delta function

In this section, we define the Dirac delta function in Eq. 2.10 and Eq. 3.4 used by the decay process and the orientation process respectively. We state the orientation case here. The case for the decay process is the same or simpler.

Let us assume an image defined by $\mathbf{I} \equiv \{\mathbf{x} : \mathbf{x} \in [0, A] \times [0, B]\}$ and an orientation set $\Theta \equiv \{\theta : \theta \in [0, 2\pi)\}$. For a given closed region Ω surrounded by its boundary

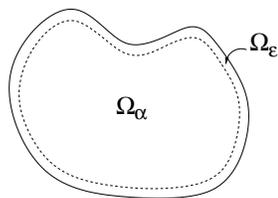


Figure A.1. A 2-D Dirac-delta function defined on Ω . It is partitioned into two regions Ω_ϵ and $\Omega_\alpha \equiv \Omega - \Omega_\epsilon$. ■

curve $\partial\Omega$ (assuming $\partial\Omega \subset \Omega$), we define the 2-D Dirac-delta function. First of all, let us partition the region Ω into two parts Ω_ϵ and Ω_α by *

$$\Omega_\epsilon \equiv \{\mathbf{x} : d(\mathbf{x}, \partial\Omega) \leq \epsilon\},$$

and $\Omega_\alpha \equiv \Omega - \Omega_\epsilon$. On Ω , a piecewise constant function D is defined as

$$D(\mathbf{x}) = \begin{cases} 1/|\Omega_\epsilon|, & \text{if } \mathbf{x} \in \Omega_\epsilon \\ 0, & \text{if } \mathbf{x} \in \Omega_\alpha \equiv \Omega - \Omega_\epsilon \end{cases}$$

where $|\Omega_\epsilon|$ indicates area of the region Ω_ϵ . We have

$$\int_{\Omega} D(\mathbf{x}) d\mathbf{x} = \int_{\Omega_\epsilon} D(\mathbf{x}) d\mathbf{x} = \int_{\Omega_\epsilon} \frac{1}{|\Omega_\epsilon|} d\mathbf{x} = 1.$$

A 2-D Dirac-delta function $\delta_2(\mathbf{x})$ or simply as $\delta(\mathbf{x})$ is stated as

$$\delta(\mathbf{x}) \equiv \lim_{\epsilon \rightarrow 0} D(\mathbf{x}),$$

and its integral is given by

$$\int_{\Omega} \delta(\mathbf{x}) d\mathbf{x} = \int_{\partial\Omega} \delta(\mathbf{x}) d\mathbf{x} \equiv \lim_{\epsilon \rightarrow 0} \int_{\Omega} D(\mathbf{x}) d\mathbf{x} = 1, \quad (\text{A.1})$$

to make as if the limit and the integral sign can be exchanged from each other. Right now, let us see this definition is meaningful to our applications.

*The distance function d is defined as $d(\mathbf{x}, \partial\Omega) = \min\{d(\mathbf{x}, \mathbf{x}_1) : \mathbf{x}_1 \in \partial\Omega\}$. The 2-D Euclidean distance function is applied.

For a binary input, an intensity function I characterizing Ω , we define the hypothesis function $\sigma_0(\mathbf{x}; \theta)$ as

$$\sigma_0(\mathbf{x}; \theta) = \begin{cases} \mathbf{u}_\theta \cdot \mathbf{n} & \text{if } \mathbf{u}_\theta \cdot \mathbf{n} \geq 0 \wedge \mathbf{x} \in \partial\Omega \\ 0 & \text{if } \mathbf{u}_\theta \cdot \mathbf{n} < 0 \vee \mathbf{x} \in \Omega - \partial\Omega \end{cases}$$

and based on it, a function $\Lambda(\mathbf{x}; \theta)$ is defined by

$$\Lambda(\mathbf{x}; \theta) = \begin{cases} \sigma_0(\mathbf{x}; \theta) \delta(\mathbf{x}) & \text{if } \mathbf{u}_\theta \cdot \mathbf{n} \geq 0 \wedge \mathbf{x} \in \partial\Omega \\ \lambda & \text{if } \mathbf{u}_\theta \cdot \mathbf{n} < 0 \vee \mathbf{x} \in \Omega - \partial\Omega \end{cases}$$

The operation “ \cdot ” denotes the conventional inner product in \mathbf{R}^2 . The vector \mathbf{u}_θ and $\mathbf{n}(\mathbf{x})$ respectively denote the unit vector $(\cos \theta, \sin \theta)$ and the unit inward normal on $\mathbf{x} \in \partial\Omega$. For the case that a smooth intensity function $I(\mathbf{x})$ is available, we can define $\sigma_0(\mathbf{x}; \theta)$ as

$$\Lambda(\mathbf{x}; \theta) = \begin{cases} \frac{1}{M_e} \left| \frac{\partial I(\mathbf{x})}{\partial \mathbf{u}_\theta} \right| \delta(\mathbf{x}) & \text{if } \mathbf{x} \in \partial\Omega \\ \lambda & \text{if } \mathbf{x} \in \Omega - \partial\Omega \end{cases}$$

The constant M_e is defined by $\max\{|\frac{\partial I(\mathbf{x})}{\partial \mathbf{u}_\theta}| : \mathbf{x} \in \mathbf{I}, \theta \in \Theta\}$, evaluated on a compact set. In either case, we would like to discuss the integral of $\Lambda(\mathbf{x}; \theta)$ on $\Omega \times \Theta$, for two parts $\partial\Omega \times \Theta$ and $(\Omega - \partial\Omega) \times \Theta$. The inside part is

$$\int_{(\Omega - \partial\Omega) \times \Theta} \Lambda(\mathbf{x}; \theta) d\mathbf{x} d\theta = 2\pi \lambda |\Omega|. \quad (\text{A.2})$$

The “border part” can be estimated by

$$\int_{\partial\Omega \times \Theta} \Lambda(\mathbf{x}; \theta) d\mathbf{x} d\theta \leq \int_0^{2\pi} \int_{\partial\Omega} 1 \cdot \delta(\mathbf{x}) d\mathbf{x} d\theta = \int_0^{2\pi} 1 d\theta = 2\pi, \quad (\text{A.3})$$

which gives an upper bound of the integral. Furthermore, let us assume, at each point \mathbf{x} of the curve $\partial\Omega$, the normal direction to the curve $\partial\Omega$ is always the gradient direction of the intensity function I at point \mathbf{x} . The lower bound of the integral can be given as by

$$\begin{aligned} \int_{\partial\Omega \times \Theta} \Lambda(\mathbf{x}; \theta) d\mathbf{x} d\theta &\geq \int_{\partial\Omega} \int_{-\frac{\pi}{2}}^{\frac{\pi}{2}} m_g \cos \theta \delta(\mathbf{x}) d\theta d\mathbf{x} \\ &= m_g \int_{\partial\Omega} 2\delta(\mathbf{x}) d\mathbf{x} = 2m_g, \end{aligned} \quad (\text{A.4})$$

given $m_g = \min\{\|\nabla I(\mathbf{x})\|/M_e : \mathbf{x} \in \partial\Omega\}$, the (normalized) minimal value of length of the gradient vectors on the boundary $\partial\Omega$. We can assume $m_g > 0$. It gives $0 < \int_{\partial\Omega \times \Theta} \Lambda(\mathbf{x}; \theta) d\mathbf{x} d\theta \leq 2\pi$. Therefore it is meaningful to include the data-fitting integral in our variational functionals, as in Eq. 2.9 and Eq. 3.6.

Appendix B

Decay Diffusion Process, Random

Walk and Discrete Settings *

To write down the discrete form of the energy functional either Eq. 2.9 or Eq.2.9, first, let us discretize the information and represent the input data σ_0 , the function Λ and the diffusion field σ as column vectors by the raster scanning order. That is, for an $M \times N$ image as an index set $\mathbf{I} \equiv \{k = 1, \dots, K = N \cdot M\}$ containing the shape Ω , the input vector is given by $\sigma_0(k) : \mathbf{I} \rightarrow \{-1, 0, 1\}$ and the function Λ is $\Lambda(k) : \mathbf{I} \rightarrow \{\lambda, 1/\epsilon\}$ for a small constant ϵ called delta function coefficient where

*The conversation in this section will deal with the most general case happened in our text. It covers discussions for Sec. 2.2 and Chapter 5. Also, we will continue to use the same set of notations here in discrete case.

$1/\epsilon$ stands for the function value of $\delta_2(\mathbf{x})$ if $\mathbf{x} \in \partial\Omega$ in discrete case. We can even make a more general case. A new function $e(k, k')$ records edge information with the pixel k and one of its neighbors k' . To be precise, we set $0 \leq e(k, k') \leq 1$ to be the intensity edge between pixel k and k' , normalized by the largest intensity edge on \mathbf{I} . Follow this, an overloaded discrete edge function $e(k)$ can be defined as $e(k) = \max_{k' \in \mathbf{N}_k} e(k, k')$. We define a generalized σ_0 as

$$\sigma_0(k) = \begin{cases} e(k), & \text{if } k \in F(\partial\Omega) \\ -e(k), & \text{if } k \in G(\partial\Omega) \\ 0, & \text{if } k \in \mathbf{I} - \partial\Omega \end{cases} \quad (\text{B.1})$$

where $F(\partial\Omega)$ and $G(\partial\Omega)$ respectively indicate “figure” and “ground” side of the edges $\partial\Omega$, which sit on opposite sides of the edges. Also,

$$\Lambda(k) = \begin{cases} \frac{1}{\epsilon} \cdot e(k), & \text{if } k \in \partial\Omega \\ \lambda. & \text{if } k \in \mathbf{I} - \partial\Omega \end{cases} \quad (\text{B.2})$$

The solution or the diffusion field is given as usual,

$$\sigma(k) : \mathbf{I} \rightarrow \mathbf{R}. \quad k \in \mathbf{I} \quad (\text{B.3})$$

We write down the discrete decay diffusion energy functional (compare to Eq. 2.9)

by finite difference method as

$$E_{\text{decay}}(\sigma|\sigma_0) = E_{\text{data}}(\sigma|\sigma_0) + E_{\text{smooth}}(\sigma), \quad (\text{B.4})$$

where

$$E_{\text{data}}(\sigma|\sigma_0) = \sum_{k \in \partial\Omega} \Lambda(k)(\sigma(k) - \sigma_0(k))^2 + \sum_{k \notin \partial\Omega} \lambda \cdot \sigma^2(k), \quad (\text{B.5})$$

and

$$E_{\text{smooth}}(\sigma) = \sum_{k=1}^K \sum_{k' \in \mathbf{N}_k} (1 - e(k, k'))(\sigma(k) - \sigma(k'))^2. \quad (\text{B.6})$$

The Equivalence between Energy Minimization and Matrix Inversion

The solution to this equation can be derived by casting it into a matrix form, which is

$$E_{\text{decay}}(\sigma|\sigma_0) = (\sigma, \mathbf{D}\sigma) - 2(\sigma, \mathcal{L}\sigma_0) + C(\sigma_0). \quad (\text{B.7})$$

where “ (\cdot, \cdot) ” denotes the inner product in \mathbf{R}^K . The diagonal matrix is given by $\mathcal{L}_{kk} = \Lambda(k)$ and the symmetric and band limited matrix \mathbf{D} has the following structure

- i. The diagonal elements $\mathbf{D}_{kk} = \Lambda(k) + \mu(k, \mathbf{N}_k)$,
- ii. The non-zero off-diagonal elements are $\mathbf{D}_{kk'} = -\mu(k, k') = -(1 - e(k, k'))$
for $k' \in \mathbf{N}_k$,

for the edge function $e(k, k')$ with $\mu(k, \mathbf{N}_k) = \sum_{k' \in \mathbf{N}_k} \mu(k, k')$. Let us assume $\mu(k, \mathbf{N}_k) > 0, \forall k = 1, \dots, K$. We rewrite it as

$$E_{\text{decay}}(\sigma|\sigma_0) = (\sigma - \sigma^*, \mathbf{D}(\sigma - \sigma^*)) + C_1(\sigma_0), \quad (\text{B.8})$$

where σ^* is the solution of the equation $\mathbf{D}\sigma = \mathcal{L}\sigma_0$. It tells that the minimum is achieved when $\sigma = \sigma^*$, where

$$\sigma^* = \mathbf{D}^{-1}\mathcal{L}\sigma_0. \quad (\text{B.9})$$

For a simple case where we consider only the computation on Ω and the edge functions $e(k)$, $e(k, k')$ choose only binary numbers, this equation gives a discrete form of Eq. 2.11. For a pixel k on $\partial\Omega$, we have

$$\mu(k, \mathbf{N}_k) - \sum_{k' \in \mathbf{N}_k} \sigma(k') = \frac{1}{\epsilon}(1 - \sigma(k)) \geq 0.$$

The left side is bounded, so we can make $\sigma(k)$ as close to 1 as we want by choosing a small enough ϵ . The edge blocking occurred in Eq. B.6 does not play a role here.

Numerical Computation

We use the conjugate gradient method to compute the solution. This steepest descent method is built directly on the energy Eq. B.7. For further reading about the conjugate gradient method, please refer to [84] and [83].

B.0.1 Random Walk Formulation

The matrix \mathbf{D} can be written as

$$\mathbf{D} = \mathfrak{D}(\mathbf{I} - \mathbf{Q}), \quad (\text{B.10})$$

with \mathcal{D} being the diagonal matrix with elements $\mathcal{D}_{kk} = \Lambda(k) + \mu(k, \mathbf{N}_k)$ ($\mathcal{D}_{kk'} = 0$ for $k \neq k'$). Therefore \mathbf{Q} must have four off-diagonals non-zero and positive i.e.,

$$\mathbf{Q}_{kk'} = \begin{cases} \frac{\mu(k, k')}{\Lambda(k) + \mu(k, \mathbf{N}_k)}, & \forall k' \in \mathbf{N}_k \\ 0. & \text{otherwise} \end{cases} \quad (\text{B.11})$$

Note that in each row of \mathbf{Q} , we have $\sum_{j=1}^K \mathbf{Q}_{ij} < 1$ if the decay coefficient $\lambda > 0$.

When $\lambda = 0$, we can also directly prove that $\sum_{n=1}^{\infty} \mathbf{Q}^n$ is bounded, but we postpone the proof till further discussions. The inverse of \mathbf{D} can be given by

$$\mathbf{D}^{-1} = \left(\sum_{n=1}^{\infty} \mathbf{Q}^n \right) \mathcal{D}^{-1}, \quad (\text{B.12})$$

and so,

$$\sigma^* = \left(\sum_{n=1}^{\infty} \mathbf{Q}^n \right) g, \quad (\text{B.13})$$

with $g = \mathcal{D}^{-1} \mathcal{L} \sigma_0$, i.e., $g_k = \frac{\Lambda(k)}{\mathcal{D}_{kk}} \cdot \sigma_0(k)$. We can think g as our “new” input data.

The matrix \mathbf{Q} is the “transient” part of an “absorbing Markov chain” \mathbf{M} , where \mathbf{M} is

$$\mathbf{M} = \begin{bmatrix} 1 & \mathbf{0} \\ R & \mathbf{Q} \end{bmatrix}, \quad (\text{B.14})$$

with a $1 \times K$ zero vector $\mathbf{0}$ and R being a $K \times 1$ matrix with elements

$$R_k = 1 - \sum_j \mathbf{Q}_{kj} = \frac{\Lambda(k)}{\Lambda(k) + \mu(k, \mathbf{N}_k)}, \quad (\text{B.15})$$

so that $\sum_{j=0}^{K+1} \mathbf{M}_{ij} = 1, \forall i$. The stochastic matrix \mathbf{M} is of size $K + 1$ by $K + 1$. It is clear that

$$\mathbf{M}^n = \begin{bmatrix} 1 & \mathbf{0} \\ R' & \mathbf{Q}^n \end{bmatrix},$$

where $R' = \sum_{\alpha=0}^n \mathbf{Q}^\alpha R$. \mathbf{M} is an absorbing Markov chain in the sense that if one starts with any state s_0 , by successively applying \mathbf{M} , one ends up in the state $s_{\text{final}} = (1, 0, 0, \dots, 0)$, i.e., $s_{\text{final}} = \lim_{n \rightarrow \infty} s_0 \mathbf{M}^n$ (we are applying the matrix to the vector to the “left”). This is clear from the fact that $\lim_{n \rightarrow \infty} \mathbf{Q}^n = 0$. This is why \mathbf{Q} is the transient part of \mathbf{M} . The interpretation of the state $s = (\pm 1, 0, \dots, 0)$, is of “death” of the random walk, since the random walk jumped to the zero-th coordinate that is not a pixel in the lattice. Moreover, once it reaches the zero-th coordinate it never leaves, i.e., $s = (\pm 1, 0, 0, \dots, 0)$ is an eigenstate of \mathbf{M} or $s \mathbf{M} = s$ (or equivalently, to the “right” s^T is an eigenstate of \mathbf{M}^T , with $\mathbf{M}^T s^T = s^T$. This is the absorbing state.

Random Walk View

Let us clarify the random walk interpretation of the matrices.

$(\mathbf{Q}^n)_{kj}$ - probability that a random walk that started at pixel k will reach pixel j in the n -th step.

R_k - probability that a random walk will vanish at pixel k (move to the zero-th state coordinate).

$R'_k = (\sum_{\alpha=0}^n \mathbf{Q}^\alpha R)_k$ - probability that the random walk vanished at pixel k in n steps.

$(\sum_{n=1}^{\infty} \mathbf{Q}^n)_{k,j}$ -probability that the random walk started at pixel k and reached pixel j in any number of steps.

The last statement gives a reason why $\sum_{k=1}^n \mathbf{Q}^n$ is bounded no matter λ is 0 or not. To speak of the random walk interpretation, in one step, for example, the random walk starting at k has probability non-zero, $\mathbf{Q}_{k,j} = \frac{\mu(k,j)}{\Lambda(k) + \mu(k, \mathbf{N}_k)}$, to move to one of its $j \in \mathbf{N}_k$ neighbors and with probability $R_k = \frac{\Lambda(k)}{\Lambda(k) + \mu(k, \mathbf{N}_k)}$ to be vanished. A random walk path is a path which starts from the pixel k , travels around the image, ends up a pixel j and vanishes there with probability R_j . Thus, we conclude this section with the random walk view interpretation of the solution σ^* (Eq. B.9 or Eq. B.13) :

The solution $\sigma^(k)$ corresponds to the expected value of σ_0 at the first vanishing point of the random walk starting at the pixel k .*

The vanishing point can be either on the boundaries $\partial\Omega$ or not on the boundary. The probability of one path W_k starting at k , going through $k = k_0, \dots, k_n$ and vanishes at k_n can be written as

$$\mathbf{P}(W_k) = R_{k_n} \prod_{i=0}^{n-1} \mathbf{Q}_{k_i, k_{i+1}} = R_{k_n} \prod_{i=0}^{n-1} \frac{\mu(k_i, k_{i+1})}{\Lambda(k_i) + \mu(k_i, \mathbf{N}_{k_i})}. \quad (\text{B.16})$$

Usually, for the simple case where $e(k)$, $e(k, k')$ take only binary numbers, we can expect that once a particle reach $\partial\Omega$, it is likely to be vanished there if we choose

small enough ϵ to make the vanish probability $R_k = 1/(1 + \epsilon\mu(k, \mathbf{N}_k))$, $k \in \partial\Omega$ as close to 1 as we want. Thus, we can provide a simpler form for $\mathbf{P}(W_k)$ as

$$\mathbf{P}(W_k) = R_{k_n} \prod_{i=0}^{n-1} \mathbf{Q}_{k_i, k_{i+1}} \sim \prod_{i=0}^{n-1} \mathbf{Q}_{k_i, k_{i+1}} = (\lambda + 4)^{-n}, \quad (\text{B.17})$$

if the path touch boundary only once and vanishes there at k_n . Under this approximation, *with or without edge blocking is no longer an issue*. All the paths touch boundaries more than once can simply be ignored as truncation of higher order terms, compared to previous type walks. For other paths stop[†] at $k \in \mathbf{I} - \partial\Omega$, i.e., the paths never reach any part of the boundary, we have

$$\mathbf{P}(W_k) = R_{k_n} \prod_{i=0}^{n-1} \mathbf{Q}_{k_i, k_{i+1}} = \frac{\lambda}{\lambda + 4} (\lambda + 4)^{-n}. \quad (\text{B.18})$$

Keep remember that a smaller λ provides less decay.

Let us run all the paths backward, and consider only the path starting from $\partial\Omega$, we can write another version of interpretation for σ^* :

The solution $\sigma^(k)$ correspond to the summation of probability of multiple source random walks, where each random walk starts from a boundary pixel $i \in \partial\Omega$ and arrives (not “stops”) at k . The walk carries the weight $g_i = \frac{\Lambda(i)}{\Lambda(i) + \mu(i, \mathbf{N}_i)} \cdot \sigma_0(i)$ or*

[†]For random walks, we will use “stops” and “arrives” at certain location to specify whether we need to multiply the vanishing probability which only applied to the case of stopped walks.

for a simpler case as $g_i = \frac{1}{1+\epsilon\mu(i, N_i)}$ from a pixel $i \in \partial\Omega$ (i.e., corners tend to weight more).

When both sides Ω and $\mathbf{I} - \Omega$ are considered simultaneously, the weight carried by those random walks can be either positive or negative depends on the sign of σ_0 . The calculation is therefore done by summing all those “signed probability”. Here the summation cares only the walks starting from a boundary pixel $k \in \partial\Omega$. I.e., for the simple case where $e(k)$, $e(k, k')$ take only binary values, under this interpretation, we need to consider only those paths as in Eq. B.17 and paths as in Eq. B.18 can be ignored, if we choose a small enough ϵ .

Probability Interpretation

For the last two interpretations, we consider the solution as an expectation or a summation. It is interesting to view the result as a probability distribution. To see this, we need a linear transform from σ to p by

$$p(k) = \frac{1}{2}(1 + \sigma(k)),$$

as we have used in Eq. 2.12. Let us discuss the case with $\lambda = 0$ first, with $p_0(k)$ being either 1 or 0 for $k \in \partial\Omega$ (i.e., the inducers $\sigma_0(k) = 1 / -1$), assuming $e(k) = 1$, $\forall k \in \partial\Omega$. Again, we assume that the walk hitting boundaries more than once will vanish very likely and can be ignored. Therefore, the solution p^* , similar to σ^* , is an

expectation and can be written as

$$\begin{aligned}
p^*(k) &= 1 \cdot \sum_{i \in F(\partial\Omega), t} \mathbf{P}(W(t) = i, \text{ stops at } i \mid W(0) = k) \\
&\quad + 0 \cdot \sum_{i \in G(\partial\Omega), t} \mathbf{P}(W(t) = i, \text{ stops at } i \mid W(0) = k) \\
&\sim \sum_{i \in F(\partial\Omega), t} \mathbf{P}(W(t) = i, \text{ stops at } i \text{ and visits } \partial\Omega \text{ for the first time} \mid W(0) = k) \\
&= \mathbf{P}(W \text{ visits } F(\partial\Omega) \text{ before it visits } G(\partial\Omega) \mid W(0) = k) \tag{B.19}
\end{aligned}$$

as $F(\partial\Omega)$ and $G(\partial\Omega)$ indicating “figure” and “ground” side of the edges $\partial\Omega$. We can make $p^*(k)$ as close to the probability $\mathbf{P}(W_k \text{ visits } F(\partial\Omega) \text{ before } G(\partial\Omega))$ as we want, if we choose small enough ϵ ,

When the decay coefficient λ is not zero, we have an extra term

$$\frac{1}{2} \cdot \sum_{i \in \Omega - \partial\Omega, t} \mathbf{P}(W(t) = i, \text{ stops at } i \text{ without touching } \partial\Omega \mid W(0) = k).$$

It is for the case where a walk stops at interior pixels $i \in \Omega - \partial\Omega$. We can say that for half of the chance, the location will be figure and will be background for the other half of the chance, for it lacks of knowledge to commit. The same probability interpretation for p^* that we just described can stand still. We conclude it as

The solution $0 \leq p^(k) \leq 1$, derived by a linear transformation from $1 \leq \sigma^*(k) \leq 1$, is an approximation to the probability $\mathbf{P}(W_k \text{ visits } F(\partial\Omega) \text{ before it visits } G(\partial\Omega))$. The approximation can be as good as we want, if a small enough ϵ is provided.*

This interpretation suggests us to use the entropy criteria to measure the shape con-

vexity, as we described in Sec. 2.2.2 or Eq. 2.12 & Eq. 2.13.

B.1 Decay Process and Convexity

Let us examine why our model can catch convexity. We will check the example in Fig. B.1 where the regions Ω_a and $\Omega_b = \mathbf{I} - \Omega_a$ share a common boundary Γ . A simpler version of the energy functional in Eq. B.4 where $e(k)$ and $e(k, k')$ take only binary numbers will be discussed. For pixels p and q , we use \overline{pq} to indicate the Euclidean distance between them. The shortest legal path between p and q within the shape Ω_a (or Ω_b) is indicated by $d_a(p, q)$ (or $d_b(p, q)$).

In Fig. B.1, given a source pixel p (along the boundary) and given a pixel q at the convex side of the figure, there is a pixel q' equally distant to p (Euclidean distance) in the concave side (There are many pixels q' with such property). In the concave region the shortest legal walk between p and q' traverse a distance $d(p, q') > \overline{pq'}$ while in the convex region, the shortest legal walk between p and q always traverse a distance $d(p, q) = \overline{pq} = \overline{pq'}$.

By the result of random walk in last section, the contribution to the the final solution $\sigma^*(q)$ and $\sigma^*(q')$ of the source p to q and q' is directly dependent on $d(p, q)$ and $d(p, q')$ respectively. This is because with the decay term, only the pixels with shorter distance can be reached by the random walk. The longer is the path to reach

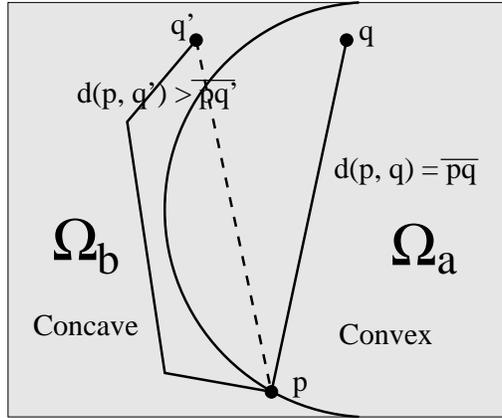


Figure B.1. Consider all the walks starting at the boundary pixel p . In the concave region, the shortest random walk from p to q' is given by $d(p, q') > \overline{pq'}$ while on the convex side, the shortest random walk to q is $d(p, q) = \overline{pq} = \overline{pq}$. The shorter are the random walks the more support is obtained by the inducers, otherwise the random walks tend to decay. Thus, concave regions get weaker response. ■

a pixel the more likely it will vanish (yielding the decay). More precisely, from the Markov theory the probability of a particle vanishing when going in the shortest path from p to q is exactly $\sum_{s=1}^{d(p,q)} (\mathbf{Q}_{p,a(s)}^s R_a)$, where the sum is over all pixels a in the shortest path from p to q , parameterized by $s = 1, \dots, d(p, q)$. Thus, the longer is the path, the more terms in the sum over a , and the smaller is the probability to reach q . Therefore, a point q in a convex side will be reached by more random walks than a point q' in the concave side. This implies that $\sigma^*(q)$ will be more defined to 1 or -1

(to figure or ground) than $\sigma^*(q')$ and the entropy will satisfy $\mathcal{S}^*(q) < \mathcal{S}^*(q')$. Thus, convexity is encouraged by the entropy criteria.

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