
Object-Based and Image-Based Image Representations

The representation of spatial objects and their environment is an important issue in applications of computer graphics, game programming, computer vision, image processing, robotics, pattern recognition, and computational geometry (e.g., [91, 1636, 1637, 1811]). The problem also arises in building databases to support them (e.g., [1638]). We assume that the objects are connected¹ although their environment need not be. The objects and their environment are usually decomposed into collections of more primitive elements (termed *cells*) each of which has a location in space, a size, and a shape. These elements can either be subobjects of varying shape (e.g., a table consists of a flat top in the form of a rectangle and four legs in the form of rods whose lengths dominate their cross-sectional areas), or they can have a uniform shape. The former yields an *object-based* decomposition; the latter yields an *image-based*, or *cell-based*, decomposition. Another way of characterizing these two decompositions is that the former decomposes the objects, while the latter decomposes the environment in which the objects lie. This distinction is used commonly in computer graphics to characterize algorithms as being either object-space or image-space, respectively [622].

Each of the decompositions has its advantages and disadvantages. They depend primarily on the nature of the queries that are posed to the database. The most general queries ask *where*, *what*, *who*, *why*, and *how*. The ones that are relevant to our application are *where* and *what* [1642]. They are stated more formally as follows:

1. Feature query: Given an object, determine its constituent cells (i.e., their locations in space).
2. Location query: Given a cell (i.e., a location in space), determine the identity of the object (or objects) of which it is a member, as well as the remaining constituent cells of the object (or objects).

Not surprisingly, the queries can be classified using the same terminology that we used in the characterization of the decomposition. In particular, we can either try to find the cells (i.e., their locations in space) occupied by an object or find the objects that overlap a cell (i.e., a location in space). If objects are associated with cells so that a cell contains the identity of the relevant object (or objects), then the feature query is analogous to retrieval by contents while the location query is analogous to retrieval by location. As we will see, it is important to note that there is a distinction between a location in space and the address where information about the location is stored, which, unfortunately, is often erroneously assumed to be the same.

¹ Intuitively, this means that a d -dimensional object cannot be decomposed into disjoint subobjects so that the subobjects are not adjacent in a $(d - 1)$ -dimensional sense.

The feature and location queries are the basis of two more general classes of queries. In particular, the feature query is a member of a broader class of queries described collectively as being *feature-based* (also *object-based*), and the location query is a member of a broader class of queries described collectively as being *location-based* (also *image-based* or *cell-based*). In computational geometry, the location query is known as the *point location query* or *problem* (e.g., [1511, 196] and Sections 2.1.3.2 and 2.1.3.3). In other domains, it is also commonly referred to as a *point query*, in which case we must be careful to distinguish it from its narrower and broader definitions. In particular, in database applications, a point query has a narrower definition—it is really an exact match query as it seeks to determine if the data point corresponding to the location is in the database (e.g., [1046] and the introduction to Chapter 1), and thus whenever possible we use the term *exact match query* in this context. On the other hand, in computer graphics applications, the point query has a broader definition and is often synonymous with the related “pick” operation (e.g., [622]), which is used to find an object that contains a given location l , and if there is none, then to find the nearest object to l . In order to avoid overloading the term *point query* in this chapter we use the term *nearest object query* when a satisfactory response includes the nearest object should no object exist that contains l .² Therefore, in this chapter, the term *point query* has the same meaning as in Chapter 3, where the response is restricted to the objects that contain l . The class of location-based queries includes the numerous variants of the *window query*, which retrieves the objects that cover an arbitrary region (often rectangular). All of these queries are used in several applications including geographic information systems (e.g., [72, 1645]) and spatial data mining (e.g., [1953]).

The most common representation of the objects and their environment is as a collection of cells of uniform size and shape (termed *pixels* and *voxels* in two and three dimensions, respectively) all of whose boundaries (with dimensionality one less than that of the cells) are of unit size. Since the cells are uniform, there exists a way of referring to their locations in space relative to a fixed reference point (e.g., the origin of the coordinate system). An example of a location of a cell in space is a set of coordinate values that enables us to find it in the d -dimensional space of the environment in which it lies. Once again, we reiterate that it should be clear that the concept of the *location* of a cell in space is quite different from that of the *address* of a cell, which is the physical location (e.g., in memory, on disk), if any, where some of the information associated with the cell is stored. This distinction between the location in space of a cell and the address of a cell is important, and we shall make use of it often.

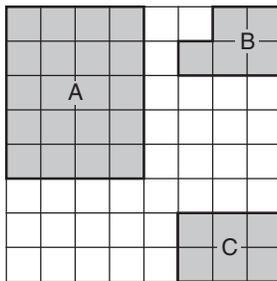


Figure 2.1
Example collection of three objects
and the cells that they occupy.

In most applications (including most of the ones that we consider here), the boundaries (i.e., edges and faces in two and three dimensions, respectively) of the cells are parallel to the coordinate axes. In our discussion, we assume that the cells making up a particular object are contiguous (i.e., adjacent) and that a different, unique value is associated with each distinct object, thereby enabling us to distinguish between the objects. Depending on the underlying representation, this value may be stored with the cells. For example, Figure 2.1 contains three two-dimensional objects, A, B, and C, and their corresponding cells. Note that, although it is not the case in this example, objects are allowed to overlap, which means that a cell may be associated with more than one object. Here we assume, without loss of generality, that the volume of the overlap must be an integer multiple of the volume of a cell (i.e., pixels, voxels, etc.).

The shape of an object o can be represented either by the interiors of the cells making up o , or by the subset of the boundaries of those cells making up o that are adjacent to the boundary of o . In particular, interior-based methods represent an object o by using the locations in space of the cells that make up o , while boundary-based methods represent o by using the locations in space of the cells that are adjacent to the boundary of o .

² A more general term is *nearest neighbor query*, which is discussed in a more general context and in much greater detail in Sections 4.1–4.3 of Chapter 4.

In general, interior-based representations make it very easy to calculate properties of an object such as its mass and, depending on the nature of the aggregation process, to determine the value associated with any point (i.e., location) in the space covered by a cell in the object. On the other hand, boundary-based representations make it easy to obtain the boundary of an object.

Regardless of the representation that is used, the generation of responses to the feature and location queries is facilitated by building an index (i.e., the result of a sort) either on the objects or on their locations in space, and implementing it using an access structure that correlates the objects with the locations. Ideally, we want to be able to answer both types of queries with one representation. This is somewhat tricky, but, as we will show, it is doable.

The rest of this chapter is organized as follows. Section 2.1 discusses interior-based representations. These are the most prevalent, and thus this discussion forms the main part of the chapter. Section 2.2 discusses boundary-based representations. The discussion in these sections shows how some of the representations can be made more compact by aggregating similar elements. These elements are usually identically valued contiguous cells (possibly adjacent to identically oriented boundary elements) or even objects that, ideally, are in proximity. The representations can be made even more compact by recording only the differences between the elements. The use of difference-based compaction methods is the subject of Section 2.3. Many of the representations that we have described are based on the principle of recursive decomposition of the underlying space (as well as the aggregation of the results of the decomposition). Therefore, we conclude in Section 2.4 with a brief historical overview of this principle and some of its early uses.

As stated earlier, most of our discussion assumes that the objects can be decomposed into cells whose boundaries are parallel to the coordinate axes. Nevertheless, we discuss the representation of objects with boundaries that are hyperplanes (i.e., straight lines and planar faces for two- and three-dimensional objects, respectively), as well as objects with arbitrary boundaries (e.g., edges of arbitrary slope, nonplanar faces). In this case, the representation of the underlying environment is usually, but not always, based on a decomposition into cells whose boundaries are parallel to the coordinate axes.

2.1 Interior-Based Representations

In this section, we focus on interior-based representations. Section 2.1.1 examines representations based on collections of unit-size cells. An alternative class of interior-based representations of the objects and their environment removes the stipulation that the cells that make up the object collection be a unit size and permits their sizes to vary. The varying-sized cells are termed *blocks* and are the subject of Section 2.1.2. The representations described in Sections 2.1.1 and 2.1.2 assume that each unit-size cell or block is contained entirely in one or more objects. A cell or block cannot be partially contained in two objects. This means that either each cell in a block belongs to the same object or objects, or that all of the cells in the block do not belong to any of the objects. Section 2.1.3 permits both the blocks and the objects to be arbitrary polyhedra while not allowing them to intersect, although permitting them to have common boundaries. Section 2.1.4 permits a cell or block to be a part of more than one object and does not require the cell or block to be contained in its entirety in these objects. In other words, a cell or a block may overlap several objects without being completely contained in them. This also has the effect of permitting the representation of collections of objects whose boundaries do not coincide with the boundaries of the underlying blocks and also permitting them to intersect (i.e., overlap). Section 2.1.5 examines the use of hierarchies of space and objects that enable efficient responses to both the feature and location queries.

Exercises

1. Given a $[6^3]$ tiling such that each side of an atomic tile has a unit length, compute the three adjacency distances from the centroid of an atomic tile.
2. Repeat Exercise 1 for $[3^6]$ and $[4^4]$, again assuming that each side of an atomic tile has a unit length.
3. Suppose that you are given an image in the form of a binary array of pixels. The result is a square grid. How can you view this grid as a hexagonal grid?

2.1.1.2 Ordering Space

There are many ways of laying out the addresses corresponding to the locations in space of the cells, each having its own mapping function. Some of the most important ones for a two-dimensional space are illustrated in Figure 2.5 for an 8×8 portion of the space and are described briefly below. To repeat, in essence, what we are doing is providing a mapping from the d -dimensional space containing the locations of the cells to the one-dimensional space of the range of index values (i.e., integers) that are used to access a table whose entries contain the addresses where information about the contents of the cells is stored. The result is an ordering of the space, and the curves shown in Figure 2.5 are termed *space-filling curves* (e.g., [1603]).

Choosing among the space-filling curves illustrated in Figure 2.5 is not easy because each one has its advantages and disadvantages. Below, we review a few of their desirable properties, show how some of the two-dimensional orderings satisfy them, and provide some exercises to help you to explore them further:

- The curve should pass through each location in space once and only once.
- The mapping from the higher-dimensional space to the integers should be relatively simple and likewise for the inverse mapping. This is the case for all but the Peano-Hilbert order (Figure 2.5(d)). For the Morton order (Figure 2.5(c)), the mapping is obtained by interleaving the binary representations of the coordinate values of the location of the cell. The number associated with each cell is known as its *Morton number*. The Gray order (Figure 2.5(g)) is obtained by applying a Gray code (see footnote 35 in Section 1.6 of Chapter 1) to the result of bit interleaving, and the double Gray order (Figure 2.5(h)) is obtained by applying a Gray code to the result of bit interleaving the Gray code of the binary representation of the coordinate values. The U order (Figure 2.5(i)) is obtained in a similar manner to the Morton order, except for an intermediate application of $d - 1$ “exclusive or” (\oplus) operations on the binary representation of selected combinations of the coordinate values prior to the application of bit interleaving [1192, 1698] (see Exercise 7). Thus, the difference in cost between the Morton order and the U order in d dimensions is just the performance of additional $d - 1$ “exclusive or” operations. This is in contrast to the Peano-Hilbert order, where the mapping and inverse mapping processes are considerably more complex (see Exercises 5 and 6).
- The ordering should be stable. This means that the relative ordering of the individual locations is preserved when the resolution is doubled (e.g., when the size of the two-dimensional space in which the cells are embedded grows from 8×8 to 16×16) or halved, assuming that the origin stays the same. The Morton, U, Gray, and double Gray orders are stable, while the row (Figure 2.5(a)), row-prime (Figure 2.5(b)), Cantor-diagonal (Figure 2.5(e)), and spiral (Figure 2.5(f)) orders are not stable (but see Exercises 1 and 2). The Peano-Hilbert order is also not stable, as can be seen by its definition (but see Exercise 3). In particular, in two dimensions, the Peano-Hilbert order of resolution $i + 1$ (i.e., a $2^i \times 2^i$ image) is constructed by taking the Peano-Hilbert curve of resolution i and rotating the NW, NE, SE, and SW quadrants by 90°

Section 2.1

INTERIOR-BASED REPRESENTATIONS

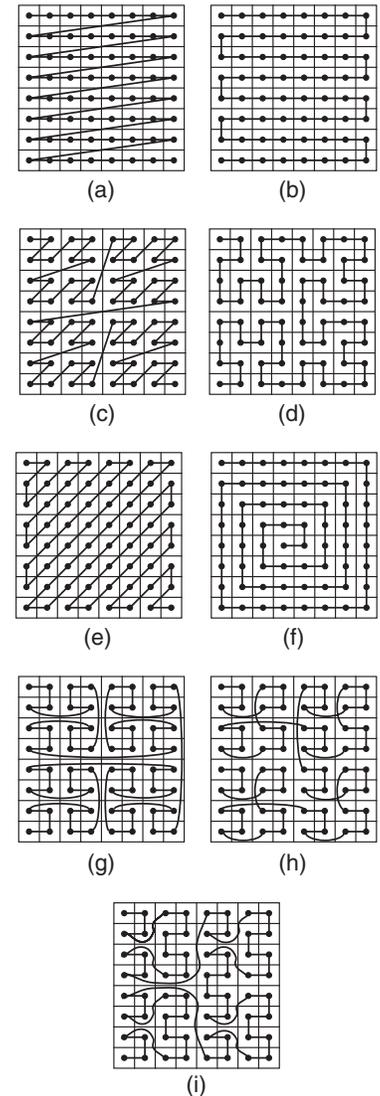


Figure 2.5

The result of applying several different space-ordering methods to an 8×8 collection of cells whose first element is in the upper-left corner: (a) row order, (b) row-prime order, (c) Morton order, (d) Peano-Hilbert order, (e) Cantor-diagonal order, (f) spiral order, (g) Gray order, (h) double Gray order, and (i) U order.

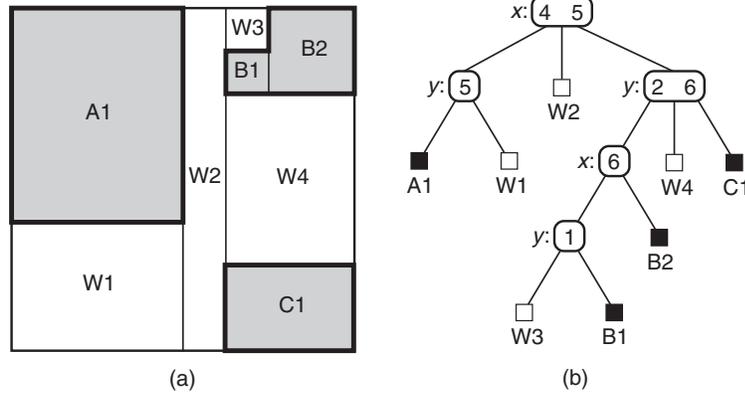
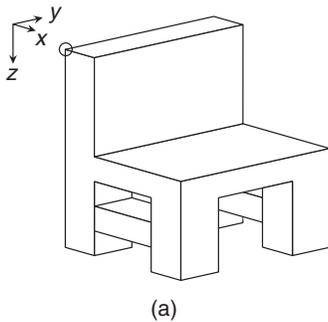
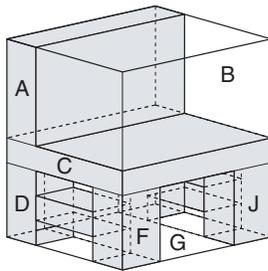


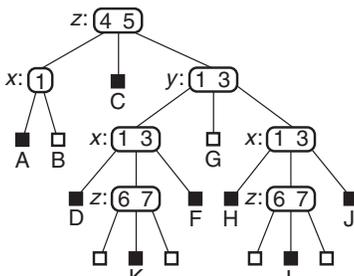
Figure 2.28 (a) Block decomposition for the puzzletree corresponding to the collection of objects and cells in Figure 2.1 and (b) the corresponding tree access structure. The nonleaf nodes are labeled with the partition axes and the partition values.



(a)



(b)



(c)

Figure 2.29 (a) A three-dimensional chair, (b) the block decomposition for its puzzletree, and (c) its corresponding tree access structure. The nonleaf nodes are labeled with the partition axes and the partition values, assuming an origin at the upper-left corner of the chair.

collection of photos). In some sense, these alternative region-representation techniques resemble the imposition of gridlike partitions at the various levels of the directory.

The X-Y tree was developed as a page representation for regions in optically scanned documents, and its presentation focused on issues related to its use in this application [1334]. However, Nagy and Seth do point out in passing that the X-Y tree could be extended to three and more dimensions, which could be useful for dealing with multipage documents as well as, possibly, with other applications. In contrast, the puzzletree was developed as an object representation with a detailed discussion of the ramifications of its use in the representation of both two- and three-dimensional objects. As our interest in this section is more in object representations, the discussion in the rest of this section follows the presentation of Dengel and hence is usually in terms of the puzzletree, especially in the context of objects, although almost all that we say is equally applicable to the X-Y tree and also to the original formulation of the treemap.

Figure 2.28(a) is the block decomposition for the puzzletree for Figure 2.1, and Figure 2.28(b) is the corresponding tree access structure. Notice that the puzzletree was created by compressing the successive initial partitions on $x = 4$ and $x = 5$ at depths 0 and 1, respectively, and likewise for the successive partitions on $y = 6$ and $y = 2$ at depths 2 and 3, respectively, in Figure 2.27. Nonleaf nodes are labeled with the name of the axis of the partition and the location of the partition line. We have labeled the blocks corresponding to object O as O_i and the blocks that are not in any of the objects as W_i , using the suffix i to distinguish between them in both cases.

The fact that the puzzletree compresses all successive partitions on the same dimension means that for $d = 2$ the partitions cycle through the different dimensions so that a different dimension is tested at each level of the decomposition. Note that for $d \geq 3$ a requirement that the partitions cycle through the various dimensions before being permitted to repeat may prevent us from being able to use certain more perceptually appealing block combinations. For example, consider the three-dimensional four-legged chair in Figure 2.29(a) whose corresponding puzzletree has the block decomposition given in Figure 2.29(b) and the tree access structure given in Figure 2.29(c). Here, we see the impossibility of cycling through all of the dimensions in an alternating manner.

The puzzletree is motivated by a desire to overcome the rigidity in the shape, size, and position of the blocks that result from the bintree (and to an equivalent extent, the region quadtree) partitioning process (because of its regular decomposition). In particular, in many cases, the decomposition rules ignore the homogeneity present in certain regions on account of the need to place the partition lines in particular positions as well as a possible limit on the number of permissible partitions along each dimension at each

This decomposition process is applied recursively to $P_{k,L}$ and $P_{k,R}$ and terminates upon encountering empty sets.

Two items are worthy of further note. First, a polygonal face, say p_i , may be contained in both the left and right subtrees of a node in the BSP tree. Such a situation arises when p_i 's plane intersects the plane of one of its ancestor nodes in the BSP tree. Second, the plane of a polygonal face is assumed to extend beyond the boundary of the polygon (i.e., it spans the entire underlying block). For example, the plane of the root's polygonal face partitions the entire scene. Moreover, for each subtree rooted at T , the polygonal face associated with T , say p_T , is extended so that p_T partitions the entire space of T .

One of the drawbacks of the BSP tree is that its shape is heavily dependent on the order in which the polygonal faces are processed and on the polygonal faces that are selected to serve as the partitioning planes. In the worst case, the BSP tree looks like a chain. Furthermore, the decomposition may be such that a polygonal face is contained in many subtrees. Such a situation arises in complex nonconvex scenes and will lead to large BSP trees. Nevertheless, it can be somewhat alleviated by choosing the root polygonal face more carefully at each stage of the BSP tree construction process.

For example, one heuristic is to choose the polygonal face, say M , in the set that splits the minimum number of the remaining polygonal faces in the set (see Exercise 5 for another approach). Fuchs, Abram, and Grant [657] have found that, in practice, there is no need to try out all possibilities to determine M . Instead, they follow a suggestion of Kedem [657] and just select a small subset of the polygonal faces at random, say S , to serve as candidates and then choose the polygonal face in S that splits the minimum number of the remaining polygonal faces in the set. In two dimensions, for n line segments, this method yields a BSP tree of expected size $O(n \log n)$ and can be constructed in $O(n^2)$ time [1475]. The execution time can be reduced to $O(n \log n)$ with the same size bound by using an approach that does not yield an autopartition [1475]. In three dimensions, for n polygonal faces that are triangles, the best-known method yields an autopartition BSP tree with expected size $O(n \log^2 n)$ and can be constructed in $O(n \log^3 n)$ time [18], although there exist sets of data for which the size is $\Omega(n^2)$ [1475]. Naylor observed that, in practice, three-dimensional BSP trees were usually of size $O(n \log n)$ [1353, 1354]. An analogous approach of avoiding a check of all possibilities was used by Samet [1618] in selecting a node to replace the deleted root of a point quadtree (see Section 1.4.1.2 of Chapter 1).

To illustrate the differences in the storage requirements of the BSP tree, we use a collection of line segments instead of polygonal faces because it is easier to visualize what is happening. Each line segment has a direction. We say that its "positive" side is the one to the right of the direction of the line segment. When the line segment is treated as a separator between two halfspaces, we say that the "positive" side is the one whose equation has the form $a \cdot x + b \cdot y + c \geq 0$, and the "negative" side is the one whose equation has the form $a \cdot x + b \cdot y + c < 0$.

For example, consider the four line segments labeled A, B, C, and D in Figure 2.37. Suppose that the BSP tree is constructed by letting B be the root. The positive subset consists of D, while the negative subset consists of A, C, and D. D appears in both subsets because D crosses B when it is extended and thereby serves as a separator between two halfspaces. Now, let us build the BSP tree for the negative subset. Letting C be the root, we find that A is in the negative subset, and D is in the positive subset. The resulting BSP tree is shown in Figure 2.38(a) and partitions the plane into six regions as shown in Figure 2.38(b).

If, in the first step, we would have chosen C instead of B to be the root of the BSP tree, then the negative subset would consist of A, and the positive subset would consist of B and D. Now, let us build the BSP tree for the positive subset. Letting D be the root, we find that the positive subset is empty, and B is in the negative subset. The resulting BSP tree is

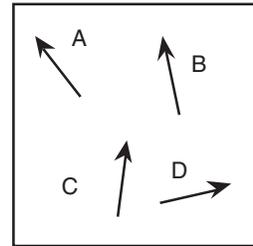


Figure 2.37
Example set of four line segments.

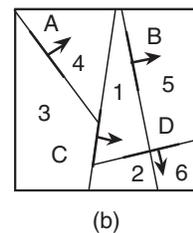
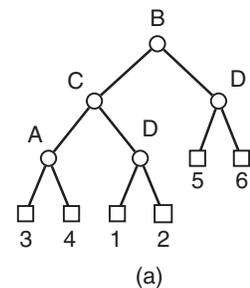


Figure 2.38
(a) The BSP tree corresponding to Figure 2.37 when B is the root and
(b) the partition induced by it.

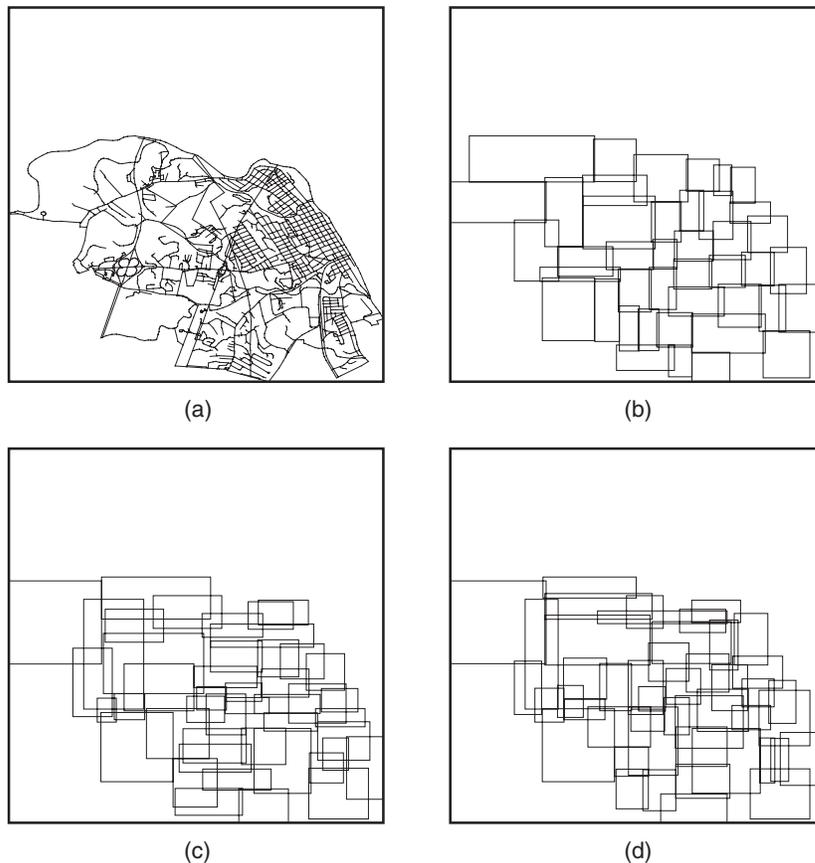


Figure 2.79
(a) A sample collection of 1,700 line-shaped objects and the space decomposition resulting from the use of (b) an R*-tree, as well as R-trees that make use of (c) linear cost and (d) quadratic cost node-splitting algorithms. All trees are order ($m = 20$, $M = 50$).

In particular, the original R-tree node-splitting algorithms [791] determine seeds, while the R*-tree algorithm determines a split axis and an axis split value. The bounding boxes of the objects are redistributed about these seeds and axis, respectively. At this point, it is important to reemphasize that the motivation for these redistribution strategies is to avoid the exhaustive search solution that looks at all possible partitions.

The R*-tree redistribution method first sorts the boundaries of the bounding boxes along each of the axes and then uses this information to find the split axis a (with respect to the minimum average perimeter of the bounding boxes of the resulting nodes) and split position (with respect to the minimal overlap once the split axis was chosen). This is a heuristic that attempts to approximate the solution to the d -dimensional problem (i.e., optimal partitioning with minimal coverage or overlap) with an approximation of the optimal one-dimensional solution along one of the axes (see Exercise 20). Intuitively, the validity of this approximation would appear to decrease as d (i.e., the dimensionality of the underlying space) increases since more and more splits are eliminated from consideration. However, the opposite conclusion might be true as it could be argued that although the number of eliminated splits grows exponentially with d , the majority of the eliminated splits are bad anyway. This is a problem for further study.

The remaining changes involving forced reinsertion and intelligent object insertion could have also been used in the R-tree construction algorithms. In particular, although the original R-tree definition [791] opts for minimizing coverage in determining the subtree into which an object is to be inserted, it does leave open whether minimizing

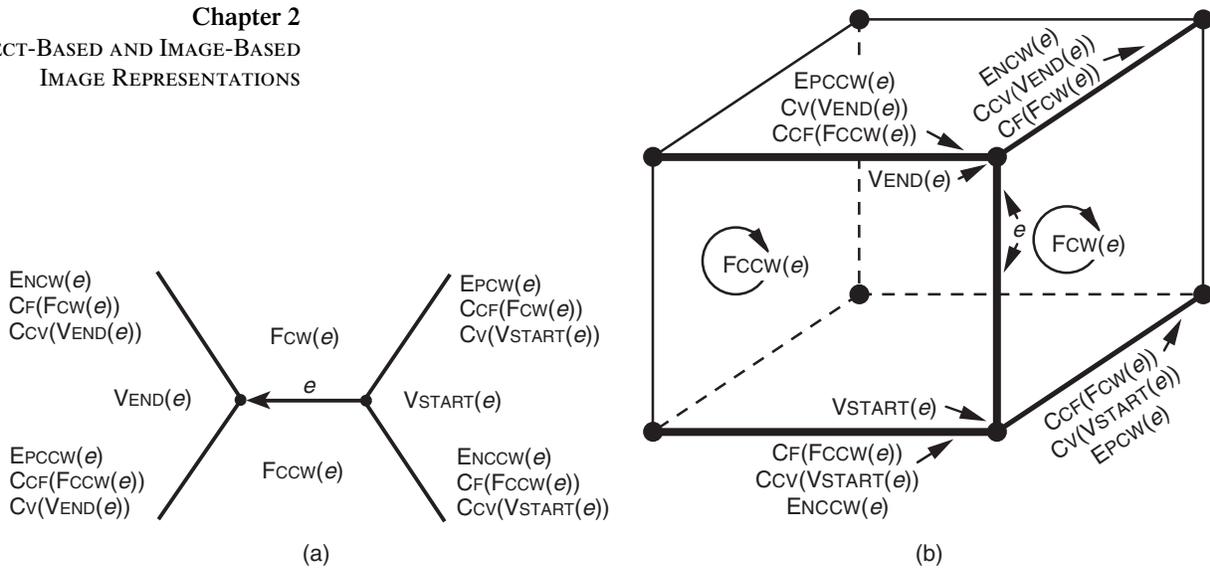


Figure 2.89
(a) Example of an edge e and its four wings and (b) the physical interpretation of e and its wings when e is an edge of a parallelepiped represented by the winged-edge data structure.

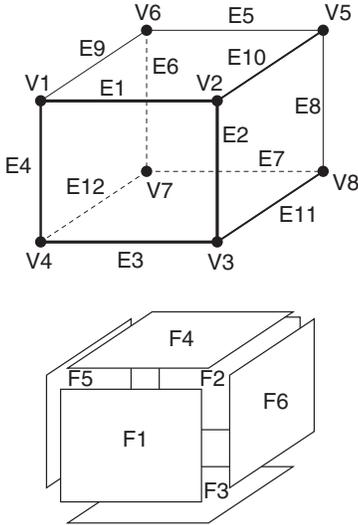


Figure 2.90
Sample parallelepiped with vertices (V1–V8), edges (E1–E12), and faces (F1–F6).

- Edge-edge relation: the preceding edges ($CCV(VSTART(e))$ and $CCV(VEND(e))$) and the next edges ($CV(VSTART(e))$ and $CV(VEND(e))$) incident at the two vertices, thereby incorporating the vertex-edge relation, as well as the face-edge relation when using an alternative interpretation of the wings of the edge.

As an example of the winged-edge data structure, consider the parallelepiped in Figure 2.89(b), whose individual vertices, edges, and faces are labeled and oriented according to Figure 2.90. One possible implementation of a winged-edge representation for it is given by tables VERTEXEDGETABLE and FACEEDGETABLE in Figures 2.91 and 2.92, which correspond to the partial vertex-edge and face-edge relations, respectively, and the collection of edge records that make up the edge-edge relation given by Figure 2.93. Observe that VERTEXEDGETABLE and FACEEDGETABLE are really indexes (i.e., access structures) that enable an efficient response to queries on the basis of the value of a given vertex v or face f , such as finding all of the edges incident at v or the edges that f comprises, respectively, in both the clockwise and counterclockwise orders. This information is accessed by the field EDGE as the tables may contain additional information, such as the actual x , y , and z coordinate values of the vertex in the case of VERTEXEDGETABLE.

VERTEX v	X	Y	Z	EDGE
V1	X1	Y1	Z1	E1
V2	X2	Y2	Z2	E2
V3	X3	Y3	Z3	E3
V4	X4	Y4	Z4	E4
V5	X5	Y5	Z5	E5
V6	X6	Y6	Z6	E6
V7	X7	Y7	Z7	E7
V8	X8	Y8	Z8	E8

FACE f	EDGE
F1	E1
F2	E5
F3	E11
F4	E9
F5	E4
F6	E8

Figure 2.91
VERTEXEDGETABLE[v].

Figure 2.92
FACEEDGETABLE[f].

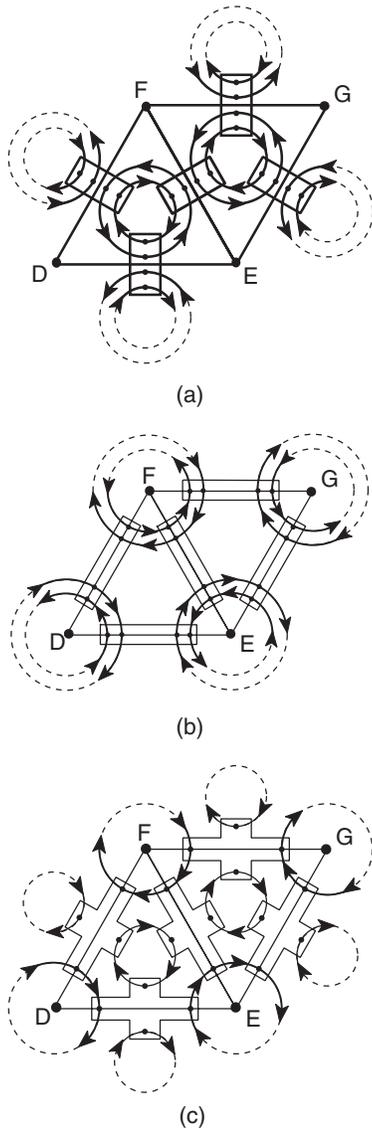


Figure 2.94
The physical interpretation of the (a) winged-edge-face, (b) winged-edge-vertex, and (c) quad-edge data structures for a pair of adjacent faces of a simple object. Assume an implementation that links the next and preceding edges in clockwise order for faces in (a), for vertices in (b), and next edges in clockwise order for both faces and vertices in (c).

$f = FCCW(e)$, and the next edge is $CCFCCW(e)$. This process terminates when we encounter the initial value of e again. For example, extracting the edges of face F1 in Figure 2.90 in clockwise order yields E1, E2, E3, and E4. The execution time of EXTRACTEDGESOFFACE is proportional to the number of edges in f as each edge is obtained in $O(1)$ time. This is a direct consequence of the use of FACEEDGETABLE, without which we would have had to find the first edge by a brute-force (i.e., a sequential) search of the edge-edge relation. Similarly, by making use of VERTEXEDGETABLE to obtain an edge incident at vertex v , we can extract the edges incident at v in time proportional to the total number of edges that are incident at v as each edge can be obtained in $O(1)$ time (see Exercise 2).

```

1  procedure EXTRACTEDGESOFFACE( $f, CWFlag$ )
2  /* Extract the edges making up face  $f$  in clockwise (counterclockwise) order
   if flag  $CWFlag$  is true (false). */
3  value face  $f$ 
4  value Boolean  $CWFlag$ 
5  pointer edge  $e, FirstEdge$ 
6   $e \leftarrow FirstEdge \leftarrow EDGE(FACEEDGETABLE[f])$ 
7  do
8    output  $e$ 
9    if  $CWFlag$  then
10      $e \leftarrow$  if  $FCW(e) = f$  then  $CFFCW(e)$ 
11     else  $CFFCCW(e)$ 
12     endif
13   else  $e \leftarrow$  if  $FCW(e) = f$  then  $CCFFCW(e)$ 
14     else  $CCFFCCW(e)$ 
15     endif
16   endif
17   until  $e = FirstEdge$ 
18 enddo

```

The above interpretations are not the only ones that are possible. Another interpretation, among many others, which finds much use, interprets the four wings in terms of the next edges at each of the faces that are adjacent to e and the next edges incident at each of the two vertices that make up e . In this case, we have combined the interpretations of the wings $CF(FCW(e))$ and $CF(FCCW(e))$ as used in the winged-edge-face data structure with the interpretations of the wings $CV(VSTART(e))$ and $CV(VEND(e))$ as used in the winged-edge-vertex data structure. The result is known as the *quad-edge data structure* [767]. It keeps track of both the edges that make up the faces in the clockwise direction and the edges that are incident at the vertices in the clockwise direction.

The quad-edge data structure is of particular interest because it automatically encodes the dual graph, which is formed by assigning a vertex to each face in the original graph and an arc to each edge between two faces of the original graph. In other words, we just need to interpret the cycles through the edges around the vertices in the original graph as faces in the dual graph and the cycles through the edges that the faces comprise in the original graph as vertices in the dual graph. In addition, the exterior face, if one exists, in the original graph is also assigned a vertex in the dual graph, which is connected to every face in the original graph that has a boundary edge. This makes the quad-edge data structure particularly attractive in applications where finding and working with the dual mesh is necessary or useful. For example, this is the case when the mesh corresponds to a Voronoi diagram whose dual is the Delaunay triangulation (DT). We discuss this further in Section 2.2.1.4. Another advantage of the quad-edge data structure over the winged-edge-face and winged-edge-vertex data structures is that, in its most general formulation, the quad-edge data structure permits making a distinction between the two sides of a surface, thereby allowing the same vertex to serve as the two endpoints of an edge, as well as allowing dangling edges, and so on.

From the above, we see that the winged-edge-face, winged-edge-vertex, and quad-edge data structures are identical in terms of the information that they store for each

Exercise

1. Show how the half-edge-vertex data structure (i.e., the vertex variant of the half-edge data structure) is closely related to the counterclockwise variant of the corner lath data structure. In this case, recall from Exercise 11 of Section 2.2.1.2 that, given half-edge L , the vertex variant of the half-edge data structure is defined in terms of $CV(L)$ and $CCV(L)$.

2.2.1.4 Voronoi Diagrams, Delaunay Graphs, and Delaunay Triangulations

In Sections 2.2.1.2 and 2.2.1.3, we pointed out that an advantage of the quad-edge interpretation of the winged-edge data structure over the other interpretations (e.g., the winged-edge-face and winged-edge-vertex) is that the quad-edge data structure is a self-dual. As an example of where this duality is useful, consider the Voronoi diagram [1941] (also known as a Thiessen polygon [1870] and a Dirichlet domain [497]) and its dual, the Delaunay graph (e.g., [1424]), as shown in Figure 2.110. Briefly, given a set of points (termed *sites*), a Voronoi diagram is a partition of the plane into regions so that each point in a region is closer to its region's site than to any other site. These regions are termed *Voronoi regions* or *Voronoi cells*. For example, Figure 2.110(a) is the Voronoi diagram for the given set of points shown with solid circles, while the vertices of the diagram are shown with hollow circles.

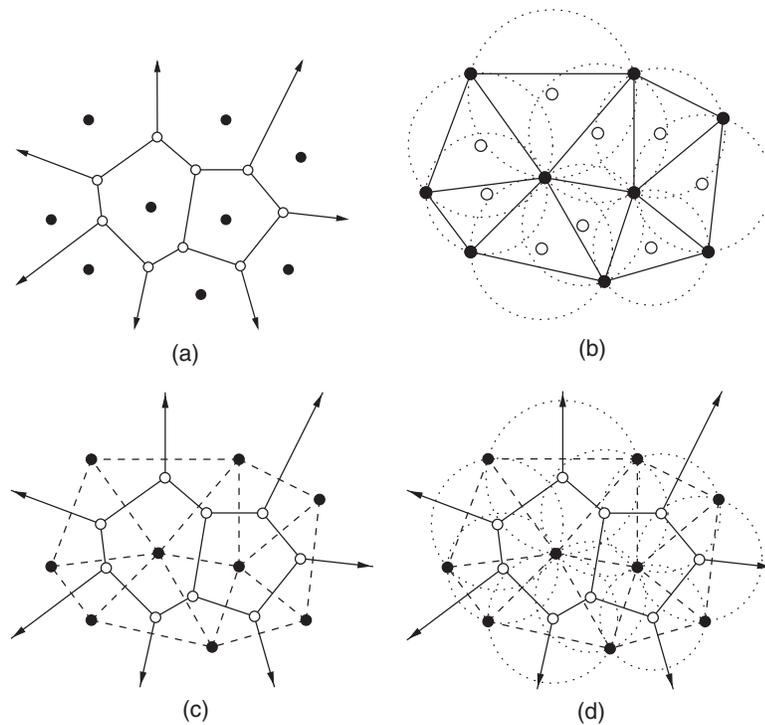


Figure 2.110
 (a) Example Voronoi diagram, (b) its dual Delaunay triangulation and the corresponding circumcircles (shown with broken lines), (c) the result of superposing the Delaunay triangulation (shown with broken lines) on the Voronoi diagram, (d) the result of superposing the Delaunay triangulation (shown with broken lines) and the corresponding circumcircles (shown with broken lines) on the Voronoi diagram. The hollow circles denote the centers of the circumcircles and the vertices of the Voronoi diagram. The arrowheads on some of the edges of the Voronoi diagram correspond to “infinite” edges that meet at some infinite vertex.

There are a number of ways of measuring the approximation error (e.g., relative, absolute, average), although no specific one is prescribed [655]. If the approximation error function is a relative one, then the sampling is more frequent around the object boundary, where the distance field is close to zero and hence the sampled distance field should follow the distance field more closely. On the other hand, if the approximation error function is an absolute one, then the approximation error threshold determines how far the sampled distance field can deviate from the actual distance field based on an absolute value. In this case, the sampling will be more uniform across the entire space.

If the application is only concerned with the distance field around the boundary of the object, then we can increase the approximation error threshold as the distance of the sampled points from the boundary of the object increases. The effect of such an approach is similar to a relative approximation error function. However, the problem is that such a solution tends to oversample at the boundary of the object. Hence, we can combine the two and use an absolute approximation error e_a if the sampled point p is sufficiently close to the boundary of the object, say within d of the boundary of the object, and use a relative approximation error e_r otherwise. Such an approximation error function e_p is given by

$$e_p = \min(e_a, e_r \times |d_p|)$$

which we characterize as an *adaptive approximation error*.

Being able to vary the approximation error at different locations is one of the main reasons for the attractiveness of the adaptively sampled distance field. This can be seen by observing that the distance field undergoes a change of sign at the boundary of the object as this is its zero-crossing. These changes are important and may require a relatively low approximation error threshold for their proper handling. On the other hand, the distance field undergoes rapid changes (i.e., points of discontinuity of the first derivative) at points on the skeleton (recall the definition in Section 2.1.2.2) of the object as the identity of the nearest object changes at these locations. In other words, the normal of the distance field surface is discontinuous at the object's skeleton, thereby requiring much decomposition in the skeleton's neighborhood when a low approximation error threshold is used, regardless of whether it is relative or absolute. On the other hand, there is little need for decomposition at the skeleton, and thus adaptively varying the approximation error threshold so that it is low around the boundary of the object and high at the neighborhood of the skeleton is very useful. In essence, an adaptive sampling of the distance field based on the values of the distance is being done, and hence the rationale for the name adaptively sampled distance field.

Regardless of how the approximation error threshold is measured, even highly curved edges and faces can be efficiently represented using adaptively sampled distance fields. In particular, since trilinear (bilinear) interpolation can represent curved surfaces (edges) well, cells with smoothly curved faces (edges) do not require many levels in the adaptively sampled distance field hierarchy, and thus much decomposition is only concentrated at corners, cusps, and sharp edges.

For example, Figure 2.122(a) is the adaptively sampled distance field for the two-dimensional object in Figure 2.121(a) using a relative approximation error threshold of 0.9. Notice the similarity in the decomposition to the edge quadtree in Figure 2.121(b, c). This is not surprising as it is a direct result of our observation of the duality of these representations.

It is interesting to observe that the distance field is somewhat of an indirect representation of the shape of an object as it describes the object by associating information with the parts of the space that are not occupied by it rather than the parts of the space that are occupied by it. In addition, note that, although the concept of distance is also used in the definition of a medial axis transformation (MAT) and a quadtree medial axis

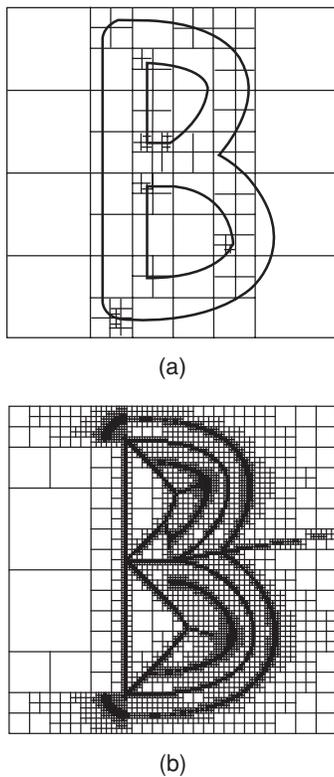


Figure 2.122
The adaptively sampled distance field for the two-dimensional object in Figure 2.121(a) using a relative approximation error threshold of (a) 0.9 and (b) 0.01.

trast to raster data) and has led them to be called *vector quadtrees* (and *vector octrees* in three dimensions) [1666, 1667].

The PM_2 quadtree is particularly useful for storing polygonal maps that correspond to triangulations or, more generally, to polygonal maps where all vertices have degree 3 (e.g., nondegenerate Voronoi diagrams where no four sites are cocircular) as the space requirements are often considerably smaller than those of the PM_1 quadtree, while not being much larger than the PM_3 quadtree. This is especially true for the Delaunay triangulation, which satisfies the minimum angle property and thus maximizes the minimum interior angle, thereby implying that the constituent triangles are as close as possible to being equiangular (recall properties 2 and 6 of the Delaunay triangulation in Section 2.2.1.4). This can be seen in Figure 2.124(a–c), which shows the block decompositions induced by the PM_1 , PM_2 , and PM_3 quadtrees, respectively, for the Delaunay triangulation given in Figure 2.110(b) in Section 2.2.1.4, where the corresponding Voronoi diagram is shown using broken lines. Notice the difference in the maximum depth of the decomposition between the PM_1 and PM_2 quadtrees, while there is no difference in this example between the PM_2 and PM_3 quadtrees.

In fact, in such a case, instead of storing with each leaf block b the set of edges that pass through b , we use an alternative implementation where each leaf block b stores the vertex, if any, that is common to the edges that pass through b [453]. If only one edge passes through b , then one of its vertices is arbitrarily chosen to be stored. In any case, regardless of how many edges pass through b , the vertices are usually part of an auxiliary data structure (e.g., a triangle table or some other variant of the winged-edge data structure) that keeps track of the topological properties of the polygonal map. The auxiliary structure serves to facilitate a number of operations that involve visiting adjacent triangles that are necessary for updating the polygonal map in a consistent manner so that it remains a triangulation of a particular type or even operations such as point location. For another related application of the PM_2 quadtree, see the discussion of the approximate Voronoi diagram (AVD) [799] in Section 4.4.5 of Chapter 4. In particular, see the (3,0)-AVD [97, 98], which, for two-dimensional polygonal maps that correspond to Voronoi diagrams, yields an identical partition of the underlying space into blocks as does the PM_2 quadtree.

Exercises

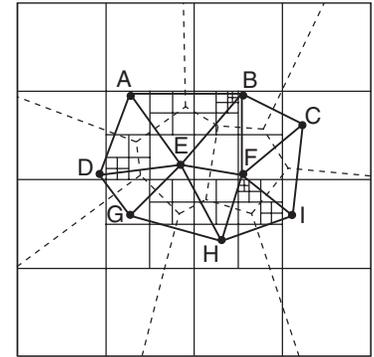
Exercises 1–6 assume the following implementation for the PM quadtree. The basic entities are vertices and edges. Each vertex is represented as a record of type *point*, which has two fields called XCOORD and YCOORD that correspond to the x and y coordinate values, respectively, of the point. They can be real or integer numbers, depending on implementation considerations such as floating-point precision.

An edge is implemented as a record of type *line* with four fields, P1, P2, LEFT, and RIGHT. P1 and P2 contain pointers to the records containing the edge's vertices. LEFT and RIGHT are pointers to structures that identify the regions that are on the two sides of the edge. We shall use the convention that LEFT and RIGHT are with respect to a view of the edge that treats the vertex closest to the origin as the start of the edge.

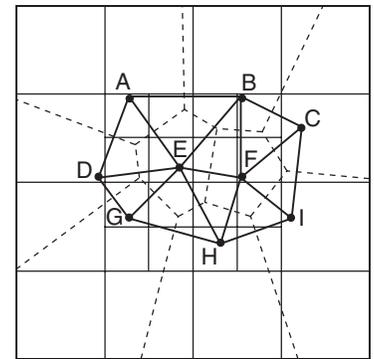
Each node in a PM quadtree is a collection of q -edges, which is represented as a record of type *node* containing seven fields. The first four fields contain pointers to the node's four children corresponding to the directions (i.e., quadrants) NW, NE, SW, and SE. If P is a pointer to a node, and I is a quadrant, then these fields are referenced as $CHILD(P, I)$. The fifth field, NODETYPE, indicates whether the node is a leaf node (LEAF) or a nonleaf node (GRAY). The sixth field, SQUARE, is a pointer to a record of type *square*, which indicates the size of the block corresponding to the node. It is defined for both leaf and nonleaf nodes. It has two fields CENTER and LEN. CENTER points to a record of type *point* that contains the x and y coordinate values of the center of the square. LEN contains the length of a side of the square that is the block corresponding to the node in the PM_1 quadtree.

DICTIONARY is the last field. It is a pointer to a data structure that represents the set of q -edges that are associated with the node. Initially, the universe is empty and consists of no

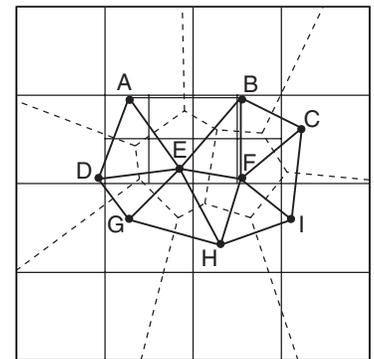
Section 2.2 BOUNDARY-BASED REPRESENTATIONS



(a)



(b)



(c)

Figure 2.124
The block decompositions induced by the (a) PM_1 , (b) PM_2 , and (c) PM_3 quadtrees for the Delaunay triangulation given in Figure 2.110(b) of the set of sites $\{A, B, C, D, E, F, G, H, I\}$ whose Voronoi diagram is given in Figure 2.110(a) and shown here using broken lines.

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REPRESENTATIONS

The newly added points that do not lie on existing edges represent new samples. Thus, we must also obtain an elevation value for them. There are two choices for some of them since they fall at the midpoints of edges between the midpoints of both a horizontal and a vertical edge. We choose to take the average of the two interpolated values. For example, v is obtained by taking the average of the values obtained by applying linear interpolation to the elevation values at L and t and the elevation values at u and r . The same technique is applied to obtain the elevation value at z once we have obtained the elevation values at x , y , and w .

At this point, let us briefly compare the two different quaternary hierarchical decompositions. As we saw, both methods suffer from the alignment problem when we try to fit a surface through their vertices. However, as we have demonstrated, this problem can be overcome through the addition of the appropriate triangles and rectangles. For the triangular decomposition, we obtained a planar surface because we were able to use a planar triangulation. For the rectangular decomposition, we obtained a nonplanar surface because it is impossible to fit a plane through the four vertices of each rectangle.

It could be argued that a nonplanar surface is a better approximation of a nonplanar surface than a planar one, thereby making the rectangular decomposition more attractive than the triangular decomposition. Of course, we could also fit a nonplanar surface through the vertices of the triangles. However, it is well-known that the more sample points that are used in a surface patch, the better is the approximation of the underlying surface. Thus, such reasoning implies that the rectangular decomposition is preferable to the triangular decomposition.

An argument could also be made that the planar approximation of the surface provided by the triangle is preferable to the nonplanar approximation provided by the rectangle. This is especially true if ease of interpolation is an issue. In fact, we could also obtain a planar approximation using rectangles by splitting each rectangle into two triangles by adding a diagonal edge between two opposing vertices. There are two choices as shown in Figure 2.154. As long as the four vertices are not coplanar, one will result in a ridgelike planar surface (Figure 2.154(a)), while the other will result in a valleylike planar surface (Figure 2.154(b)). We usually choose the diagonal edge whose angle is the most obtuse.

Neither of the above solutions to the alignment problem that employ a quaternary hierarchical decomposition (whether they use a triangular or rectangular decomposition) yields a conforming mesh because not all of the polygons that make up the mesh have the same number of vertices and edges. Nevertheless, the quaternary hierarchical decomposition that makes use of a rectangular quadtree can form the basis of a conforming mesh and overcome the alignment problem. The idea is to apply a planar triangulation to the result of transforming the underlying rectangular quadtree to a restricted quadtree [1453, 1771, 1772, 1940]. The *restricted quadtree* has the property that all rectangles that are edge neighbors are either of equal size or of ratio 2:1. Such a quadtree is also termed a *1-irregular quadtree* [124], a *balanced quadtree* [204], and a *1-balanced quadtree* [1312]. A somewhat tighter restriction requires that all vertex neighbors either be of equal size or of ratio 2:1 (termed a *0-balanced quadtree* [1312]). It should be clear that a 0-balanced quadtree is also a 1-balanced quadtree but not vice versa (see Exercise 4).

Given an arbitrary quadtree decomposition, the restricted quadtree is formed by repeatedly subdividing the larger nodes until the 2:1 ratio holds. For example, Figure 2.155(a) is the restricted quadtree corresponding to the rectangular decomposition in Figure 2.152(b). Note that the SE quadrant of Figure 2.152(b) has to be decomposed once, in contrast to twice when building the hierarchy of rectangles in Figure 2.153(b). This method of subdivision is also used in finite-element analysis (e.g., [1147]) as part of a technique called *h-refinement* [1013] to refine adaptively a mesh that has already been analyzed, as well as to achieve element compatibility.

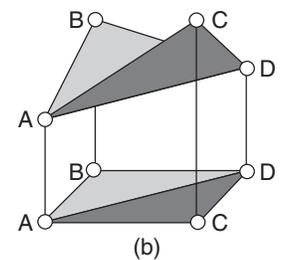
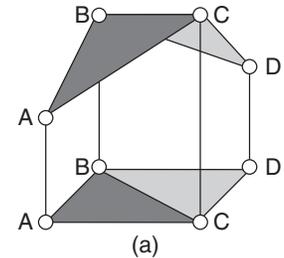


Figure 2.154
 Example illustrating the ambiguity that results when each rectangle is split into two triangles by adding a diagonal edge between (a) vertices B and C and (b) vertices A and D.

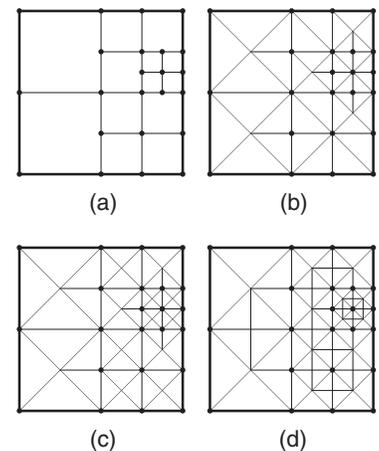


Figure 2.155
 (a) Restricted quadtree corresponding to the rectangular decomposition in Figure 2.152(b) and the result of triangulating it using a (b) two-triangle rule, (c) four-triangle rule, and (d) eight-triangle rule.