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# THE ARC TREE: AN APPROXIMATION SCHEME TO REPRESENT ARBITRARY CURVED SHAPES 

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Memorandum No. UCB/ERL M87/76
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# The Arc Tree: An Approximation Scheme 

## To Represent Arbitrary Curved Shapes

## 1. Introduction

The exact representation of curved geometric objects in finite machines is only possible if the objects can be described by finite mathematical expressions. Typical examples for such objects are paraboloids or ellipses, which can be described by functional equations such as $x^{2} / a^{2}+y^{2} / b^{2}=1$. Many applications, however, especially in computer vision and robotics, do not fit this pattern. The objects to be represented are rather arbitrary in shape, and some approximation scheme has to be employed to represent the data. Any finite machine can only store an approximate representation of the data with limited accuracy. In particular, the answer to any query is based on this approximate representation and may therefore be approximate as well.

Of course, the initial description of a curved object, coming from a camera, a tactile sensor, a mouse, or a digitizer may already be an approximate description of the real object. In most practical applications, this description will be a sequence of curve points or a spline, i.e. a piecewise polynomial function that is smooth and continuous. To support set, search, and recognition operators, however, it is more efficient to represent the data by a hierarchy of detail [Hopc87], i.e. a hierarchy of approximations, where higher levels in the hierarchy correspond to coarser approximations of the curve. Geometric operators can then be computed in a hierarchical manner: algorithms start out near the root of the hierarchy and try to answer the given query at a very coarse resolution. If that is not possible, the resolution is increased
where necessary. In other words, algorithms "zoom in" on those parts of the curve that are relevant for the given query.

In this paper, we develop this theme of hierarchy of detail, focusing on the arc tree, a balanced binary tree that serves as an wproximation scheme to represent arbitrary curved shapes. Section 2 gives a definition of the arc tree and an algorithm to obtain the arc tree representation of a given curve. Section 3 generalizes the concept of the arc tree to include related approaches such as Ballard's strip trees [Ball81] and Bezier curves [Bezi74, Pavl82]. Sections 4 and 5 show how to use arc trees to perform point queries and set operations, such as union or intersection. Both sections also discuss the performance of our implementation. Section 6 outlines how to embed arc trees into an extended database system such as POSTGRES [Ston86b], and section 7 contains a summary and our conclusions.

## 2. Definition

A curve is a one-dimensional continuous point set in $d$-dimensional Euclidean space $\mathbf{E}^{\mathbf{d}}$. For simplicity, we restrict this presentation to the case $d=2$. The generalization to arbitrary $d$ is straightforward. A curve is open if it has two distinct endpoints, otherwise it is called closed; see figure 1 for some examples. As mentioned in the introduction, in practical applications, curves are usually given as a polygonal path, i.e. a sequence of curve points, or as a spline, i.e. a piecewise polynomial function that is smooth and continuous.

The arc tree scheme approximates curves by a sequence of polygonal paths. Let the curve $C$ have length $l$ and be defined by a function $C(t):[0,1] \rightarrow \mathbf{E}^{2}$, such that the


Figure 1: A closed and two open curves
length of the curve from $C(0)$ to $C\left(t_{0}\right)$ is $t_{0} l$. The $k$-th approximation $C_{k}$ ( $k=0,1,2 \ldots$ ) of $C$ is a polygonal path consisting of $2^{k}$ line segments $e_{k, i}\left(i=1 . .2^{k}\right)$, such that $e_{k, i}$ connects the two points $C\left(\frac{i-1}{2^{k}}\right)$ and $C\left(\frac{i}{2^{k}}\right)$. Each edge $e_{k, i}$ can be associated with an arc $a_{k, i}$ of length $l / 2^{k}$, which is a continuous subset of $C$. $C\left(\frac{i-1}{2^{k}}\right)$ and $C\left(\frac{i}{2^{k}}\right)$ are the common endpoints of $e_{k, i}$ and $a_{k, i}$. For $k \geq 1$, each $k$-th approximation is a refinement of the corresponding ( $k-1$ )-th approximation: the vertex set of the $(k-1)$-th approximation is a true subset of the vertex set of the $k$-th approximation. See figure 2 for an example.


Figure 2: A 0th, 1st and 2nd approximation of a curve

More formally, the $k$-th approximation of $C$ is defined by a piecewise linear function $C_{k}(t):[0,1] \rightarrow \mathrm{E}^{2}$ as follows. Here, $\underline{t}$ and $\bar{t}$ denote $\frac{\left|t \cdot 2^{k}\right|}{2^{k}}$ and $\frac{\left[t \cdot 2^{k}\right]}{2^{k}}$, respectively.

$$
C_{k}(t)= \begin{cases}C(t) & t \cdot 2^{k}=0.2^{k} \\ \frac{\bar{t}-t}{\bar{t}-\underline{t}} \cdot C(t)+\frac{t-\underline{t}}{\bar{t}-\underline{t}} \cdot C(\bar{t}) & \text { otherwise }\end{cases}
$$

Then the following convergence theorem is easily proven.

Theorem 1: The sequence of approximation functions $\left(C_{k}(t)\right)$ converges uniformly towards $C(t)$.

Proof: We have to prove $\max _{0 \leq t \leq 1} d\left(C_{k}(t), C(t)\right) \underset{k \rightarrow \infty}{\rightarrow} 0$, or that for any $\varepsilon$, there is a $K$ such that for all $k>K$ and for all $t \in[0,1]$, it is $d\left(C_{k}(t), C(t)\right)<\varepsilon$. Here, $d$ denotes Euclidean distance. Let $K=\log _{2} \frac{l}{\varepsilon}$. Now assume (*) there were some $t$ and some $k>K$ such that

$$
d\left(C_{k}(t), C(t)\right) \geq \varepsilon
$$

Then we have

$$
\begin{aligned}
& d\left(C_{k}(t), C(t)\right) \geq \frac{l}{2^{K}} \\
\Rightarrow & d\left(C_{k}(t), C(t)\right)>\frac{l}{2^{k}} \\
\Rightarrow & d\left(C_{k}(t), C(t)\right)+d\left(C_{k}(t), C(t)\right)>\frac{l}{2^{k}}
\end{aligned}
$$

$\Rightarrow d(C(t), C(t))+d(C(t), C(t))>\frac{l}{2^{k}}$
This is a contradiction to the definition of the $k$-th approximation. The arc from $C(t)$ to $C(t)$ may not be longer than $l / 2^{k}$. Hence, assumption (*) is wrong which proves the theorem.

Moreover, for each approximation $C_{k}$ there is a well-defined area that contains the curve. We have

Lemma 2: Let $E_{k, i}$ denote the ellipse whose major axis is $l / 2^{k}$ and whose focal points are the two endpoints of the edge $e_{k, i}, C\left(\frac{i-1}{2^{k}}\right)$ and $C\left(\frac{i}{2^{k}}\right)$. Then the arc $a_{k, i}$ is internal to $E_{k, i}$.

Proof: (by contradiction) Let $X \in a_{k, i}$ denote a point external to $E_{k, i}$. Then

$$
d\left(X, C\left(\frac{i-1}{2^{k}}\right)\right)+d\left(X, C\left(\frac{i}{2^{k}}\right)\right)>\frac{l}{2^{k}}
$$

Thus, the length of $a_{k, i}$ would be greater than $l / 2^{k}$ which is a contradiction.
Corollary 3: The curve $C$ is internal to the area formed by the union of the bounding ellipses, $\bigcup_{i=0}^{2^{k}} E_{k, i}(k=0,1, \ldots)$.

See figure 3 for an example.


Figure 3: A curve $C$ with its 2 nd approximation $C_{2}$ and corresponding ellipses $E_{2, i}$.

The family of approximations of a given curve $C$ can be stored efficiently in a binary tree. The root of the tree contains the three points $C(0), C(1 / 2)$ and $C(1)$ and is considered on level 1 . If a tree node on level $i$ contains point $C\left(\frac{x}{2^{i}}\right)\left(x=1 . .2^{i}-1\right)$, then its left son contains point $C\left(\frac{2 x-1}{2^{i+1}}\right)$, and its right son contains point $C\left(\frac{2 x+1}{2^{i+1}}\right)$. We call this tree the arc tree of the curve $C$. The arc tree is an exact representation of $C$; each of its subtrees represents a continous subset of $C$. An inorder traversal of the first $k(k \geq 1)$ levels of the arc tree yields the vertices of the $k$-th approximation, sorted by increasing $t$. On the other hand, a breadth-first traversal of the first $k$ levels yields these vertices in an order such that the first $2^{i}+1$ vertices yielded form the $i$-th approximation of $C$. See figure 4 for an example.

In practice, only a finite number of levels of the arc tree is stored. An arc tree with $r$ levels is called an arc tree of resolution $r$. It is a balanced binary tree and it represents the 0 th through $r$-th approximation of $C$.


Figure 4: A curve with approximations and its arc tree. For a closed curve, it is $A=F$.

An arc tree of resolution $r$ can be constructed in two traversals of the given curve $C$. In the first round, one determines the length $l$ of $C$. If $C$ is a spline (or a polygonal path), $l$ can be computed using the following formula for the arc length of an analytical curve. If the curve is given by $y=f(x)$, its length between the points $P_{1}\left(x_{1}, y_{1}\right)$ and $P_{2}\left(x_{2}, y_{2}\right)$ is

$$
l=\int_{x_{1}}^{x_{2}} \sqrt{1+f^{\prime 2}(x)} d x
$$

If it is given by $x=x(t), y=y(t)$, its arc length is

$$
l=\int_{t_{1}}^{t_{2}} \sqrt{x^{2}(t)+y^{\prime 2}(t)} d t
$$

with $x_{i}=x\left(t_{i}\right)$ and $y_{i}=y\left(t_{i}\right)$. One may also attach a label to each knot of $C$ indicating the length accumulated so far. This does not require any additional computation, but it will speed up the second round. In the second round, one picks up the curve
points $C\left(\frac{i}{2^{r}}\right)\left(i \in\left\{0,1 . .2^{r}\right\}\right)$ and inserts them into the appropriate tree nodes while performing a depth-first inorder traversal of the tree.

Note that arc trees can be used to represent any given curve that can be parametrized with respect to arc length. This requirement poses no problem if the input curve is given as a polygonal path or a spline. Nevertheless, there remain problems with some curves such as fractals, for example [Mand77], or with curves that are distorted by high-frequency noise. In both cases the concept of arc length becomes somewhat meaningless and it is necessary to smooth the curve first before the parametrization can take place.

## 3. Generalization

The arc tree parametrizes the given curve by arc length and localizes it by means of bounding ellipses. At higher resolutions the number of ellipses increases, but their total area decreases, thus providing a better localization.

The arc tree can be viewed as just one instance of a large class of approximation schemes that implement Hopcroft's idea of hierarchy of detail [Hopc87]. Higher levels in the hierarchy correspond to coarser approximations of the curve. Associated with each approximation is a bounding area that contains the curve. Set and search operators are computed in a hierarchical manner: algorithms start out near the root of the hierarchy and try to solve the given problem at a very coarse resolution. If that is not possible, the resolution is increased where necessary.

In this section we will present several approximation schemes that are based on the same principle, but that use different parametrizations or bounding areas. For all of these schemes, it is fairly straightforward to obtain the representation of a given spline. Moreover, the algorithms for the computation of set and search operators are essentially the same as the ones for the arc tree, which are presented in sections 4 and 5. It is a subject of further research to conduct a detailed practical comparison of these schemes to find out which schemes are suited best for certain classes of curves.

The first modification of the arc tree concerns the choice of the ellipses $E_{k, i}$ as bounding areas. These ellipses provide the tightest possible bound but, on the other hand, ellipses are fairly complex objects, which has a negative impact on the performance of this scheme. For example, it is often necessary to test two bounding areas for intersection; if the bounding areas are ellipses, this operation is rather costly. Our implementation showed that it is in fact sometimes more efficient to replace the ellipses by their bounding circles; see section 5.1. The circles provide a poorer localization of the curve, but they are easier to handle computationally, which caused the total performance to improve. Other alternatives would be to use bounding boxes whose axes are parallel to the coordinate axes or to the axes of the ellipses. Both of these approaches, however, proved to be less effective than the bounding circles.

If the curves to be represented are polygonal paths with relatively few vertices, it is more efficient to break up the polygonal paths at their vertices rather than to introduce artificial vertices $C\left(1 / 2^{k}\right)$. If a polygonal path has $n+1$ vertices $v_{1} \ldots v_{n+1}$, it can be represented exactly by a polygon arc tree of depth $\left\lceil\log _{2} n\right\rceil$ as follows. The
root of the polygon arc tree contains the vertices $v_{1}, v_{[n / 2\rceil+1}$, and $v_{n+1}$. Its left son contains the vertex $\nu_{\lceil n / 4\rceil+1}$, its right son the vertex $v_{\lceil 3 / 4 \cdot n\rceil+1}$, and so on, until all vertices are stored. Clearly, the arc length corresponding to a node is no more implicit; it has to be stored explicitly with each node. In particular, at each node $N$ it is necessary to know the lengths of the subcurves corresponding to $N$ 's left and right subtree. An example is given in figure 5.


Figure 5: A polygon and corresponding polygon arc tree. The numbers in italics denote arc length.

It is easily seen that some of this length data is redundant. Indeed, with some care it is sufficient to store only one arc length datum per node. For this reason, the storage requirements for a polygon arc tree are only about $20 \%$ to $40 \%$ higher than for a regular arc tree of the same depth.

There are other structures that also implement some hierarchy of detail. One of them is the strip tree, introduced by Ballard [Ball81]. As the arc tree, the strip tree
represents a curve by a binary tree such that each subtree $T$ represents a continuous part $C_{T}$ of the curve. $C_{T}$ is approximated by the line segment connecting its endpoints $\left(x_{b}, y_{b}\right)$ and $\left(x_{e}, y_{e}\right)$. The root node of $T$ stores these two endpoints and two widths $w_{l}$ and $w_{r}$, thus defining a bounding rectangle $S_{T}$ (the strip) that tightly encloses the curve segment $C_{T} . S_{T}$ has the same length as the line segment $\left(\left(x_{b}, y_{b}\right),\left(x_{e}, y_{e}\right)\right)$ and its sides are parallel or perpendicular to it. See figure 6 for an example of a curve and a corresponding strip tree. Clearly, this approach requires some extensions for closed curves and for curves that extend beyond their endpoints (fig. 7).

When a strip tree is constructed for a given curve $C$, a curve segment $C_{T}$ is subdivided further until the total strip width $w_{l}+w_{r}$ is below a certain threshold. As it is a non-trivial operation to obtain the strip $S_{T}$ for every curve segment $C_{T}$, the construction of a strip tree for a given curve may be quite costly. To subdivide $C_{T}$, one can choose any point of $C_{T}$ that lies on the boundary of the corresponding strip $S_{T}$. Clearly, a strip tree is not necessarily balanced (see also figure 6) which has a negative impact on its average-case performance. Note that arc trees are balanced, which might give them an edge over strip trees in terms of average performance.

Also, a strip tree requires about twice as much space as an arc tree of same depth: each arc tree node stores a minimum of two real numbers and two pointers, whereas a strip tree node stores six real numbers and two pointers. . Note, however, that strip trees can be modified to require less storage. First, all subdivision points belong to more than one strip and are therefore stored in more than one node. The


Figure 6: A curve with strip, a hierarchy of strips, and a corresponding strip tree.
redundant data may be replaced by pointers or deleted, which may require that some of the algorithms are slightly modified. Second, rather than storing $w_{l}$ and $w_{r}$, one may just store the maximum of these two widths. The resulting strip is potentially


Figure 7: $\quad$ A curve $C$ that extends beyond its endpoints. There is no bounding box of length $l$ that contains $C$.
wider and provides a poorer localization. In both cases, some loss in performance is likely, but it will probably be minor compared to the savings in storage space.

A very different approach to implement a hierarchy of detail is based on curve fitting techniques such as Bezier curves [Bezi74] or B-splines [DeBo78]; see also [Pav182] for a good survey of these and related techniques. A Bezier curve of degree $m$ is an $m$-th degree polynomial function defined by $m+1$ guiding points $P_{1}$. . $P_{m+1}$. The curve goes through the points $P_{1}$ and $P_{m+1}$ and passes near the remaining guiding points $P_{2} \ldots P_{m}$ in a well-defined manner. The points $P_{2}$ through $P_{m}$ may be relocated interactively to bring the Bezier curve into the desired form. See figure 8 for two examples.

It can be shown that a Bezier curve lies within the corresponding characteristic polygon, i.e. the convex hull of its guiding points. Also, a Bezier curve $B$ can be subdivided into two Bezier curves $B_{1}$ and $B_{2}$ of same degree. The characteristic polygons of $B_{1}$ and $B_{2}$ are disjoint and subsets of $B$ 's characteristic polygon. They


Figure 8: Examples of Bezier polynomials with three and five guiding points.
therefore provide a better localization of $B$; see figure 9 .


Figure 9: $\quad$ A Bezier curve $B$ partitioned into two curves $B_{1}$ and $B_{2}$ with characteristic polygons.

Now we can derive a hierarchical representation of a given Bezier curve $B$ as follows. The first approximation is the edge segment connecting $B$ 's endpoints; its bounding area is given by $B$ 's characteristic polygon. The second approximation is the polygonal path connecting the endpoints of $B_{1}$ and $B_{2}$; its bounding area is the
union of the characteristic polygons of $B_{1}$ and $B_{2}$, and so on. There are various efficient subdivision algorithms to obtain $B_{1}$ and $B_{2}$ from a given $B$; see for example [Pav182], pp. 221-230.

The main problem with this approach seems to be that not every curve can be approximated well by a low-order Bezier curve. A high-order Bezier curve, however, is harder to partition and has a more complex characteristic polygon, which has an adverse impact on the performance of this scheme. In practice, complex curves are often approximated by several third-order Bezier curves. This would mean that the bounding area of the first approximation is a union of convex polygons, which is already rather complex. Further approximations are then obtained by subdivisions of each one of these polygons. Nevertheless, this approach seems very promising and should be included in a practical comparison of the various approaches to implement a hierarchy of detail.

We expect arc or strip trees to be superior to Bezier curves if the curves to be represented are initially described by a long sequence of curve points and can only be described by high-order splines or a large number of simpler splines. This is often the case if curves are input from a digitizer pad or a mouse. On the other hand, if a curve is initially given by a few simple splines, it is probably more efficient to keep this representation and use spline subdivision algorithms as described above to implement a hierarchy of detail.

B-splines can be used in a way similar to Bezier curves to implement a hierarchy of detail. For appropriate subdivision algorithms, see [Bohm84].

Certainly, there are many more possibilities to implement a hierarchy of detail as a tree structure similar to the schemes presented above. Note that in all of these schemes it is possible to trade space with time as follows. Rather than storing all lower level approximations explicitly, one could keep the source description of the curve in main memory and compute finer approximations "on the fly" when needed. This approach can be viewed as a procedural arc tree as finer approximations are defined procedurally, i.e. by means of the appropriate subdivision algorithm that computes finer approximations from coarser ones. This approach seems particularly promising for the Bezier approach where highly efficient subdivision algorithms are available. In the case of arc and strip trees, the computations to obtain finer approximations are probably too complex to be repeated at every tree traversal.

As mentioned above, the algorithms for set and search operations for these various approximation schemes are all essentially the same. In the following two sections, we give the algorithms for the arc tree scheme. In most cases, the corresponding algorithms for the other schemes are simply obtained by replacing the ellipses $E_{k, i}$ by the corresponding bounding areas, viz., the characteristic polygons for the curve fitting approaches or the strips for the strip tree.

## 4. Hierarchical Point Inclusion Test

To demonstrate the power of the arc tree representation scheme, we first show how to answer point queries on the arc tree. Given a point $A \in \mathbf{E}^{\mathbf{2}}$ and a simple ${ }^{*}$

[^0]closed curve $C$, a point query asks if $A$ is internal to the simple point set enclosed by $C, P(C)$. For simplicity, we also describe this case by stating that $A$ is internal to $C$, or that $A \in P(C)$.

The point inclusion test is performed by a hierarchical algorithm called HPOINT, which starts with some simple approximation $C_{a p p}$ of $C$. For each edge $e_{k, i}$ of $C_{\text {app }}\left(i=1 . .2^{k}\right)$, it checks if the replacement of $e_{k, i}$ by the arc $a_{k, i}$ may affect the internal/external classification of $A$. If there is no such edge $e_{k, i}$, then $A \in P\left(C_{a p p}\right)$ is equivalent to $A \in P(C) ; H P O I N T$ uses a conventional algorithm to solve the point query $A \in P\left(C_{a p p}\right)$ ? and terminates. Otherwise, HPOINT replaces each edge $e_{k, i}$, whose replacement by $a_{k, i}$ may affect $A$ 's classification, by the two edges $e_{k+1,2 i-1}$ and $e_{k+1,2 i}$. The resulting polygon is a closer approximation of $C$. HPOINT proceeds recursively with that polygon.

If the maximum resolution has been reached without obtaining a result, then the problem cannot be decided at that resolution. In fact, there are boundary points (such as $C(1 / 3)$ ) that cannot be decided at any finite resolution. There are three ways to resolve this situation: (i) the algorithm returns unclear, (ii) the algorithm considers the point a boundary point, or (iii) the arc tree is extended at its leaf nodes to include the source description of the curve; then, edges $e_{k, i}$ may eventually be replaced by $\operatorname{arcs} a_{k, i}$ to allow an exact query evaluation. For HPOINT, we choose option (ii), thus considering the boundary as having a nonzero width. In our definition of the point inclusion test, where the given point set $P(C)$ is closed, HPOINT returns $A \in P(C)$, accordingly.

We are left with the problem of how to find out quickly if the replacement of $e_{k, i}$ by $a_{k, i}$ may affect the internal/external classification of $A$. From lemma 2, we obtain

Lemma 4: Let $C_{k, i}$ denote the curve obtained from $C$ by replacing the arc $a_{k, i}$ by the straight line $e_{k, i}$. Then, if $A$ is external to $E_{k, i}$, it is $A \in P(C)$ equivalent to $\left.A \in P\left(C_{k, i}\right)\right)$.

Proof: Because $A$ is external to $E_{k, i}, A$ may not lie on or between $a_{k, i}$ and $e_{k, i}$. Therefore, the replacement of $a_{k, i}$ by $e_{k, i}$ may not affect the internal/external classification of $A$.

It is therefore sufficient to check if $A$ is internal to $E_{k, i}$. If yes, the replacement of $e_{k, i}$ by $a_{k, i}$ may affect the classification of $A$, otherwise it may not. Letting the initial approximation be $C_{0}, H P O I N T$ can be described more precisely as follows.

## Algorithm HPOINT

Input: A point $A \in \mathbf{E}^{2}$. The arc tree $T_{C}$ of a simple closed curve $C$.

Output: $A \in P(C)$ ?
(1) Set the approximation polygon $C_{a p p}$ to $C_{0}$ and $k$ to zero.
(2) For each edge $e_{k, i}\left(i \in\left\{1.2^{k}\right\}\right)$ of $C_{a p p}$ do
(2a) If $A$ is one of the endpoints of $e_{k, i}$, return true and stop.
(2b) Otherwise, if $A$ is internal to the ellipse $E_{k, i}, \operatorname{tag} e_{k, i}$.
(3) If $C_{\text {app }}$ has no tagged edges, use a conventional point inclusion algorithm to determine if $A \in P\left(C_{\text {app }}\right)$, return the result and stop.
(4) Otherwise, if $k$ is less than the maximum resolution, depth $\left(T_{C}\right)$, replace each tagged edge $e_{k, i}$ by the two edges $e_{k+1,2 i-1}$ and $e_{k+1,2 i}$, increase $k$ by one and repeat from (2).
(5) Otherwise, retun true and stop.

Step (2a) is necessary for termination if $A$ is a boundary point. Step (2b) can easily be done by computing the distances from $A$ to the two focal points of $E_{k, i}$. Step (4) can be performed by using $C$ 's arc tree in the following manner. Each edge $e_{k, i}$ is associated with the subtree whose root contains the point $C\left(\frac{2 i-1}{2^{k+1}}\right)$. Note that this is the curve point which corresponds to the center point of $e_{k, i}$ and which $e_{k+1,2 i-1}$ and $e_{k+1,2 i}$ have in common. If $e_{k, i}$ is to be replaced by $e_{k+1,2 i-1}$ and $e_{k+1,2 i}$, HPOINT obtains that point from the tree node and continues recursively on both subtrees of this node.

Steps (2) and (4) can now be performed during a top-down traversal of the arc tree. Each subtree can be processed independently of the others, which offers a natural way to parallelize the algorithm. If $C_{a p p}$ has no more tagged edges, or if the maximum resolution has been reached, the partial results are collected in a bottom-up traversal of the tree and put together to form the boundary of the final approximation polygon $C_{\text {app }}$. At this point, $A \in P(C)$ is equivalent to $A \in P\left(C_{\text {app }}\right)$. Step (4) can be performed by Shamos' algorithm, where one constructs a horizontal line $L$ through $A$
and counts the intersections between $L$ and the edges of $C_{a p p}$ that lie to the left of $A$. If the number of intersections is odd then $A$ is internal, otherwise it is external. Shamos' algorithm requires some special maintenance for horizontal edges; see [Prep85] for details.

We implemented this algorithm on a VAX 8800 and ran several experiments to see how HPOINT's time complexity correlates with the complexity of the given curve $C$ and with the location of $A$ with respect to $C$. Our running times should not be considered in absolute terms as we did not make a strong effort to optimize our programs. However, the figures are appropriate for comparative measurements. Figures 10 and 11 show our results. Here, $t$ is CPU time in $m s$, and $r$ is the resolution at which the query was decided. The dotted polygons are the $r$-th approximations of $C$, respectively.


Figure 10: $C$ is a spline with 12 knots.

Note that the use of alternative approximation schemes is unlikely to improve the performance of our algorithms. To test a given point for inclusion in a given

(a) $r=2, r=4.0$

(b) $r=4, t=6.4$

(c) $\mathrm{r}=4, \mathrm{t}=7.1$

(d) $\mathrm{r}=6, \mathrm{t}=8.9$

Figure 11: $C$ is a spline with 36 knots.
ellipse has about the same complexity as the corresponding tests for a characteristic polygon (say, a convex quadrilateral) or a strip. On the other hand, the test is somewhat easier for circles or for boxes whose axes are parallel to the coordinate axes. In both cases, however, the localization of the curve that is provided by these areas is poorer than for the bounding areas above.

Our algorithm HPOINT is an application of Hopcroft's idea of hierarchy of detail [Hopc87]. It solves the point inclusion problem by starting with a very simple representation of $C$ and introduces more complex representations only if they are required to solve the problem. The algorithm "zooms in" on those parts of $C$ that are interesting in the sense that they may change the internal/external classification of the point $A$ at a higher resolution. As our examples demonstrate, HPOINT terminates very quickly if $A$ is not close to $C$. The closer $A$ gets to $C$, the higher is the
resolution required to answer the point query. Due to a quick localization of the interesting parts of $C$, the algorithm does not show the quadratic growth in the complexity of $C$ that a worst-case analysis would predict.

## 5. Hierarchical Set Operations

In this section, we show how to detect and compute intersections, unions, and differences of one- and two-dimensional point sets. We assume that the input point sets are simple and that they are given by their arc trees or by the arc trees of their boundaries. Again, the idea is to inspect approximations of the input curves by increasing resolution and to "zoom in" on those parts of the boundaries that may participate in an intersection.

### 5.1. Curve-Curve Intersection Detection

We first show how to test two given curves $C$ and $D$ for intersection. The hierarchical algorithm HCURVES starts with simple approximations $C_{\text {app }}$ and $D_{\text {app }}$ of $C$ and $D$, respectively, and continues with approximations of higher resolutions where necessary. We have

Lemma 5: The arcs $a_{k, i}$ and $b_{k, j}$ corresponding to the edges $e_{k, i}$ of $C_{a p p}$ and $f_{k, j}$ of $D_{\text {app }}$, respectively, must intersect if the following three conditions are met:
(i) $e_{k, i}$ intersects $f_{k, j}$,
(ii) the two endpoints of $e_{k, i}$ are external to the ellipse $F_{k, j}$ corresponding to $f_{k, j}$,
(iii) the two endpoints of $f_{k, j}$ are external to the ellipse $E_{k, i}$ corresponding to $e_{k, i}$.

Proof: Any situation where all three conditions are met are topologically equivalent to the situation in figure 12.


Figure 12

The intersection of the two ellipses $E_{k, i}$ and $F_{k, j}$ is a quadrilateral ABCD with curved edges $\mathrm{AB}, \mathrm{BC}, \mathrm{CD}$, and DA. The segment of the arc $a_{k, i}$ that is interior to $A B C D$ connects some point of $A B$ with some point of $C D$. The segment of the arc $b_{k, i}$ that is interior to ABCD connects some point of BC with some point of DA. Obviously, this is not possible without an intersection of the two arc segments, which proves the lemma.

Now the algorithm HCURVES proceeds as follows. For each pair of edges, $e_{k, i}$ of $C_{\text {app }}$ and $f_{k, j}$ of $D_{\text {app }}\left(i, j \in\left\{0,1 . .2^{k}\right\}\right)$, HCURVES checks if their corresponding arcs may intersect. According to lemma 2, this can be done by testing if the corresponding ellipses $E_{k, i}$ and $F_{k, j}$ intersect. If yes, HCURVES puts tags on $e_{k, i}$ and $f_{k, j}$ and applies lemma 5 to see if the arcs must intersect. If yes, HCURVES reports an intersection and stops. After all edges $e_{k, i}$ of $C_{a p p}$ have been processed,

HCURVES checks if there are any tagged edges. If no, HCURVES reports no intersection and stops. Otherwise, HCURVES replaces all tagged edges by the corresponding edges of the next higher approximation, increases $\boldsymbol{k}$ by one, and proceeds recursively on the refined curves. If the maximum resolution has been reached and there are still tagged edges, HCURVES interprets the situation as an intersection of the boundaries and returns an intersection. More exactly, HCURVES can be described as follows.

## Algorithm HCURVES

Input: The arc trees $T_{C}$ and $T_{D}$ of two curves $C$ and $D$.

Output: $C \cap D \neq \phi$ ?
(1) Set the approximation polygons $C_{a p p}$ to $C_{0}, D_{a p p}$ to $D_{0}$, and $k$ to zero.
(2) For each pair of edges $e_{k, i}$ of $C_{a p p}$ and $f_{k, j}$ of $D_{a p p}$ do
(2a) Check if the two ellipses $E_{k, i}$ and $F_{k, j}$ intersect.
(2b) If yes, tag $e_{k, i}$ and $f_{k, j}$; if conditions (i) through (iii) in lemma 5 are met or if $e_{k, i}$ and $f_{k, j}$ share one or two endpoints, return true and stop.
(3) If there are no tagged edges, return false and stop.
(4) If $k$ is less than the maximum resolution, $\min \left(\operatorname{depth}\left(T_{C}\right), \operatorname{depth}\left(T_{D}\right)\right.$ ), replace each tagged edge $e_{k, i}$ of $C_{\text {app }}$ by the two edges $e_{k+1,2 i-1}$ and $e_{k+1,2 i}$. Similarly for each tagged edge $f_{k, j}$ of $D_{\text {app }}$. Increase $k$ by one and repeat from (2).
(5) Otherwise, the maximum resolution has been reached; return true and stop.

We implemented this algorithm on a VAX 8800 with a few slight modifications to speed up execution. First, the test if the two ellipses $E_{k, i}$ and $F_{k, j}$ intersect is replaced by a test if the two circumscribing circles of $E_{k, i}$ and $F_{k, j}$ intersect. If those do not intersect then the ellipses do not intersect either. Otherwise, we assume that the ellipses may intersect and proceed accordingly. We made several experiments with more accurate tests, such as to test bounding boxes of the two ellipses for intersection, or to test the two ellipses themselves for intersection. In every case, the execution times went up between $25 \%$ and $60 \%$. The more accurate tests required a significant amount of CPU time, but they only marginally reduced the number of tagged edges.

Second, rather than performing step (2) for each pair of edges $e_{k, i}$ of $C_{a p p}$ and $f_{k, j}$ of $D_{a p p}$, we maintain matrices to keep track which pairs of ellipses ( $E_{k, i}, F_{k, j}$ ) pass the intersection test in step (2a). Then, step (2) is executed for a pair of edges ( $e_{k, i}, f_{k, j}$ ) if and only if the ellipses $E_{k-1,\lceil i / 2\rceil}$ and $F_{k-1,\lceil j / 2\rceil}$, which correspond to their parent edges, intersect. Otherwise, it is known in advance that $E_{k, i}$ and $F_{k, j}$ do not intersect.

Figures 13 and 14 give several examples for the performance of the algorithm. Here, $r$ denotes the resolution at which the algorithm is able to decide the query, and $t$ denotes the CPU time in $m s$.


Figure 13: $C$ is a spline with 13 knots, $D$ a spline with 8 knots.


Figure 14: $C$ is a spline with 24 knots, $D$ a spline with 23 knots.

Again, it is not clear if the use of alternative approximation schemes might yield a better performance. The crucial operation in algorithm HCURVES is the test if two bounding areas intersect. In the case of circles, this is a trivial operation: two circles
intersect if the distance between their centers is no more than the sum of their radii. The corresponding tests for boxes or characteristic polygons (say, convex quadrilaterals) are about two to three times as complex.

Note that the running times do not grow quadratically with the complexity of the input curves. The example in figure 11 (b) requires a large amount of CPU time due to the fact that the two curves are quite interwoven but do not intersect. It is therefore necessary to get down to fairly high resolutions in order to determine that there is no intersection. It seems that a case like this will require a lot of computation with any other intersection detection algorithm as well.

### 5.2. Curve-Curve Intersection Computation

The intersection is actually computed by the hierarchical algorithm HCRVCRV, a variation of algorithm HCURVES. HCRVCRV does not test if two arcs must intersect. It continues recursive refinement until one of the following two conditions is met: (i) there are no more tagged edges, or (ii) the maximum resolution has been reached. In case (i), $C$ and $D$ do not intersect. In case (ii), each tagged edge of $C_{\text {app }}$ is intersected with each tagged edge of $D_{a p p}$ and the intersection points are returned.

## Algorithm HCRVCRV

Input: The arc trees $T_{C}$ and $T_{D}$ of two curves $C$ and $D$.

## Output: $C \cap D$

(1) Set the approximation polygons $C_{a p p}$ to $C_{0}, D_{a p p}$ to $D_{0}$, and $k$ to zero.
(2) For each pair of edges $e_{k, i}$ of $C_{a p p}$ and $f_{k, j}$ of $D_{a p p}$, check if the two ellipses $E_{k, i}$ and $F_{k, j}$ intersect. If yes, tag $e_{k, i}$ and $f_{k, j}$.
(3) If there are no tagged edges, return no intersection and stop.
(4) Otherwise, if $k$ is less than the maximum resolution, $\min \left(\operatorname{depth}\left(T_{C}\right)\right.$, depth $\left.\left(T_{D}\right)\right)$, replace each tagged edge $e_{k, i}$ of $C_{\text {app }}$ by the two edges $e_{k+1,2 i-1}$ and $e_{k+1,2 i}$. Similarly for each tagged edge $f_{k, j}$ of $D_{a p p}$. Increase $k$ by one and repeat from (2).
(5) Otherwise, the maximum resolution has been reached. Intersect each tagged edge $e_{k, i}$ with each tagged edge $f_{k, j}$, report all intersection points and stop.

We implemented this algorithm on a VAX 8800 with the same modifications as in the case of HCURVES. Figures 15 and 16 give two examples for the performance of the algorithm at various maximum resolutions $r . P$ is an intersection point, $d$ is the distance between $P$ and its approximation, $C_{r}$ and $D_{r}$ are $C$ 's and $D$ 's approximations at maximum resolution, and $t$ is CPU time required to compute all intersections.


Figure 15: $C$ is a spline with 13 knots, $D$ a spline with 8 knots.


Figure 16: Both $C$ and $D$ are splines with 20 knots.

Note that the running times do not increase quadratically with the number of edges, $2^{r}$, or with the complexity of the input curves. In fact, the increase in CPU time is about cubical in $r$, i.e. polylogarithmic in the number of edges. The following plot shows the increase in CPU time for both figures and for resolutions $r=2$ through $r=7$. The broken lines indicate the distance $d$ between the actual intersection point $P$ and the corresponding intersection point returned by $H C R V C R V$ at maximum resolution $r$.


Figure 17

### 5.3. Curve-Area Intersection Detection

Given the arc trees of a curve $C$ and a closed curve $D$, it is now easy to detect if $C$ intersects the point set $P(D)$. First, one employs algorithm HCURVES to check $C$ and $D$ for intersection. If the two curves do not intersect, it may be possible that $C$ is internal to $D$. This can be checked by algorithm HPOINT by testing some point of $C$ if it is internal to $D . C$ and $P(D)$ do not intersect if and only if both tests fail.

### 5.4. Curve-Area Intersection Computation

To actually compute the intersection of a curve with an area, we present the hierarchical algorithm $H C R V A R A$. Given the arc trees of a curve $C$ and a simple closed curve $D, H C R V A R A$ computes $C \cap P(D)$. The initiation and the recursion
step of HCRVARA are identical to the corresponding sections of algorithm HCRVCRV. As HCRVCRV, HCRVARA proceeds recursively until one of two conditions is met: (i) there are no more tagged edges, or (ii) the maximum resolution has been reached.

In case (i), it may be that $C$ is internal to $D$. A point query on some point of $C$ suffices to decide if that is the case. In case (ii), each tagged edge of $C_{a p p}$ is intersected with each tagged edge of $D_{\text {app }}$ and subdivided at the intersection points into disjoint edge segments. Now each edge segment of $C_{a p p}$ is either internal or external to $D_{\text {app }} . H C R V A R A$ performs a point query for some point of $C_{a p p}$ to see if it is internal or external. Starting from that point, $H C R V A R A$ performs a traversal of $C_{a p p}$ to label each edge as internal or external. The label is alternately internal or external, changing at each intersection point. Some special handling is required for edges of $C_{a p p}$ that coincide with edges of $D_{a p p} ;$ see figure 18 for an example.


Figure 18: The dotted segments of $C_{a p p}$ are internal, the broken segments external.

Finally, HCRVARA replaces all untagged internal edges of $C_{a p p}$ by the corresponding edges of maximum resolution, and returns the internal edges and edge
segments of $C_{\text {app }}$. It follows a more exact description of $\operatorname{HCRVARA}$.

## Algorithm HCRVARA

Input: The arc trees $T_{C}$ and $T_{D}$ of a curve $C$ and a simple closed curve $D$.
Output: $C \cap^{P(D)}$
(1) Set the approximation polygons $C_{a p p}$ to $C_{0}, D_{a p p}$ to $D_{0}$, and $k$ to zero.
(2) For each pair of edges $e_{k, i}$ of $C_{a p p}$ and $f_{k, j}$ of $D_{\text {app }}$, check if the two ellipses $E_{k, i}$ and $F_{k, j}$ intersect. If yes, tag $e_{k, i}$ and $f_{k, j}$.
(3) If there are no tagged edges, return no intersection and stop.
(4) Otherwise, if $k$ is less than the maximum resolution, $\min \left(\operatorname{depth}\left(T_{C}\right), \operatorname{depth}\left(T_{D}\right)\right)$, replace each tagged edge $e_{k, i}$ of $C_{\text {app }}$ by the two edges $e_{k+1,2 i-1}$ and $e_{k+1,2 i}$. Similarly for each tagged edge $f_{m, j}$ of $D_{\text {app }}$. Increase $k$ by one and repeat from (2).
(5) Otherwise, the maximum resolution has been reached. Intersect each tagged edge $e_{k, i}$ with each tagged edge $f_{k, j}$ and subdivide the edges $e_{k, i}$ at their intersection points into disjoint segments.
(6) Perform a point query for some point of $C_{a p p}$ to see if it is internal or external to $D_{a p p}$.
(7) Traverse $C_{a p p}$ and label edges as internal or external. The label is alternately internal or external, changing at each intersection point.
(8) Replace the internal untagged edges by the corresponding edges of maximum resolution.
(9) Return the internal edges and edge segments of $C_{\text {app }}$.

We implemented this algorithm on a VAX 8800 with the same modifications as in the case of HCURVES. Figures 19 and 20 give two examples for the output of the algorithm at various maximum resolutions $r$. The dotted curves are the $r$-th approximation of $D$, respectively.


Figure 19: $C$ is a spline with 10 knots, $D$ a spline with 18 knots.


Figure 20: Both $C$ and $D$ are splines with 20 knots.

Again, the running times do not increase quadratically with the number of edges, $2^{r}$, or with the complexity of the input curves. In fact, the increase in CPU time is about cubical in $r$, i.e. polylogarithmic in the number of edges. Figure 21 shows the increase in CPU time for both figures and for resolutions $r=2$ through $r=7$.


Figure 21

### 5.5. Area-Area Intersection Detection

Given the arc trees of two closed curves $C$ and $D$, it is now easy to detect if the enclosed point sets $P(C)$ and $P(D)$ intersect. First, one employs algorithm HCURVES to check $C$ and $D$ for intersection. If the two curves do not intersect, it may be possible that $C$ is internal to $D$, or vice versa. This can be checked by algorithm HPOINT by testing some point of $C$ if it is internal to $D$, and some point of $D$ if it is internal to $C$. The two areas do not intersect if and only if all tests fail.

### 5.6. Area-Area Set Operations

Given the arc trees of two closed curve $C$ and $D$, the intersection of $P(C)$ and $P(D)$ can now be computed as follows. First, one employs algorithm HCRVARA to
compute $C \cap^{P(D)}$ and $D \cap^{P(C) \text {. The resulting curves form the boundary of the }}$ intersection $P(C) \cap P(D)$. Some special handling is required for those edge segments that $C$ and $D$ have in common. HCRVARA has to be modified such that it marks these segments in its output. These segments are included in the boundary if and only if the corresponding edges of $C$ and $D$ have the same orientation; see figure 22.


Figure 22: EF is included in the boundary of $P(C) \cap P(D), \mathrm{AB}$ is not.

We implemented this algorithm on a VAX 8800 with the same modifications as in the case of HCURVES. Figures 23 and 24 give two examples for the performance of the algorithm at various maximum resolutions $r$. The broken curves are the $r$-th approximations of $C$ and $D$, respectively.


Figure 23: $C$ is a spline with 10 knots, $D$ a spline with 20 knots.


Figure 24: Both $C$ and $D$ are splines with 20 knots.

Again, the running times do not increase quadratically with the maximum resolution or with the complexity of the input curves.

To obtain the boundary of the union $P(C) \cap P(D)$, one computes those segments of $C$ that are external to $D$ and those segments of $D$ that are external to $C$. Again, the edge segments that $C$ and $D$ have in common are included if and only if the corresponding edges of $C$ and $D$ have the same orientation.

To retrieve the boundary of the difference $P(C)-P(D)$, one computes those segments of $C$ that are external to $D$ and those segments of $D$ that are internal to $C$. The edge segments that $C$ and $D$ have in common are included if and only if the corresponding edges of $C$ and $D$ do not have the same orientation.

## 6. Implementation in a Database System

As the previous sections have shown, the arc tree is an efficient scheme to represent curves. In large-scale geometric applications such as geography or robotics, is is usually most efficient to have a separate data management component and to maintain a geometric database to store a large number of geometric objects. In order to use the arc tree representation scheme efficiently in this context, it is therefore necessary to embed arc trees as complex objects in the database system. This section will discuss several ways to perform this embedding; we will restrict our analysis to relational databases.

There are three major ways to implement complex objects in an extended relational database system such as POSTGRES [Ston86b] or DASDBS [Paul87]. First,
one may organize the data of a complex object in relational form and represent the object as a set of tuples, each marked with a unique object identifier. Then the algorithms may be either programmed in an external host language with embedded query language commands [RTI84], or within the database system by means of user-defined operators [Wong85]. These approaches have been used in earlier attempts to extend relational database systems to applications in geography and robotics [Kung84, Gunt87]. Second, one supports a procedural data type to store expressions in the query language or any other programming language directly in the database. This approach is emphasized in the POSTGRES database system [Ston86a]. Third, one may define an abstract data type (ADT) with corresponding operators and abstract indices; see for example [Ston83]. The importance and suitability of ADT mechanisms for geometric data management has also been discussed by Schek [Sche86]. The following subsections will discuss these approaches in turn and evaluate their suitability to embed arc trees in a relational database.

### 6.1. The Pure Relational Approach

The traditional approach would be to represent a complex object as a set of tuples, i.e. as a relation or subrelation. For the representation of an arc tree the following database design may be used.

$$
\begin{aligned}
& \text { arctreenodes }(\text { tree-id }=\text { int, node-id }=\text { int, point- } x=\text { real, point- } y=\text { real, } \\
& \qquad \text { left-son }=\text { int, right-son }=\text { int })
\end{aligned}
$$

Then the algorithms for intersection detection and so on are coded in a generalpurpose programming language (the host language) that allows the embedding of
query commands to access the database. In the case of INGRES [Ston76], one may use, for example, EQUEL/FORTRAN [RT184].

For this approach, the relational data model as defined by Codd [Codd70] would be sufficient. It would not be necessary to extend the data model by new concepts such as special data types, and query optimization could be carried out as usual. Nevertheless, we do not believe that this approach will be very efficient. For each access to a tree node it is necessary to activate the interface between host language and the database system. In order to get the left -son node of a given node $N$, for example, it is necessary to process the following query.
range of a1, a2 is arctreenodes
retrieve (al.all)
where a1.node-id = a2.left-son
and a2.node-id $=N$

This query involves a join of the relation arctreenodes with itself. Then the resulting tuple has to be returned to the host language before the execution of the program can continue. This is a major effort to retrieve just one node, which may slow down the overall performance of our algorithms considerably.

### 6.2. Relational Data Type and User Defined Features

A variation of this approach would be to represent the arc tree as above, but to program the algorithms within the query language by means of a relational data type and user defined data types and operators [Wong85]. First, the relational data type is used to represent each arc tree as one tuple in a relation arctrees:
arctrees $($ tree-id $=$ int, nodes $=$ arctreenodes using tree-id $)$
Here, the domain nodes is of the relational data type arctreenodes. A value of this domain is the set of all tuples in arctreenodes that share the same tree-id value.

Second, the user has to define the geometric data types and operators that are needed in this context, based on the data types and operators provided by the database system. For example, one may define a data type line in two dimensions as define type line $($ phase $=$ real, dist $=$ real $)$
where phase denotes the angle between the line and the $\boldsymbol{x}$-axis, and dist is the distance between the line and the origin. Then one defines an operator intersect as

$$
\text { define operator intersect ( } l 1=\text { line, } l 2=\text { line }) \text { as } z=\text { boolean }
$$

where $\mathrm{z}=1$ if $l 1$.phase $\neq 12$.phase or 11 .dist $=12$.dist

Eventually, one will be able to program arc tree algorithms within the extended query language. Clearly, each such program $P$ that uses any of the user defined data types and operators can be mapped onto a program $\bar{P}$ in the basic query language. Then the query optimization can be performed on $\bar{P}$ in the usual manner. Moreover, there will be opportunities to perform some kind of global query optimization [Sell85] because the queries do not have to be processed one by one, as in the case of the host language approach.

One problem with this approach is that it requires the definition of a lot of data types and operators before algorithms can be coded. Also, it is not sure if the database can provide an efficient environment for the program execution. Finally, this approach does not really make use of the special properties of the arc tree and the
access paths required. The arc tree is a very regular structure, and the set of operators to be performed is very limited. Any selective access to lower level subtrees is embedded in a more complex operator, such as union or intersection, that starts out at the root of the tree and works its way down from there. Nevertheless, this approach seems to be promising and should be included in a practical performance analysis.

### 6.3. Procedure as a Data Type

Another method to support complex objects is to introduce a procedural data type; in particular, a data type query seems to be useful. This approach has first been suggested by Stonebraker [Ston84] and it is currently being implemented in POSTGRES. The procedural data type refers to components that are complex objects themselves by means of a retrieval command. This approach provides easy access to lower level components via the multiple-dot notation and provides efficient support for shared subobjects.

Consider the following POSTGRES example with two objects apple and orange and three relations polygon, circle, and line .

| name | desc |
| :--- | :--- |
| apple | retrieve (polygon.all) where polygon.id $=10$ <br> retrieve (circle.all) where circle.id $=40$ |
| orange | retrieve (line.all) where line.id $=17$ <br> retrieve (polygon.all) where polygon.id $=10$ |

Table 1: The object relation.

Clearly, the polygon 10 is a complex object that is shared by both apple and orange . To retrieve the area of the shared polygon, for example, one may use the multiple-dot notation [Zani83] as follows.
retrieve (object.desc.polygon.area) where object.name = 'apple'

In order to improve performance, it is usually useful to precompute access plans or even answers to stored queries. This precomputation step makes the query optimization somewhat more complicated, but it improves overall efficiency. As discussed in [Ston86a], the procedural data type also provides efficient support for complex objects with many levels of subobjects and complex objects with unpredictable composition.

The arc tree is certainly an object with many levels of subobjects, but it has a very regular structure and no shared subobjects. Furthermore, the set of operators to be performed is very limited, and any selective access to lower level subtrees is
embedded in a more complex operator, such as union or intersection, that starts out at the root of the tree and works its way down from there. We therefore do not believe that the procedural data type is an adequate embedding for arc trees; it is too complicated because it is too powerful. We advocate to use the simpler ADT scheme as described in the following subsection.

### 6.4. Abstract Data Types

Although the arc tree is a useful representation scheme for the most important geometric operators, it should not necessarily be visible to the user. On the contrary, all set and search operators should be executed without revealing the internal representation scheme - the arc tree - to the user. The only operator where the internal representation may be visible to the user is the rendering of approximations of the curve. But even then, it seems preferable to offer an operator that maps an abstract object of type curve and a resolution into an approximation of the curve. Note that for none of the common operators the user needs to have explicit access to subtrees or to retrieve or manipulate details of the arc tree. On the other hand, it is important to implement the algorithms for set and search operations as efficiently as possible. The algorithms are complex, and their performance should not be impeded unnecessarily by an insufficient runtime environment or an inadequate implementation language.

Because of these considerations and because of the limited number of operators, we believe that an embedding of the arc tree as an abstract data type (ADT) into an extended database system is the superior solution to the problem. An ADT is an encapsulation of a data structure (so that its implementation details are not visible to
an outside client procedure) along with a collection of related operators on this encapsulated structure. The canonical example of an ADT is a stack with related operators new, push, pop and empty.

In our case, the user is given an ADT curve; each curve is represented internally as an arc tree, but this fact is completely transparent to the user. The operators defined on curves are given in table 2. Internally, all of these operators can be implemented in a high level programming language such as LISP or C++. Because the nodes of the arc trees are accessed along the parent-child pointers of the tree, it will be useful to store nodes near their parent nodes.

| operator | operand-1 | operand-2 | result |
| :--- | :--- | :--- | :--- |
| approximation | curve | integer | curve |
| point inclusion test | curve | point | boolean |
| curve-curve intersection detection | curve | curve | boolean |
| curve-curve intersection computation | curve | curve | set of points |
| curve-area intersection detection | curve | (closed) curve | boolean |
| curve-area intersection computation | curve | (closed) curve | set of curves |
| area-area intersection detection | (closed) curve | (closed) curve | boolean |
| area-area intersection computation | (closed) curve | (closed) curve | set of (closed) curves |
| area-area union computation | (closed) curve | (closed) curve | set of (closed) curves |
| area-area difference computation | (closed) curve | (closed) curve | set of (closed) curves |

Table 2: The curve ADT.

Note that it is not necessary to define a separate data type for closed curves. Each operator that requires the input curves to be closed may just extend its type checking by a test for closedness. Operators that return sets may just be implemented as relation-valued operators (such as the common retrieve command that may return
relations as well as single tuples).

## 7. Summary and Conclusions

We presented the arc tree, a balanced binary tree that serves as an approximation scheme for curves. It is shown how the arc tree can be used to represent curves for efficient support of common set and search operators. The arc tree can be viewed as just one instance of a large class of approximation schemes that implement some hierarchy of detail. We gave an overview of several other approximation schemes that are based on the same idea, and indicated how to modify the arc tree algorithms to work with these schemes.

Several examples are given for the performance of our algorithms to compute set and search operators such as point inclusion or area-area intersection detection and computation. The results of the practical analysis are encouraging: in most cases, the computation of boolean operators such as point inclusion or intersection detection can be completed on the first four or five levels of the tree. Also, the computation of non-boolean operators such as intersection computation gives fairly good results even if one restricts the computation to the first few levels. Finally, it is described how to embed the arc tree as an abstract data type into an extended database system. It is subject of future research to conduct a more comprehensive and systematic study of these arc tree algorithms. Also, we are planning to conduct a theoretical analysis of the arc tree, and to compare the arc tree to Ballard's strip tree and Bezier curves, both theoretically and practically.

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[^0]:    * A point set is simple if it is continuous, closed and not self-intersecting. In two or more dimensions this means in particular that it has no holes.

