

# Indexing For Function Approximation

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## ABSTRACT

Simulation is one of the most powerful tools that scientists have at their disposal for studying and understanding real-world physical phenomena. In order to be realistic, the mathematical models which drive simulations are often very complex and run for a very large number of simulation steps. The required computational resources often make it infeasible to evaluate simulation models exactly at each step, and thus scientists trade accuracy for reduced simulation cost.

In this paper, we explore function approximation for a combustion simulation. In particular, we model high-dimensional function approximation (HFA) as a storage and retrieval problem, and we show that HFA defines a novel class of applications for high dimensional index structures. The interesting property of HFA is that it imposes a mixed query/update workload on the index which leads to novel tradeoffs between the efficiency of search versus updates. We investigate in detail one specific approach to HFA based on Taylor Series expansions and we analyze tradeoffs in index structure design through a thorough experimental study.

## 1. INTRODUCTION

Studying physical phenomena through computer simulation is an important method of scientific research. Application areas include studies of heat and mass transfer, fluid dynamics, combustion, evaporation and many more [1, 2]. The general methodology in these application areas is similar. Scientists first understand the physical laws that govern the observed phenomenon. These laws then drive a mathematical model that is used in simulations as an approximation of reality.

In practice scientists often face serious computational challenges. The more realistic the model, the more complex the corresponding mathematical equations. As an example, consider the simulation of a combustion process [25], the application that brought our group together. Simulation of combustion requires tracking the composition of gases in a combustion chamber and the change in their compositions

over time. The transition from one composition to the next is defined by a complex high-dimensional *transition function*. Depending on the gases studied, a composition can be described using nine dimensions for a simple Hydrogen simulation and fifty or more dimensions for a Methane simulation. A single transition step (which is an evaluation of this function) can require millions of floating point operations. On a modern processor with the optimized code of the domain scientists, a simple Hydrogen simulation of low dimensionality would run for a few hours, while a Methane simulation with higher dimensionality would require several weeks. Simulations need to be run for many different gases and different configurations for each gas. This problem is not specific to combustion simulation, but representative of a large class of scientific simulations that require repeated evaluations of computationally expensive functions that govern a physical process [14, 21, 22].

To trade accuracy for simulation time, the domain scientists do not expend the resources to evaluate the function for each transition step, but instead use cheaper *approximations*. The main approach to function approximation is to build a model  $\hat{f}$  of the function. Given a query point  $\mathbf{x}$  at which the function  $f$  must be evaluated, function approximation techniques return  $\hat{f}(\mathbf{x})$  which approximates the true value  $f(\mathbf{x})$  within a specified error tolerance. There are two main types of function approximations. Global approximation techniques use a single model to represent  $f$ . Local approximation techniques break the domain into regions, representing each region with a different model. It has been shown that local approximations work better for the class of simulations that are the focus of this paper [14, 18].

A local function approximation scheme poses two main challenges. First, we need to decide how to select appropriate regions in the domain of the function, such that  $f$  can be approximated well within each region by a model  $\hat{f}$ . Approaches based on the Taylor Series are generally accepted for approximating high-dimensional functions in this domain [20, 22, 25]. The second challenge is to efficiently store the regions such that given a query point  $\mathbf{x}$ , we can efficiently find the region responsible for  $\mathbf{x}$  in order to calculate  $\hat{f}(\mathbf{x})$ . It is this storage and retrieval part that we think the database community can make significant contributions to. The domain scientists already took the first steps by developing the ISAT method [22] for indexing of regions. In collaboration with them, we are now studying the general problem of high-dimensional function approximation (HFA) and the design of efficient index structures for it. The workload imposed on indexing structures by HFA is very different

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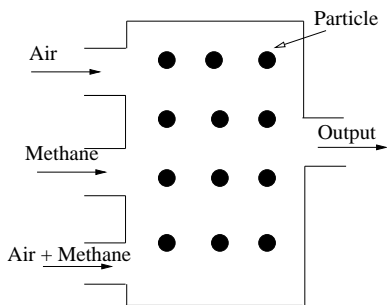


Figure 1: Combustion chamber

from the workloads studied so far in the literature [5, 10].

In summary, this paper makes the following contributions:

- We introduce the novel problem of high-dimensional function approximation as an application of high-dimensional indexing, and we describe an algorithmic framework that abstracts the salient elements of an HFA application (Sections 2 and 3).
- We give an analysis of the index design tradeoffs that this problem poses, and we identify their effects on the overall performance of HFA (Section 4).
- We perform a thorough evaluation of a set of candidate index structures from the database community for this application (Section 5).

Section 6 discusses related work and Section 7 concludes the paper.

We would like to emphasize that this paper is just the beginning of an exciting new direction of research into high-dimensional indexing, and it is this connection between an active application area and the database community that we believe is one of the major contributions of this paper.

## 2. PROBLEM FORMULATION

We first introduce the basic problem of high-dimensional function approximation and then show in Section 3 how it leads to a challenging indexing problem. We start with our case study, an example of a typical scientific application that uses HFA to improve the running time of simulations. We then formally define the resulting HFA problem.

### 2.1 Simulating Combustion

The application simulates the combustion of a hydrocarbon in a reaction chamber. The chamber has three inflows (air, the hydrocarbon being studied — Methane in the diagram, and a mixture of air and hydrocarbon) and a single outflow (see Figure 1). The gases flow into the chamber at different rates which are input parameters to the simulation. The simulation starts with a user-specified number of particles in the chamber. Each particle  $p$  has a user-specified chemical composition, which is described by its thermochemical composition vector  $\phi^p(t) = \langle Y_1^p(t), Y_2^p(t), \dots, Y_s^p(t), h \rangle$ , where  $s$  is the number of chemical species in  $p$ ,  $Y_i^p(t)$  is the mass fraction of chemical species  $i$  in particle  $p$  at time  $t$  of the simulation, and  $h$  (which is a constant) is the enthalpy of the particle [25].

Each simulation step consists of the following three phases: **(1) Inflow-Outflow.** Some of the particles in the reactor leave through the outflow and the same number of new particles enter from the inflows in ratios proportional to

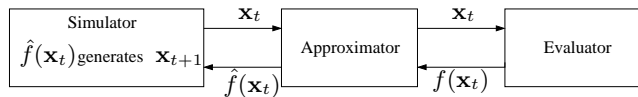


Figure 2: Application model

the rates of the inflow. **(2) Mixing.** Particles in the reactor mix with each other, generating new particle compositions. **(3) Reaction.** The particle compositions evolve due to reaction and the new compositions must be calculated for all particles.

The computationally expensive part is the reaction step, which for a particle  $p$  is described by a reaction function  $f$  that maps one thermochemical composition to the next composition. Moreover, typical simulations require  $10^8$  to  $10^{10}$  reaction function evaluations. These factors can cause simulations to run for years if the function value is calculated at each step. Thus in practice, the domain scientists accept approximations to  $f$  in order to be able to run large, complex simulations.

### 2.2 Application Model

Combustion simulation is representative of the class of applications that our methods apply to in general. Figure 2 shows the general framework. There is an application simulating some mathematical model, which we call the simulator. The simulator generates query points at which the value of a function  $f$  is required. These function values are used by the simulator to generate future query points. The application queries a function approximator for the function values. The approximator can calculate the exact value<sup>1</sup> using the function evaluator, which is an expensive operation, or it can use some algorithm to return approximate values within a user specified error tolerance. Typically, the approximator has limited knowledge about  $f$ , e.g., only previously calculated function values.

### 2.3 Problem Definition

Let us now define the function approximation problem for the above application model. We start by defining an  $\varepsilon$ -approximation of a function value.

**DEFINITION 1.** Let  $f : \mathbb{R}^m \rightarrow \mathbb{R}^n$  be a function, let  $\mathbf{x} \in \mathbb{R}^m$  and let  $\varepsilon \in \mathbb{R}$ . We say that  $\hat{f}(\mathbf{x}) \in \mathbb{R}^n$  is an  $\varepsilon$ -approximation of  $f(\mathbf{x})$  at  $\mathbf{x}$  if  $\|\hat{f}(\mathbf{x}) - f(\mathbf{x})\| < \varepsilon$ .

We can now formalize the function approximation problem as a game between two players, the simulator (application) and the function approximator. In the first round of the game, the simulator produces query point  $\mathbf{x}_1$ , and the function approximator computes  $\hat{f}(\mathbf{x}_1)$  at computational cost  $c_1$ , where  $\hat{f}$  is an  $\varepsilon$ -approximation to  $f$ . In the next round, the simulator takes  $\hat{f}(\mathbf{x}_1)$  and computes  $\mathbf{x}_2$ , the function approximator generates  $\hat{f}(\mathbf{x}_2)$  at cost  $c_2$  and so on. Note that in general  $\mathbf{x}_{i+1}$ , among other things depends on  $\hat{f}(\mathbf{x}_i)$ ,  $i = 1, \dots, n - 1$ . The game stops after  $n$  rounds. The goal of function approximation is, for a given  $\varepsilon$ , to minimize the total cost,  $\min \sum_{i=1}^n c_i$ .

In this paper we study approximators that attempt to minimize the total cost by partitioning the input domain

<sup>1</sup>Up to the accuracy of the evaluator. In practice the evaluator is a differential equation solver that introduces some error as well.

into local regions and modeling each region with some  $\hat{f}$ . The motivation is that the local regions enable cheaper approximate computation. We make this notion of local region more formal in the next definition.

**DEFINITION 2.** An  $\varepsilon$ -Local Region  $R_{f,\hat{f}}(\mathbf{x},\varepsilon) \subseteq \mathbb{R}^m$  for function  $f$  based on approximation  $\hat{f}$  at point  $\mathbf{x}$  is a maximal connected region containing  $\mathbf{x} \in \mathbb{R}^m$  such that  $\forall \mathbf{x}' \in R_{f,\hat{f}}(\mathbf{x},\varepsilon) : \hat{f}(\mathbf{x}')$  is an  $\varepsilon$ -approximation of  $f(\mathbf{x}')$ .

As a shortcut, we will often refer to an  $\varepsilon$ -Local Region  $R_{f,\hat{f}}(\mathbf{x},\varepsilon)$  for function  $f$  based on approximation  $\hat{f}$  at point  $\mathbf{x}$  as *Local Region* when the parameters are clear from the context.

There is a cost associated with finding the Local Region around a point. At the very least a function evaluation is required for the “center” of the region, together with additional computation for determining the extent of the region. Assuming that this cost is approximately the same for all regions, we can minimize total function approximation cost by finding the smallest set of Local Regions that covers all query points. The hardness of this problem is analyzed in the next section.

## 2.4 Analysis

In this section we show that the function approximation problem is hard. This applies to both its offline and its online formulation. We show hardness for an easier version of the problem, where the Local Region of  $\mathbf{x}$  is obtained for free when  $f(\mathbf{x})$  is computed. The more general case, i.e., where determining the extent of the Local Region around  $\mathbf{x}$  has some cost assigned to it as well, is therefore at least as hard.

**Offline problem:** Given a set  $X = \{\mathbf{x}_1, \dots, \mathbf{x}_n\}$  of query points, find the smallest set  $L = \{l_1, \dots, l_k\}$  of Local Regions in the data space (not limited to Local Regions around the query points), such that for each  $\mathbf{x}_i \in X$  there is an  $l_j \in L$ , which contains  $\mathbf{x}_i$ .

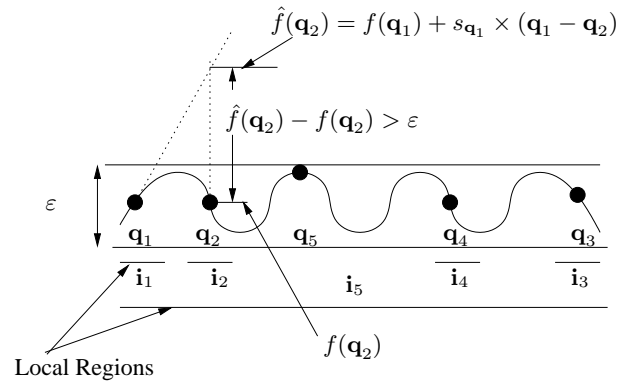
If the Local Regions are constrained to be hyper-spheres, we can show by reduction from Geometric Covering By Discs [16], that the offline problem is *NP-complete*. This implies that the more general formulation above is at least as hard.

Using a similar reduction, we can show the same hardness results even for a restricted version of the offline problem. In this restricted version, we constrain  $L$  to be a subset of  $\{R_{f,\hat{f}}(x_1,\varepsilon), \dots, R_{f,\hat{f}}(x_n,\varepsilon)\}$ , i.e., we can only choose from the Local Regions around query points.

In practice the algorithm for function approximation does not know the query points in advance. It has to solve an *online* problem, where query points are presented one-by-one.

**Online problem:** For  $i > 0$  let  $X(i) = \{\mathbf{x}_1, \dots, \mathbf{x}_i\}$  and  $L(i) = \{l_1, \dots, l_{k(i)}\}$ , where  $k(i)$  is some integer with  $k(i-1) \leq k(i)$  for all  $i > 1$ . Find the smallest set  $L(n)$ , such that the following holds for each set  $X(i)$ : Each  $\mathbf{x} \in X(i)$  is contained in some Local Region  $l \in L(i)$ .

Intuitively the set  $X(i)$  contains the query points seen until time  $i$ , and  $L(i)$  contains the Local Regions that have been materialized until time  $i$ . To be able to compute the function for query point  $\mathbf{x}_i$ ,  $\mathbf{x}_i$  has to be contained in one of the Local Regions that are available at time  $i$ . If no such Local Region exists, it has to be inserted into  $L(i)$ .



**Figure 3: Example for non-competitiveness**

A standard performance measure for online algorithms is the *competitive ratio* [6]. It measures the cost of the online algorithm for an input sequence  $(\mathbf{x}_1, \dots, \mathbf{x}_n)$  relative to an optimal offline algorithm, which knows the whole input in advance. The competitive ratio is defined as the worst (i.e., highest) ratio over all possible inputs of length  $n$ . If this ratio is independent of  $n$ , i.e., bounded by a constant  $c$ , then the online algorithm is  $c$ -competitive. E.g., if the online algorithm never materializes more than twice as many Local Regions as the optimal offline algorithm, then it is 2-competitive. Notice that it is often possible to analyze competitiveness even without knowing the optimal algorithm [6].

We can show that *there exists no deterministic online algorithm that is competitive*. Due to space constraints we only sketch the proof. The adversary can always construct a function (even if we are restricted to smooth functions), such that the online algorithm has to materialize a new Local Region for every query  $\mathbf{x}_i$ , while the offline algorithm can pick a single Local Region that contains all query points.

Figure 3 illustrates the construction. The figure shows a one-dimensional function where linear interpolations are used as  $\hat{f}$ , i.e.,  $\hat{f}$  for a Local Region is  $\hat{f}(\mathbf{a}) = f(\mathbf{x}) + s_{\mathbf{x}} \times (\mathbf{x} - \mathbf{a})$ , where  $s_{\mathbf{x}}$  is the derivative of  $f$  at  $\mathbf{x}$ . Note that the Local Regions in this case are intervals. For  $q_1, \dots, q_4$ , the Local Regions around them do not contain any other  $q_i$ , but the Local Region of  $q_5$  covers the whole domain. Since the online algorithm is deterministic, the adversary can always select a function such that all Local Regions computed by the online algorithm are around points like  $q_1, \dots, q_4$ , while the offline algorithm can choose an optimal point like  $q_5$ . This construction even works for the more restricted online problem, where materialized Local Regions have to be selected from the Local Regions defined by the query points. The adversary constructs the same function and input; it simply selects a point like  $q_5$  as the *last* query point.

The proof of non-competitiveness might appear contrived. However, without knowing the nature of the function it is impossible to rule out the case that there are “large” Local Regions that contain many “small” Local Regions. We are currently exploring what properties of a function could be taken advantage of to obtain better competitiveness.

## 3. AN ALGORITHMIC FRAMEWORK

In this section we introduce an algorithmic framework for the problem, which highlights the indexing problem in function approximation. Since the results from our analy-

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**Algorithm 1** : Framework

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**Require:** Query Point  $\mathbf{x}$ , index structure  $S$ 

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1: if  $\exists \langle R_{f,\hat{f}}(\mathbf{x}', \varepsilon), \hat{f} \rangle \in S$  such that  $\mathbf{x} \in R_{f,\hat{f}}(\mathbf{x}', \varepsilon)$  then
2:   Compute  $\mathbf{y} = \hat{f}(\mathbf{x})$ 
3: else
4:   Compute  $\mathbf{y} = f(\mathbf{x})$ 
5:   Update( $S, \mathbf{x}, f(\mathbf{x})$ )
6: end if
7: return  $\mathbf{y}$ 

```

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**Algorithm 2** : Update

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**Require:**  $S, \mathbf{x}, f(\mathbf{x})$ 

```

1: Add new  $\langle R_{f,\hat{f}}(\mathbf{x}, \varepsilon), \hat{f} \rangle$  to  $S$ 

```

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sis in 2.4 are discouraging, we start with algorithms that we abstracted from the approach of the domain scientists: A greedy heuristic (Section 3.1) and its refinements (Section 3.2). We then summarize the instantiation of choice of this framework by the domain scientists (Section 3.3).

### 3.1 A Greedy Heuristic

Since the problem is hard and there is no hope for a competitive online algorithm, we use a simple greedy strategy as shown in Algorithm 1. The algorithm maintains an index structure  $S$ , which contains the Local Regions around previously evaluated query points. Given a new query point  $\mathbf{x}$ , it first tries to find a Local Region that contains  $\mathbf{x}$  (Lines 1-2). If the point lies in some Local Region, an approximate value of  $f(\mathbf{x})$  is calculated and returned. If the point does not lie in any indexed region, then the algorithm has to compute  $f(\mathbf{x})$  (Line 4). It then updates the index based on the knowledge of  $f(\mathbf{x})$  as it belongs to a part of the domain that is yet to be indexed (Line 5). In the simple version of the algorithm, the update routine creates a new Local Region containing  $\mathbf{x}$  and inserts it into the index (Algorithm 2).

### 3.2 Practical Constraints

In practice, it is often impossible to accurately compute the Local Region ( $R_{f,\hat{f}}(\mathbf{x}, \varepsilon)$ ) around a point. We will see why this is the case in Section 3.3. Usually an initial (conservative) guess of the Local Region is first obtained and inserted into the index. We denote these approximate Local Regions as  $\hat{R}_{f,\hat{f}}(\mathbf{x}, \varepsilon)$ . As the simulation proceeds and larger portions of the domain are seen, better approximations of the existing Local Regions in  $S$  are obtained. This calls for a modified update operation in Algorithm 1. Algorithm 3 is an update stub which replaces Algorithm 2.

In our initial algorithm, if the function was evaluated because no existing region contained the query point, then a new Local Region was created and inserted into the index. The new update stub on the other hand first checks to see if the current query point can be part of any existing Local Region (Line 1). If such regions exist, then it finds the regions that can include  $\mathbf{x}$  and updates them (Lines 2-6). Finally, only if no existing region can include  $\mathbf{x}$ , then a new Local Region is initialized and inserted into  $S$  in Line 8. Updating existing Local Regions is usually more beneficial than adding new ones because it reduces the total number of Local Regions. We will also see later that updating an existing region is cheaper than creating a new one.

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**Algorithm 3** : Update

---

**Require:**  $S, \mathbf{x}, f(\mathbf{x})$ 

```

1: if  $\exists \langle \hat{R}_{f,\hat{f}}(\mathbf{x}', \varepsilon), \hat{f} \rangle \in S$  :  $\mathbf{x}$  can be included in  $\hat{R}_{f,\hat{f}}(\mathbf{x}', \varepsilon)$  then
2:   for all  $\langle \hat{R}_{f,\hat{f}}(\mathbf{x}', \varepsilon), \hat{f} \rangle \in S$  do
3:     if  $\mathbf{x}$  can be included in  $\hat{R}_{f,\hat{f}}(\mathbf{x}', \varepsilon)$  then
4:       Update  $\langle \hat{R}_{f,\hat{f}}(\mathbf{x}', \varepsilon), \hat{f} \rangle$  to include  $\mathbf{x}$ 
5:     end if
6:   end for
7: else
8:   Add new  $\langle \hat{R}_{f,\hat{f}}(\mathbf{x}, \varepsilon), \hat{f} \rangle$  to  $S$ 
9: end if

```

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## 3.3 An Instantiation

In practice finding a representation of the Local Regions of a function is not easy. In this section we review a method based on the Taylor Series [22]. This method, commonly used by scientists finds the Local Regions in two steps. It first creates an initial approximation, which is then refined over time.

### 3.3.1 Initializing Local Regions

Under fairly general conditions a function  $f(\mathbf{x}+\mathbf{a})$  can be expanded using the Taylor Series as

$$f(\mathbf{x}+\mathbf{a}) = \sum_{j=0}^k \left[ \frac{1}{j!} (\mathbf{a} \cdot \nabla_{\mathbf{x}})^j f(\mathbf{x}) \right] + \phi_k(\mathbf{x}, \mathbf{a}), \quad (1)$$

where  $\nabla_{\mathbf{x}}$  is the gradient  $[\frac{\partial}{\partial x_1} \frac{\partial}{\partial x_2} \dots \frac{\partial}{\partial x_m}]$  and the error  $\phi_k$  is  $O(|\mathbf{a}|^k)$ , i.e.,  $\lim_{h \downarrow 0} \frac{\phi_k(\mathbf{x}, h\mathbf{a})}{h|\mathbf{a}|^k} = 0$ .

The Taylor Series provides us with a simple mechanism for function approximation. Given the value of  $f$  at a point  $\mathbf{x}$ , the value at any point  $\mathbf{x}+\mathbf{a}$  “near”  $\mathbf{x}$  is usually approximated using the first few terms of the summation in Equation 1. For example, using the first term only, we get a constant approximation  $\hat{f}_{0,x}$  of  $f$  as follows:

$$f(\mathbf{x}+\mathbf{a}) \approx \hat{f}_{0,x}(\mathbf{x}+\mathbf{a}) \equiv f(\mathbf{x}) \quad (2)$$

Similarly, the first two terms give the following linear approximation  $\hat{f}_{1,x}$

$$f(\mathbf{x}+\mathbf{a}) \approx \hat{f}_{1,x}(\mathbf{x}+\mathbf{a}) \equiv f(\mathbf{x}) + (\mathbf{a} \cdot \nabla_{\mathbf{x}})f(\mathbf{x}) \quad (3)$$

The errors of the above approximations can be obtained from the remaining terms in the summation of Eq. 1, i.e., those terms not used in the approximation. For small values of  $|\mathbf{a}|$ , high-order terms in Eq. 1 are dominated by low-order terms and are therefore commonly ignored. The domain scientists only use the single lowest-order term to estimate the approximation error. More precisely, the approximation quality requirement is defined for the constant approximation as

$$\| (\mathbf{a} \cdot \nabla_{\mathbf{x}})f(\mathbf{x}) \| < \varepsilon, \quad (4)$$

and similarly for the linear approximation

$$\| \frac{1}{2!} (\mathbf{a} \cdot \nabla_{\mathbf{x}})^2 f(\mathbf{x}) \| < \varepsilon. \quad (5)$$

Equation 5 for a high dimensional function is the equation of a tensor and hence, except under special conditions, it is computationally infeasible to compute the Local Region

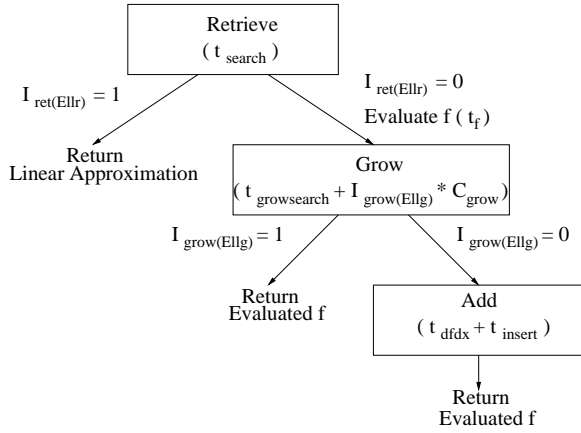


Figure 4: Block diagram

defined by it. However, it can be shown that Eq. 4 defines a hyper-ellipsoid around  $\mathbf{x}$  [22]. Therefore, using a constant approximation, the local region around  $\mathbf{x}$  is a hyper-ellipsoid.

### 3.3.2 Growing

The linear approximation is preferred over the constant approximation, because it tends to generate much larger Local Regions. Unfortunately, since the region defined by Eq. 5 is difficult to compute, we have to start out with the more conservative (and hence smaller) region defined by Eq. 4. Since we know that the true Local Region is much larger, we use the grow operation to extend the initial region over time as follows. Consider a query point  $\mathbf{x}$  and an ellipsoid  $e$  around it. Suppose, there exists another query point  $\mathbf{x}'$  such that  $\mathbf{x}'$  lies just outside  $e$  but  $\hat{f}_{1,x}(\mathbf{x}')$  is an  $\varepsilon$ -approximation. Then  $\mathbf{x}'$  is assumed to be part of the Local Region of  $\mathbf{x}$ ; therefore  $e$  is grown to a larger ellipsoid that contains  $\mathbf{x}'$ . This simple heuristic of growing has been found to work well in practice [20]. For applications with stricter error guarantees, growing can be further controlled by using domain specific information like the maximum allowable size of ellipsoids or it may even be turned off and other function specific methods may be used to find the true Local Regions.

### 3.3.3 Final Algorithm

Instantiating the framework with ellipsoids as the Local Regions is straightforward. The algorithm performs the following high level operations on a query point  $\mathbf{x}$ .

**Retrieve:** The algorithm first tries to find an ellipsoid that contains  $\mathbf{x}$  (Line 1 of Algorithm 1).

**Grow:** If the retrieve fails then the algorithm attempts to grow existing ellipsoids in the index to include  $\mathbf{x}$  (Lines 1-6 of Algorithm 3).

**Add:** If both the retrieve and the grow fail a new ellipsoid derived from the constant approximation is initialized and inserted into the index (Line 8 of Algorithm 3).

## 4. INDEXING PROBLEM

We now turn to the indexing problem in function approximation, which produces a challenging workload for the operations on index  $S$  in Algorithms 1, 2 and 3. The retrieve requires the index to support fast lookups. The grow requires

both a fast lookup to find growable ellipsoids and then an efficient update process once an ellipsoid is grown. Finally, an efficient insert operation is required for the add step. Also, past decisions about growing and adding affect future performance of the index, therefore the algorithm produces a query/update workload that is not common in traditional indexing applications.

A straightforward implementation of the algorithm introduced in the previous section would search in the index for an ellipsoid containing the query point until it finds an ellipsoid or has established that no such ellipsoid exists. The grow similarly would try to find all ellipsoids that can be grown and finally the add is performed if these operations fail. Our initial experiments showed that this implementation can result in poor performance.

This observation brings us to the most interesting aspect of the indexing problem in function approximation: It presents a very different framework in which indexes must be evaluated. Traditionally, the performance of index structures has been measured simply in terms of the cost of a search and in some cases update. There are two distinct cost factors in the function approximation problem. First, there are the costs associated with the search and update operations on the index. Second, there are costs of the function approximation application which include function evaluations and ellipsoid operations. Since, the goal of function approximation is to minimize the total cost of the simulation, all these costs must be accounted for when evaluating the performance of an index. We will see that a principled analysis leads to the discovery of novel tradeoffs. These tradeoffs produce significant and different effects on different index structures. This makes indexing for function approximation a challenging problem.

## 4.1 Costs

This section introduces a cost model for the function approximation algorithm. We use the cost model to qualitatively explore the tradeoffs in the indexing problem. The formulation of a quantitative cost model for optimization purposes has limited use. This is because the benefits from an operation can be determined only in the future after the operation has been done. As a simple example, consider the grow operation. The benefit from the grows cannot be estimated accurately until the actual grown ellipsoids are known, which requires the grow operation to be performed. Therefore, we do not explore a true quantitative cost model further in the paper. Table 1 is a summary of the commonly referenced variables in this section and the next.

The total cost ( $C_{\text{tot}}$ ) of processing a query  $\mathbf{q}$  can be expressed as

$$C_{\text{tot}} = t_{\text{search}} + I_{\text{ret}} \times t_{\text{la}} + (1 - I_{\text{ret}}) \times C_{\text{miss}} \quad (6)$$

Note that the the costs in Eq. 6 vary from one query to the next. However, the dependence on  $\mathbf{q}$  has been dropped for ease of notation.  $t_{\text{search}}$  is the cost of searching the index for an ellipsoid containing  $\mathbf{q}$ . This is essentially the cost of the retrieve step.  $I_{\text{ret}}$  is an indicator which is 1 if there exists an ellipsoid containing  $\mathbf{q}$  in the index and 0 otherwise.  $t_{\text{la}}$  is the cost of calculating the linear approximation on a successful retrieve.  $C_{\text{miss}}$  is the cost incurred if  $\mathbf{q}$  does not result in a retrieve operation. A miss results in either a grow or an add.  $C_{\text{miss}}$  can therefore be written as

$$C_{\text{miss}} = t_f + t_{\text{growsearch}} + I_{\text{grow}} \times C_{\text{grow}} + (1 - I_{\text{grow}}) \times C_{\text{add}} \quad (7)$$

Name	Description
$C_{\text{tot}}$	Total cost of a query
$C_{\text{miss}}$	Total cost of a miss
$C_{\text{grow}}$	Total cost of a grow
$C_{\text{add}}$	Total cost of an add
$I_{\text{ret}}$	Indicator variable for successful retrieve
$I_{\text{grow}}$	Indicator variable for successful grow
$t_{\text{search}}$	Index search cost during Retrieve
$t_f$	Cost of a function evaluation
$t_{\text{la}}$	Cost of a linear approximation
$t_{\text{growsearch}}$	Index search cost during grow
$t_{\text{inellipsoid}}$	Cost of checking if ellipsoid contains point
$t_{\text{grow}}$	Cost to grow ellipsoid
$t_{\text{update}}$	Index update cost
$t_{\text{dfdx}}$	Cost of a derivative evaluation
$t_{\text{insert}}$	Index insertion cost
$\text{Ellr}$	Max. number ellipsoids examined for Retrieve
$\text{Ellg}$	Max. number ellipsoids examined for Grow
$N_{\text{fpos}}$	Number of false positives during Retrieve
$N_{\text{found}}$	Number of growable ellipsoids found
$N_{\text{growmax}}$	Max. number of grows allowed
$N_{\text{grown}}$	Number of ellipsoids grown

**Table 1: Index of variables**

Ellipsoid Operation	Cost
$t_f$	2000
$t_{\text{dfdx}}$	1200
$t_{\text{grow}}$	10
$t_{\text{la}}$	1
$t_{\text{inellipsoid}}$	1
Usual search cost	1

**Table 2: Relative costs of the ellipsoid operations**

The cost of a grow comprises two parts.  $t_{\text{growsearch}}$  is the cost associated with searching for growable ellipsoids.  $C_{\text{grow}}$  is the cost of actually growing the ellipsoids and updating the index.  $I_{\text{grow}}$  is an indicator which is 1 if there is some ellipsoid in the index that can be grown and 0 otherwise. Finally, if no ellipsoids were grown, an ellipsoid is added at a cost of  $C_{\text{add}}$ .

Table 2 describes typical ratios between costs of the application in the combustion problem, which is a typical example of an application that our methods apply to. The table also lists the commonly observed cost of searching for an ellipsoid in an index. It is important to note that for most indexes the costs of the application are more expensive than the index operations.

## 4.2 Effects And Tradeoffs

In the previous section we outlined the cost of processing a query. Here we will analyze the different components of  $C_{\text{tot}}$ . Figure 4 displays the cost associated with each high-level operation.

**Retrieve:** The first component of  $C_{\text{tot}}$  is  $t_{\text{search}}$ . In most high dimensional index structures the ellipsoid containing a query point is usually not the first ellipsoid found. The index ends up looking at a number of ellipsoids before finding “the right one” (Section 5.1). The additional ellipsoids that are examined by the index are called *false positives*, the number of which is denoted by  $N_{\text{fpos}}$ . Taking  $N_{\text{fpos}}$  into account, we can rewrite  $t_{\text{search}}$  as

$$t_{\text{search}} = (N_{\text{fpos}} + 1) \times (t_r + t_{\text{inellipsoid}}).$$

For each false positive the algorithm pays to search and retrieve the ellipsoid from the index ( $t_r$ ) and to check if the ellipsoid contains the query point ( $t_{\text{inellipsoid}}$ ). In practice using an iterator for the search could lead to different values of  $t_r$  for each false positive. However, this is not important for this qualitative study and hence we ignore such effects to keep the analysis simple.

In traditional indexing problems, if an object that satisfies the query condition exists in the index, then finding this object during search is mandatory. Therefore,  $N_{\text{fpos}}$  is a fixed property of the index. However, the function approximation problem provides the flexibility to tune  $N_{\text{fpos}}$ , because we can evaluate the function if the index search was not successful. The number of false positives can be tuned by limiting the number of ellipsoids examined during the retrieve step. We denote this parameter by  $\text{Ellr}$ .  $\text{Ellr}$  places an upper bound on the number of false positives for a query. Taking this parameter into account, the total cost of processing a query can be rewritten as

$$C_{\text{tot}} = t_{\text{search}} + I_{\text{ret}(\text{Ellr})} \times t_{\text{la}} + (1 - I_{\text{ret}(\text{Ellr})}) \times C_{\text{miss}} \quad (8)$$

$I_{\text{ret}(\text{Ellr})}$  is 1 if an ellipsoid containing the query point is found by the index *before Ellr ellipsoids are examined*, and 0 otherwise. Notice that for a given query,  $I_{\text{ret}} \geq I_{\text{ret}(\text{Ellr})}$ . Tuning  $\text{Ellr}$  introduces the following tradeoffs:

**Effect 1:** Decreasing  $\text{Ellr}$  restricts the number of ellipsoids examined during the retrieve for a given query. This effectively reduces the number of false positives, therefore decreasing  $t_{\text{search}}$ .

**Effect 2:** Decreasing  $\text{Ellr}$  decreases the probability that  $I_{\text{ret}(\text{Ellr})} = 1$  for a given query  $\mathbf{q}$ , thereby lowering the probability of a query resulting in a retrieve. This is because there might be an ellipsoid containing  $\mathbf{q}$  in the index but it is not found before  $\text{Ellr}$  ellipsoids are examined. Reducing the probability of retrieve increases the probability of an expensive miss operation.

**Effect 3:** The previous tradeoffs demonstrated the effects of decreasing  $\text{Ellr}$  on the probability of a retrieve for a given query. There is another tradeoff unique to the problem. Misses that result from decreasing  $\text{Ellr}$  can grow and add ellipsoids. These grows and adds index new parts of the domain and also change the overall structure of the index. Both of these affect the probability of retrieves for future queries.

**Grow:** The next major cost component is the cost of the grow operation:  $t_{\text{growsearch}} + I_{\text{grow}} \times C_{\text{grow}}$ . In the first part of the grow process the index is traversed to find ellipsoids that can be grown. Every ellipsoid in the index is a candidate for growing. Checking an ellipsoid for growing involves a cost of retrieving it from the index ( $t_{\text{rg}}$ ) and then checking to see if the ellipsoid can be grown ( $t_{\text{la}}$ ). Therefore, most indexes prune the search space for finding growable ellipsoids using some domain information or heuristic. Just like the retrieve operation, the cost incurred in searching for growable ellipsoids can be tuned by restricting the number of ellipsoids examined for growing ( $\text{Ellg}$ ). Taking this parameter into account,  $C_{\text{miss}}$  can be rewritten as  $C_{\text{miss}} =$

$$t_f + t_{\text{growsearch}} + I_{\text{grow}(\text{Ellg})} \times C_{\text{grow}} + (1 - I_{\text{grow}(\text{Ellg})}) \times C_{\text{add}}.$$

$I_{\text{grow}(\text{Ellg})}$  is 1 if at least one growable ellipsoid is found before  $\text{Ellg}$  ellipsoids are examined, and 0 otherwise.

The cost of searching for growable ellipsoids can be writ-

ten as

$$t_{\text{growsearch}} = \text{Ellg} \times (t_{\text{rg}} + t_{\text{la}}).$$

Tuning Ellg introduces the following tradeoffs:

**Effect 4:** Decreasing Ellg decreases  $t_{\text{growsearch}}$  because fewer ellipsoids are examined for growing.

**Effect 5:** Decreasing Ellg decreases the number of ellipsoids examined for growing and hence the number of growable ellipsoids that are found ( $N_{\text{found}}$ ). Therefore, restricting Ellg also limits  $N_{\text{grown}}$  (the number of ellipsoids that are finally grown). The effects associated with  $N_{\text{grown}}$  are described in Effects 6 and 7.

After finding the ellipsoids that can be grown, the next step of the grow operation is to actually grow the ellipsoids, which costs  $C_{\text{grow}}$ . There is another tuning parameter,  $N_{\text{growmax}}$ , which represents the maximum number of ellipsoids that are allowed to be grown during a grow operation. Hence the number of ellipsoids that are actually grown,  $N_{\text{grown}}$ , is only  $\min\{N_{\text{found}}, N_{\text{growmax}}\}$ . For each ellipsoid that is grown during the grow step, the algorithm incurs a cost to grow the ellipsoid ( $t_{\text{grow}}$ ) and update the index ( $t_{\text{update}}$ ).

$$C_{\text{grow}} = N_{\text{grown}} \times (t_{\text{grow}} + t_{\text{update}})$$

$N_{\text{grown}}$  has the following effects on the cost of the algorithm:

**Effect 6:** Lower values of  $N_{\text{grown}}$  decreases  $C_{\text{grow}}$ . This can affect the simulation time significantly because  $t_{\text{grow}}$  and  $t_{\text{update}}$  can be expensive.

**Effect 7:** Larger values of  $N_{\text{grown}}$  increase the fraction of the domain that is covered by ellipsoids. Therefore,  $I_{\text{ret}}$  changes from 0 to 1 for future query points that lie in the newly covered part of the domain.

**Effect 8:** For  $N_{\text{grown}} > 1$  the false positive rate of the index can increase, because the grown ellipsoids overlap (they all cover the new query point). This in turn might negatively affect  $I_{\text{ret}(\text{Ellr})}$ . The reason for this is that the higher false positive rate can result in failed retrieves due to the search limit imposed by Ellr.

**Add:** The last cost component is the cost of adding an ellipsoid. It includes the cost of finding the derivative of the function ( $t_{\text{dfox}}$ ), which is costly (see Table 2), and inserting the new ellipsoid into the index ( $t_{\text{insert}}$ ). The derivative is needed to estimate the initial ellipsoid and for computing the linear approximation [22]. A new ellipsoid is added only if the retrieve and grow both fail. Therefore there is no direct way of controlling the number of adds.

**Effect 9:** Lowering the effort spent on the retrieve and grow can cause the number of add operations to increase. This can be undesirable because the add operation is expensive and a newly added ellipsoid is a conservative approximation of a Local Region. Adds also increase the index size.

In summary, the algorithm provides us with a set of tunable parameters, namely the number of ellipsoids examined during retrieve (Ellr), the number of ellipsoids examined during grow (Ellg), and the maximum number of ellipsoids allowed to be grown ( $N_{\text{growmax}}$ ). Each parameter can have different effects on  $C_{\text{tot}}$ . What makes the problem interesting is that these effects often move in opposite directions. Moreover, tuning affects indexes differently and to varying degrees, which makes it necessary to analyze each index individually. In the next section we will demonstrate the effects of these

parameters on the performance of different index structures when used in the function approximation algorithm.

## 5. EXPERIMENTS

In the previous section we introduced the tuning parameters of the function approximation problem and we identified nine qualitative effects these parameters could have on the runtime. In this section we study the corresponding tradeoffs for a concrete instance of the problem and different index structures.

All experimental results are for a Methane combustion simulation. In the simulation, the number of species was set to  $s = 31$ , i.e., the thermochemical composition vector has 32 dimensions. There are 100 particles in the combustion chamber; at each time step a single particle enters the reaction step. The simulation was run for  $6 \times 10^6$  timesteps, thereby generating  $6 \times 10^6$  query points for function evaluation. The error tolerance, unless otherwise noted, was set to  $\varepsilon = 5 \times 10^{-5}$ . All reported measurements are wall-clock time for an execution on a Windows XP machine with a 2.4Ghz processor and 2GB of memory.

### 5.1 Candidate Index Structures

Our goal in this paper is *not* to find the best index for function approximation. In fact because of the diversity of function approximation applications in terms of their cost structure, dimensionality, and locality of access, the existence of a single best index is unlikely. For this reason we selected a very diverse set of indexes, without attempting a comparison of all existing ones. Another criterion for selection for this initial study was to pick only well understood index structures with predictable behavior, rather than highly optimized and complex indexes.

We chose a candidate from each of the different classes of commonly used indexes, namely linear scan, spatial partitioning, balanced index for points and balanced index for extended objects. In the simulations we studied, the indexes were small enough to fit in main memory, therefore we limited our attention to in-memory performance. Extending our experiments to other indexes and I/O performance is part of our future work.

**Bounding Box Rtree (Bbox Rtree):** An obvious choice for function approximation is an index that can manage the Local Regions, i.e., the ellipsoids in our case. The most well-known data structure with this functionality is the Rtree, a balanced multidimensional generalization of the B-tree which can handle both point and hyper-rectangular objects. There exists a large number of Rtree-variants (see Section 6); as a representative we selected the robust R\*-tree [3]. Bbox Rtree indexes the axis-parallel minimum bounding boxes of the ellipsoids, using the standard R\*-tree algorithms. The retrieve operation finds leaf objects (minimum bounding boxes of ellipsoids) that contain the query point. Then it needs to verify that the corresponding ellipsoid also contains the query point. Growing of ellipsoids is implemented by a deletion, followed by an insertion. Growable ellipsoids are found by performing a nearest-neighbor (NN) search on the bounding boxes.

**Point Rtree:** Managing objects with extent is far more challenging than handling points [10]. We can map our problem to a point-indexing problem by only indexing the center points of the ellipsoids in an Rtree. The Point Rtree does not have to deal with overlapping leaf objects (bounding

boxes of *inner* nodes can still overlap) and growing does not require an index update, because the center of an ellipsoid is not modified by the grow operation. Unfortunately, without any information about the dimensions of the ellipsoid, the index has no way of pruning search—as long as the ellipsoid is large enough, even a center point far away from the query could be relevant. Intuitively based on the Taylor Series, the Euclidean distance between query point and ellipsoid center should be correlated with the probability that the query point is within the ellipsoid. We therefore implemented both the retrieve and the search for growable ellipsoids as a NN-query.

**Binary Tree:** This is a binary space partitioning tree [10], which was introduced for the ISAT function approximation problem [22]. The Binary Tree indexes the centers of the ellipsoids by recursively partitioning the space with cutting planes. Leaf nodes of the tree correspond to ellipsoid centers and non-leaf nodes represent cutting planes. During the retrieve step, the index is traversed from the root by following the subtree corresponding to the side of the cutting plane that the query point lies on. Like the Point Rtree, the Binary Tree requires no update when an ellipsoid is grown, because it only indexes the ellipsoid center points. A more detailed discussion of the index can be found in Section 5.2.2.

**MRU List + Rtree:** For high dimensional data, it has been shown that a simple linear scan often outperforms any sophisticated indexing technique [27]. We therefore include a list-based data structure. This simple structure has the advantage that if there is locality of access, we can directly apply existing cache-replacement policies. The MRU List stores the ellipsoids ordered by their most recent access. The retrieve operation simply scans the list, starting with the most recently used object. To improve the search for growable ellipsoids, we index the ellipsoids with a “secondary” point Rtree. This tree is identical to the Point Rtree described above, but it is not used for the retrieve operation. Notice that the leaf objects also contain a pointer to the corresponding ellipsoid in the MRU list.

## 5.2 Tradeoffs: Detailed Analysis

We examined the tradeoffs for all candidate index structures. Due to space constraints, we only discuss the two best-performing indexes in detail and report the overall performance for the others. Somewhat to our surprise, the Binary Tree and the simple MRU List + Rtree have the best performance after tuning. If we do not tune any index, i.e., set  $\text{Ellr} = \text{Ellg} = N_{\text{growmax}} = \infty$ , the Bbox Rtree clearly outperforms the Binary Tree. However, the Binary Tree benefits much more from tuning. This will be discussed in more detail in this section.

For our experiments we do not examine the effect of the  $N_{\text{growmax}}$  parameter. Recall that we grow  $N_{\text{grown}} = \min\{N_{\text{found}}, N_{\text{growmax}}\}$  ellipsoids, i.e.,  $N_{\text{growmax}}$  limits the number of finally grown ellipsoids if there are too many growable candidates. However, we currently do not have any meaningful way of preferring one grow candidate over another, therefore we set  $N_{\text{growmax}} = \infty$ . Notice also that  $\text{Ellg}$  directly limits  $N_{\text{found}}$ , and therefore we can control  $N_{\text{grown}}$  through  $\text{Ellg}$ . Hence in our experiments we only study the effect of two tuning parameters:  $\text{Ellr}$  and  $\text{Ellg}$ .

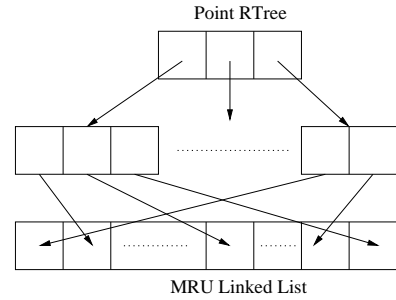


Figure 5: MRU List + Rtree

### 5.2.1 MRU List + Rtree

The MRU List + Rtree index (Figure 5) uses the list for the retrieve operations. The list contains pointers to the ellipsoids. During the retrieve step, it is scanned, starting with the most recently used ellipsoid. This continues until either an ellipsoid is found that contains the query point, or  $\text{Ellr}$  ellipsoids have been examined unsuccessfully. If a containing ellipsoid is found, it is moved to the front of the list.

As mentioned earlier, the Rtree improves the performance of the search for growable ellipsoids during the grow step. It indexes the ellipsoid centers (see Figure 5 for an illustration). Instead of scanning the list for growable ellipsoids, we perform a NN-search for the query point to find grow candidates sorted by the Euclidean distance to the ellipsoid centers. The search terminates once the  $\text{Ellg}$  nearest neighbors have been examined. All growable ellipsoids are grown. Since their center points do not change, we do not need to update the Rtree. The linked list is updated by moving all grown ellipsoids to the front.

An add operation adds the new ellipsoid to the front of the list. Its center point, together with a pointer to the ellipsoid object, is inserted into the Rtree.

We first examine the effect of tuning  $\text{Ellr}$ , the number of ellipsoids in the list that are examined during the retrieve step. For this experiment, we held the grow search limit constant at  $\text{Ellg} = 3000$ . Limiting the grow search to 3000 nearest neighbors was found to be enough to find all growable ellipsoids. We report the total number of retrieves (Figure 6) and the total cost of the simulation, also broken down into retrieve and miss (grow and add) cost (Figure 7). As we increase  $\text{Ellr}$ ,  $t_{\text{search}}$  increases in accordance with Effect 1 (Figure 7). At the same time the number of retrieves increases (Figure 6), because we are searching further, thereby reducing  $C_{\text{miss}}$  (Effect 2). As we increase  $\text{Ellr}$ , Effect 2 initially causes the simulation time to decrease. However, at some point further increasing  $\text{Ellr}$  will only add very few additional retrieves, hence the total number of retrieves asymptotes and the increased effort in searching does not pay off any more. At this point Effect 1 causes the overall simulation time to increase slightly.

Next we examine the effect of tuning  $\text{Ellg}$ , the number of nearest neighbors examined for growing. In this experiment there were no restrictions placed on  $\text{Ellr}$ . The Rtree examines ellipsoids for growing in nearest neighbor order. Therefore, as we start to increase  $\text{Ellg}$ , larger ellipsoids are grown and the domain is indexed more aggressively. The number of retrieves increases (Figure 8) and the number



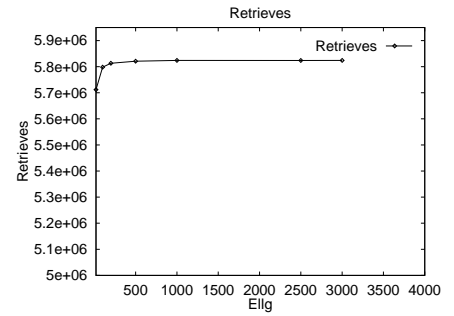
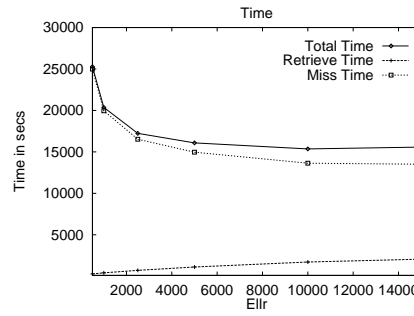
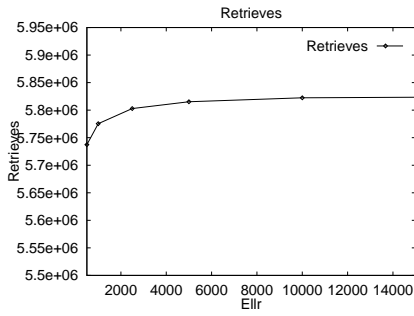


Figure 6: Number of retrieves vs. Ellr (MRU List + Rtree)

Figure 7: Time vs. Ellr (MRU List + Rtree)

Figure 8: Number of retrieves vs. Ellg (MRU List + Rtree)

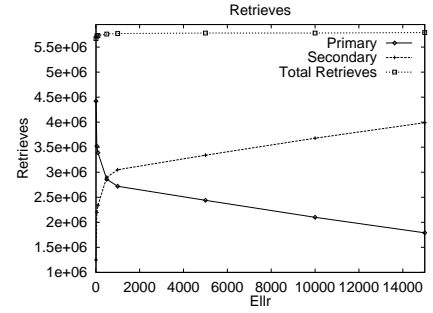
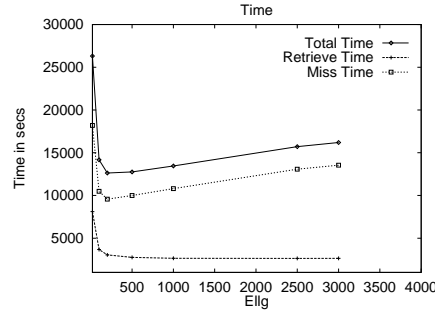
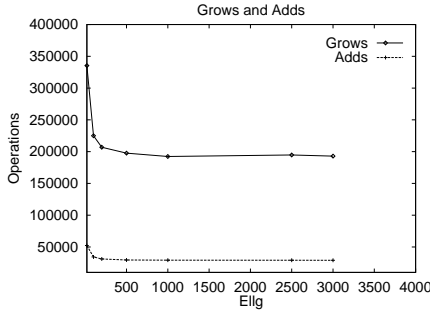


Figure 9: Number of grows, add vs. Ellg (MRU List + Rtree)

Figure 10: Time vs. Ellg (MRU List + Rtree)

Figure 11: Number of retrieves vs. Ellr (Binary Tree)

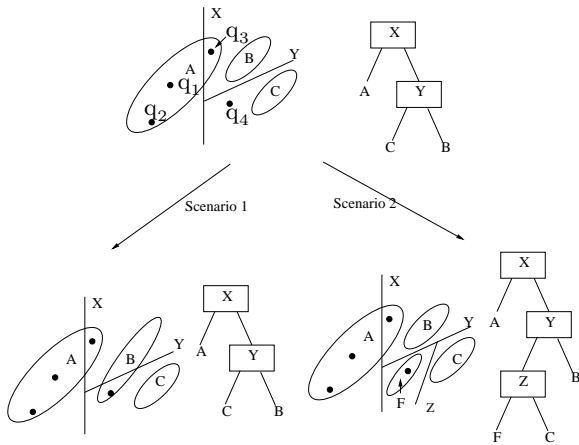


Figure 12: Binary Tree

of misses decreases (Figure 9). The total simulation time (Figure 10) therefore decreases (Effect 7). Increasing Ellg increases the chances of finding at least one growable ellipsoid, hence reduces the number of adds (Effect 9). Reduction in the number of adds causes the list size to decrease for larger values of Ellg. This accounts for the decrease in  $t_{\text{search}}$  as we start increasing Ellg. As we go on increasing Ellg, the number of retrieves asymptotes (Figure 8) because no additional ellipsoids are found for growing. There is no increase in retrieve time. However, the total simulation time then increases because  $t_{\text{growsearch}}$  increases with Ellg caused by Effect 4 (Figure 10).

### 5.2.2 Binary Tree

The Binary Tree partitions the data space recursively, using cutting planes [22]. It might be unbalanced, i.e., leaves can be at different depths. Figure 12 shows an example tree with three ellipsoids  $A, B, C$  and two cutting planes  $X$  and  $Y$ . For now we focus on the tree in the top half of the figure, together with the corresponding view of the data space showing the cutting planes and ellipsoids.

We illustrate the retrieve step with query point  $q_2$ . The retrieve starts at the root, checking on which side of hyperplane  $X$  the query point lies. The search continues recursively with the corresponding subtree, the left one in our example. When we reach a leaf node, we test if the ellipsoid in the leaf contains the query point. In the example,  $A$  contains  $q_2$ , therefore we have found a containing ellipsoid. This process, i.e., when the traversal from root to leaf is successful, will be denoted as a *Primary Retrieve* [22].

Notice that ellipsoids can straddle cutting planes, e.g.,  $A$  covers volume on both sides of cutting plane  $X$ . If ellipsoids are straddling planes, then the Primary Retrieve can result in a false negative. For example,  $q_3$  lies to the right of  $X$  and so the Primary Retrieve fails even though there exists an ellipsoid  $A$  containing it. To overcome this problem the Binary Tree performs a *Secondary Retrieve* if the Primary fails. The main idea of the Secondary Retrieve is to explore the “neighborhood” around the query point by examining “nearby” subtrees. In the case of  $q_3$ , the failed Primary Retrieve ended in leaf  $B$ . Nearby subtrees are explored by moving up a level in the tree and exploring the other side of the cutting plane. Specifically, we first examine  $C$  (after moving up to  $Y$ ,  $C$  is in the unexplored subtree). Then the search would continue with  $A$  (now moving up another level

to  $X$  and accessing the whole left subtree). This process continues until a containing ellipsoid is found, or Ellr ellipsoids have been examined unsuccessfully.

The search for growable ellipsoids proceeds in exactly the same way as a Secondary Retrieve, starting where the failed Primary Retrieve ended. Assume that in the example in Figure 12, ellipsoid  $B$  can be grown to include  $\mathbf{q}_4$ , but  $C$  and  $A$  cannot. After the retrieve failed, the grow operation first attempts to grow  $C$ . Then it continues to examine  $B$ , then  $A$  (unless  $\text{Ellg} < 3$ ).  $B$  is grown to include  $\mathbf{q}_4$ , as shown on the bottom left (Scenario 1). Growing of  $B$  made it straddle hyper-plane  $Y$ . Hence, for any future query point near  $\mathbf{q}_4$  and “below”  $Y$ , a Secondary Retrieve is necessary to find containing ellipsoid  $B$ , which is “above”  $Y$ .

The alternative to growing  $B$  is illustrated on the bottom right part of Figure 12 (Scenario 2). Assume  $\text{Ellg} = 1$ , i.e., after examining  $C$ , the grow search ends unsuccessfully. Now we add a new ellipsoid  $F$  with center  $\mathbf{q}_4$  to the index. This is done by replacing leaf  $C$  with an inner node, which stores the hyper-plane that best separates  $C$  and  $F$ . The add step requires the expensive computation of  $F$ , but it will enable future query points near  $\mathbf{q}_4$  to be found by a Primary Retrieve.

As we can see from this example, tuning parameter  $\text{Ellg}$  affects the Binary Tree in its choice of scenario 2 over 1. Furthermore, this choice, i.e., performing an add instead of a grow operation, reduces  $N_{\text{fpos}}$  for future queries, but adds extra-cost for the current query. The experiments show that this tradeoff has a profound influence on the overall simulation cost. We will also see that the effect of the tuning parameters is very different for the Binary Tree as compared to the MRU List + Rtree.

We first study the effect of varying  $\text{Ellr}$ , which limits the number of ellipsoids examined during the Secondary Retrieve phase. For this experiment we set  $\text{Ellg} = \infty$ . It can be seen from Figure 13 that as we increase  $\text{Ellr}$ ,  $t_{\text{search}}$  goes up (Effect 1). This increase in the retrieve time is accompanied by a reduction in miss time, which is caused by the improved total number of retrieves (hence fewer misses) due to the more aggressive Secondary Retrieves (Effect 2, see Figure 11).

One of the most interesting observations from Figure 13 is the super-linear increase in the time for successful retrieves, which starts dominating the total simulation time. Figure 11 reveals the explanation: As we increase  $\text{Ellr}$ , Secondary Retrieves (and hence also  $N_{\text{fpos}}$ ) are increasing, because we are searching the index more extensively. Therefore we are reducing the number of add operations, ultimately causing the Primary Retrieve rate to decrease (Effect 3). At the same time, the average cost of a Secondary Retrieve also increases, because the search proceeds further in the tree. These two effects together—increase in number of Secondary Retrieves and in average cost per Secondary Retrieve—create the super-linear trend of the retrieve time with increasing  $\text{Ellr}$ .

Lastly, we examine the effect of varying  $\text{Ellg}$ , the number of ellipsoids examined for growing, while setting  $\text{Ellr} = \infty$ . As we start increasing  $\text{Ellg}$ , because of Effect 7 the total number of retrieves increases slightly (Figure 14). Therefore there are fewer misses, which results in lower miss cost and better total simulation time (see Figure 15). Note the initial drop in retrieve time in Figure 15. The reason is that  $t_{\text{search}}$  includes the cost of *all* searches, including unsuccessful ones. A better retrieve rate therefore also reduces the total retrieve

**Table 3: Total simulation time in seconds**

Index Type	Value of $\varepsilon$		
	0.0005	0.00005	0.00004
Binary Tree (tuned)	1073	10181	13100
MRU List + Rtree	1125	14000	19920
Bbox Rtree	1201	14700	20850
Random Projection Rtree	1378	15800	22051
Binary Tree (default)	1344	29186	31200
FIFO List + Rtree	2164	33770	42900
Point RTree	10431	> 44000	-
Ellipsoidal Rtree	14328	> 44000	-

cost.

Figure 14 shows that as we increase  $\text{Ellg}$ , Primary Retrieves are being replaced by Secondary Retrieves, while the overall number of successful retrieves stays fairly constant for larger values of  $\text{Ellg}$ . This is because increasing  $\text{Ellg}$  is replacing adds with grow operations, which as we discussed earlier increases  $N_{\text{fpos}}$  and is a manifestation of Effect 8. The explanation of the super-linear increase in retrieve time is similar to that described for tuning  $\text{Ellr}$ . As we increase  $\text{Ellg}$  the miss cost also increases slightly (see Figure 15) because of Effects 4 and 6. Overall the total simulation time first decreases because of the dominance of Effect 7, but later starts to increase because of Effects 4 and 8.

### 5.3 Tradeoffs: The Big Picture

It is evident from the detailed analysis in Section 5.2 that the tuning parameters can have a significant effect on the performance of the indexes and the overall function approximation algorithm. We performed a similar analysis for the other index structures and selected the best parameter setting for each of them accordingly. Table 3 lists the overall running time of the Methane combustion simulation; the times are for the indexes *after tuning*, unless explicitly stated otherwise. We report times for different values of  $\varepsilon$ , because the index size increases with lower error tolerance and hence smaller ellipsoids.

The tuned Binary Tree performs significantly better than the Binary Tree with default parameter settings ( $\text{Ellr} = \text{Ellg} = \infty$ ). In fact, it outperforms all competitors. The “natural” index for this problem, the Bbox Rtree, performs well, but is 20-40% slower than the tuned Binary Tree. We established that the cause for this difference was the ability of the Binary Tree to achieve a large number of Primary Retrieves because it partitions the space, rather than searching through levels of overlapping bounding boxes. Careful tuning can bias the Binary Tree toward a high rate of Primary Retrieves, with little reduction in overall retrieval rate. On the other hand, tuning had comparatively little effect on the Bbox Rtree. The overlap of bounding boxes at all levels of the tree resulted in large numbers of false positives during search. We note here that the difference in performance of the Bbox Rtree and the Binary Tree is not due to the update costs of the Rtree. We have found in our experiments that Effect 6 does not significantly affect performance because very few ellipsoids actually grow during a single grow step.

The dramatic difference between the FIFO List and MRU List indexes is caused by locality in the combustion simulation. Both index structures are identical; they only differ in the order of the ellipsoids in the list. MRU sorts by most recent access, while FIFO maintains the ellipsoids in the order in which they were added.

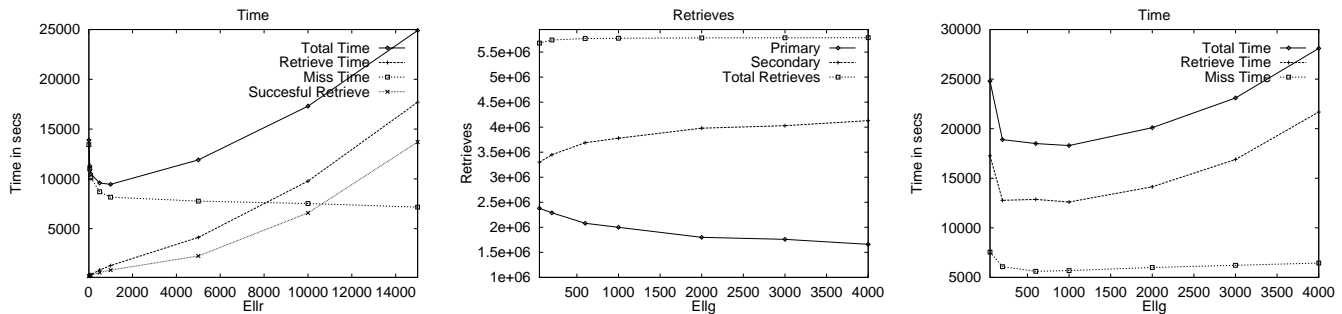


Figure 13: Time vs. Ellr (Binary Tree) Figure 14: Number of retrieves vs. Ellg (Binary Tree) Figure 15: Time vs. Ellg (Binary Tree)

A surprising result was the poor performance of the Point Rtree. Since it does not know the spatial extent of the ellipsoids, we implemented the retrieve operation with an NN-query to find the “best” ellipsoids early on. Unfortunately because of the limited pruning power and the high cost of the NN-search, the Point Rtree was not more successful than scanning the FIFO List. The MRU List + Rtree essentially uses the same Rtree during the grow step to find grow candidates. Hence the performance difference between it and the Point Rtree approach is mostly due to the poor retrieve performance of the Point Rtree.

We also experimented with two extensions of the Bbox Rtree to explore ways to improve its performance. Both are motivated by the problem that in high dimensions, hyper-rectangular bounding boxes are only poor approximations of ellipsoids. The bounding boxes contain a large fraction of “dead space”, i.e., volume that is outside the ellipsoid, which creates many false positives during search.

The **Random Projection Rtree** addresses the problem by projecting all ellipsoids onto a fixed set of  $k$  randomly selected lines. This transforms a  $d$ -dimensional ellipsoid into a  $k$ -dimensional hyper-rectangle in the transformed space defined by the projection lines. We can now use a standard Rtree to index the objects. By using larger numbers of projections, we can achieve a tighter bounding polyhedron around an ellipsoid, at the cost of more expensive index operations in the higher-dimensional space. The results for  $k = 60$  showed the expected lower false positive rates (compared to Bbox Rtree), but slightly worse overall performance (see Table 3) because of higher dimensionality. A detailed study of Random Projection Rtrees is part of our future work.

We can also reduce dead space by using ellipsoids as the bounding shape at all tree levels. The corresponding **Ellipsoid Rtree** performed very poorly, because of the high cost of basic index operations like testing if a point is within a bounding ellipsoid or splitting nodes and computing the new bounding ellipsoids, which is done approximately.

## 6. RELATED WORK

The ISAT function approximation approach was first introduced by Pope [22]. It is one of the most widely used techniques for function approximation in the scientific community and is now a part of Fluent’s CFD package [2]. Machine learning and data mining research have extensively studied the problem of automatically learning unknown functions [13]. By treating the known function values as a train-

ing sample, machine learning techniques can be used for function approximation. For the combustion simulation, neural networks have been proposed and used [14]. However, there has been very little work on studying function approximation as an indexing problem. Pope [22] and later Veljkovic et al. [26] propose new index structures for combustion simulation. Our work is the first principled analysis of the indexing problem.

A large variety of index structures have been proposed by the database and computational geometry communities, and their suitability for the function approximation problem needs to be studied. Work prior to 2001 is surveyed in [5] and [10]. In the following we discuss a few selected indexes, which are most related to the ones studied in this paper.

The Rtree [12] is a commonly used multidimensional index in the database community. It is a balanced data structure based on axis-parallel bounding boxes and can manage both point and extended objects. It is thus a natural choice for indexing Local Regions for function approximation. Several variants of the Rtree have been proposed, e.g., R\*-tree [3], R+-tree [24], and Xtree [4]. The goal of most improvements is to reduce the overlap of bounding boxes in tree nodes, which is a major factor in degrading performance for high-dimensional data. The SS-Tree [28] takes this a step further by using spheres as bounding regions. It is therefore a good candidate for managing spherical or ellipsoidal regions of accuracy.

To avoid overlap of bounding regions, some index structures partition the space, e.g., the Binary Space Partitioning (BSP) tree [10]. The Binary Tree used in our experiments is an adaptation of this index structure.

It has been shown that in high dimensions linear scans are sometimes faster than complex index structures, especially when data is accessed on disk. The VA-file [27] improves the performance of linear scans by quantizing the space. A simple approach based on scanning files at different resolutions has been shown to outperform sophisticated bulk-loaded Rtrees [23].

The Random Projection Rtree in this paper was motivated by work on using projections for containment queries [7] and approximate nearest-neighbor queries [17]. Multidimensional problems can be mapped to lower dimensions by hashing [11, 15]. Other common approaches to combat the curse of dimensionality are dimensionality reduction and principal component analysis, e.g., used by the TV-tree [19], the  $\Delta$ -tree [8] and the VA+ file [9].

## 7. CONCLUSIONS AND FUTURE WORK

In this paper we introduced the function approximation problem. We showed its hardness and how it motivates an interesting indexing problem. A principled analysis of the indexing problem led to the discovery of novel tradeoffs which have a significant and varied impact on different index structures.

This is the first in-depth study of the general indexing problem in high-dimensional function approximation and there are several avenues for future research. (1) Function approximation can provide a new and non-traditional benchmark for comparing indexes. (2) We have so far evaluated tradeoffs when parameter values are fixed throughout the simulation. Performance could be improved considerably by varying parameters adaptively. For example, growing and adding could be turned off towards the end of the simulation. (3) The Random Projection Rtree provides a new direction in the design of index structures for ellipsoids. (4) It is also interesting to compare this approach of function approximation to the algorithms used by the machine learning community.

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