

Unified data structures to optimize solving a complex of interrelated geometric problems

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Abstract—The paper is devoted to the development of an efficient algorithmic tools for solving a set of interrelated computational geometry problems. To do this, a single algorithmic environment with unified data structures is created, which allows to implement complex use cases efficiently with respect to computational resources. We build the environment on the basis of the “divide and conquer” strategy. Once a convex hull is key to a set of computational geometry problems, we offer a special weighted concatenable queue data structure to maintain it. The data structure is implemented in a form of a binary tree. **This allows to perform procedures for a set of tasks within no more than $O(\log n)$ time.** Furthermore we offer a way to execute the algorithms both sequentially and in parallel. In the future the algorithmic environment can be improved to support other computational models with similar integral properties for solving problems. As an example, the Voronoi Diagram or the Delaunay Triangulation can be considered.

Index Terms—Algorithmic tools, Computational geometry, Interrelated problems set, Single algorithmic environment, A weighted concatenable queue

I. INTRODUCTION

The current situation in the field of research. The convex hull problem arises not only in many practical applications, but also is an core tool for solving many others computational geometry problems.

There are implicit relations between such problems. Currently, the state of the art approach for solving such sets of problems is the single algorithmic environment, which takes advantage of those implicit relations to create a final software product. An example to such approach may be the usage of a generic data structures for solving different problems. Thus, the algorithmic environment allows to reduce complex problems to some basic computations.

This article introduces a new approach for solving the convex hull problem on a static set of points. Here the divide and conquer strategy is used as a base for the algorithm and the actual convex hull is maintained with a concatenable queue during the computations. The resulting algorithm has the worst case complexity of $O(n \log n)$, whereas the subroutine of merging 2 convex hull has a complexity of $O(\log n)$, where n is the number of points.

Henceforth a notion of a single algorithmic environment is introduced that formalizes the divide and conquer algorithm from a programmatic standpoint. The formalization allows to

create a level of abstraction upon an actual implementation of an algorithm and therefore perform computations sequentially and in parallel on a more generic level.

Relevance consist in a large specter of application problems that computational geometry allows to solve with its methodology. Implementation of those algorithms enables in an single algorithmic environment allows to use them in the development of modern technology and apply them on large datasets.

The goals of this article is to design an optimal algorithms for solving the convex hull problem on a static set of points and to create a single algorithmic environment on the divide and conquer principle which allows to execute its algorithms both sequentially and in parallel.

Tasks:

- 1) To systematize and analyze existing approaches to solve the aforementioned problem.
- 2) Create a formal description of a new algorithm.
- 3) Formally prove its computational complexity.
- 4) Implement the algorithm and create.
- 5) Create a single algorithmic environment which allows to execute computational geometry algorithms based on divide and conquer principle sequentially and in parallel.
- 6) Describe practical use cases of the developed software.

Object of research is methods of convex hull computation, applying the divide and conquer strategy for designing computational geometry algorithms, programmatic tools for executing computations in parallel. To achieve the goals the following **methods of research** are used: collection the information, information analysis, formalizing of the collected data and method of proving the scientific hypothesis.

The application scope of the results described in this article includes designing of dentures and corsets, where it is necessary to solve a large set of computational geometry problems on large inputs.

II. CONVEX HULL ALGORITHM

A. Previous work

The notion of a convex hull is simple. For a set of points S in a k -dimensional space it is a smallest convex set, that comprise S . The problem of building such a set has a long

history. In practice to solve such problem, means to find a subset of S , which can a "skeleton" for the convex hull.

Despite its simplicity, the definition mentioned earlier does not allow to build on its basis an algorithm. To find a solution, it is needed to introduce more notions that will address the nature of the problem in a more detailed way and allow to solve it in a constructively.

It is differentiated between two types of points set in this case: a static and a dynamic one. In the first case all points are known in advance. The algorithms for solving that kind of problem are well researched and described in [1]–[3]. In the latter case an algorithm should be able to address the problem on a new points being added to the initial set and some others being removed from it after a certain period of time. Initially a set of points can either contain points or be empty.

The first known algorithm for solve such problem was introduces by Preparata in [4]. Convex hull is maintained in a form of an AVL tree, that allows to perform operations of adding new points form it. The split and finding of tangents is performed in optimal time of $O(\log n)$. The major drawback of this method is that there is no mean to remote points from the set.

Soon after Overmars and van Leeuwen described a method of how a concatenable queue can be used to maintain the convex hull [5]. This, as well as special data structure that is based on a balanced binary tree and maintain the information gathered during previous queries, allows to support fully dynamic sets of points and perform a add or remove query $O(\log^2 n)$ in the worst case.

B. Concatenable queue

Concatenable queue is an Abstract Data Type (ADT), that supports following operations:

- ADD_ELEMENT();
- REMOVE_ELEMENT();
- GET_MINIMUM();
- CONTAINS();
- SPLIT();
- MERGE().

By default the elements in a concatenable queue are kept i a certain predefined order [6].

Here the concatenable queue is implemented as a binary $B+$ tree. Its elements are divided into non-leaf and leaf elements. The leaf elements contain all data kept in a tree. Every element has a pointer to its left and right child element. For the leaf elements those pointers point to the elements left and right neighbors or *null* if the element is utmost in the tree. Additionally every element keep a pointer to the largest element in its left sub-tree, which allows to perform binary search [6]. An example of such tree is given on fig. 1. The height of earch element is kept for the balancing process during the split and merge operations. It is measured as a maximum amount of steps it is possible to do in order to reach a leaf element.

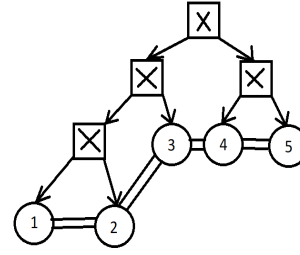


Fig. 1. Example of a concatenable queue.

From now we will go into details on how this data structure is implemented. So on the list. 1 its formal definition is provided:

```
class Node:
    int data
    int height

    Node left
    Node right
    Node leftSubtreeMax

    boolean isLeaf

class ConcatenableQueue:
    Node root
    Node minNode
    Node maxNode
```

Listing 1. Structure of a concatenable queue vertex

The contains operations uses binary search by the tree elements so its complexity is $O(\log n)$, where n hereafter denotes the numbers of elements in the queue.

Algorithm of insertion an element in a queue looks like as follows. First, the position of a new element is searched. Then the connections between adjacent leaf elements are broken to insert a new element. Going back a new non-leaf element is created - parent for the new element and one of its neighbor. The algorithm is formally described on the list. 2.

The *updateHeight* subroutine updates the value of an element height, the *createLeafBetween* subroutine breaks the connection between adjacent elements and insertion of a new one.

The *insert* operation. On the first step we find out if *leftSubtreeMax* point to an element with greater value than the value to be inserted e . If so, then if current element is a leaf, a new element is created between the current element and its left neighbor. Otherwise the search proceeds on the left sub-tree of the current element. The case, when *leftSubtreeMax* is smaller than the value e is analogous. The procedure ends

```

Node insert(Node node, int e):
Node result = nil
if e <= node.leftSubtreeMax.data:
if node.isLeaf:
if e == node.data:
node.data = e
else:
Node createdLeaf =
createLeafBetween(e, node.left, node)
result =
Node(createdLeaf, createdLeaf, node)
else:
node.left = insert(node.left, e)
else:
if node.isLeaf:
Node createdLeaf =
createLeafBetween(e, node, node.right)
result = Node(node, node, createdLeaf)
else:
// go left and update node.right pointer
node.right = insert(node.right, e)

if result == nil:
result = node

updateHeight(result)
return result

```

Listing 2. Queue element insertion algorithm

with updating the height on a newly created element. The element is returned as its result value.

Since on every step we perform a constant amount of work, the complexity of the procedure is $O(h) = O(\log n)$, where h hereafter denotes the height of the tree.

The operation of removing an element from the queue is performed analogously.

The split operation is a bit more complex. As an input the corresponding procedure receives current element and value based on which the split is performed. As a result two independent queues are formed. The value, by which the split has been performed, belongs to the left sub-queue. The procedure is described formally on the list. 3.

```

split(Node node, int e,
ConcatenableQueue leftQueue,
ConcatenableQueue rightQueue):
if node.isLeaf
leftQueue.root = node
leftQueue.maxNode = node

rightQueue.minNode = node.right
cut(node)
else:
if e == node.leftSubtreeMax.data:

```

```

leftQueue.root = node.left
leftQueue.maxNode = node.leftSubtreeMax

rightQueue.root = node.right
rightQueue.minNode = node.leftSubtreeMax.right

cut(node.leftSubtreeMax)
else if e < node.leftSubtreeMax.data:
split(node.left, e, leftQueue, rightQueue)
rightQueue.root =
concatenateNodes(rightQueue.root,
node.right, node.leftSubtreeMax)
else:
split(node.right, e, leftQueue, rightQueue)
leftQueue.root =
concatenateNodes(node.left,
leftQueue.root, node.leftSubtreeMax)

```

Listing 3. Queue split algorithm

Here the *concatenateNodes* procedure is used. It performs concatenation of two arbitrary elements and uses their heights to balance the resulting queue. Its implementation is described further in the work. The *cut* procedure breaks connection between two adjacent leaf elements in a queue and therefore is trivial. On the first step in the *split* operation we check, is the current element is a leaf. If so, its connection are broken and the value of *maxNode* is updated for the left queue as well as the value of *minNode* for the right queue. If element is not a leaf, then the procedure continues of either left or right sub-tree. Here a special corner cases is considered, where *leftSubtreeMax* contain the dividing value. Then the analogous action to the usual search are performed.

The algorithm of the *concatenateNodes* procedure is described on the list. 4.

```

Node concatenateNodes(Node leftNode,
Node rightNode, Node leftSubtreeMax) {
if leftNode == nil:
return rightNode
else if rightNode == nil:
return leftNode
else if leftNode.height == rightNode.height:
Node result = Node(leftSubtreeMax,
leftNode, rightNode)
updateHeight(result)
return result
else if leftNode.height < rightNode.height:
rightNode.left = concatenateNodes(leftNode,
rightNode.left, leftSubtreeMax)
updateHeight(rightNode)
return rightNode
else
leftNode.right =
concatenateNodes(leftNode.right,
rightNode, leftSubtreeMax)
updateHeight(leftNode)

```

```
return leftNode
```

Listing 4. Merging two queues using heights

First, we consider corner cases where one of the elements is *null*. This is required to ensure correctness of the recursion. Then, if the left element is higher than the right, one step down is taken for the left element. If the right element is higher - we take a step down for it. If the heights are equal, the joining point is found and a new element must be created. At each step, it is necessary to update the height of current element as it changes.

We begin by analyzing the complexity of the *split* procedure by determining the complexity of the *concatenateNodes* procedure. At each iteration, the step is performed either to the left son of the current element or to the right. The execution of the recursive procedure is finished by merging two elements. Since each step moves us down one level and a constant amount of work is performed for each level, the total complexity of the *concatenateNodes* procedure is $O(h) = O(\log n)$.

The *split* procedure uses the *concatenateNodes* function as a subroutine. The complexity a *split* call is equal to the complexity of *concatenateNodes*. The number of recursive calls *split* for one split is $\log n$, so the total complexity of the procedure $\log^2 n$.

The merge operation of two queues is reduced to the clamping of their root vertices by the *concatenateNodes* procedure, so its complexity is $O(\log n)$.

C. Precomputation

In the pre-processing stage, the "inner" points, which lie on a horizontal or vertical line, are removed from the set. Formally, the removal criterion is formulated as follows. For $a = (x_a, y_a)$ we denote $x(a) = x_a$, $y(a) = y_a$. Let points a_1, a_2, \dots, a_k lie on one horizontal line and $x(a_1) < x(a_2) < \dots < x(a_k)$. Then, by the criterion, the points a_2, a_3, \dots, a_{k-1} must be removed. Similarly for the vertical case.

Consider an algorithm that, in a sorted array, for each group of identical elements, deletes all but the first and the last (if the repetition is more than two). The description of the algorithm is given in Listing 5.

```
removeDuplicated(int[] points) {
    int n = points.length

    if n == 0:
        return

    it1 = 1
    it2 = 0

    while it2 < n:
        if it2 + 1 > 0 && it2 + 1 < n - 1:
            int previous = points[it2 - 1]
            int current = points[it2]
            int next = points[it2 + 1]
```

```
if previous != current || current != next:
    points[it1] = current
    if it1 < n-1:
        ++it1
    else if it2 + 1 == 0:
        if it1 < n-1:
            ++it1
            ++it2
        else:
            points[it1] = points[it2]

    if it1 < n-1:
        while it1 + 1 != points.length:
            delete points[n - 1]
```

Listing 5. Algorithm of removing duplicates in a sorted array

The idea behind the algorithm is to use two pointers to delete repetitions by overwriting their place with other non-repeating element. In this way, additional memory usage can be reduced to a constant value. The algorithm makes one pass through the array. At each step, a constant amount of work is performed to decide whether to delete the current item. Given this, the complexity of the above algorithm is $O(n)$.

To perform the above pre-processing, it is needed to:

- 1) Sort points by y (if y coordinates are equal, the x coordinates are compared).
- 2) Delete repetitions by y coordinate using the described algorithm.
- 3) Sort points by x (if x coordinates are equal, the y coordinates are compared).
- 4) Delete repetitions by x coordinate using the described algorithm.

D. The divide and conquer principle

At the stage of splitting the initial problem, the current set of points is divided into left and right parts of equal size. Since a list is used to store the points, this operations can be completed by $O(1)$ using the formula

$$M_{i,j} = \frac{i+j}{2} \quad (1)$$

The recursion stops when there are no more than 3 points in the set.

For the base case, the set of points can have 2 or 3 elements. In the first case, as shown in fig. 2, the highest of two points forms the upper part of the hull, and the lower - the lower part.

To consider the base case of 3 points, we introduce an additional notion of the tangent slope given by two points on the plane. For arbitrary points a_1, a_2 such that $x(a_1) < x(a_2)$ slope is denoted as λ :

$$\lambda(a_1, a_2) = \frac{y(a_2) - y(a_1)}{x(a_2) - x(a_1)} \quad (2)$$

According to this definition, the possible cases for points a_1, a_2, a_3 are given in table. 1 and shown in fig. 3:

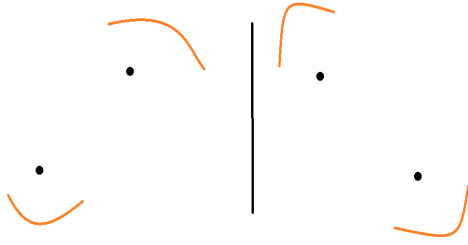


Fig. 2. Base case for 2 points

TABLE I
POSITION OF 3-POINTS BASE CASE

$\lambda(a_1, a_2) > \lambda(a_2, a_3)$	$y(a_1) < y(a_2)$	upper sub-hull	lower sub-hull
false	false	a_1, a_3	a_2
false	true	a_1, a_3	a_2
true	false	a_2, a_3	a_1
true	true	a_1, a_2	a_3

E. The merge step

[5] provides an algorithm for merging two convex hulls. The essence of the algorithm is to maintain the hull with the help of two concatenated queues. The hull is divided into two parts. In this article, the separation occurs from top to bottom. Then, using the two-pointer technique and the cases described in [5], the proper tangent line is searched. It remains to split the two queues at the 2 found vertices that form the tangent and to merge the remaining parts. An example of performing such a procedure is shown in fig. 4.

It remains to consider the corner cases that arise when performing the merging. The first of these cases is related to the ambiguity of the position of the utmost points in the described representation of the convex hull. The leftmost point of the left hull and the rightmost point of the right hull must belong to the upper parts of the view before finding the tangent line, because otherwise such tangent may be found incorrectly.

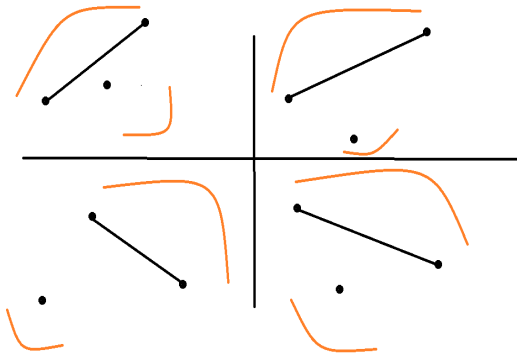


Fig. 3. Base cases for a set of 3 points

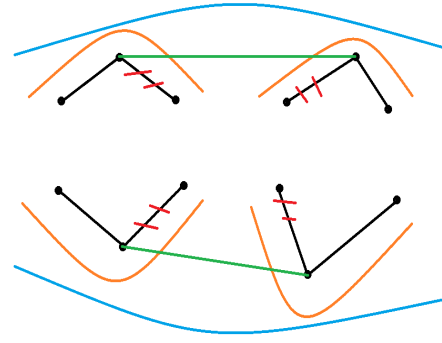


Fig. 4. Merging step of two hulls

An example of such incorrect search is given in fig. 5.

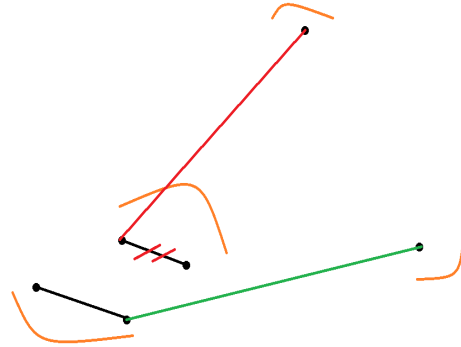


Fig. 5. Example of an incorrect position of the utmost left in the left sub-hull

To avoid such a situation, it is necessary to move the indicated points of their upper parts before merging the sub-hulls.

For the rightmost point of the left hull and the leftmost point of the right hull we have the following cases. Similarly to the previous argument, they must be transferred to the upper parts of the hulls. And after merging these points must be transferred to the lower parts of the hull, if they do not belong to the created upper part of the final hull. Otherwise, the formed hull may be incorrect. An example of such case is shown in fig. 6.

After combining the parts of the convex hulls, another corner case might take place. The search for the tangent for the upper parts of the hulls does not take into account the position of the lower parts and vice versa. As a result, the upper and lower parts of the final hull may not form a coherent structure. An example of such a situation is shown in Fig. 7.

To avoid such a situation, it is necessary to perform the step of cutting off the redundant parts of the formed lower sub-hull. Then searching for the left and right pivoting vertices in the concatenable queue is performed. After that, the queue is splits over the found vertices.

The cases corresponding to the left search are shown in fig. 8, and those corresponding to the right - in fig. 9. Fig. 10 shows correctly constructed convex hull.

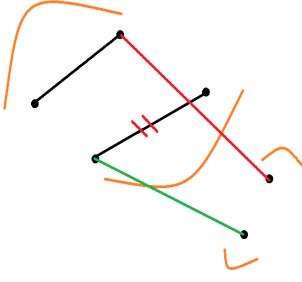


Fig. 6. Example of a convex hull for a wrong position of the utmost left points of the left sub-hull

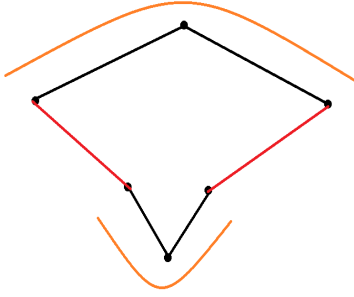


Fig. 7. An example of a non-integral hull after merging along the reference lines

F. Algorithm complexity

Theorem II.1. *The complexity of the described convex hull construction algorithm for a static set of points is $O(n \log n)$ with sequential execution.*

Proof. We will argue the complexity of the algorithm by listing the complexities of the main stages it consists of.

- Precomputation $O(n \log n)$.
- Recursive descent and splitting the set into 2 parts $O(1)$.
- Recursive ascent and merging parts of the convex hull $O(\log n)$.

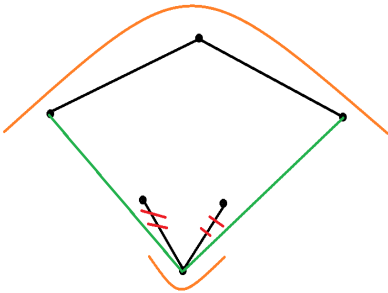


Fig. 8. Correctly constructed convex hull

- Transfer of the utmost points to upper parts of convex hulls $O(\log n)$.
- Finding the tangent for the upper parts of the hulls $O(\log n)$.
- Splitting and merging the upper parts $O(\log^2 n)$.
- Moving the utmost points to the bottom of the hulls $O(\log n)$.
- Finding the tangent for the upper parts of the hulls $O(\log n)$.
- Splitting and merging the upper parts $O(\log^2 n)$.
- Merging the lower parts of the hull $O(\log n)$.
- Normalization of the obtained lower part $O(\log n)$.

The complexity of pre-processing directly depends on the complexity of the sort algorithm. Using fast sorting algorithms [7, page 159], this procedure can be performed in the optimal time of $O(n \log n)$.

To estimate the complexity of the recursive procedure for constructing a convex hull, we make the following equation:

$$T(n) = 2T\left(\frac{n}{2}\right) + O(\log^2 n)$$

According to the well-known result from the theory of algorithmic complexity we have that the solution of this equation is:

$$T(n) = O(n)$$

Thus, taking into account the pre-processing, we get the total complexity of the algorithm $O(n \log n)$ \square

Theorem II.2. *The complexity of the recursive convex hull construction is $O(\log^3 n)$ when executed concurrently on $\frac{n}{2}$ processors.*

Proof. The recursion tree has a height of no more than $\log n$ levels. At the lowest level, the number of subtasks created is $\frac{n}{2}$. Thus, each subtask takes not more than $\frac{n}{2}$ time.

Next, $O(\log^2 n)$ work is performed at each level. Having the height of the recursion tree, we get the total complexity of the algorithm

$$T(n) = O(\log^3 n)$$

\square

III. SINGLE ALGORITHMIC ENVIRONMENT

A. Divide and conquer algorithm interface

The next goal of this work is to build a single algorithmic environment. The construction of such an object requires the combination of an algorithmic database together with the necessary data structures. It will allow to incorporate all implemented algorithms together, without necessity to know their internal implementation. In fact, it is necessary to create an interface of generic algorithm based on divide and conquer strategy, which will then be used on a specific set of input data.

We start creating the generic interface by listing its components:

- Precomputation.
- Splitting task into sub-tasks.
- Solving sub-tasks.
- Merging obtained results.
- Checking, if a given input data is a base case for the algorithm.
- Solving the base case.

Each of these components will represent a function of the future interface. Here there is a clear separation between the input type for the algorithm and the type of result it returns. These two types should be parameters of the algorithm model to be built. One should also pay attention to the types of functions input and output types of the interface.

A large number of computational geometry algorithms, such as computing minimal spanning tree, the Delaunay triangulation, the Voronoi diagram and the convex hull accept the list of points. However, it is not possible limit the type of input to the algorithm interface in a single algorithmic environment. The reason for this is that some algorithms can use the results of other algorithms as input data. A well known example is the construction of a Delaunay triangulation given the Voronoi diagram represented in a special data structure [9, pp. 209].

Listing 6 shows the constructed algorithm model:

```
interface DaCAlgorithm[IT, OT]:
  boolean isBaseCase(IT input)
  int inputSize(IT input)
  OT solveBaseCase(IT input)
  OT merge(OT first, OT second)
  Pair[IT, IT] divide(IT input)
  IT precompute(IT input)
```

Listing 6. Algorithm model based on the divide and conquer principle

B. Sequential and parallel execution

Although this model very accurately describes the class of algorithms, it does not make it possible to solve the problem directly by having input data, which in fact enables to the algorithm from how it will be executed. In this part of the article, principles of sequential and parallel execution are considered.

When executing sequentially an algorithm, its individual subtasks computed one by one. Such algorithm is shown in Listing 7:

```
OT solveRecursively(IT input,
  DaCAlgorithm[IT, OT] algorithm):
  if algorithm.isBaseCase(input):
    return algorithm.solveBaseCase(input)

  Pair[IT, IT] p = algorithm.divide(input)
  IT left = p[0]
  IT right = p[1]

  OT leftResult = solveRecursively(left)
  OT rightResult = solveRecursively(right)
```

```
return algorithm.merge(leftResult, rightResult)
```

Listing 7. Sequential execution algorithm

In parallel execution, it is necessary to take into account that the individual subtasks can be calculated independently, which significantly speeds up the execution of the algorithm.

To construct the concurrent algorithm, we use the following parallel computation abstraction *computeInParallel(function1, function2)*, which runs the functions *function1* and *function2* in parallel.

Listing 8 contains the main steps of this implementation.

```
OT compute(IT input,
  DaCAlgorithm[IT, OT] algorithm):
  if algorithm.isBaseCase(input):
    return algorithm.solveBaseCase(input)

  Pair[IT, IT] p = algorithm.divide(input)
  IT left = p[0]
  IT right = p[1]

  OT leftResult, rightResult =
    computeInParallel(solveRecursively(left),
      solveRecursively(right))

  return algorithm.merge(leftResult, rightResult)
```

Listing 8. Parallel execution algorithm

From the standpoint of the practical implementation the performance of such algorithm was improved by introducing a limit on the size of subtasks that can be calculated in parallel.

C. Performance comparison

A number of algorithm performance measurements were performed for different input sizes and the average number of recursive tasks per thread. The results are shown in Table. 2 and 3.

TABLE II
SEQUENTIAL EXECUTION PERFORMANCE

Number of points	1000000	10000000	100000000
	52.207 (ms)	648.621 (ms)	1302.485 (ms)

TABLE III
PARALLEL EXECUTION PERFORMANCE

NosptNop	1000000	10000000	100000000
20	52.434 (ms)	458.921 (ms)	969.243 (ms)
30	53.615 (ms)	472.127 (ms)	845.707 (ms)
40	52.797 (ms)	464.418 (ms)	907.498 (ms)
50	53.189 (ms)	474.804 (ms)	905.952 (ms)

CONCLUSION

As a result of the work, existing methods for constructing a convex hull were analyzed and a new algorithm was proposed to solve this problem for a static set of points.

A unified algorithmic environment was also created on the basis of divide and conquer algorithms, which allows to develop efficient implementation of such algorithms quickly and enables to execute them both sequentially and in parallel. The described object was implemented in Java software using standard library tools. Additionally, a tool was developed to visualize the results of the convex hull construction.

The implemented of data structures and other elements does not use other libraries. This makes it independent and easy to use.

The main advantages of the developed algorithm are the optimization of the preprocessing stage and the effective application of such data structures as the concatenable queue, which allows to spend minimum time for the merging step of the convex hulls construction. In the preprocessing stage constant memory usage is achieved.

The performance comparison of the developed algorithm for both types of execution allows to conclude that it has high degree of parallelism. The speedup was 31% percent in the best case. It is worth noting that the use of parallel execution is only necessary for very large input data.

The architecture of the created algorithmic environment makes it easy to expand its capabilities: add new as well as modify existing algorithms, implement the necessary data and extend the programmatic representation of the algorithm based on the divide and conquer scheme. This flexibility of the software product is achieved by using the modular principle in its design. In the future, it is necessary to investigate the possibility of performing parallel preprocessing. The basis for this research is given in [8].

The results of this work can be incorporated in the development of modern high-precision 3D models. The concept of a single algorithmic environment has proven to be highly relevant and effective for constructing a software for solving complex problems of computational geometry. This, along with the ease of extending the product created, makes research in this area extremely relevant.

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