# On logarithmic-factor speedup algorithms for some classic problems

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January 7, 2025

#### Abstract

This survey discusses techniques that speed up the time complexity of algorithms for certain problems (mainly those under the scope of fine-grained complexity) by a logarithmic factor, and provides a brief overview of works aimed at improving the lower bound. Four main problems are discussed: boolean matrix multiplication (introducing bit-parallelization and the method of four Russians), longest common subsequence, 3-sum, and all pairs shortest path.

# 1 Introduction

Among the history of algorithmic improvements for specific problems, one type of speedup is often overlooked compared to optimizations on polynomial order. Typically, we are concerned with problems such as, "Can we reduce the time from  $O(n^2)$  to  $O(n \log n)$ ?" or "Is it possible to remove the log n factor from  $O(n \log n)$ ?".

In this survey, however, we focus on algorithms that run in  $O\left(\frac{n^k}{\text{polylog}(n)}\right)$  time, providing a logarithmic speedup compared to the trivial algorithms. Our discussion is based on the word RAM model— $w = \Omega(\log n)$  bits can be processed simultaneously, where n is the problem size. The techniques discussed are essentially complicated variations of the idea proposed by [ADKF70], mainly bit parallelization and preprocessing all possible scenarios of the locality of the unoptimized algorithm model.

This technique can also be applied to other problems not covered in this survey but worth mentioning. Using the method of four Russians, the lowest common ancestor (LCA) problem, range minimum query (RMQ) problem (see [BFC00]), level ancestor (LA) problem (see [BFC04]), and other related problems (see [Jin23]), have been optimized to O(n) preprocessing and O(1) query time, removing the log *n* factor from the common algorithms.

We consider these problems under the category of fine-grained complexity theory. Fine-grained complexity focuses on determining the exact polynomial order of time complexity achievable by the best algorithm for a given problem. Based on the following conjectures, a class of problems has been shown to be unlikely to have an algorithm with time complexity below a specific polynomial order bound (known as the conditional lower bound), via fine-grained reductions, which is analogous to polynomial-time reductions used to prove NP-hardness:

- 1. Strong Exponential Time Hypothesis (SETH): kSAT cannot be solved in  $O(2^{(1-\epsilon)n})$  time.
- 2. APSP cannot be solved in  $O(n^{3-\epsilon})$  time.
- 3. 3SUM cannot be solved in  $O(n^{2-\epsilon})$  time.

The introduction to fine-grained complexity presented above is based on [Bri21]. We will not discuss the hardness results further in the subsequent sections, but present algorithms that optimize the algorithms for LCS, 3SUM, and APSP by a logarithmic factor, which do not violate the conjectures mentioned above. We will first discuss BMM to illustrate the basic idea of optimization.

In researching the papers and algorithms, the content of [Li23] was particularly helpful and is itself a high-quality survey of this topic.

### 2 Preliminaries

We analyze time complexity based on the word RAM model with word size w. We assume that equality, comparison, and addition between elements can be performed in O(1) time. If not specified,  $A_i$  ( $A_{i,j}$ ) denotes the *i*-th (row and *j*-th column) element of sequence (matrix) A, and indices start from 1. We will not omit the log log n factor in complexity unless  $\hat{O}$  is used.

### 3 The boolean matrix multiplication problem

**Problem 1** (BMM). Given two  $n \times n$  boolean matrices A, B, compute  $n \times n$  matrix C, such that

$$C_{i,j} = \bigvee_{k=1}^{n} \left( A_{i,k} \wedge B_{k,j} \right)$$

This problem is weaker than the general matrix multiplication, which has an  $O(n^{\omega})$  algorithm, with  $\omega \approx 2.37$  so far ([Wik24]). Here, however, we focus only on combinatorial algorithms.

[ADKF70] introduced a groundbreaking algorithm, widely known as "the method of four Russians". It works as follows: partition A into  $n \times d$  and B into  $d \times n$  smaller matrices, and consider optimizing the multiplication of such a pair, namely  $A_1$  and  $B_1$ .



Figure 1: The partitioning.

Denote the rows of  $A_1$  as  $a_1, \dots, a_n$  and the rows of  $B_1$  as  $b_1, \dots, b_d$ . Although we need to compute  $a_1B, \dots, a_nB$ , there are only  $2^d$  possibilities among them, so we can precompute these results by recursion. For a row vector a such that  $a_t = 1$ , we have

$$aB = (a \setminus t)B \vee b_t,$$

where  $a \setminus t$  means setting  $a_t$  to 0, and  $\vee$  represents the "or" operation applied element-wise to the two row vectors. Since we use the word RAM model, precomputation can be done in  $O\left(2^d \frac{n}{w}\right)$  time,

and using them to compute  $A_1B_1$  takes  $O\left(\frac{n^2}{w}\right)$  time. Finally, taking the  $\vee$  of all the products requires  $O\left(\frac{n^3}{wd}\right)$  time. Choosing  $d = \log n$ , the overall time becomes  $O\left(\frac{n^3}{w\log n}\right)$ . After decades, [BW09] presented an algorithm based on advanced graph theory techniques with

After decades, [BW09] presented an algorithm based on advanced graph theory techniques with time complexity  $O\left(\frac{n^3(\log\log n)^2}{(\log n)^{9/4}}\right)$ , and [Cha15] subsequently gave an algorithm with time complexity  $O\left(\frac{n^3(\log\log n)^3}{(\log n)^3}\right)$  using a relatively simple analysis. Then, [Yu15] introduced an algorithm with  $\hat{O}\left(\frac{n^3}{(\log n)^4}\right)$  time complexity. The current best algorithm, presented by [AFK<sup>+</sup>24], runs in  $\frac{n^3}{2^{\Omega((\log n)^{1/7})}}$  time, which is a speedup strictly stronger than any polylogarithmic factor.

### 4 The longest common subsequence problem

**Problem 2** (LCS). Given two sequences A and B of length n, find the length of the longest common subsequence (LCS) of A and B. A subsequence of a sequence is obtained by removing zero or more elements from it, while preserving the order of the remaining elements. Here, we assume the range of  $A_i$  and  $B_i$  is a finite set  $\Sigma$ .

The common  $O(n^2)$  algorithm is a dynamic programming approach: let A[l, r] denote the consecutive subsequence  $(a_l, \dots, a_r)$  of A. We need to calculate  $f_{i,j}$ —the LCS of A[1, i] and B[1, j]—for every  $0 \le i, j \le n$ . The recurrence is given by:

$$f_{i,j} = \begin{cases} 0, & i = 0 \lor j = 0\\ \max\{f_{i-1,j}, f_{i,j-1}\}, & i, j \ge 1 \land A_i \ne B_j\\ f_{i-1,j-1} + 1, & i, j \ge 1 \land A_i = B_j \end{cases}$$

Thus,  $f_{n,n}$  gives the final answer. A practical  $O\left(\frac{n^2}{w}\right)$  algorithm is presented in [CIPR01], which exploits the parallelized bitwise operations of the word RAM model. The brief description of [CIPR01]'s algorithm is as follows:

Let  $d_{i,j} = f_{i,j} - f_{i,j-1}$ . We can show that  $0 \le d_{i,j} \le 1$  by considering the meanings of  $f_{i,j}$  and  $f_{i,j-1}$ , so  $(d_{i,1}, \dots, d_{i,n})$  can be seen as a boolean vector, denoted as  $d_i$ . We want to compute  $d_i$  from  $d_{i-1}$ .

We calculate  $d_{i,j}$  from j = 1 to n. The transition can be divided into three stages:

- 1.  $f_{i,j-1} = f_{i-1,j-1}$ , and  $A_i \neq B_j$  or  $d_{i-1,j} = 1$ . Simply  $f_{i,j} = f_{i-1,j}$  and  $d_{i,j} = d_{i-1,j}$ .
- 2.  $f_{i,j-1} = f_{i-1,j-1}$ , and  $A_i = B_j$  and  $d_{i-1,j} = 0$ .  $f_{i,j}$  will increase,  $f_{i,j} = f_{i-1,j} + 1$  and  $d_{i,j} = 1$ . 3.  $f_{i,j-1} = f_{i-1,j-1} + 1$ . It must be  $d_{i,j} = 0$ . Also  $f_{i,j} = f_{i-1,j} + 1$  until  $d_{i-1,j} = 1$ .

In conclusion, as long as stage 2 is triggered, it will continue in stage 3 until  $d_{i-1,j} = 1$ , after which it returns to stage 1, which is simply copying the elements of  $d_{i-1}$ . The entire process can be interpreted as a combination of addition and bitwise operations if  $d_i$  is viewed as the binary representation of an integer. Let  $\bar{d}_i = \text{not } d_i$ , the "chain effect" can be seen as the carry bit of the addition process. We have

$$\bar{d}_i = \left[\bar{d}_{i-1} + \left(\bar{d}_{i-1} \operatorname{and} m_{A_i}\right)\right] \operatorname{or} \left(\bar{d}_{i-1} \operatorname{and} \bar{m}_{A_i}\right)$$

where  $m_c = ([B_1 = c], \dots, [B_n = c])$  and  $\bar{m}_c = \text{not } m_c$ . Both  $m_c$  and  $\bar{m}_c$  can also be viewed as integers.

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Figure 2: An example, A = aaaaababbccc, B = bcbcacbbacba, i = 10. Blue, green and orange&red grids represent stage 1, 2, 3 respectively.

Since the involved operations can be performed in  $O\left(\frac{n}{w}\right)$  time, the overall time complexity is  $O\left(\frac{n^2}{w}\right)$ , and the memory complexity is O(n).

[MP80] uses a similar technique to the method of four Russians, leading to a speedup by a  $\log^2 n$  factor, under the assumption that  $|\Sigma| = O(1)$ . For arbitrary  $|\Sigma|$ , [BFC08] achieves  $O\left(\frac{n^2(\log\log n)^2}{\log^2 n}\right)$ , and [Gra16] improves it to  $O\left(\frac{n^2\log\log n}{\log^2 n}\right)$ .

The basic idea of [MP80] is to divide the table of d into  $y \times y$  blocks, with adjacent blocks sharing borders. For a block with  $i \in [i'y + 1, (i' + 1)y]$  and  $j \in [j'y + 1, (j' + 1)y]$ , it can be fully determined as long as its top row, leftmost column, and A[i'y + 1, (i' + 1)y] and B[j'y + 1, (j' + 1)y]are given. If  $|\Sigma| = C \ge 2$ , then  $2y(1 + \log C)$  bits are needed to represent this information.

We can precompute the result (only the bottom row and rightmost column of d of the block are needed for subsequent calculation) for all these  $\Theta(2^{2y}C^{2y})$  possibilities in  $O(2^{2y}C^{2y}y^2)$  time, and then compute the values of all blocks in  $O(l \cdot n^2/y^2)$  time, with lookups from the precomputed results done in  $l = \Theta\left(\frac{y}{\log n}\right)$  time. Taking  $y = \left\lfloor \frac{\log n}{4\log C} \right\rfloor = \Theta(\log n)$  leads to l = O(1), and the overall time complexity becomes  $O\left(\frac{n^2}{\log^2 n}\right)$ .

For the case where  $|\Sigma|$  is not constant, [Gra16] designs a more delicate structure: consider A[i'y+1, (i'+1)y], a snippet of length y at a time. Relabel the elements of B by equality relation with elements of A[i'y+1, (i'+1)y] so that a block of d with dimensions  $x_1 \times x_2$ , where the first dimension is fixed, can now be determined using  $x_1 + x_2(1 + \log(y+1))$  bits. Precompute all the possibilities for each snippet. The overall time complexity is:

$$O\left(\frac{n^2}{y}\log y + \frac{n}{x_1}2^{x_1+x_2}(y+1)^{x_2}x_1x_2 + \frac{n^2}{x_1x_2} \cdot \frac{(x_1+x_2)\log y}{\log n}\right)$$

Taking  $y = \frac{\log^2 n}{2}$ ,  $x_1 = \frac{\log n}{4}$ ,  $x_2 = \frac{\log n}{4 \log \log n}$ , the resulting time complexity will be  $O\left(\frac{n^2 \log \log n}{\log^2 n}\right)$ . No better algorithm is known.

#### 5 The 3-sum problem

**Problem 3** (3SUM). *Given a set*  $A \subseteq U$ *, determine whether*  $\exists a, b, c \in A$  *such that* a + b + c = 0*.* 

An  $O(n^2)$  algorithm is based on the two-pointers technique. First, sort the elements of A, denoted as  $a_1, \dots, a_n$ , in increasing order. We can run the following algorithm:

#### Algorithm 1 Brute force for 3SUM

for  $i \leftarrow 1$  to n do  $k \leftarrow n$ for  $j \leftarrow i$  to k do while  $k > j \land a_i + a_j + a_k > 0$  do  $k \leftarrow 1$ if  $a_i + a_j + a_k = 0$  then Report finding  $(a_i, a_j, a_k)$ Report not found

We focus on  $U = \mathbb{R}$ . [GP18] developed several faster algorithms based on this method, initially presenting a slow algorithm but with only  $O(n^{3/2}\sqrt{\log n})$  decision tree complexity. The main idea is to partition  $a_1, \dots, a_n$  into groups of size g, namely  $A_1, \dots, A_{\lceil n/g \rceil}$ , where  $A_i = \{a_t \mid (i-1)g < t \le ig\}$ , and then keep the first loop. Instead of enumerating j and k as single elements, we enumerate them as two groups, and check if  $-a_i \in A_{j,k} = A_j + A_k = \{x + y \mid x \in A_j, y \in A_k\}$ .

In the following analysis, we denote  $A_j(x)$  as the x-th element of  $A_j$ . We assume there are no duplicate elements in the definitions of  $A_j$  and  $A_{j,k}$ , as ties can be broken by perturbation or by taking the index into account.

This algorithm uses binary search to speed up the checking process to  $O(\log g)$ , resulting in a total complexity of  $O\left(\frac{n^2 \log g}{g}\right)$ . Sorting each  $A_{j,k}$  is then required. The number of possible orders of  $A_{j,k}$  has an upper bound of  $(g^2)!$ . Consider enumerating these permutations, denoted as  $\pi_1, \pi_2 : [g^2] \to [g]$ , which corresponds to  $A_j(\pi_1(t)) + A_k(\pi_2(t))$  being the *t*-th smallest element. A permutation sorts some  $A_{j,k}$  correctly iff for all *t*,

$$A_j(\pi_1(t)) + A_k(\pi_2(t)) < A_j(\pi_1(t+1)) + A_k(\pi_2(t+1))$$
  
$$\iff A_j(\pi_1(t)) - A_j(\pi_1(t+1)) < A_k(\pi_2(t+1)) - A_k(\pi_2(t)).$$

Thus, we can regard  $(A_j(\pi_1(1)) - A_j(\pi_1(2)), \dots, A_j(\pi_1(g^2 - 1)) - A_j(\pi_1(g^2)))$  and  $(A_k(\pi_2(2)) - A_k(\pi_2(1)), \dots, A_k(\pi_2(g^2)) - A_k(\pi_2(g^2 - 1)))$  as two  $g^2$ -dimensional points (red and blue). Now finding all the  $A_{j,k}$ s that can be sorted by  $(\pi_1, \pi_2)$  is equivalent to finding all pairs consisting of a red and a blue point, where every coordinate of the red point is less than that of the blue point.

For this problem, [GP18] presents a divide-and-conquer algorithm. If the input consists of n d-dimensional points, it runs in  $O\left(\left(\frac{2^{\epsilon}}{2^{\epsilon}-1}\right)^d n^{1+\epsilon} + \text{output size}\right)$  time for all  $\epsilon > 0$ . We omit the details here. Taking  $\epsilon = \frac{1}{2}$ , the sorting step takes  $O\left(\left(g^2\right)!\left(2+\sqrt{2}\right)^{g^2}\left(\frac{2n}{g}\right)^{3/2}+\frac{n^2}{g^2}\right)$  time, since each  $A_{j,k}$  will be correctly sorted once. For the binary search part, since the query entries  $\left[-a_i \in A_{j,k}\right]$  are known before sorting, we can simply process the queries right after  $A_{j,k}$  is found sorted. The time complexity remains  $O\left(\frac{n^2 \log g}{g}\right)$ .

Taking  $g = \frac{1}{2}\sqrt{\frac{\log n}{\log \log n}}$ , the dominating term for the sorting time will be  $\frac{n^2}{g^2}$ , which results in a time complexity of  $O\left(\frac{n^2(\log \log n)^{3/2}}{(\log n)^{1/2}}\right)$ . By [Fre76a], there are actually only  $O(g^{8g})$  possible orders of all  $A_{j,k}$ , so taking  $g = \frac{\sqrt{\log n}}{2}$  reduces the time complexity to  $O\left(\frac{n^2 \log \log n}{(\log n)^{1/2}}\right)$ .

Further optimization from [GP18] reduces the time complexity to  $O\left(\frac{n^2(\log \log n)^{2/3}}{(\log n)^{2/3}}\right)$  by preventing the sorting of the entire  $A_{j,k}$ . This optimization narrows down the region to the space between two contours. A randomized algorithm based on the same approach runs in  $O\left(\frac{n^2(\log \log n)^2}{\log n}\right)$ time. [KLM18] further lowers the decision tree complexity to  $O(n \log^2 n)$ , and the SOTA algorithm [Cha19] runs in  $O\left(\frac{n(\log \log n)^{O(1)}}{\log^2 n}\right)$  time using a geometric approach. These algorithms are more complicated, so we do not delve into the details here.

#### 6 The all-pairs shortest path problem

**Problem 4** (APSP). Given a directed graph G = (V, E) with |V| = n, and assuming there is a weighted edge between every ordered pair of vertices, where the weights are nonnegative real numbers, find the shortest path between each pair of vertices.

The basic  $O(n^3)$  algorithm for this problem is the Floyd-Warshall algorithm [Flo62], which essentially computes the closure of the adjacency matrix. In this case, the addition and multiplication operations in the definition of matrix multiplication are replaced by minimum and addition operations. This is known as min-plus matrix multiplication. [AH74] states that:

The time necessary to compute the closure of a matrix of nonnegative reals, with operations MIN and + playing the role of addition and multiplication of scalars, is of the same order as the time to compute the product of two matrices of this type.

Thus, we only need to focus on the min-plus matrix multiplication problem.

**Problem 5** ((min,+) MM). Given two  $n \times n$  real matrices A, B, calculate matrix C, such that

$$C_{i,j} = \min_{k=1}^{n} \{A_{i,k} + B_{k,j}\}.$$

The first breakthrough in this problem was made by [Fre76b], who developed a decision tree with  $O(n^{5/2})$  complexity and an algorithm with time complexity  $O\left(\frac{n^3(\log \log n)^{1/3}}{(\log n)^{1/3}}\right)$ . The algorithm starts by dividing matrix A into  $b \times m$  stripes and matrix B into  $m \times b$  stripes. The original problem can then be reduced to performing  $\frac{n^3}{b^2m}$  multiplications between  $b \times m$  and  $m \times b$  matrices, alongside  $O\left(\frac{n^3}{m}\right)$  min operations. For a  $b \times m$  matrix  $A_1$  and an  $m \times b$  matrix  $B_1$ , multiplying them needs comparisons between  $a_{i,r} + b_{r,j}$  and  $a_{i,s} + b_{s,j}$  for every  $1 \le i, j \le b$  and  $1 \le r < s \le m$ , where  $a_{\cdot,\cdot}$  and  $b_{\cdot,\cdot}$  denotes the elements of  $A_1$  and  $B_1$ . We can use transposition of terms, to change the values needed for comparison to  $a_{i,r} - a_{i,s}$  and  $b_{s,j} - b_{r,j}$ . So we just sort  $\{a_{i,r} - a_{i,s}\} \cup \{b_{s,j} - b_{r,j}\}$ for every r, s to obtain  $\binom{m}{2}$  lists  $\{L_{r,s}\}$ , and no more information is needed. Sorting requires  $O(m^2b \log b)$  comparison in total, so we can precompute a decision tree of this depth, to obtain the result of all matrix multiplications efficiently, i.e. we can directly know  $\operatorname{argmin}_{k=1}^m \{a_{i,k} + b_{k,j}\}$ without enumerating through k. [Fre76b] further notes that the information complexity can be reduced. Consider a 2mbdimensional space, where each dimension corresponds to an element of  $A_1$  or  $B_1$ . We construct  $\binom{m}{2}\binom{2b}{2} = \Theta(m^2b^2)$  hyperplanes:  $a_{i,r} - a_{i,s} = a_{i',r} - a_{i',s}$ ,  $b_{s,j} - b_{r,j} = b_{s,j'} - b_{r,j'}$ , and  $a_{i,r} - a_{i,s} = b_{s,j} - b_{r,j}$ . Then, each possible order corresponds to a region. According to [Buc43], an *r*-dimensional space can be partitioned into at most  $O(rn^r)$  regions by *n* hyperplanes, so the information complexity is reduced to  $O(mb \log mb)$ .

[Fre76a] provides an algorithm to construct a decision tree for sorting a sequence of length n, under the constraint that only the orders in a subset  $\Gamma$  of n-permutations are possible. The tree has depth  $\log |\Gamma| + O(n)$ , and its construction can be done in polynomial time in  $|\Gamma|$ . In this case,  $n = \Theta(m^2 b)$ , so the depth is  $O(m^2 b)$ , and the construction time is  $\exp(O(mb \log mb))$ .

Now if  $\Gamma$  can be obtained in time T, then the total time will be

$$O\left(T + \exp(O(mb\log mb)) + n^3\left(\frac{m}{b} + \frac{1}{m}\right)\right).$$

To obtain  $\Gamma$ , we consider sampling at least one point from each region. It can be proven that such points can all be obtained by solving the intersections of hyperplanes:  $a_{i,r} - a_{i,s} =$  $a_{i',r} - a_{i',s} \pm 1$ ,  $b_{s,j} - b_{r,j} = b_{s,j'} - b_{r,j'} \pm 1$ ,  $a_{i,r} - a_{i,s} = b_{s,j} - b_{r,j} \pm 1$ ,  $a_{i,r} = 0$ , and  $b_{j,r} = 0$ . For the intersections, we only need to calculate the  $\{L_{r,s}\}$ s and remove duplicates. Since there are  $O((m^2b^2)^{2mb})$  intersections, T will also be  $\exp(O(mb\log mb))$ .

Finally, we need to choose m and b such that the  $\exp(O(mb\log mb))$  term doesn't dominate, and the term  $\frac{m}{b} + \frac{1}{m}$  is minimized. By choosing  $m = c \frac{(\log n)^{1/3}}{(\log \log n)^{1/3}}$  and  $b = m^2$  for a sufficiently small constant c, we get the time complexity:

$$O\left(\frac{n^3(\log\log n)^{1/3}}{(\log n)^{1/3}}\right).$$

For further improvements, [Tak92] works under the same framework and achieves a time complexity of  $O\left(\frac{n^3(\log\log n)^{1/2}}{(\log n)^{1/2}}\right)$ . The result of  $O\left(\frac{n^3}{\log n}\right)$  was achieved in [Cha08], while [HT12] achieves  $O\left(\frac{n^3\log\log n}{(\log n)^2}\right)$ . The current SOTA result [Wil14], based on tools from circuit complexity, runs in time  $\frac{n^3}{2^{\Omega((\log n)^{1/2})}}$ .

Here we shall mention another very common problem closely related to the problem above:

**Problem 6** ((min,+) Conv). Given two real sequences A, B of length n, compute sequence C of length n, such that

$$C_i = \min_{j+k=i} \{A_j + B_k\}.$$

We make indices start from 0 for alignment. This problem can be reduced to (min,+) MM. By [BCD<sup>+</sup>06], we can construct the following multiplication between a  $\frac{n}{d} \times d$  and a  $d \times n$  matrix:

$$\begin{bmatrix} A_0 & A_1 & \cdots & A_{d-1} \\ A_d & A_{d+1} & \cdots & A_{2d} \\ \vdots & \vdots & \ddots & \vdots \\ A_{n-d} & A_{n-d+1} & \cdots & A_{n-1} \end{bmatrix} \begin{bmatrix} B_0 & B_1 & \cdots & B_{n-1} \\ +\infty & B_0 & \cdots & B_{n-2} \\ \vdots & \vdots & \ddots & \vdots \\ +\infty & +\infty & \cdots & B_{n-d} \end{bmatrix} = \begin{bmatrix} P_{0,0} & P_{0,1} & \cdots & P_{0,n-1} \\ P_{1,0} & P_{1,1} & \cdots & P_{1,n-1} \\ \vdots & \vdots & \ddots & \vdots \\ P_{\frac{n}{d}-1,0} & P_{\frac{n}{d}-1,1} & \cdots & P_{\frac{n}{d}-1,n-1} \end{bmatrix},$$

$$C_i = \min_{j=0}^{\lfloor i/d \rfloor} \{P_{j,i-jd}\}$$

Take  $d = \sqrt{n}$ . If  $n \times n \pmod{(\min, +)}$  matrix multiplication can be done in  $\mathsf{M}(n) = \frac{n^3}{\mathsf{R}(n)}$  time, then the convolution can be done in  $O(\sqrt{n}\mathsf{M}(\sqrt{n})) = O\left(\frac{n^2}{\mathsf{R}(\sqrt{n})}\right)$  time.

# 7 Conclusion

From the algorithms discussed above, we can observe that those following the method of four Russians generally work by identifying the gap between information complexity and actual computational complexity, and try to precompute the results of all possible cases. However, since the number of possibilities grows exponentially with n, they can only divide the input into small groups or pieces and make trade-offs, seeking a balance between precomputation and actual computation. The decision tree complexity cannot be fully realized.

While different approaches to breaking down the model or varying the processing order may lead to significant improvements, most of the algorithms require a careful analysis of information complexity—that is, counting the possible cases. As a result, combinatorics, information theory, and more advanced mathematical tools are increasingly used. Some SOTA methods even rely on tools from circuit complexity, and surprisingly, shaving logs is also related to computation hardness. [AHWW16] shows that

If Edit Distance or LCS on two binary sequences of length n can be solved in  $O(n^2/\log^c n)$  time for every c > 0, then  $\mathsf{NTIME}[2^{O(n)}]$  does not have non-uniform polynomial-size log-depth circuits.

Finally, regarding parallelized bitwise operations, actually every algorithm implicitly or explicitly uses this property. It's needed for encoding and addressing the possibilities, or at least for performing basic operations like additions. Algorithms for BMM and LCS (the first one) explicitly leverage parallelization, and these are the two algorithms actually practical. Other algorithms have large constant factors. To prevent precomputation time from exceeding the desired complexity, the group/block sizes need to be kept very small, and the optimization will barely be effective or may even lead to deoptimization.

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