HAKUBI RESEARCHERS' ACTIVITIES IN ICR

Hakubi Project: Fosterage and Support of Young Researchers, Kyoto University **Research Topic**

Algorithmic Graph Theory with Applications to Bioinformatics



Host Laboratory Laboratory of Mathematical Bioinformatics Host Professor AKUTSU, Tatsuya

Program-Specific Assoc Prof JANSSON, Jesper (Ph D)

Outline of Research

One of my research topics this year is fast matrix multiplication. Given two square matrices A and B of size (n x n) with nonnegative integer entries, the naive algorithm for computing the matrix product AB runs in $O(n^{3})$ time. There exist algorithms that run in substantially subcubic time, e.g., a very recent one due to F. Le Gall uses $O(n^{2}.3728639)$ time, and a major open question in Theoretical Computer Science is whether it can be done in quadratic time. We have developed a new technique based on interpreting matrices as 3D histograms. To multiply A and B, we decompose their 3D histograms into 3D blocks which are then manipulated in a pairwise manner using the interval tree data structure. This leads to an $O^*(n^{2} + rs)$ -time algorithm for matrix multiplication, where r and s denote the minimum number of 3D blocks into which A and B can be partitioned, respectively. In other words, whenever A and B admit a partition into a small number of 3D blocks, our algorithm is very efficient.