

Weak Kernels and Their Applications

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Traditional NP-completeness theory [3, 8, 7] and FPT algorithms theory [4, 6] focus on *decision* problems. Valiant first raised the question on whether *search* problems and decision problems are equivalent [9], which leads to a small branch in traditional complexity theory. The notable conclusion is that even for problems in NP, they are not equivalent, i.e., under a complexity assumption, there is an associated search problem ρ in NP which cannot be reduced to its corresponding decision problem [2].

In the FPT theory, *kernelization* (also viewed as *data reduction*), is about reducing problem size and is one of the most fundamental techniques in the area [4, 5]. In this work, we formalize a folklore concept and formally define *weak kernels* for (NP-hard) *search* problems, which is about search space reduction and stands as a new generic technique for designing FPT algorithms. (The idea of designing FPT algorithms for search problems was used by Bosma et al. as early as in 2003 [1].) We show that weak kernels are different from the (traditional) kernels for decision problems, by exhibiting an example out of P such that its decision version has no kernel while the equivalent search problem has a weak kernel.

As this work was motivated by applications in computational biology, we show a few applications of weak kernels on some of these problems, for which a traditional kernelization seems hard to apply. The three applications are: Sorting by Minimum Unsigned Reversals, Minimum Co-Path Set and Sorting by Minimum Unsigned DCJ Distance. Among them, we present the first FPT algorithm for the famous Sorting by Minimum Unsigned Reversals problem. While the three applications are mainly from genome analysis, hopefully new applications will follow later on.

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