# High-Performance and Mobile Computing Platform for Drug Comparisons on NVIDIA Jetson TK1

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### A. Background and Motivation

Compound comparison is an important task for computational chemistry. In most of works, molecules can be represented as fingerprints and *SMILES*, and then the work of compound comparison can be seen as the *string comparison*. However, it will be time-consuming when comparing with a large amount of compounds, such as ZNIC and GDB-13 database.

#### **B.** Goal and Method

We proposed a GPU-based parallel algorithm for multiple compound comparisons (O2A and A2A) on NVIDIA Jetson TK1. In this algorithm, the goal is to compare two sets of compounds listed as *Query* and *Database* at first, and then find the compounds in *Database* with more than 0.85 Tanimoto coefficient for each compound in *Query*.

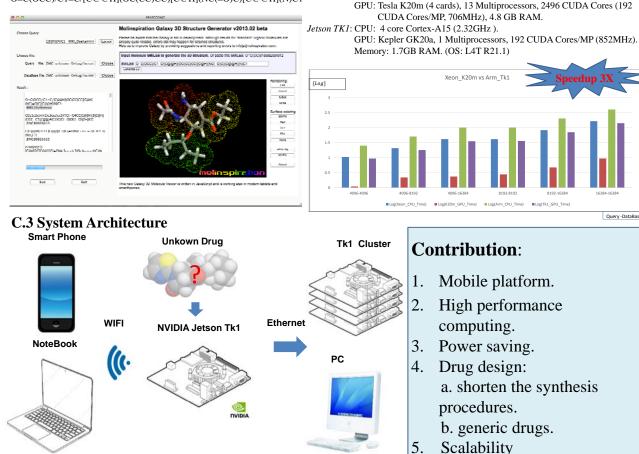
For each compound in *Query* and *Database*, it should be fragmented into a set of *q*-Lingos, respectively. a *preprocessing phase* is designed to do this procedure on CPU. After this phase, *a GPU implementation of comparison phase* is designed in order to accelerate the computation speed. In our algorithm, a compound in *Query* will be used to compare with all compounds in *Database* by all threads on GPUs when executing the kernel function once. A load-balancing strategy by considering four LINGO types can be used to accelerate the computation speed for single GPU and multiple GPUs. All of compounds in *Database* with more than 0.85 Tanimoto coefficient for each compound in *Query* are reported in the *output phase* on CPU.

### **C. Results**

#### C.1 User interface

*Oseltamivir (Tamiflu)* SMILE:

O=C(OCC)/C1=C/[C@@H](OC(CC)CC)[C@H](NC(=O)C)[C@@H](N)C



**C.2 Experimental Results** 

(s1) ten thousand compounds in *Query* and *Database*, respectively, (s2) thirty thousand compounds in *Query* and *Database*, respectively.

Machine 1: CPU: Intel Xeon E5-2650 2.0GHz, 32 processors, 125.87GB RAM.

Two test sets:

Environment:

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