# GPU Parallelization of Massive Sample-size Survival Analysis

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## Introduction

#### Large-scale Observational Data

- Observational databases have millions of individuals [1] with thousands of patient characteristics and up to 10 years of data per life [2].
- Resource for comparative effectiveness and safety study
- Survival analysis is a main statistical method in comparative effectiveness and safety study.
- Problem: computational burden.

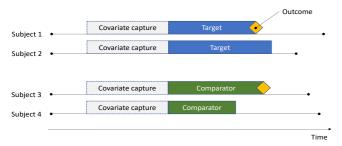


Figure 1: Comparative effectiveness study with cohort design.

#### **GPU Parallelization**

- Graphics processing units (GPUs) contain thousands of processor cores that can apply the same numerical operations simultaneously to elements of large data arrays under a "Single Instruction, Multiple Threads" (SIMT) programming paradigm.
- GPUs are relatively inexpensive, easy-to-use hardware that offers impressive potential for speeding up computations.

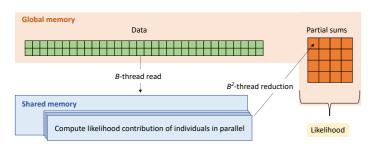


Figure 2: Massive parallelization strategy for computing the likelihood.

## Methods

### Cox Proportional Hazards Model

- · The hazard model formula depends on a baseline survival function and a set of explanatory variables.
- Parameter estimation of the Cox proportional hazards model follows from the logpartial likelihood.

$$h(t|\boldsymbol{\beta}) = h_0(t|\boldsymbol{\beta}) \exp\left(\boldsymbol{\beta}^T \boldsymbol{x}\right)$$

$$l_{partial}(\pmb{\beta}) = \sum_{i=1}^n \delta_i \left\{ \pmb{\beta}^T \mathbf{x}_i - \log \left[ \sum_{k \in R(Y_i)} \exp \left( \pmb{\beta}^T \mathbf{x}_k \right) \right] \right\}$$

#### Fine-Gray Sub-distribution Proportional Hazards Model

- The Fine-Gray model generalizes the Cox proportional hazards model to competing risks time-to-event data that consists of more than one type of events.
- Competing risks arise when individuals can experience more than one type of event and the occurrence of one type of event will prevent the occurrence of the others.

#### Massive Parallelization for Parameter Estimation with Prefix Sums and Reduction

- We identify prefix sums [3] and reductions in log-partial likelihood of Cox model and Fine-Gray model due to the cumulative structure of the risk set.
- We avoid unnecessary memory transactions by fusing prefix sums and reductions operations in likelihood calculations into a single kernel.
- We minimize data movements by exploiting the sparsity of the design matrix.

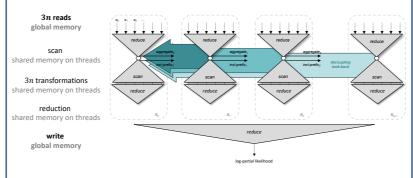


Figure 3: Fused kernel for maximum likelihood estimation

# Results



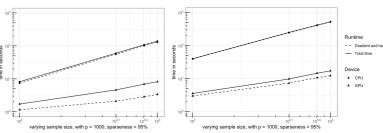


Figure 4: GPU vs CPU runtimes for Cox model (left) and Fine-Gray model (right).

#### Antihypertensive drug classes study

- Dataset: thiazide or thiazide-like diuretics (THZ) and angiotensin-converting enzyme inhibitors (ACEi) cohort study, reproduction of [4] in CCAE dataset.
- Cohort size: 1,065,745 hypertension patients and 7891 covariates.
- Outcome of interest: major cardiovascular events.
- GPU parallelization reduce the time of parameter estimation from 16 hours on multicore CPU to less than one hour.

## Conclusions

- We implement the massive parallelization of the Cox proportional hazards model and Fine-Gray model by using NVIDIA's CUB library for parallel computing of prefix sums and exploiting the sparsity of data.
- By saving data movement and clever manipulation of likelihood structure, our parallelization significantly reduces the runtime of large-scale survival analysis.

#### References

[1] G. Hripcsak, P. B. Ryan, J. D. Duke, N. H. Shah, R. W. Park, V. Huser, M. A. Suchard, M. J. Schuemie, F. J. DeFalco, A. Perotte, et al., "Characterizing treatment pathways at scale using the ohdsi network," Proceedings of the National Academy of Sciences, vol. 113, no. 27, pp. 7329-7336, 2016.

[2] M. A. Suchard, S. E. Simpson, I. Zorych, P. Ryan, and D. Madigan, "Massive parallelization of serial inference algorithms for a complex generalized linear model," ACM Transactions on Modeling and Computer Simulation (TOMACS), vol. 23, no. 1, p. 10, 2013.

[3] D. Merrill and M. Garland, "Single-pass parallel prefix scan with decoupled look-back," NVIDIA, Tech. Rep. NVR-

[4] M. A. Suchard, M. J. Schuemie, H. M. Krumholz, S. C. You, R. Chen, N. Pratt, C. G. Reich, J. Duke, D. Madigan, G. Hripcsak, et al., "Comprehensive comparative effectiveness and safety of first-line antihypertensive drug classes: a systematic, multinational, large-scale analysis," The Lancet, vol. 394, no. 10211, pp. 1816-1826, 2019.