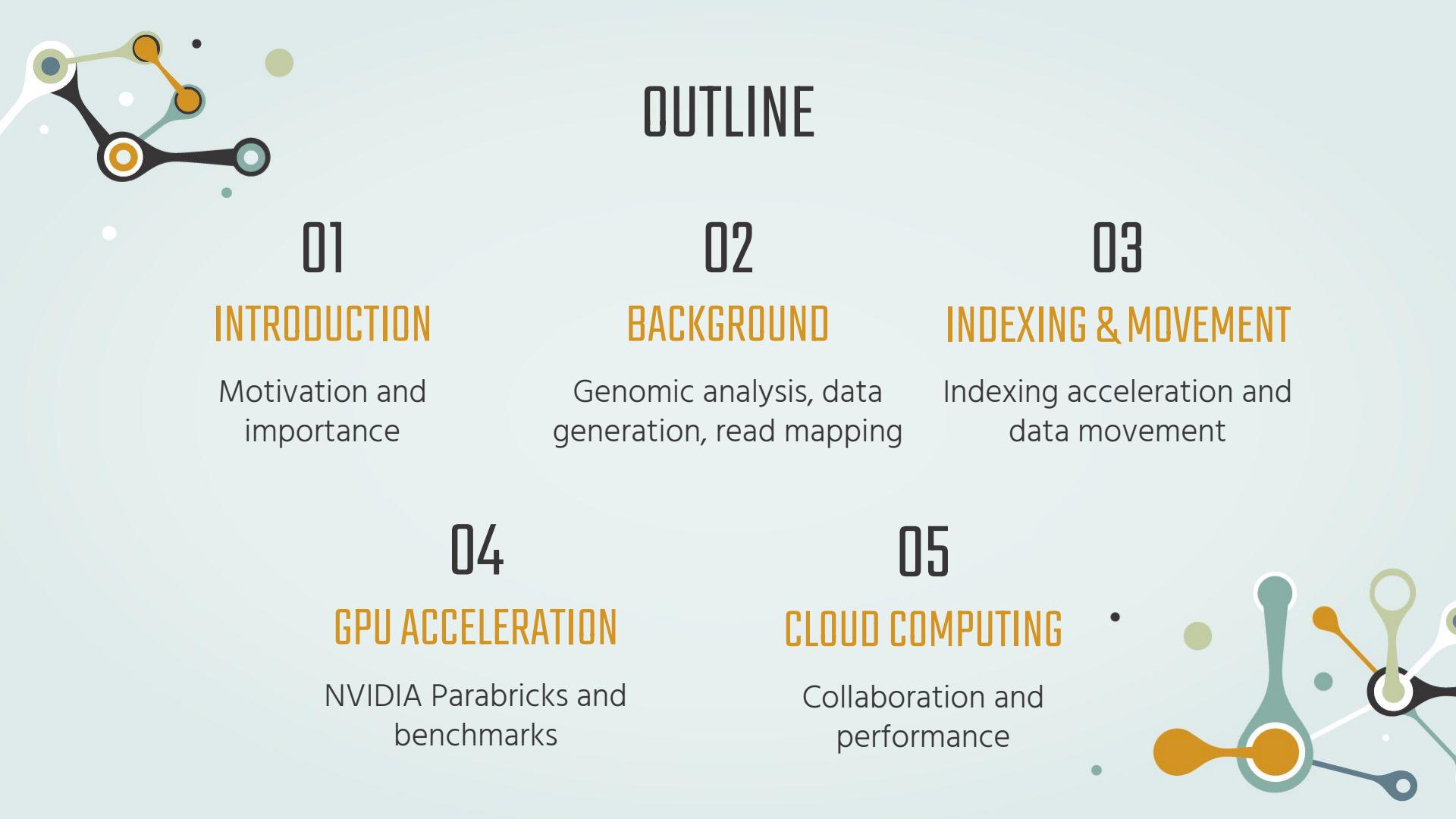


High-Performance Computing in Precision Medicine

Maksym Turkot



OUTLINE

01

INTRODUCTION

Motivation and importance

02

BACKGROUND

Genomic analysis, data generation, read mapping

03

INDEXING & MOVEMENT

Indexing acceleration and data movement

04

GPU ACCELERATION

NVIDIA Parabricks and benchmarks

05

CLOUD COMPUTING

Collaboration and performance



01

INTRODUCTION



Introduction

- HPC in Precision Medicine
- Shifting Bottleneck in Genomic Research
- Acceleration Techniques
 - Algorithmic (indexing)
 - Hardware (GPUs)
- Cloud Computing in Genomic Workflows

BACKGROUND

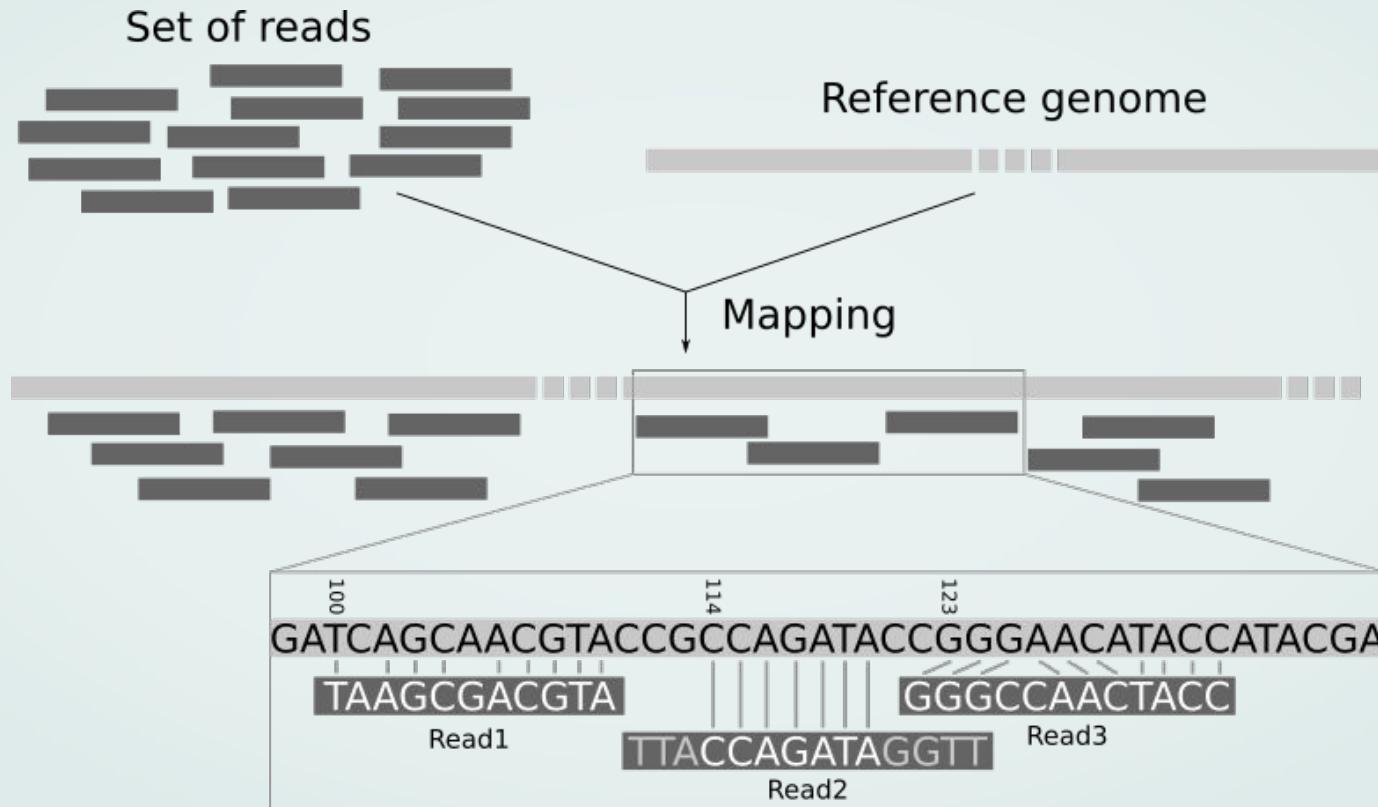


Obtaining Genomic Data

- Generation
 - Machine-based, sequencer
- Download
 - Prepared data
- Simulation
 - Mimic sequencer. Useful



Read Mapping

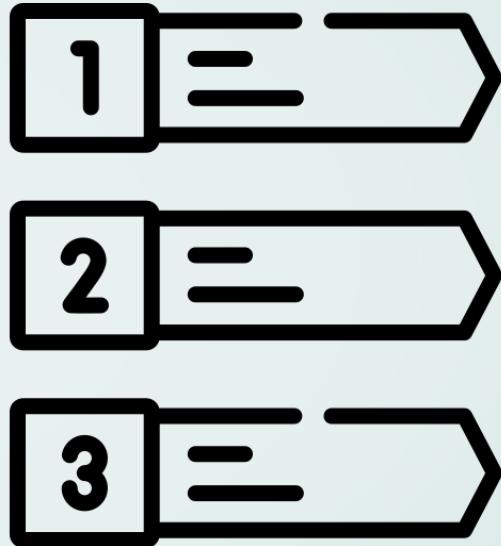


INDEXING & MOVEMENT



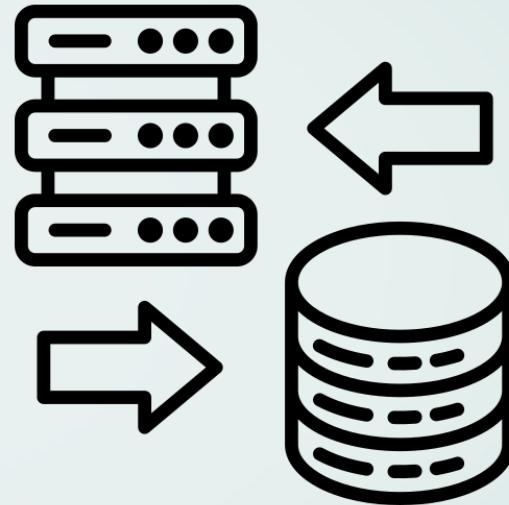
Indexing and Seeding Acceleration

- Indexing and Seeding
- Indexing Process
- Impact of Seed Characteristics
- Tradeoff and Improvements



Data Movement Reduction

- Memory-Intensive Tasks
- Processing-in-Memory (PIM)
- Alternative Approaches
- 3D ReRAM-based Memory



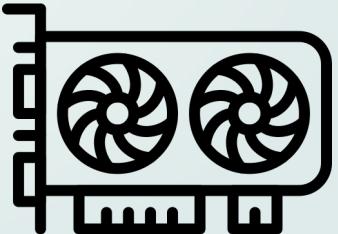
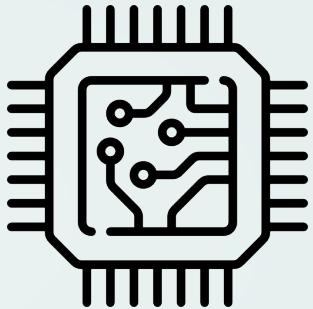


04

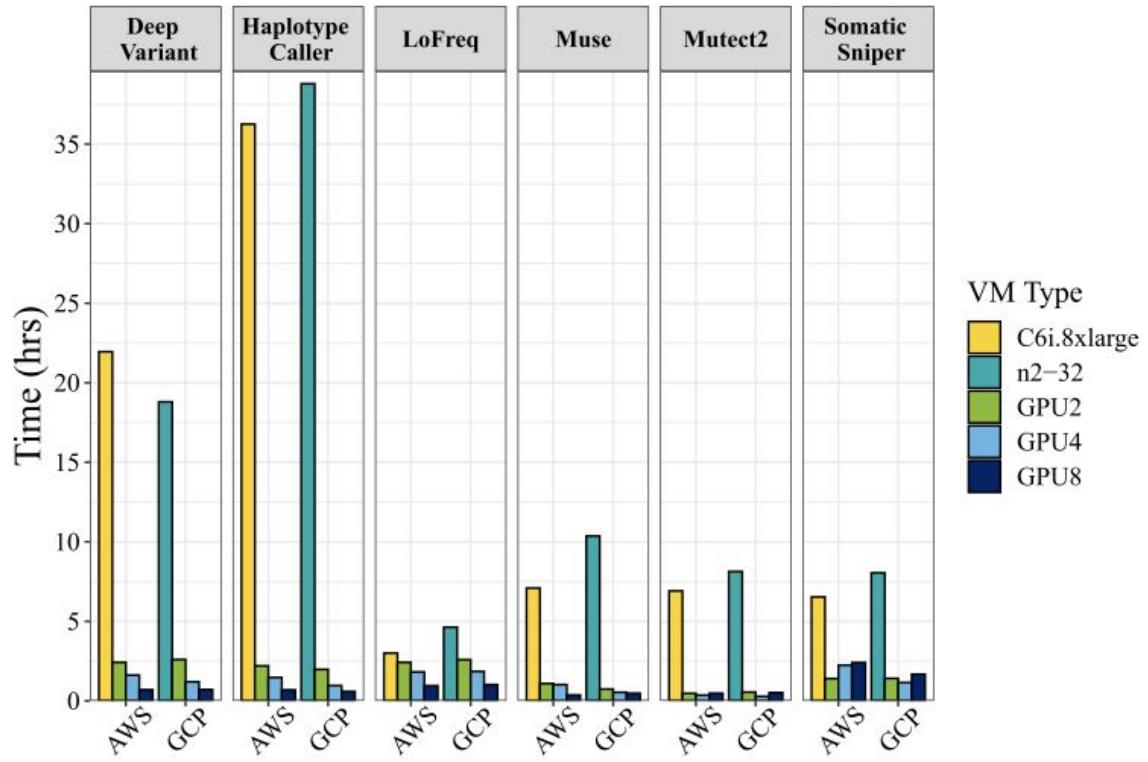
GPU ACCELERATION

GPU Acceleration

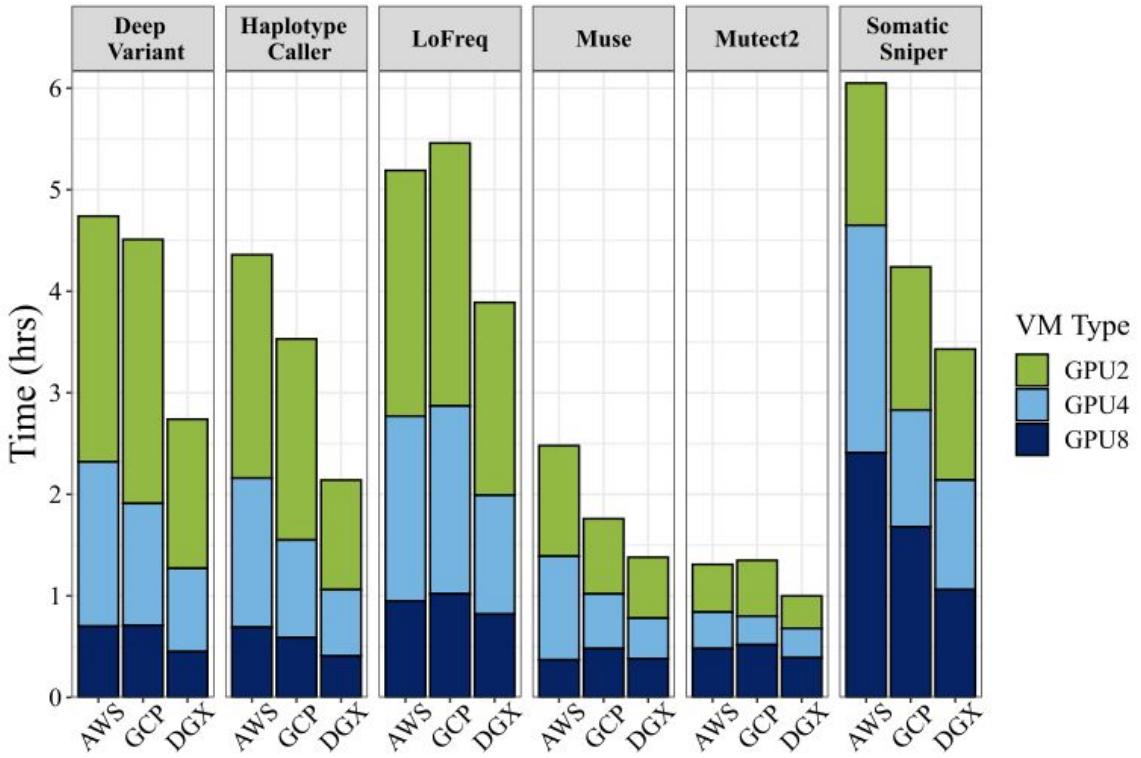
- Data Analysis Bottleneck
- NVIDIA Parabricks
- Varying CPU performance
- Massive GPU acceleration



Runtimes of All Cloud-Based Analyses



GPU Runtimes Across Platforms



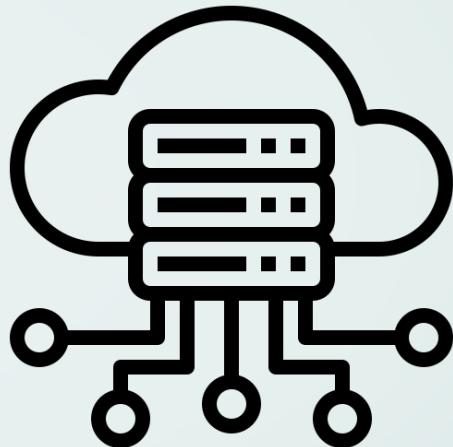


05

CLOUD COMPUTING

Cloud Computing

- An Invaluable Tool
- Reproducibility & Accessibility
- Simplified setup
- Collaboration
- Advantage in genome work
- Scaling





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Thank You!

Question time

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