

LISA WU WILLS, DUKE UNIVERSITY

**GENESIS: A COMPOSABLE HARDWARE
ACCELERATION FRAMEWORK FOR GENOME
ANALYSIS**



My Research Goal #1: **Design and deploy highly efficient domain-specific hardware accelerators to advance state-of-the-art emerging applications.**

My Research Goal #2: **Democratize end-to-end accelerated system development and deployment while leveraging hardware acceleration to advance emerging application domains in the interdisciplinary area of computer architecture and healthcare.**

FPGA ACCELERATED INDEL REALIGNMENT IN THE CLOUD

[HPCA 2019]



LISA WU, DAVID BRUNS-SMITH, FRANK NOTHAFT, QIJING HUANG, SAGAR KARANDIKAR, HOWARD MAO, BRENDAN SWEENEY, KRSTÉ ASANOVIC, DAVID PATTERSON, AND ANTHONY JOSEPH

GENESIS: A HARDWARE ACCELERATION FRAMEWORK FOR GENOMIC DATA ANALYSIS [ISCA 2020, IEEE MICRO TOP PICK 2021]



서울대학교
SEOUL NATIONAL UNIVERSITY

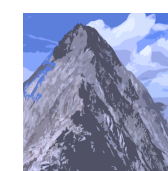


Berkeley
UNIVERSITY OF CALIFORNIA



Duke
UNIVERSITY

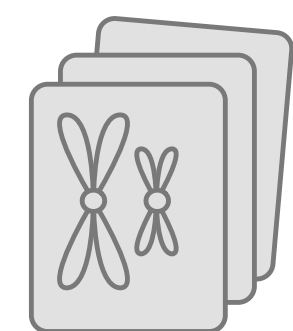
TAE JUN HAM, DAVID BRUNS-SMITH, BRENDAN SWEENEY, YEJIN LEE, SEONG HOON SEO, U GYEONG SONG, YOUNG H OH, KRSTÉ ASANOVIC, JAE W LEE, AND LISA WU WILLS



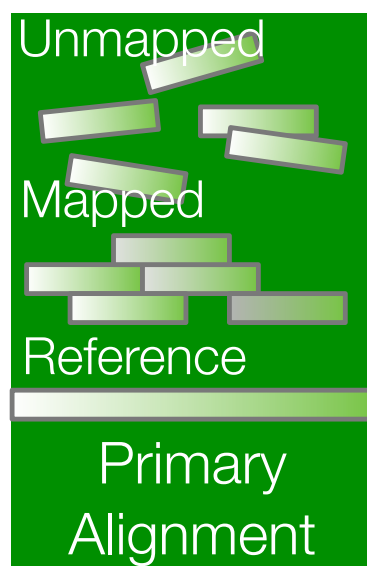
GOAL OF GENOMIC ANALYSIS:

Identify the **nucleotide differences (or variants)** between an individual genome and the reference genome at a given position, with **acceptable accuracy**.

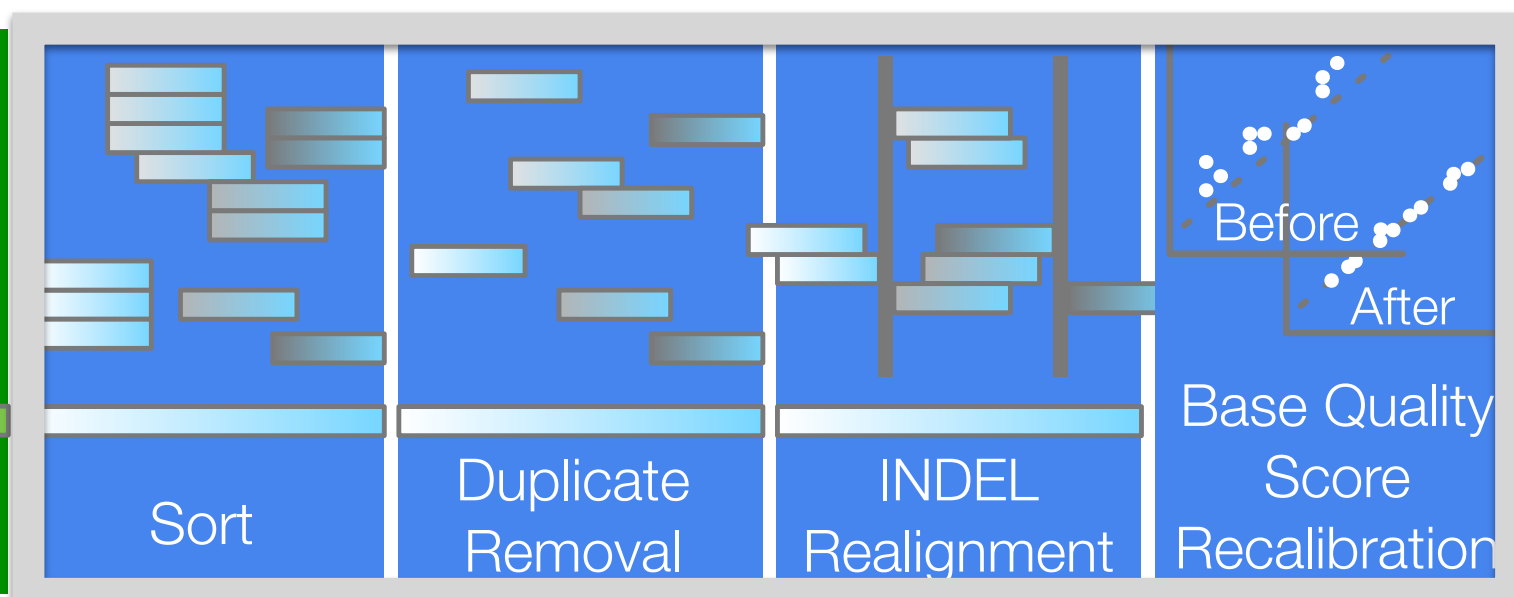
ALIGNMENT REFINEMENT IS THE SLOWEST PIPELINE



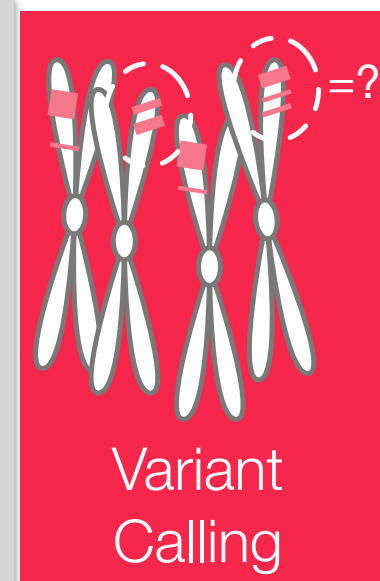
Raw
Sequence
Reads



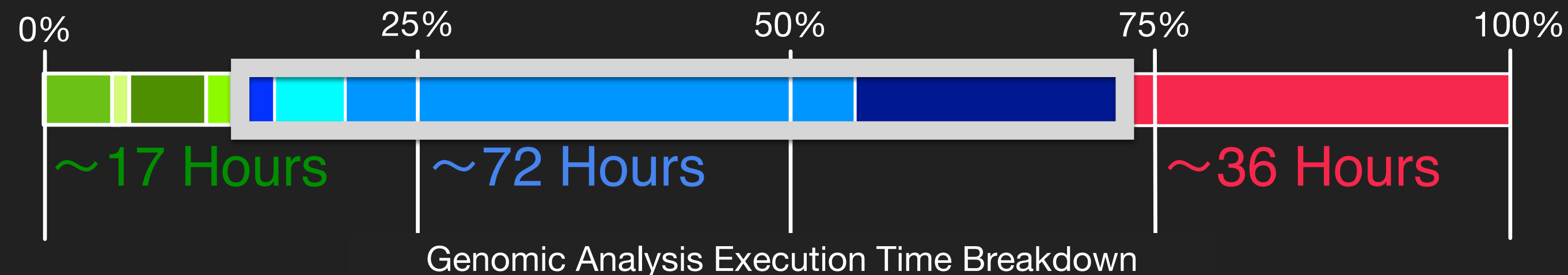
Read Mapping
14%



Alignment Refinement
58%



Variant Calling
28%



WHAT DOES DATABASE ANALYTICS AND GENOMIC ANALYSIS HAVE IN COMMON?

Generic Data Manipulations (i.e. aggregation, sorting)!



Domain-Specific Language: SQL+



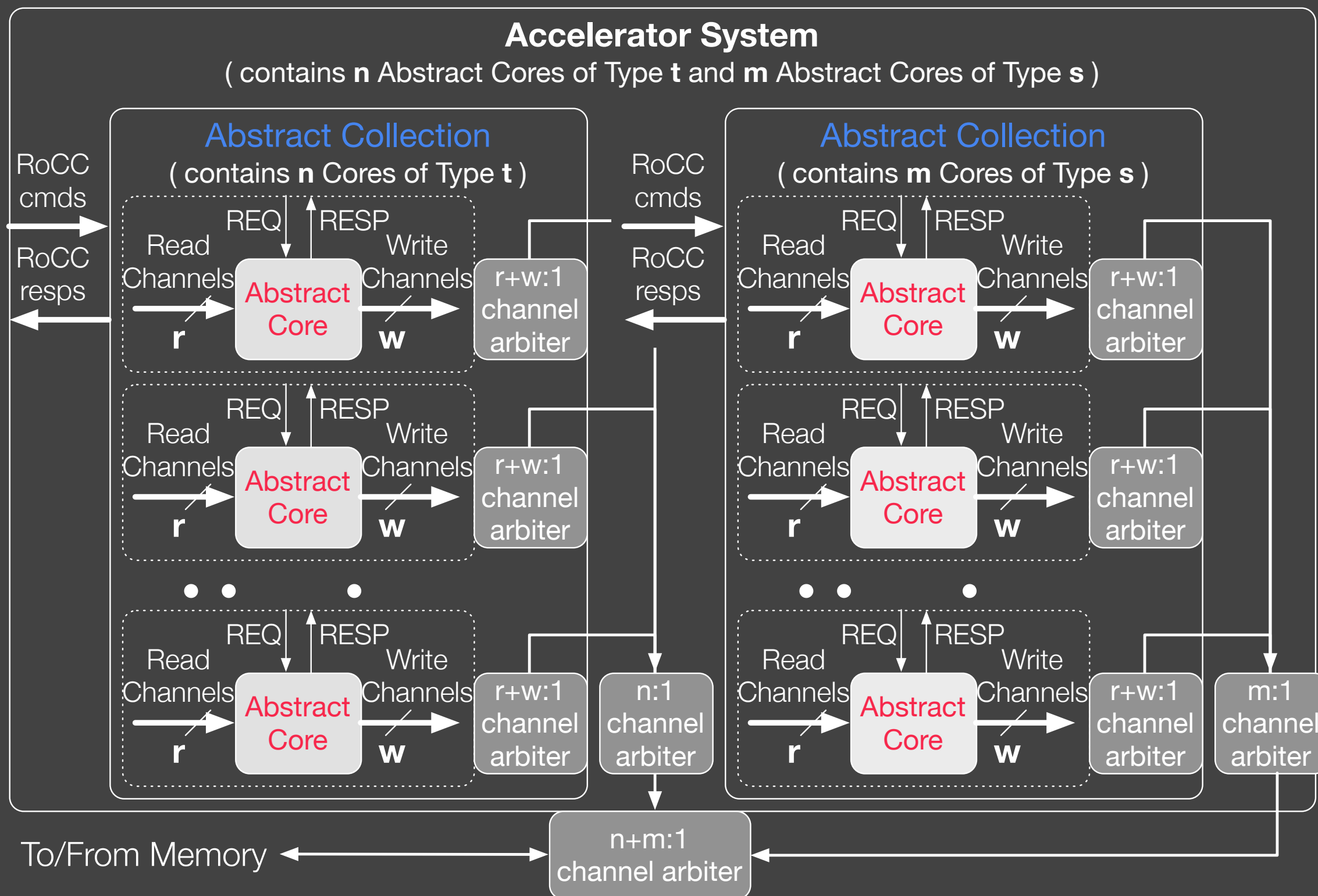
Genomic Hardware Library



Accelerator Composer

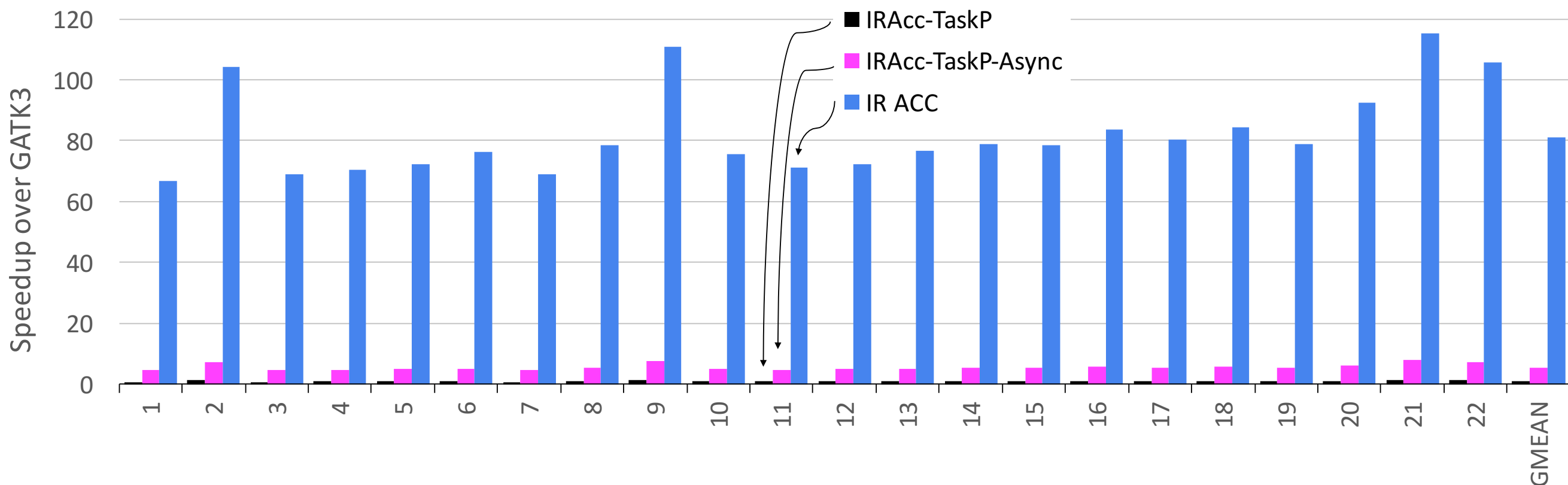


Accelerator Composer Design Templates



OUR ACCELERATED IR SYSTEM PERFORMS **81X** BETTER THAN SOFTWARE RUNNING 8 THREADS

Hardware Accelerated INDEL Realignment Performance Compared Against Software



42 hours —> roughly 30 minutes

IMPACTS/SIGNIFICANCE

ACCELERATING **COMMON PRIMITIVES ACROSS DOMAINS** ALLOWS THE SHARING, REUSING, AND COMPOSITION OF ACCELERATED SYSTEMS ACROSS DOMAINS, LOWERING DEVELOPMENT EFFORT.

IMPACTS/SIGNIFICANCE

LEVERAGE AN **ALREADY-STANDARDIZED LANGUAGE** AS THE DSL AND CONSTRUCT **PRIMITIVE OPERATORS** THAT DIRECTLY MAP **SOFTWARE PRIMITIVES** TO **HARDWARE BLOCKS** PRODUCES EFFICIENT ACCELERATED SYSTEMS.

IMPACTS/SIGNIFICANCE

THIS DEVELOPMENT METHODOLOGY CAN BE
ADOPTED FOR VARIOUS DOMAINS BEYOND DATABASE
AND GENOMIC ANALYTICS.

