

PAUL SCHERRER INSTITUT



Filip Leonarski :: MX Data :: Swiss Light Source

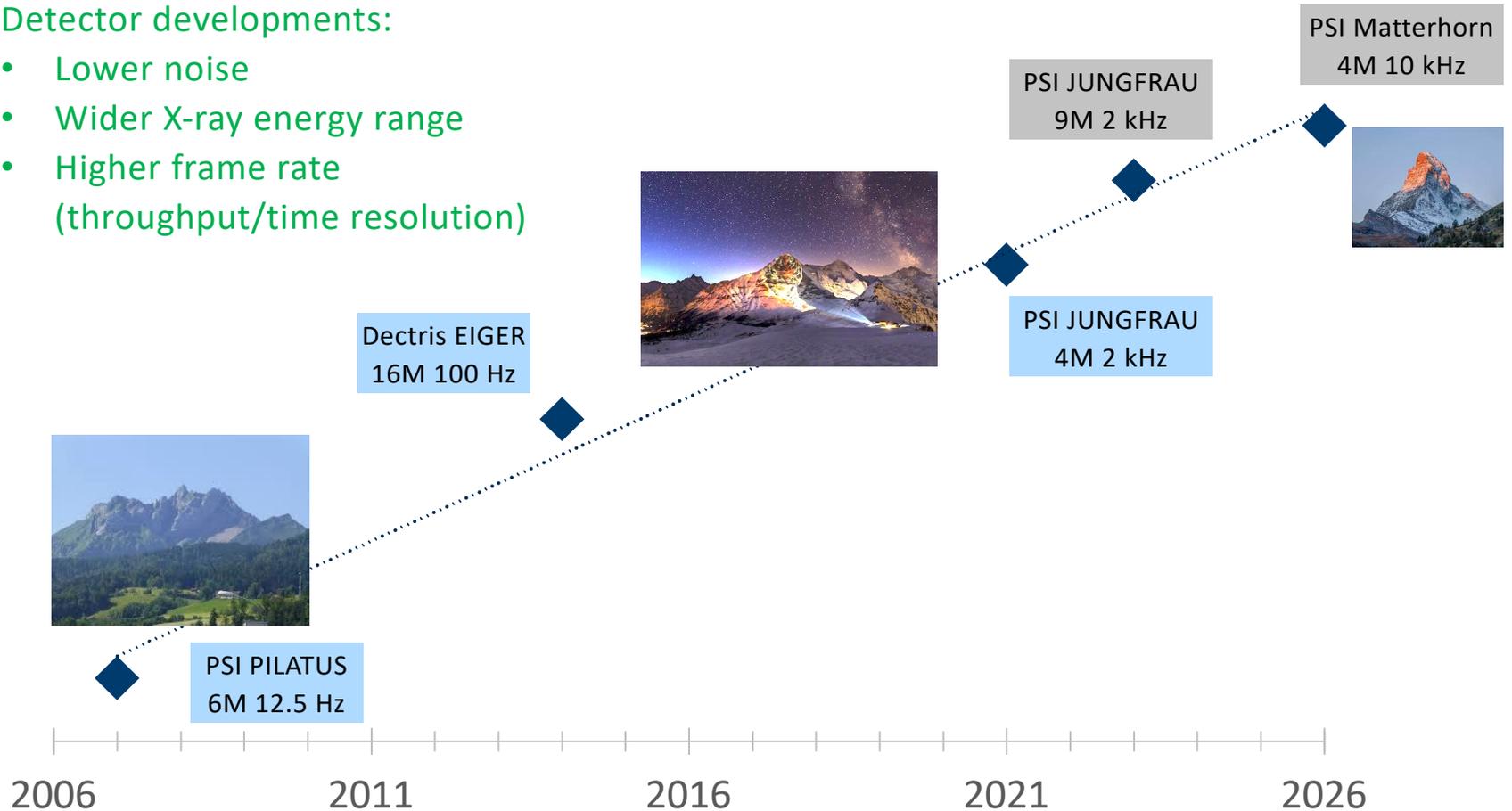
Jungfrauoch: kilohertz data acquisition and on-the-fly image analysis for MX beamlines

LEAPS Innov WP7 Seminar
13 April 2022

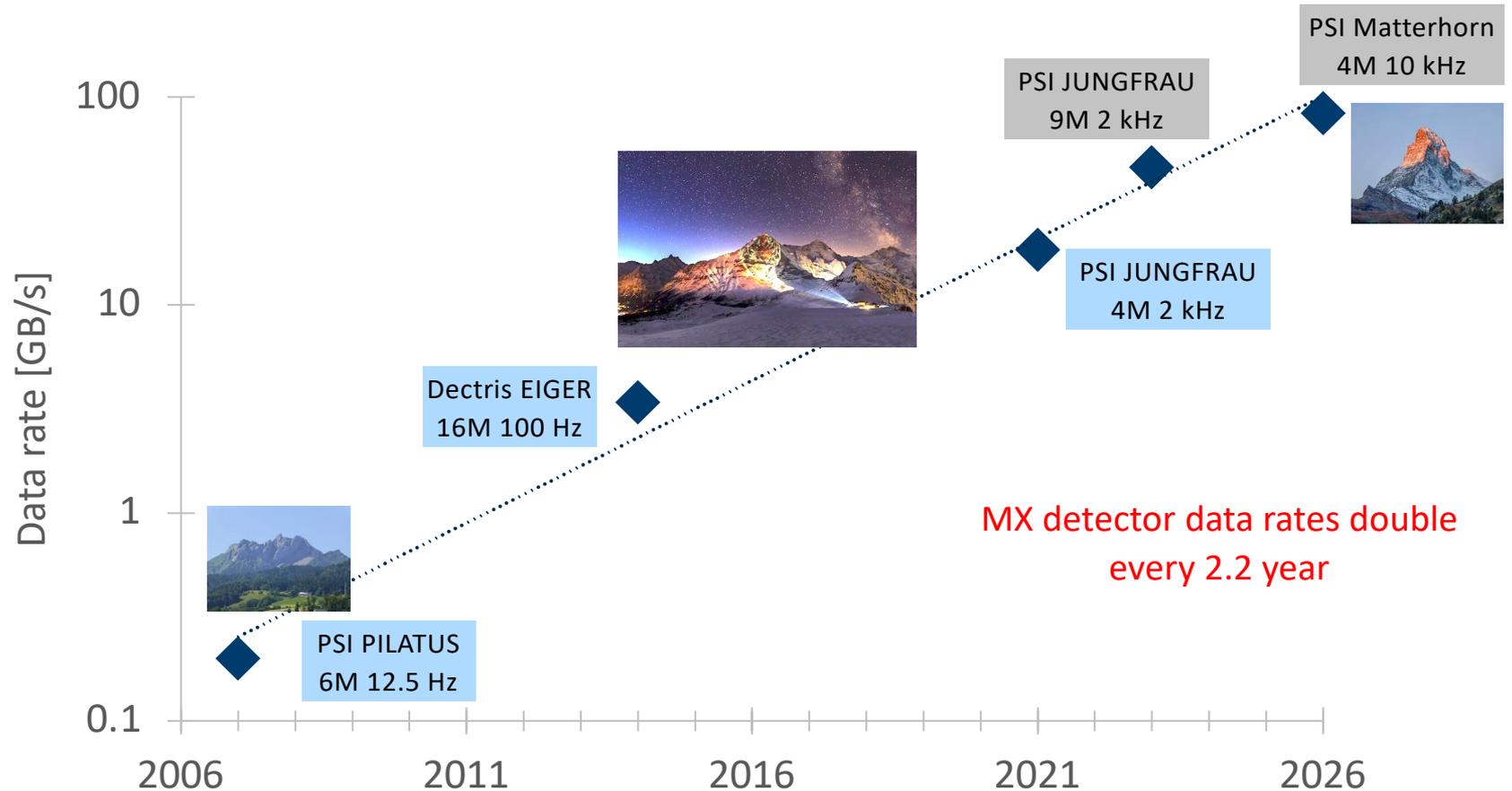
MX development is driven by X-ray detectors

Detector developments:

- Lower noise
- Wider X-ray energy range
- Higher frame rate (throughput/time resolution)



Detector development is challenging for IT infrastructure

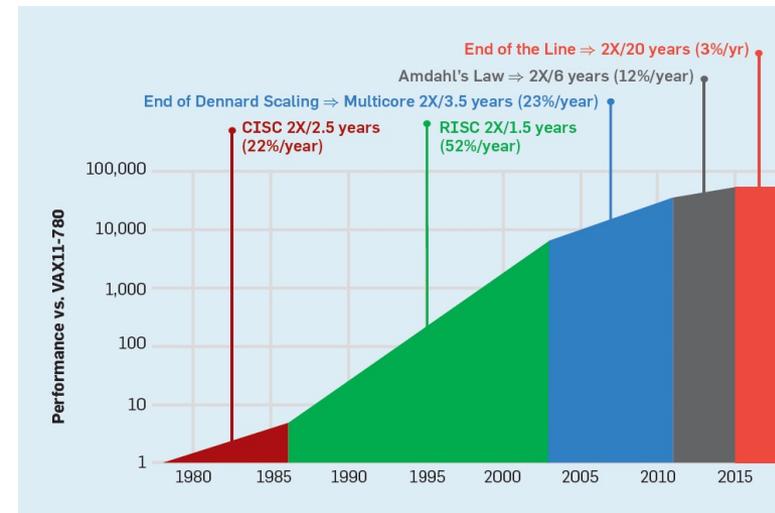


MX detector data rates double every 2.2 year

“JUNGFRAU detector for brighter x-ray sources: Solutions for IT and data science challenges in macromolecular crystallography”
 Leonarski F., et al, *Structural Dynamics* (2020)

Solution to data challenge

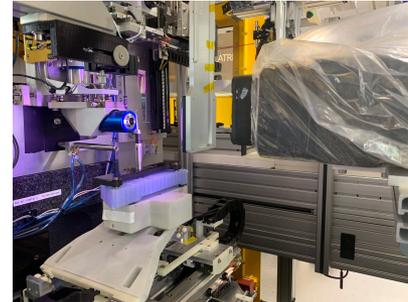
- Computing challenge
 - Mainstream computing is not growing exponentially anymore
 - Massive parallelization (multiple servers) of software-based solution possible, but not sustainable in long run
 - Data movement (memory/network) is the main issue
- Development needed
 - **Reduce data close to source**
 - Combine acquisition and on-the-fly processing
 - Use task-specific computing architectures



Hennessy & Patterson
doi:10.1145/3282307

Science application: JUNGFRAU detector for synchrotron MX

- JUNGFRAU applications
 - Native-SAD (3-6 keV)
 - High-flux (time resolved/high throughput)
 - Room temperature
- JUNGFRAU operational requirements
 - To reach >95% duty cycle at a synchrotron source, one needs to operate JUNGFRAU at kHz frame rates (at least internally)
 - Raw ADUs need to be converted to photons for analysis and compression



Test at **VMXi Diamond Light Source (UK)**

nature|methods

ARTICLES

<https://doi.org/10.1038/s41592-018-0143-7>

Fast and accurate data collection for macromolecular crystallography using the JUNGFRAU detector

Filip Leonarski¹, Sophie Redford¹, Aldo Mozzanica¹, Carlos Lopez-Cuenca¹, Ezequiel Panepucci¹, Karol Nass¹, Dmitry Ozerov¹, Laura Vera¹, Vincent Olieric¹, Dominik Buntschu¹, Roman Schneider¹, Gemma Tinti¹, Erik Froejdh¹, Kay Diederichs², Oliver Bunk¹, Bernd Schmitt^{1*} and Meitian Wang^{1*}

SwissFEL



10 μ s

SLS



980 μ s

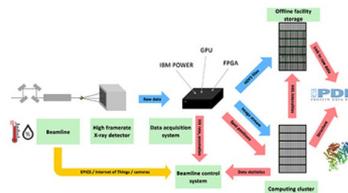
A. Mozzanica

Jungfrauoch development story

- 2017: JUNGFRAU detector demonstration for synchrotron MX
- 2018: Jungfrauoch backend development started (technology exploration)
- 2019: Collaboration with IBM started
- 2020: First operation after 6 months of development at X06SA
- 2020: Tested JF 4 Mpixel 1 kHz at BL-1A, Photon Factory KEK (native-SAD in helium at 4 keV; rotation crystallography)
- 2021: Tested JF 4 Mpixel 2 kHz at BioMAX, MAX IV (time resolved serial crystallography)



Test at **BL-1A Photon Factory KEK (JP)**



Open . Jan 1, 2020 . 1 Citations

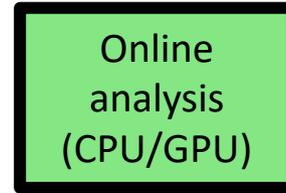
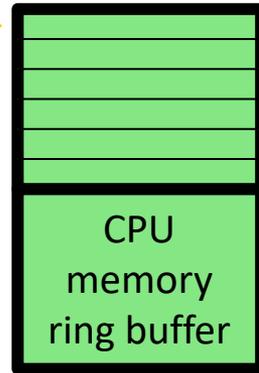
JUNGFRAU detector for brighter x-ray sources: Solutions for IT and data science challenges in macromolecular crystallography

Filip Leonarski, Aldo Mozzanica, Martin Brückner, Carlos Lopez-Cuenca, Sophie Redford, Leonardo Sala, Andrej Babic, Heinrich Billich, Oliver Bunk, Bernd Schmitt and Meitian Wang

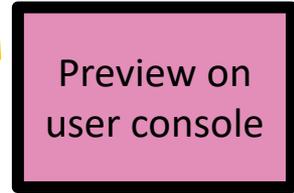
Structural Dynamics 7, 014305 (2020); <https://doi.org/10.1063/1.5143480>

Jungfrauoch general architecture

Eiger Mönch Jungfrau



Work in progress

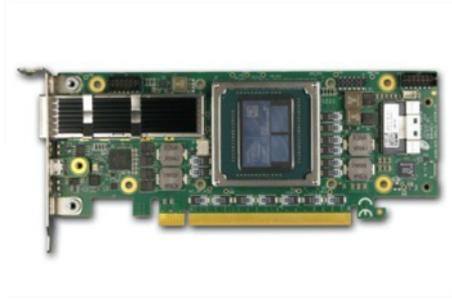


1 card = 12 GB/s
(~2.5 Mpixel JF)
4 cards possible
(up to 48 GB/s)



Jungfrauoch server (IBM IC922/POWER9)

FPGA board with OpenCAPI interface



- Data acquisition
- Initial data analysis
- Pre-compression
(2.5 Mpixel/board for JF)

Ethernet
UDP/IP

Dark
current

Conversion

Pixel mask

Strong
pixel finder

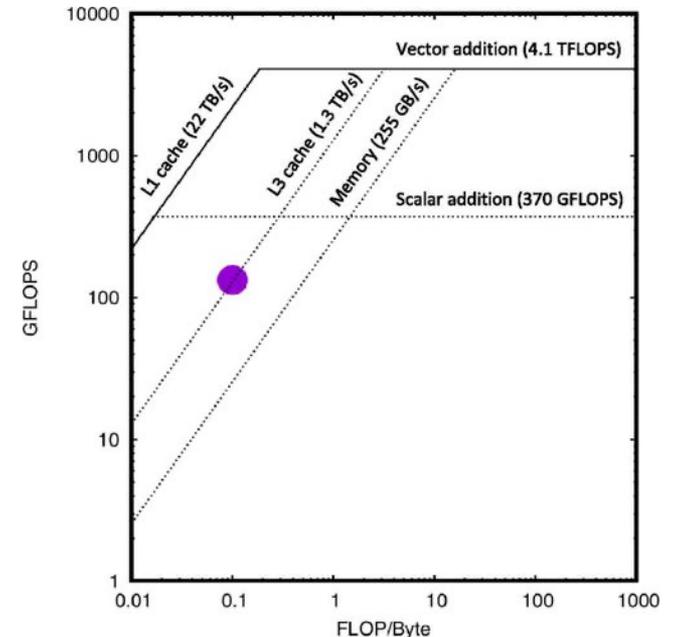
Bitshuffle

Memory
writer

Functionality described in C++ and
transcompiled with high-level
synthesis to hardware description
language



- Network
 - Ethernet/IP/UDP stack
 - ARP and ICMP for integration
 - Can be reused for different UDP data source
- Pixel conversion for JUNGFRAU
 - 460 GB/s high-bandwidth memory is crucial (memory BW problem on CPU)
 - Pedestal tracking possible
- Pre-compression/compression
 - Bitshuffle filter trivial on FPGA
 - Any transformations need to be done before (multipixel/spot finding)
 - Compression must happen on CPU (at the moment)



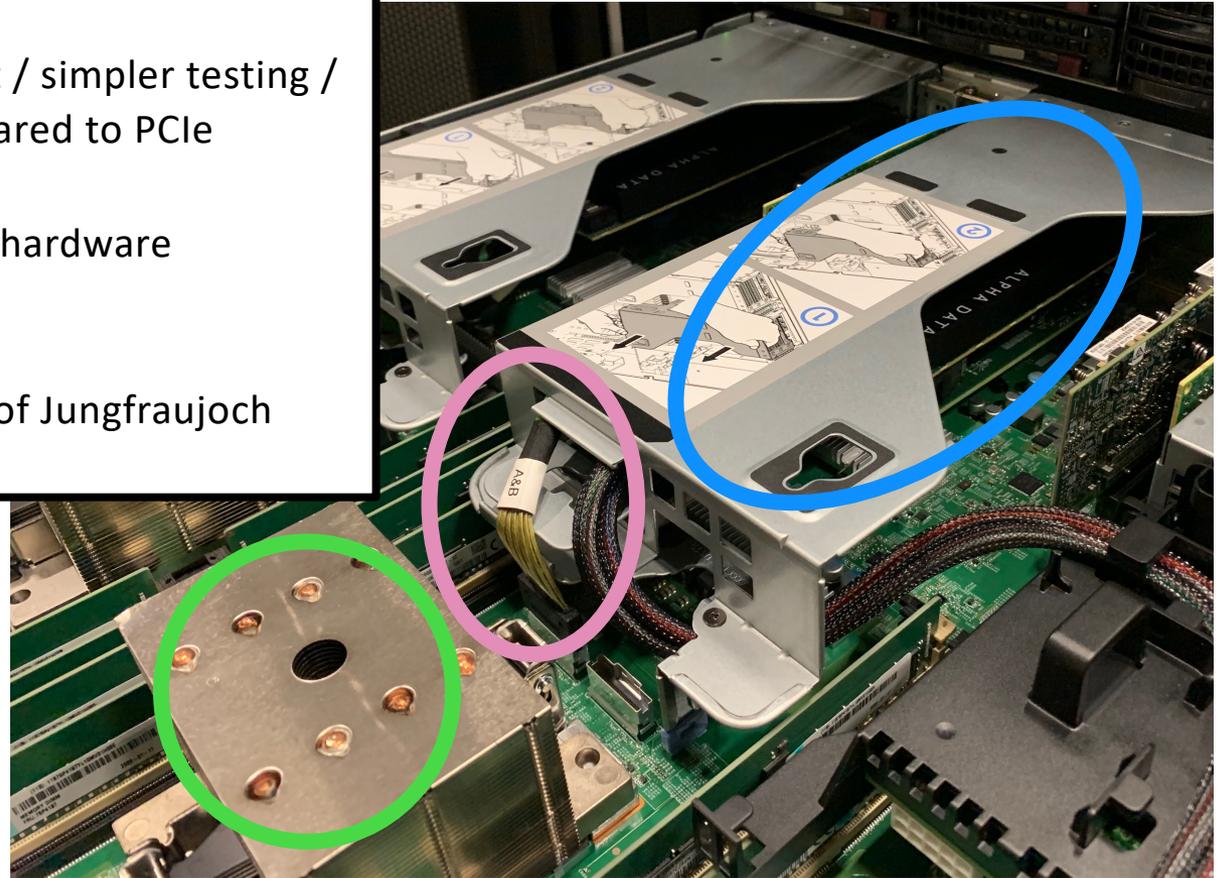
CPU performance (roofline plot)

See: “JUNGFRAU detector for brighter x-ray sources: Solutions for IT and data science challenges in macromolecular crystallography” Leonarski et al. Structural Dynamics (2019)

<https://doi.org/10.1063/1.5143480>

OpenCAPI interface

- Coherent interconnect => FPGA has access to virtual memory, no kernel level development necessary
- Cheaper development / simpler testing / better safety as compared to PCIe
- Requires IBM POWER hardware (or Intel CXL soon)
- Experimental version of JungfrauJoch works with PCIe



**POWER9
CPU**

**OpenCAPI
cable**

**FPGA
board**

Jungfraujoch FPGA, CPU and GPU

- Host memory writing
 - Work request/work completion mechanism
 - Unit of transfer 1 frame/module:
 - 500k pixel = 1 MiB
 - For OpenCAPI: standard user-space mmap buffer, no special HW allocation, just 128b alignment
 - For PCIe: kernel-space physically continuous buffer

- Spot finding
 - COLSPOT algorithm from XDS
 - CPU, GPU and FPGA implementations
 - FPGA is possible for real-time full frame rate performance, but constrained and inflexible
 - GPU is optimal for 100-200 Hz (online feedback)
 - CPU for even lower frame rate



IBM POWER9



Nvidia Tesla T4

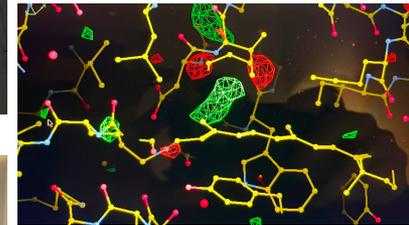
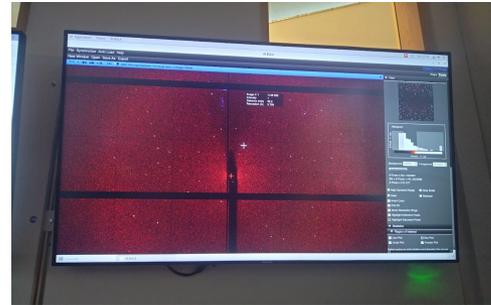
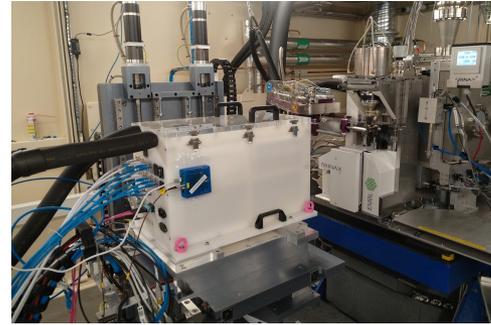
High-level synthesis => continuous integration

- For HLS code written in C++ initial tests can be done on CPU
 - GCC + few extra headers
 - 20 minutes to get wide coverage
 - Possible FPGA HLS code + receiver CPU code in single test
- Hardware simulation for deadlocks and (rare) HLS compiler issues
 - Cadence Xcelium
 - IBM provides OpenCAPI simulation environment => host code is the same as for real hardware test
 - 2-3 hours for simple data acquisition tests (3 images)

The screenshot displays a GitHub Actions pipeline interface for the repository 'leonarski_f / Jungfrauojch'. The pipeline is named 'Pipelines' and has 589 runs. The interface includes buttons for 'Run Pipeline', 'Clear Runner Caches', and 'CI Lint'. Below the buttons, there are filters for 'All', 'Finished', 'Branches', and 'Tags'. A search bar labeled 'Filter pipelines' is present. The main content is a table of pipeline runs with columns for Status, Pipeline, Triggerer, Commit, Stages, and a time/duration indicator. All four runs shown are 'passed' and have a 'latest' badge.

| Status | Pipeline | Triggerer | Commit | Stages | Time/Duration |
|--------|----------|----------------|----------|------------------------|------------------------|
| passed | #1981 | bl1a-march2... | 635804ad | Added tools to calc... | 00:15:28 4 days ago |
| passed | #1980 | bl1a-march2... | c38c6b4a | Minor modifications... | 00:15:28 5 days ago |
| passed | #1979 | fpga_refact... | 347224a2 | Minor modifications... | 00:15:30 5 days ago |
| passed | #1978 | fpga_refact... | 7244d4a1 | Refactored FPGA C ... | 09:10:54 5 days ago |

- Collaboration between MAX IV and PSI (MX + Detector + Bio)
- JF 4Mpixel from PSI + MAX IV computing infrastructure with Jungfraujoch
- Continuous measurements up to 2 kHz (17 GB/s) frame rate on protein targets
 - Consecutive 8 minute runs of 1 million frames at 2 kHz (not tried longer)
- 1 ms time resolution for biological KR2 sample with laser trigger
- Online conversion and compression reduced initial 17 GB/s to below 4 GB/s (GPFS limit)



- Task-specific architectures (like FPGA or GPU) are important tools to handle high data rates in coming years
- Design with FPGA is not difficult with proper tools (coherent interconnects & high-level synthesis)



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