Al-for-Scienceを加速しその先へと導く初のエクサスケールスーパーコンピューター

(The First Exascale Supercomputer Accelerating Al-for-Science and Beyond)

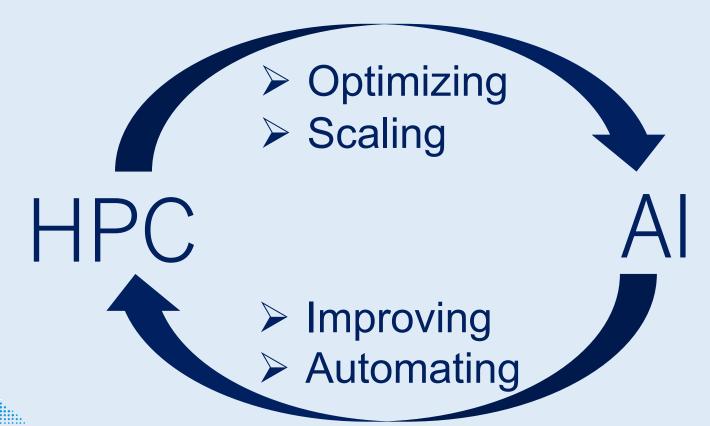
モハマド ワヒブ (Mohamed WAHIB) 理研 計算科学研究センター (RIKEN-CCS) 高性能人工知能システム研究チームリーダー





Prelude

- > AI/ML is not magic
 - > Important to distinguish hype from real prospect
- > There is prospect for AI/ML in HPC



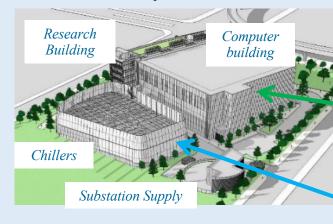




Fugaku Supercomputer



423 km (263 miles) west of Tokyo





Computer room 50 m x 60 m = 3,000 m²

Electric power up to 37 MW

Water cooling system

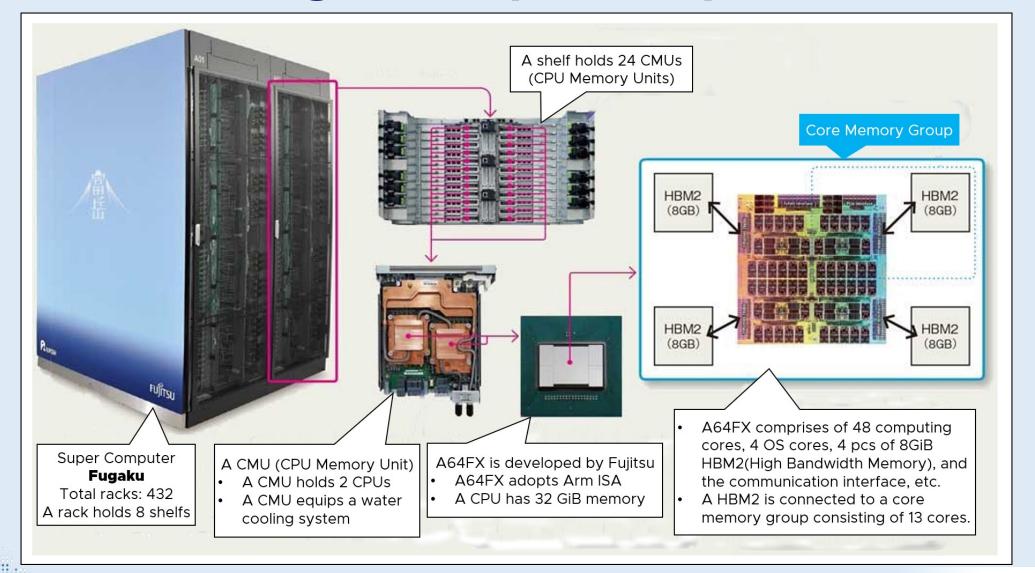
Gas-turbine co-generation 5 MW x 2







Fugaku Supercomputer









Fugaku Research Highlights (FY21)

Achieved world's first five titles

(4 consecutive terms + new ML Perf HPC 1st place)

In four HPC performance rankings (Top500, HPCG, HPL-AI, Graph500), Fugaku won four titles consecutively from June 2020. In November 2021, also awarded first place in ML Perf HPC, a new overall performance evaluation of AI processing.

Fugaku's high overall performance in a wide range of fields, as well as its ability to make a significant contribution to the realization of Society 5.0/SDGs.

Gordon Bell Special Prize

Fight against COVID-19

Successfully developed a detailed and quantitative COVID-19 droplet and aerosol dispersion model using "Fugaku" for the first time in the digital transformation of infectious disease epidemiology. Visualizing arised awareness of the importance of understanding droplet and aerosol infection changing behaviour not only in Japan, but also around the world.

ITU-AJ Special Achievement Award

Data Assimilation Research

(Prediction of Sudden downpours, COVID-19 infection)
Using big data from weather radar, a real-time of ultra-fast
precipitation forecasting, was conducted in the Tokyo area
using Fugaku during the Tokyo Olympic and Paralympic
Games.

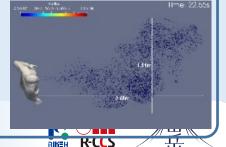
Data assimilation methods developed in numerical weather forecasting were applied to the forecasting of COVID-19 infections

"GENESIS" new version released

Molecular dynamics (MD) simulations

A new version of GENESIS, optimized for 'Fugaku' by co-design, more than 125 times faster and with many new features, has been released as free software in 2020. Work on the dynamic

structure of spike proteins on the surface of COVID-19 has analized successfully. RIKEN EIHO Award (RIKEN Significant Achievement Award)



Al on Fugaku

- > World 2nd fastest Supercomputer (Top500: June 2022)
 - > 1st on HPCG and Graph500
- Fujitsu A64FX Arm v8.2-A + SVE CPUs (32GB HBM memory)
- Creating DL ecosystem for Fugaku:
 - https://github.com/dl4fugaku







Scaling Al/ML on Fugaku

- Past and on-going effort on scaling ML on Fugaku goes into benchmarks
- MLPerf (<u>https://mlcommons.org/en/</u>)
 - Subcategory: MLPerf Training HPC

Area	Benchmark	Dataset	Quality Target	Reference Implementation Model
Scientific	Climate segmentation	CAM5+TECA simulation	IOU 0.82	DeepCAM
Scientific	Cosmology parameter prediction	CosmoFlow N-body simulation	Mean average error 0.124	CosmoFlow
Scientific	Quantum molecular modeling	Open Catalyst 2020 (OC20)	Forces mean absolute error 0.036	DimeNet++





Scaling Al/ML on Fugaku

PERFORMANCE METRICS (TIME TO SOLUTION IN MINUTES) FROM SUBMISSIONS IN CLOSED AND OPEN DIVISIONS

Division	System	Submission	Software	#Processors	#Accelerators	Parallelism [†]	CosmoFlow	DeepCAM
Closed	Piz Daint	Piz-Daint-128	TensorFlow 2.2.0	128	128	2 s/1 GPU	461.01	-
	Piz Daint	Piz-Daint-256	TensorFlow 2.2.0	256	256	2 s/1 GPU	327.01	-
	ABCI	ABCI-1024	PyTorch 1.6.0	512	1,024	2 s/1 GPU		11.71
	ABCI	ABCI-512	TensorFlow 2.2.0	256	512	1 s/1 GPU	34.42	12
	Fugaku	Fugaku-512	TensorFlow 2.2.0 + Mesh TensorFlow	512	-	1 s/1 CPU	268.77	-
	Fugaku	<mark>Fugaku</mark> -8192	TensorFlow 2.2.0 + Mesh TensorFlow	8,192	-	1 s/16 CPUs	101.49	-
	Cori-GPU	Cori-GPU-64	PyTorch 1.6.0	16	64	2 s/1 GPU	-	139.29
	Cori-GPU	Cori-GPU-64	TensorFlow 1.15.0	16	64	1 s/1 GPU	364.73	-
	Cori-KNL	Cori-KNL-512	TensorFlow 1.15.2	512	-	1 s/1 CPU	536.06	-
	HAL	HAL-64	TensorFlow 1.15.0	32	64	1 s/1 GPU	265.59	-
	Frontera-RTX	Frontera-RTX-64	TensorFlow 1.15.2	32	64	1 s/1 GPU	602.23	-
Open	ABCI	⋆ABCI-1024	PyTorch 1.6.0	512	1,024	2 s/1 GPU	=:	10.49
	ABCI	★ABCI-2048	TensorFlow 2.2.0	1,024	2,048	1 s/1 GPU	13.21	-
	<mark>Fugaku</mark>	∗ <mark>Fugaku</mark> -16384	TensorFlow 2.2.0 + Mesh TensorFlow	16,384	-	1 s/4 CPUs	30.07	-
	Cori-KNL	*Cori-KNL-1024	TensorFlow 1.15.2	1,024	-	1 s/1 CPU	419.69	-



AI-for-Science on Fugaku*

by R-CCS Research Teams & Priority application programs (collected by Kento Sato)



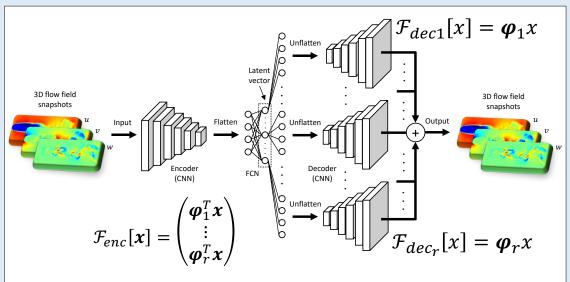


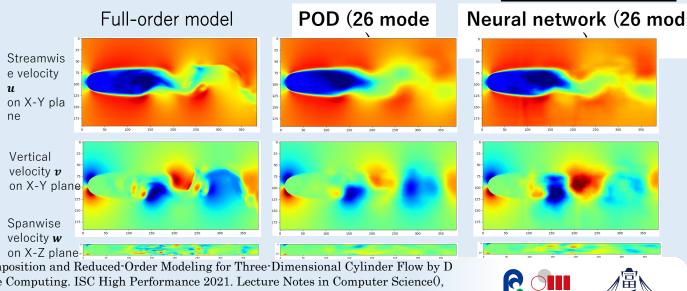
^{*} To Appear: Book Chapter "The First Exascale Supercomputer Accelerating Al-for- Science and Beyond" Satoshi Matsuoka, Mohamed Wahib, Aleksandr Drozd, Kento Sato

Nonlinear Dimensionality Reduction for Three-dimensional Flow Field Around Circular Cylinder with Distributed Parallel Machine Learning on Fugaku (Kazuto Ando et al. [1])

- Mode decomposition is an approach used to decompose flow fields into physically important flow structures known as modes, POD (Proper Orthogonal Decomposition) is a powerful method
- MD-CNN-AE [1] extended to support **3-D flow**
 - MD-CNN-AE [2]: A neural network for a mode decomposition method for **2-D flow** field around circular cylinder using a convolutional neural network
- A hybrid parallelism method combining the distribution of network structure (model parallelism) and the training data (data parallelism) using up to 10,500 nodes on Fugaku was employed for learning.

Schematics of 3D-extended version of MD-CNN-AE



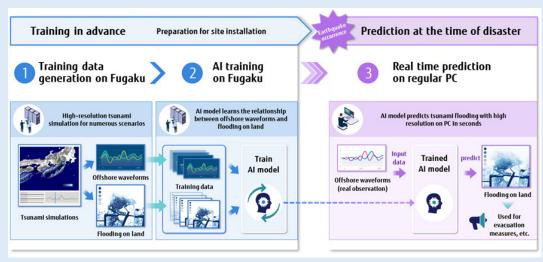


[1] Ando, K., Onishi, K., Bale, R., Tsubokura, M., Kuroda, A., Minami, K. (2021). Nonlinear Mode Decomposition and Reduced-Order Modeling for Three-Dimensional Cylinder Flow by D istributed Learning on Fugaku. In: Jagode, H., Anzt, H., Ltaief, H., Luszczek, P. (eds) High Performance Computing, ISC High Performance 2021. Lecture Notes in Computer Science(), vol 12761. Springer, Cham. https://doi.org/10.1007/978-3-030-90539-2 8

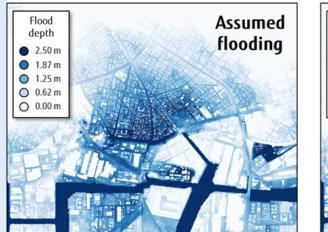
2 T. Murata, K. Fukami, and K. Fukagata, "Nonlinear mode decomposition with convolutional neural networks for fluid dynamics," J. Fluid Mech., vol. 882, A13., 2020, doi:10.1017/jfm.

Development of NN for High-resolution, Real-Time Tsunami Flood Prediction (Fumihiko Imamura group [1])

- Tsunami simulations to generate training data
 - Training Input data: Tsunami waveform in offshore areas
 - Training Output data: Flooding conditions in coastal areas
- Training an AI model to predict flooding condition in coastal areas from Tsunami wave format in offshore
- → This approach makes it possible to accurately and rapidly obtain detailed flooding forecast before landfall of Tsunami







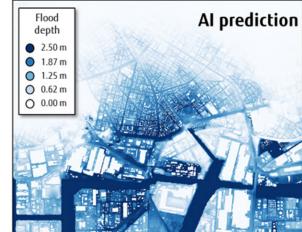


Fig 2. Comparison between anticipated flooding (tsunami source model created by Cabinet Office of Japan with tripled wave heights) of Nankai Trough Megathrust Earthquake and prediction results of newly developed AI







Elucidation of the Cause and Diversity of Cancer using Large-scale Data Analysis and Al Tech. [1]

- Even if a patient receives a targeted cancer drug therapy, the appearance of drug-resistant cancer cells represents an ongoing threat to full remission
 - The mechanism for how certain cancers become drug resistant remains unclear
- Fujitsu implemented parallel conditional and causal algorithms
 - Utilizing Fujitsu's "Wide Learning" AI technology to extract combinations of potential genes relating to the emergence of drug resistance based on statistical information

metastasis of cancer.

Also, maximizing computational performance with the supercomputer Fugaku

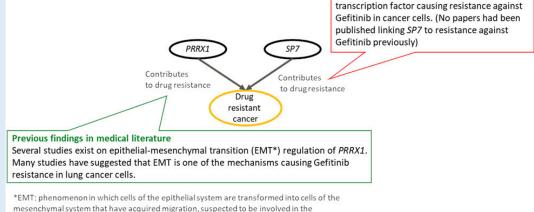
Fugaku, Fujitsu and TMDU were able to search the entire human genome for conditions and causality

within a single day and determine the genes that cause resistance

to drugs used to treat lung cancer

Gene names and expression levels	Indications regarding genes in existing medical literature	Possible findings from existing medical literature		
Low expression level of ZNF 516	Contributes to the suppression of EGFR expression. Suppresses cell proliferation by inhibiting the EGFR pathway*	A low expression level of <i>ZNF 516</i> may indicate activation of the EGFR pathway targeted by Gefitinib.		
Low expression level of E2F6	<i>E2F1</i> and <i>E2F2</i> stimulate cell proliferation by cycling the cell cycle pathway. <i>E2F6</i> inhibits this function.	A low expression of <i>E2F6</i> may promote cell proliferation (not covered by the EGFR pathway).		
Low expression level of EMX1	Known to suppress the Wnt pathway (which promotes cell cycle pathways).	A low expression level of <i>EMX1</i> may not suppress the Wnt pathway and may promote cell proliferation (not covered by the EGFR pathway).		

^{*} mechanism that activates cell growth when stimulated by a biomolecule called epidermal growth factor (EGF)







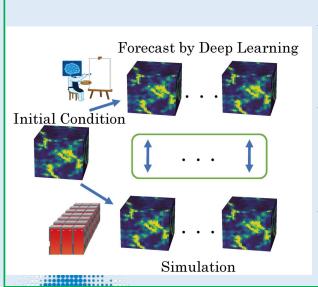
SP7 is suggested as a further potential



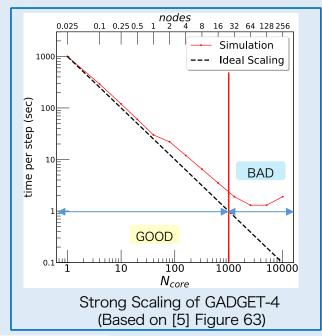
[1] (Press release) Fujitsu Limited, Tokyo Medical and Dental University," Fujitsu and Tokyo Medical and Dental University leverage world's fastest sup ercomputer and AI technology for scientific discovery to shed light on drug resistance in cancer treatment, March 7, 2022 (https://www.fujitsu.com/global/ about/resources/news/press-releases/2022/0307-01.html)

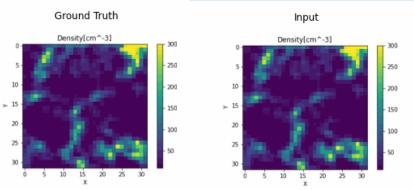
Forecasting the Expansion of Supernova Shells Using Deep Learning toward High-Resolution Galaxy Simulations (Kaiya Hirashima et al. [1])

- Goal: achieve the highest-resolution galaxy simulation which can resolve individual stars on the supercomputer Fugaku.
- One of the bottlenecks is the calculation including small scall phenomena (e.g., supernovae)
- Just using massively parallel computers cannot solve it.
- And algorithm that uses deep learning to resolve the bottlenecks.
 - → The deep learning model to forecast the expansion of supernova shells



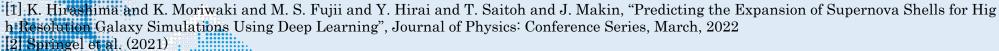
- They applied the framework for video predictions to forecasting the results of simulations
- Prediction of spatial features: CNN
- Prediction of temporal changes: LSTM
- Training data
 - Input: 1 image, Density distribution just before the explosion
 - Output: 19 images, Spatiotemporal changes in density after the explosion
- Training & Prediction
 - The training takes several days using an NVIDIA GeForce RTX 3090 GPU.
- The estimation itself takes only a few second.











Other AI-based Scientific Achievements (Fugaku is not used)

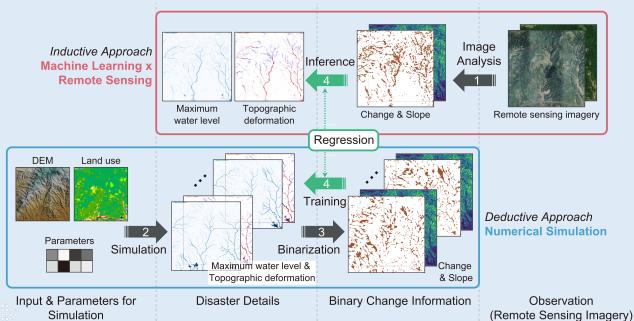
by R-CCS research team & Priority application programs





Remote Sensing by Deep Learning From Simulated Data for Flood and Debris-Flow Mapping for flood disaster damage estimation (Naoto Yokoya et al., 2020 [1])

- Yokoyama et al. proposed a framework that estimates the inundation depth (maximum water level) and debris-flow-induced topographic deformation
 - From remote sensing imagery by integrating deep learning and numerical simulation.
 - A water and debris-flow simulator generates training data for various artificial disaster scenarios.
- They showed that regression models based on Attention U-Net and LinkNet architectures trained on such synthetic data can predict the maximum water level and topographic deformation from a remote sensing-derived change detection map and a digital elevation model
- The proposed framework has an inpainting capability, thus mitigating the false negatives that are inevitable in remote sensing image analysis qualitatively



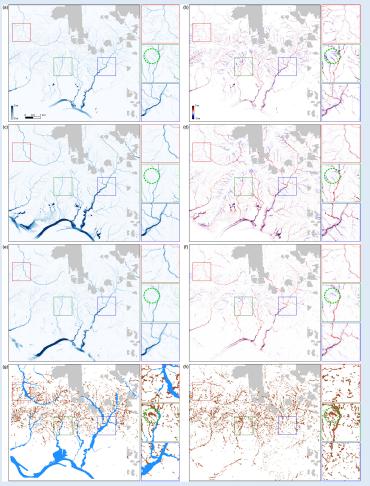


Fig. 8. (a), (c), and (e) Maximum water level and (b), (d), and (f) topographic deformation of (a) and (b) reference simulation, (c) and (d) average simulation and (e) and (f) our method with (g) reference map of (blue) flood and (brown) debris-flow extent and (h) binary change detection map. Mask for clouds an no data is shown in gray

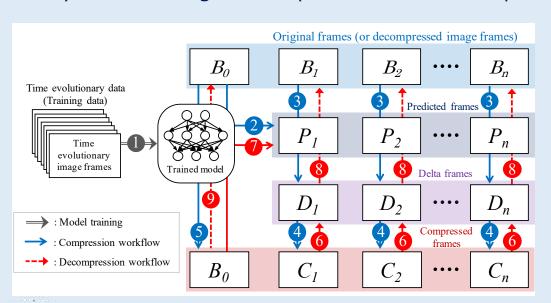
RIKEN R-C



[1] Yokoya, Naoto & Yamanoi, Kazuki & He, Wei & Baier, Gerald & Adriano, Bruno & Miura, Hiroyuki & Oishi, Satoru. (2020). Breaking Limits of Remote Sensing by Deep Learning From Simulated Data for Flood and Debris-Flow Mapping. IEEE Transactions on Geoscience and Remote Sensing. PP. 1-15. 10.1109/TGRS.2020.3035469.

Compression of Time Evolutionary Image Data through Predictive Deep Neural Networks (Rupak Roy et al., 2021 [1])

- Roy et al. proposed new AI-drive data compressor (TEZIP) for time evolutionary data
 - We train PredNet to learn how pixels move and how fast by inputting a number of time evolutional frames
 - When compressing frames, we predict future frames from original base data, compute delta values and apply series of encoding
 - → We only store (1) base frame data and (2) compressed data
- They achieved higher compression ration compared to existing video encoder (Zstd, HFYU, FFV1, x.265)



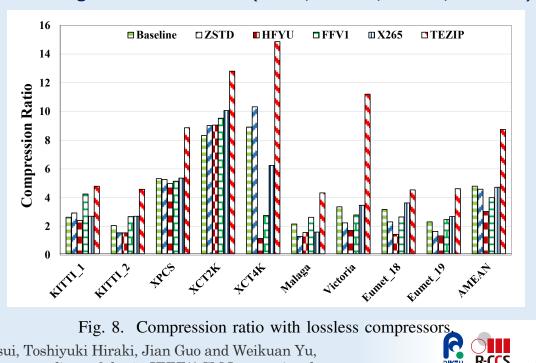


Fig. 1. Workflows of TEZIP (de)compression







Automatic Quantum Circuit Encoding of a given Arbit (Tomonori Shirakawa et al. [1])

- Quantum computer can in principle treat an exponentially large amount of day
 practice to represent classical data such as a classical image into a quantum
- Inspired by a tensor network method, they have proposed a quantum-classical hybrid algorithm to construct an
 optimal quantum circuit for classical data that is represented as a quantum state by the amplitude encoding
- The proposed algorithm employs as an objective function the absolute value of fidelity $F = \langle 0 | \hat{C}^{\dagger} | \Psi \rangle$, which is maximized iteratively to construct an optimal quantum circuit \hat{C} with controlled accuracy.

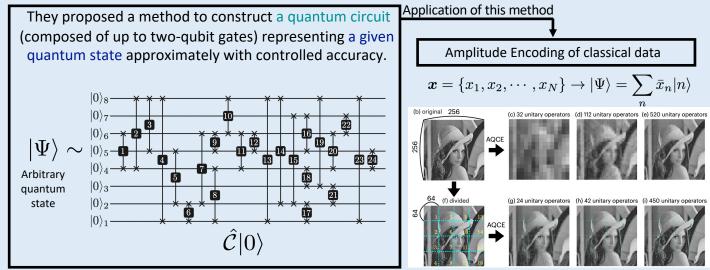


Fig 1: Quantum circuit encoding of a gray scale. (a) Labeling of two-dimensional classical data (with 8×8 pixels, as an example). (b) Original picture with 256×256 pixels. (c)-(e) Pictures reconstructed by decoding the quantum circuit states. (f) Original picture divided into 16 pieces. (g)-(i) Pictures reconstructed by decoding each quantum circuit states.

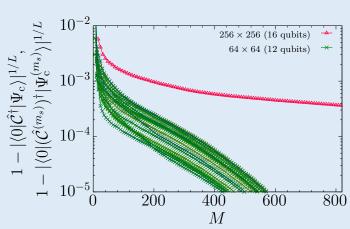


Fig 2: Fidelity per site between the quantum state qubits representing the original picture in Fig. 1(b) [the msth s egment of the original picture in Fig.1(f)] and the quantum circuit state indicated by red triangles (green crosses)





Priority Application

(g) 24 unitary o

Other AI researches by Priority Application Projects (Fugaku is not used)

- Comprehensive gene network analysis by XAI for uncovering molecular mechanism of disease (Heewon Park et al., 2020 [1])
 - Although various approaches were developed to gene network analysis, comprehensive interpretation of the massive networks remains a challenge
 - They conducted comprehensive analysis of the massive gene networks with the use of explainable artificial intelligence (XAI) approaches
 - DeepTensor and Tensor Reconstruction-based Interpretable Prediction (TRIP), proposed by Maruhashi et al. 2018; 2020
 - The use of the XAI enables us to overcome limitation of existing gene network analysis, i.e., narrow angle in the large-scale gene network analysis, and this leads to a better understanding of molecular interplay involved to disease
- Machine learning by neural network for quantum many-body solver and experimental data analysis (Yusuke Nomura et al., 2021 [2])
 - Quantum many-body problems are known to show NP hard difficulty
 - We have developed a solver by using Boltzmann machines to approximate solutions accurately and efficiently
 - It was applied to a challenging problem of a quantum spin model and established the existence of quantum spin liquid phase, which bears a long-ranged quantum entanglement. On another application of machine learning, we have extracted electron self-energy, which is hidden in the photoemission experimental data. It has established the existence of prominent resonance peaks which are responsible for the high-temperature superconductivity.

Other AI researches by Priority Application Projects (Fugaku is not used)

- Facilitating ab initio configurational sampling of multicomponent solids using an on-lattice neural network model and active learning (Kasamatsu et al., 2020 [1])
 - Kasamatsu et al. proposed a scheme for ab initio configurational sampling in multicomponent crystalline solids using Behler-Parinello type neural network potentials (NNPs)
 - The NNPs are trained to predict the energies of relaxed structures from the perfect lattice with configurational disorder instead of the usual way of training to predict energies as functions of continuous atom coordinates.
 - Training set bias is avoided through an active learning scheme
 - This enables bypassing of the structural relaxation procedure which is necessary when applying conventional NNP approaches to the lattice configuration problem
 - This idea is demonstrated on the calculation of the temperature dependence of the degree of A/B site inversion in MgAl2O4, which is a multivalent system requiring careful handling of long-range interactions
 - The present scheme may serve as an alternative to cluster expansion for 'difficult' systems, e.g., complex bulk or interface systems with many components and sublattices that are relevant to many technological applications today
- Density functional theory from supervised learning (Ryo Nagai et al. 2021 [2])
 - Density functional theory (DFT) is the standard electronic structure theory and is widely used as a basis for materials design
 - DFT is based on the Hohenberg-Kohn theorem that there is a one-to-one correspondence between particle density and energy, a relationship that should be machine learnable, but the lack of this relationship makes the accuracy limited
 - They recently proposed a method to establish this relationship using neural network methods
 - They found that the DFT developed in this study can improve the accuracy not only for molecular systems, for which the training set can be obtained by a very accurate quantum chemical method, but also for solids, for which the training set is unavailable but theoretically derived physical conditions work alternatively
 - This paves a systematic way for further accuracy and will serve as a tool for developing the ultimate material database

[1] Kasamatsu, Shusuke and Motoyama, Yuichi and Yoshimi, Kazuyoshi and Matsumoto, Ushio and Kuwabara, Akihide and Ogawa, Takafumi, "Facil itating {Yit ab initio} configurational sampling of multicomponent solids using an on-lattice neural network model and active learning, 10.48550/ARXI



Highlight of Efforts in Bridging AI <-> HPC in R-CCS (Fugaku)

by R-CCS High Performance Artificial Intelligence Systems Research Team





R-CCS HPAIS Team

- > PI: Mohamed Wahib
- > Senior Research Scientists: Jun Igarashi
- > Research Scientists: Aleksandr Drozd, Emil Vatai
- Visiting Scientists: Rio Yokota (TokyoTech), Balazs Gerofi (Intel)
- > Interns: Soyturk (Koc U., Turkey), Zhang (Hokkaido U), Pucceti (SNS)





Jun Igarashi



Aleksandr Drozd



Emil Vatai



Rio Yokota



Balasz Gerofi



Muhammet Soyturk



Enzhi Zhang



Giovanni **Pucceti**

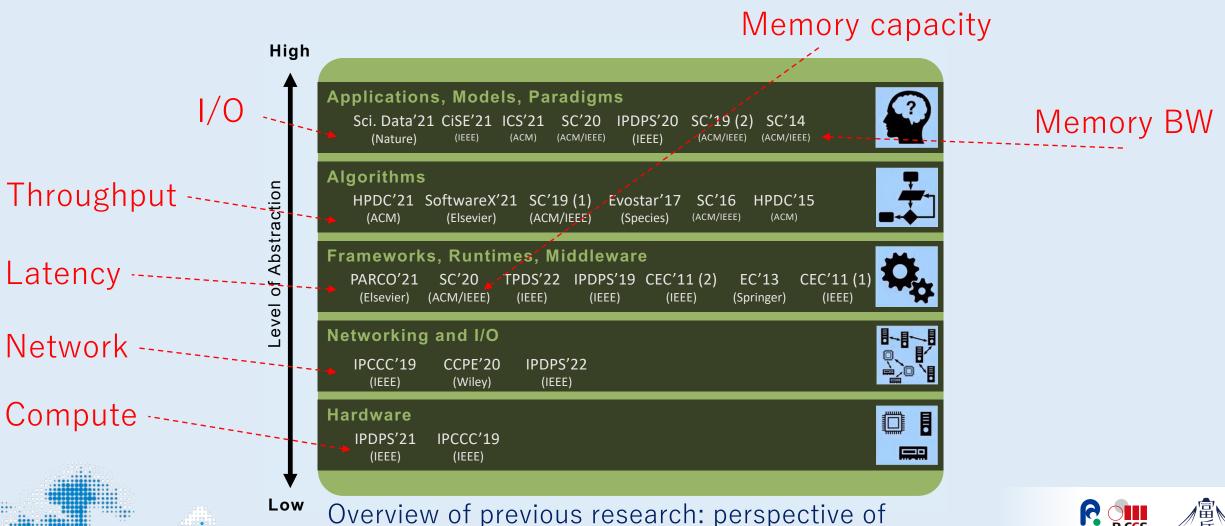






Bottlenecks in Scaling Al

Unlike traditional HPC applications, Al workloads have multiple bottlenecks

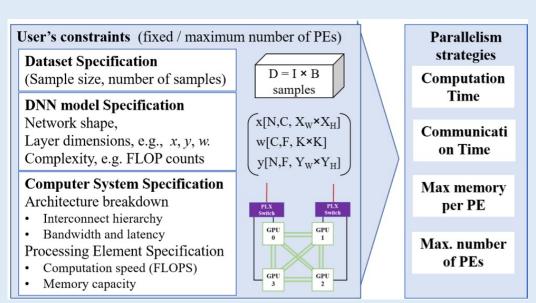


AI+HPC computing stack

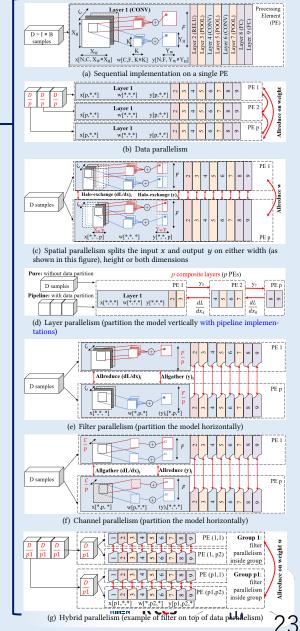
ParDNN: An Oracle for Characterizing and Guiding Large-Scale Training of Deep Neural Networks

- > Different forms of model parallelism are emerging
- ➤ We provide a model-driven analysis and a utility to help in detecting the limitations and bottlenecks of different parallelism approaches at scale

Overview of ParDNN





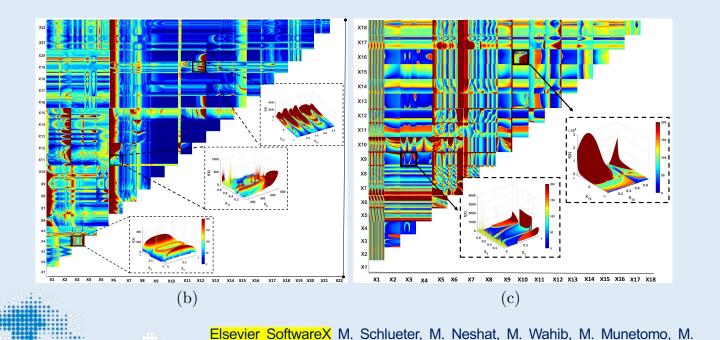


HPC → AI

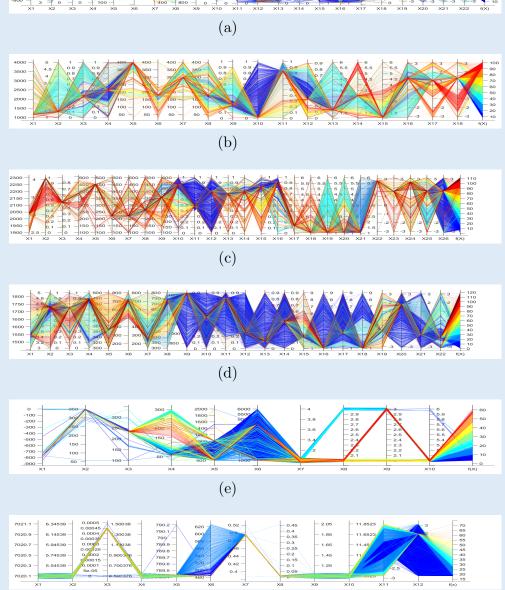
GTOPX Space Mission Benchmarks

- **➤ Landscape Analysis for ML**
 - ➤ Important to understand importance of parameters
 - **≻**Space mission benchmarks

June 2021



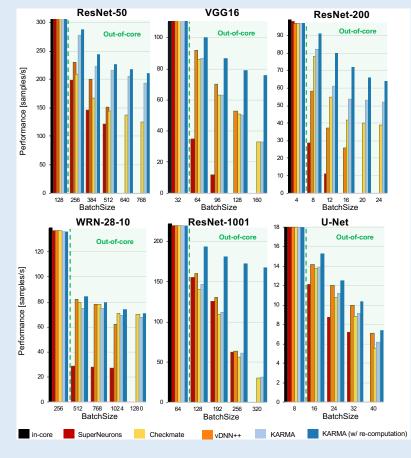
Wagner: GTOPX Space Mission Benchmarks Elsevier SoftwareX Volume 14,



Scaling Distributed Deep Learning Workloads beyond the Memory Capacity with KARMA

- **≻**Concurrency-driven out-of-Core
 - > Capacity-based interleaved with recompute
- **>1.52x over state-of-the-art (single GPU)**
- >First out-of-core to support multi-GPU
 - > Heterogeneity and careful orchestration
- **≻Outperforming DP+MD with out-of-core**
 - > Experiments with up to 2,048 GPUs

Performance on V100. For all figures, only the first reported mini-batch size (x-axis) fits in memory







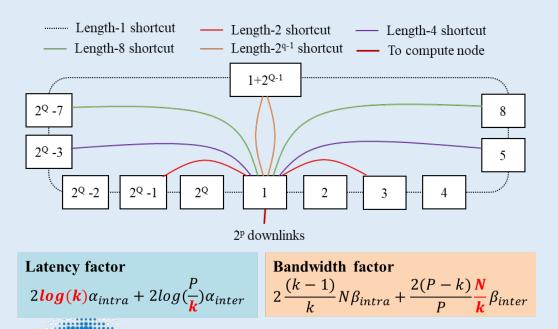


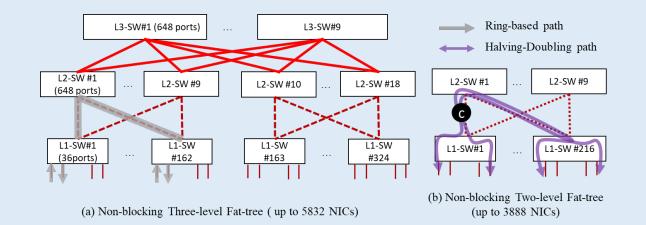
Optimizing Network Architecture for Deep Learning

Problem: Number of involved GPUs (*P*) becomes too big, 1000s GPUs.

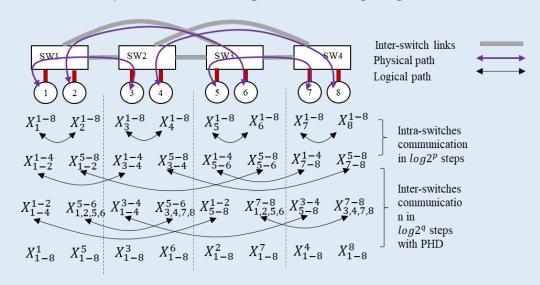
- Ring-based algorithm $O(P)\alpha$: latency factor increase.
- ightharpoonup Halving-Doubleling algorithms O(logP) α : network congestion of communication.

Proposal: Distributed Loop Network topology





Pipelined Halving – Doubling Algorithm



CCGRID'21 Truong Thao Nguyen, Mohamed Wahib: An Allreduce Algorithm and Network Co-design for Large-Scale Training of Distributed Deep Learning, In proceedings of the IEEE/ACM International Symposium on Cluster, Cloud and Internet Computing 2021

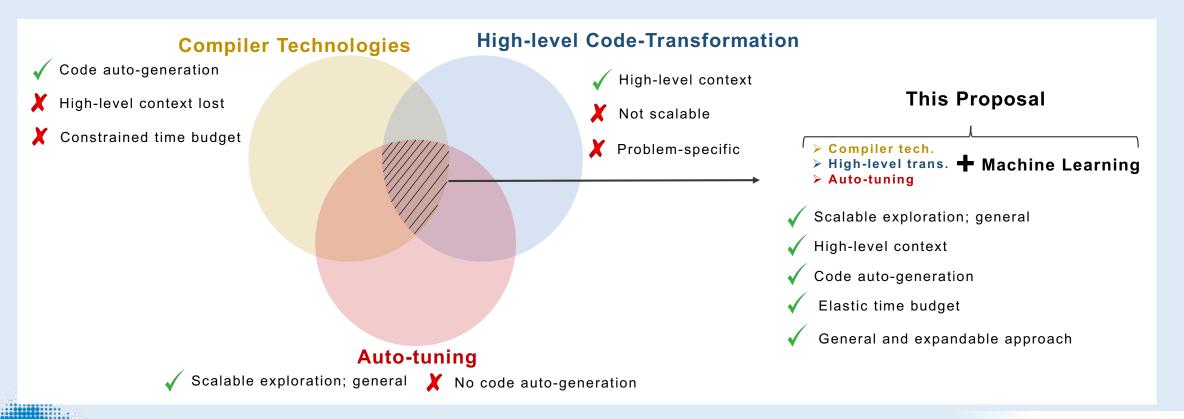






Al for Scientific Codes

Use Al for code auto generation to aid scientific programmers in producing HPC programs.







Fugaku is driving innovation in bridging HPC <-> AI, and will continue to do!

Thank you.

We are hiring!





