



Recent Activities in CCS

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Oakforest-PACS



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Center for Computational Sciences



U. Tokyo

東京大学情報基盤センター

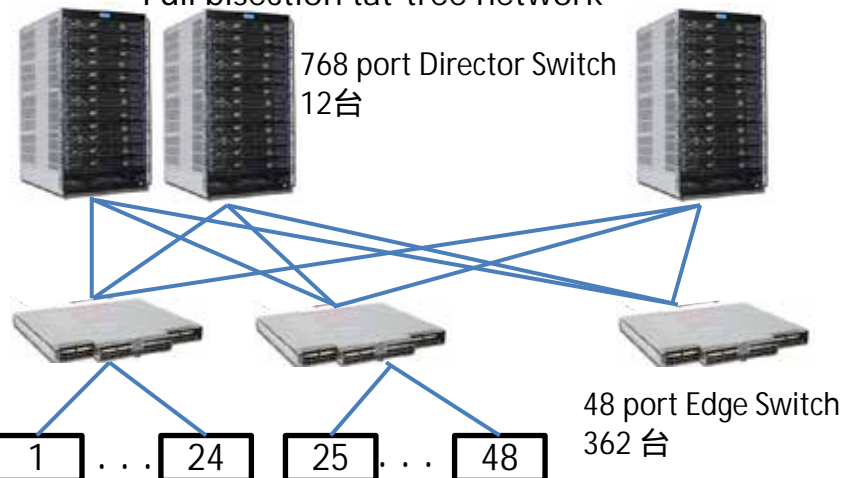
INFORMATION TECHNOLOGY CENTER, THE UNIVERSITY OF TOKYU



Ø Peak Performance 25 PFLOPS
Ø LINPACK 13.55PFLOPS

Intel® Omni-Path Architecture

Full bisection tat-tree network



FUJITSU
Node (Fujitsu PRIMERGY CX1640 M1)
CPU Intel Xeon Phi 7250 (Knights Landing)



DataDirect
NETWORKS

26 PB Parallel file system , 940 TB High-performance file cash system

Oakforest-PACS ranks #1 in IO 500 list

Oakforest-PACS Storage System

Parallel file system (DataDirect Networks ES14KX)

File cache system (Infinite Memory Engine IME14KX): Improving the storage performance.

IO-500 benchmark in 2017

The file cache system achieved 742 GiB/s^[1] for file-per-process write access that parallel processes access their own file.

and 600 GiB/s for single-shared-file write access that parallel processes access a single shared file but a different position.



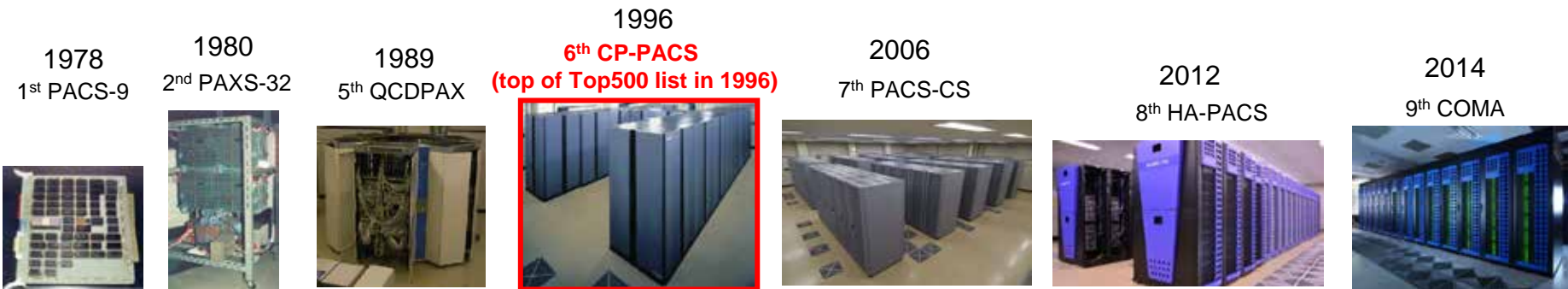
OFP file cache system
IME14KX

Development of Massively Parallel Computer Systems in CCS

- 1977 research begins (by Hoshino, Kawai)
- 1978 1st machine
- 1996 CP-PACS (top of Top500)
- 2006 7th machine PACS-CS
- 2012 8th machine HA-PACS

CP-PACS

- First large-scale general-purpose MPP system in Japan
- Development supported by “Research of Field Physics with Dedicated Parallel Computers” funded by the Ministry of Education of the Japanese Government.
- ranked as No. 1 system in the November 1996 Top 500 List.
- Collaboration by physicists and computer scientists
- Collaboration with industry, and released as Hitachi SR2201



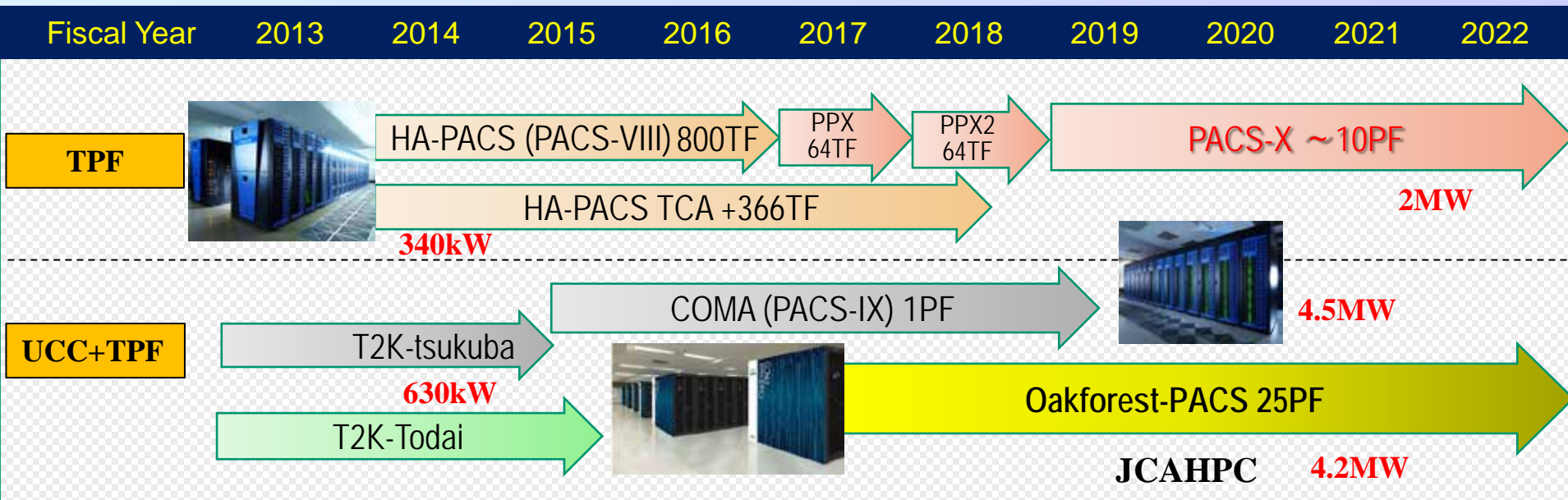
Year	System	Performance
1978	PACS-9 (PACS I)	7 KFLOPS
1980	PACS-32 (PACS II)	500 KFLOPS
1983	PAX-128 (PACS III)	4 MFLOPS
1984	PAX-32J (PACS IV)	3 MFLOPS
1989	QCDPAX (PACS V)	14 GFLOPS
1996	CP-PACS (PACS VI)	614 GFLOPS
2006	PACS-CS (PACS VII)	14.3 TFLOPS
2012	HA-PACS (PACS VIII)	1.166 PFLOPS
2014	COMA (PACS IX)	1.001 PFLOPS

2007
FIRST
(Hybrid Simulator)



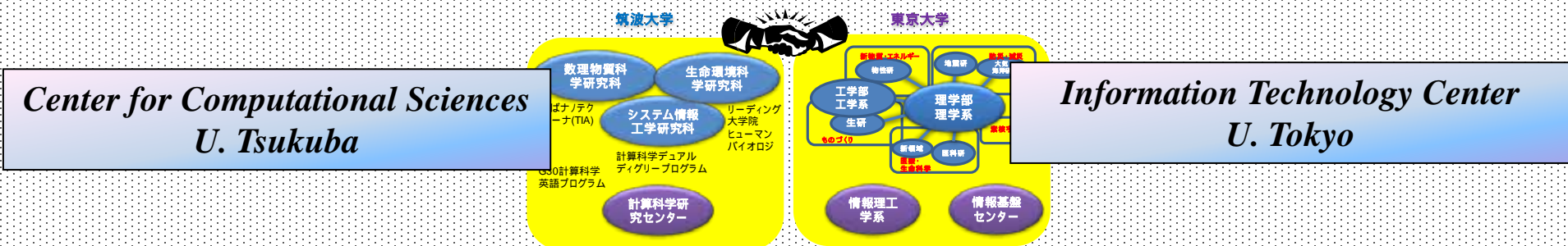
36 TFLOPS

Timeline of Computing Systems in CCS



TPF = Technology Path-Forward Machine
UCC = Upscale Commodity Cluster Machine

U Joint Center for Advanced High Performance Computing (JCAHPC)



U Feasibility Study toward Exa-scale Computing (National Project)

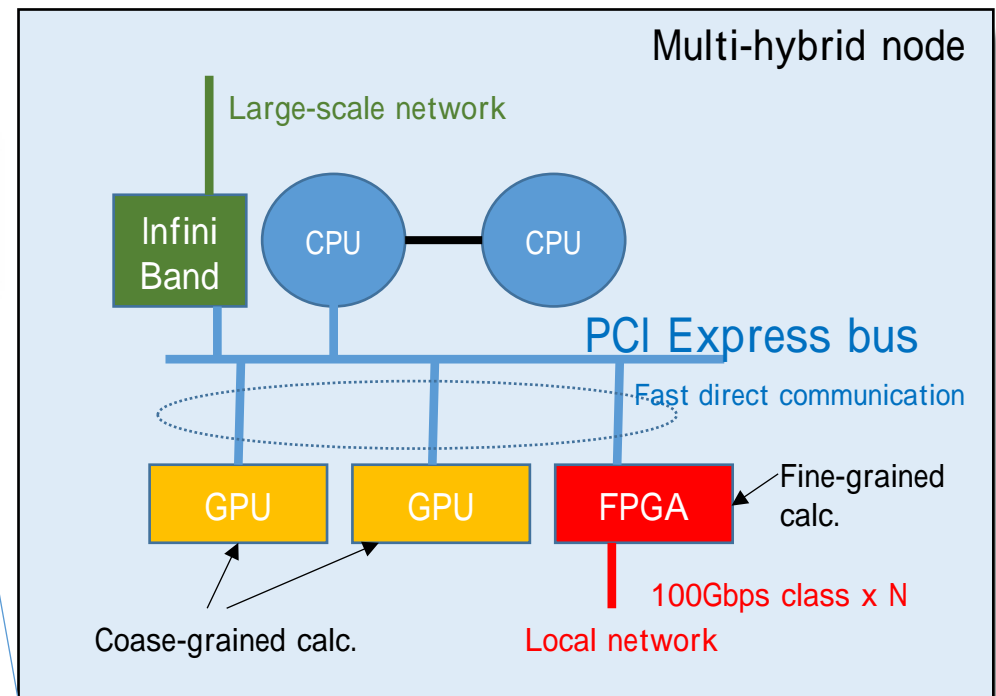
PACS-X: Hybrid Accelerator System

- Ø GPU&FPGA are embedded with PCIe bas
- Ø Local network system and off-loading with FPGA
- Ø Acceleration of hot spots with GPU

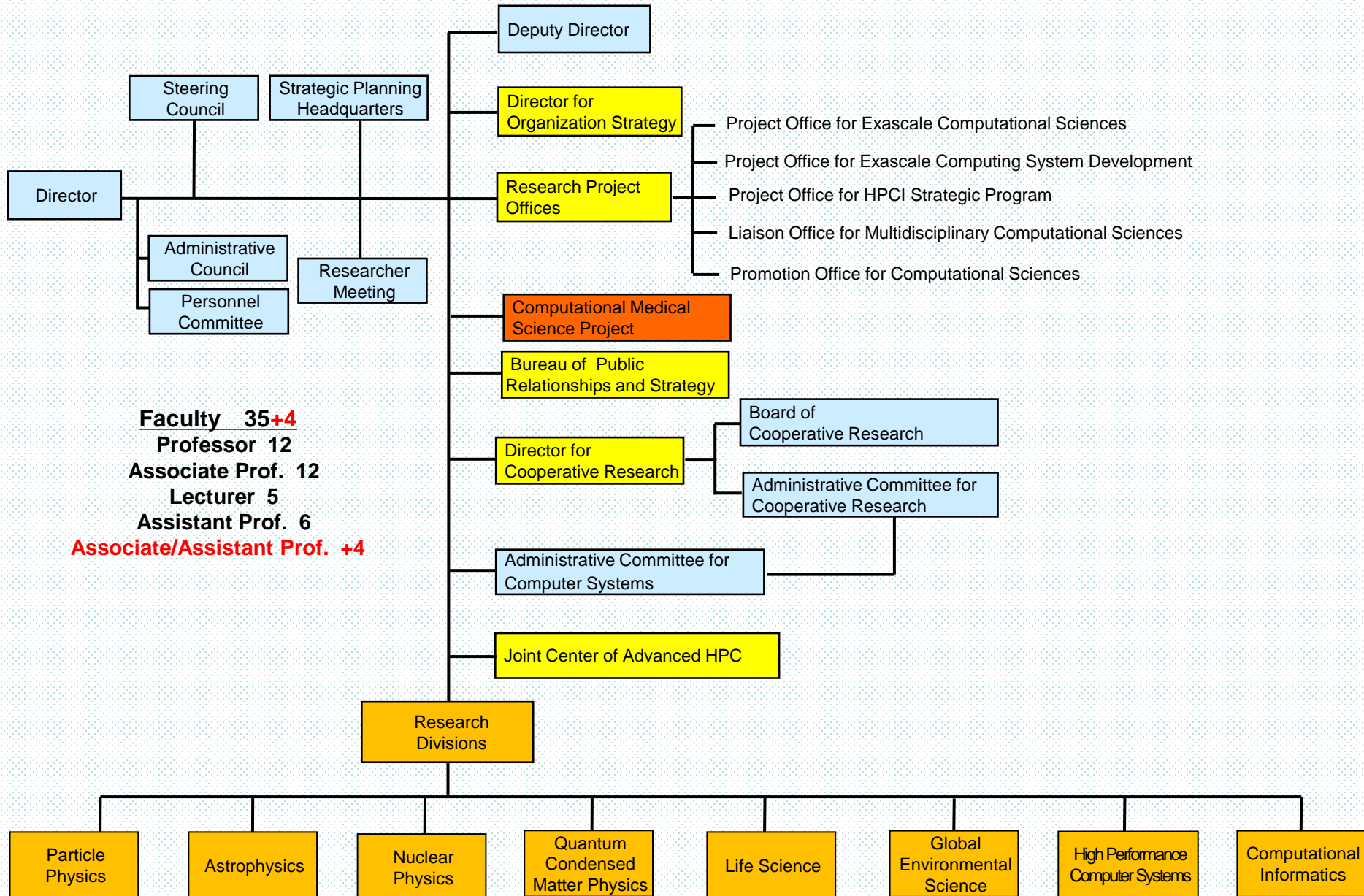


Pre-PACS-X (PPX)

- Ø PPX1(6nodes), PPX2(7nodes)
- Ø Optimization of off-loading with FPGA



Organization of CCS





Development of Novel Scientific Fields



Computational Medical Science Project

Development of medical technology by computational science in cooperation with medicine, physics, biology, mathematics, data science, and image processing technology.

Computational Biomolecular
Medical Science

**Collaborations: Medicine,
Biology, and Physics**

Drug discovery, detection of SNP,
and physical mechanism analysis
by bioinformatics and simulations

Sleep Big Data Analytics &
Automatic Sleep Diagnosis

**Collaborations: IIS,
C-AIR, CCR**

System for automatic analysis of
EEG (electroencephalogram) in
sleep with machine learning and
deep learning

Life Science

Data Science

Computational
Medical Science

Computational
Media

Computational
Physics

3DCG Virtual Surgery

**Collaborations: Medicine,
Engineering, Design**

Development of surgical operation support system using
computational surgery in cooperation with Medical, System
Information Engineering, Information Media researches

Computational Optical
Bioimaging

**Collaborations: Biomedical optics,
Mathematics**

Development of non-invasive near-infrared computed
tomography in cooperation with biomedical optics, radiation
transfer physics, and mathematical inverse problem.

Industry

Computational Biomolecular Medical Science

Main Target

- Medium size drugs
- Covalent bond-type drugs
- Single Nucleotide Polymorphism

Current Problems

- Diversity in Structures (Vast Conformations)
- Reaction Control (First-principles Calculations)
- Fast Gene Analysis

Drug discovery needs long time (10 ~ 15 year) and a lot of fund (\$1billion). But the success ratio is less than 1/20000

We need a breakthrough for in silico drug discovery, i.e. detection of target, design of drug and proteins & etc.



Fusion of Biology, Medicine, and Physics to solve these problems

Evolution

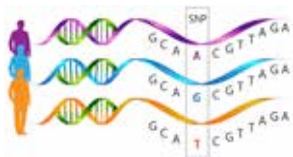


Prof. Y. Inagaki
Department of Biology

Computational
Biology



Next-generation
sequencer

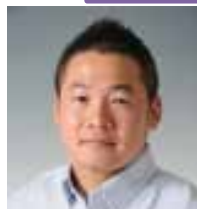


Fast Single Nucleotide
Polymorphism detection

Phylogenetic tree analysis by GPU
Mitochondrial protein detection by supercomputer

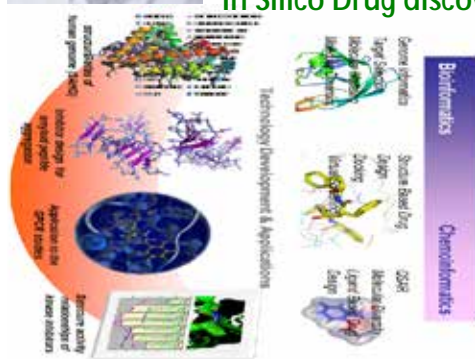
Gene analysis
SNP analysis

Drug Simulation



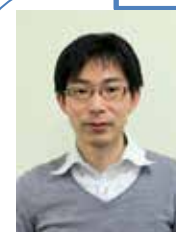
Prof. T. Hirokawa
Department of Medicine
& AIST

Bioinformatics,
In Silico Drug discovery



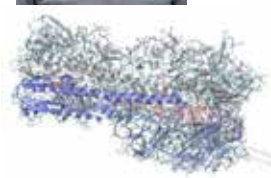
In Silico drug discovery project by AMED
since 2017.

First-principle



Prof. Y. Shigeta
Department of Physics

Biophysics, First-principles
calculations



Influenza HA3
First-principles Calculation



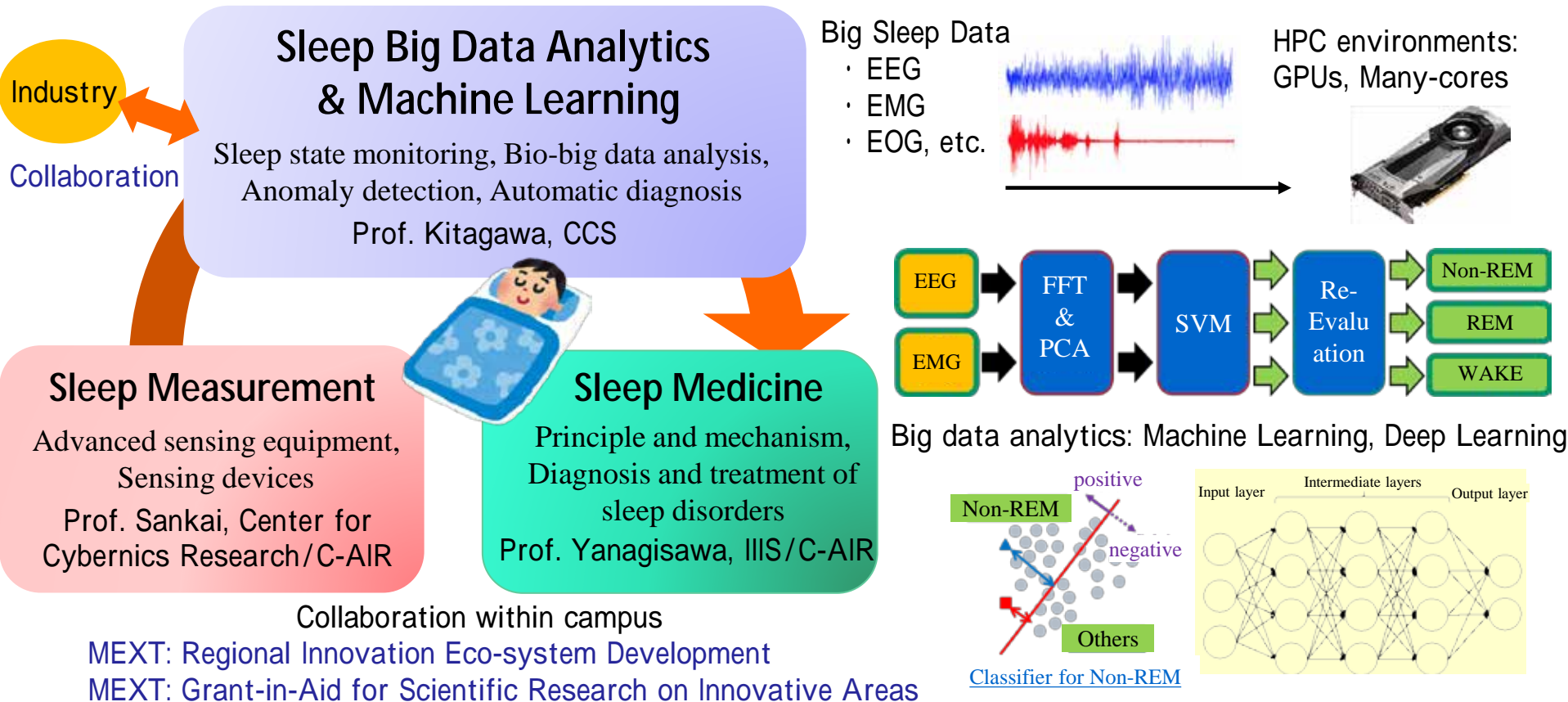
Molecular dynamics
of FtsZ In cell fission

Fragment molecular orbital by GPU architecture
Efficient sampling method by Molecular Dynamics

Post K Computer Project by MEXT
since 2015

Sleep Big Data Analytics & Automatic Sleep Diagnosis

- Sleep plays an important role in daily life of people. Sleeping problems could trigger various diseases.
- There has not been any means for conveniently measuring the sleep state as accurately as medical doctors and technicians.
- Integrate bio-big data analysis, machine learning, and new sensing technology to realize automatic analysis and diagnosis of sleep.

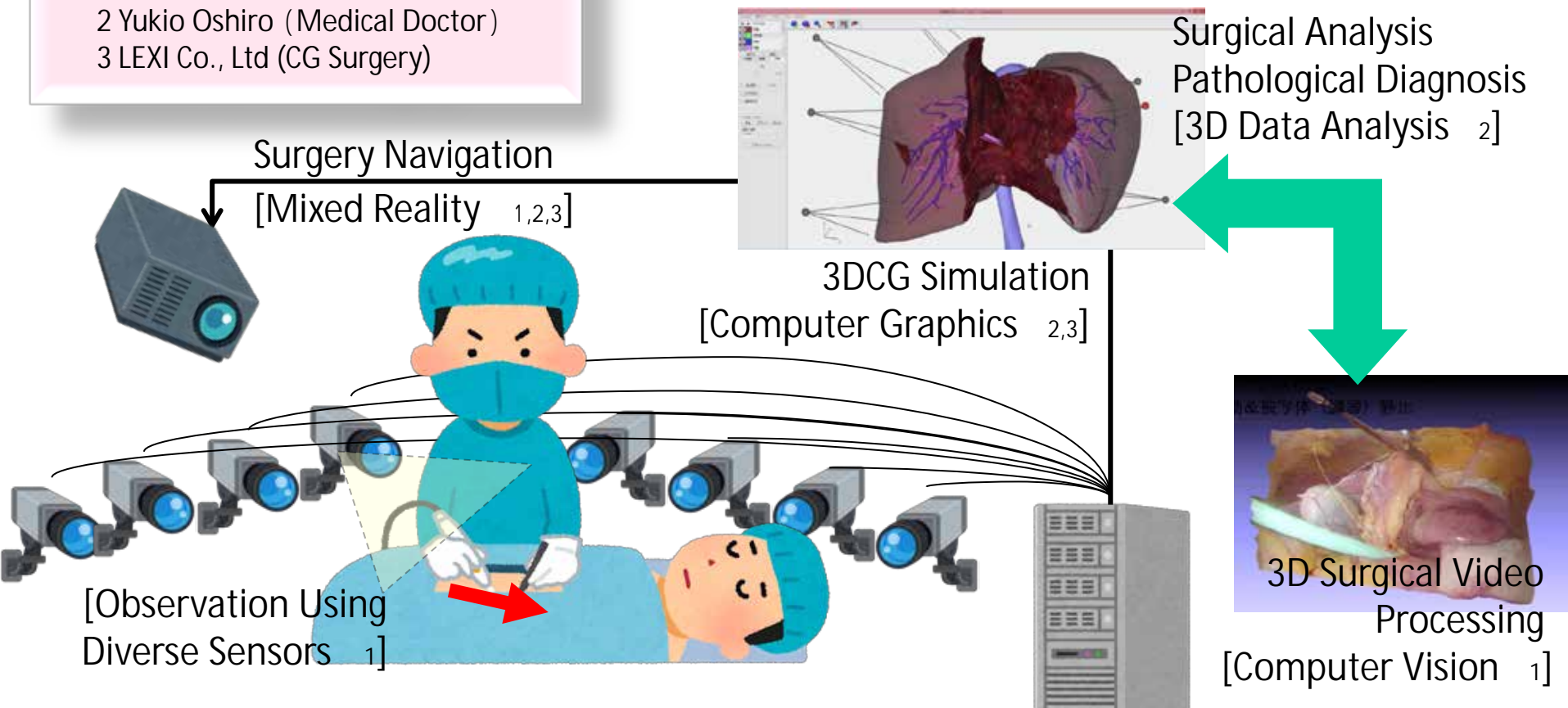


3DCG Virtual Surgery

- **Improving instruction level:** Integration of 3D CG simulation and onsite navigation.
- **Solving surgeon shortage:** VR remote operation, advanced skill passing.
- **Boosting medical education:** Evaluation of surgical procedures and proficiency.

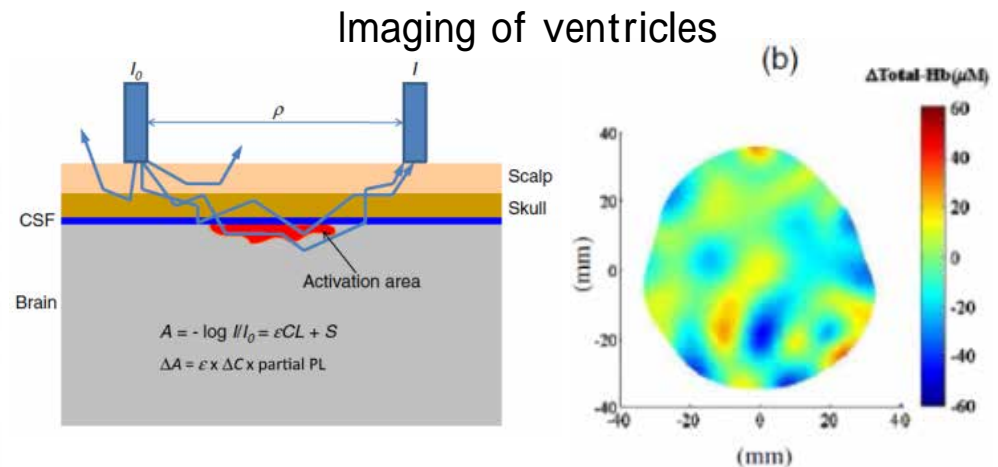
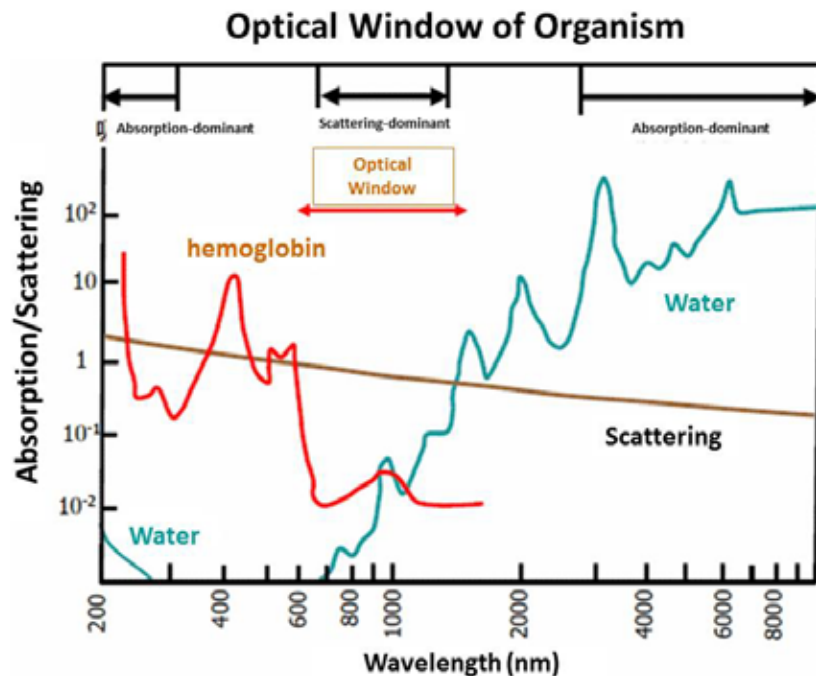
Collaboration Team

- 1 Itaru Kitahara (CCS)
- 2 Yukio Oshiro (Medical Doctor)
- 3 LEXI Co., Ltd (CG Surgery)

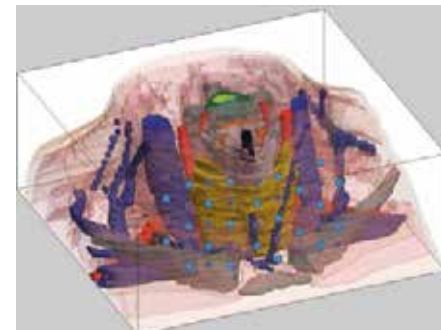


Computational Optical Bioimaging

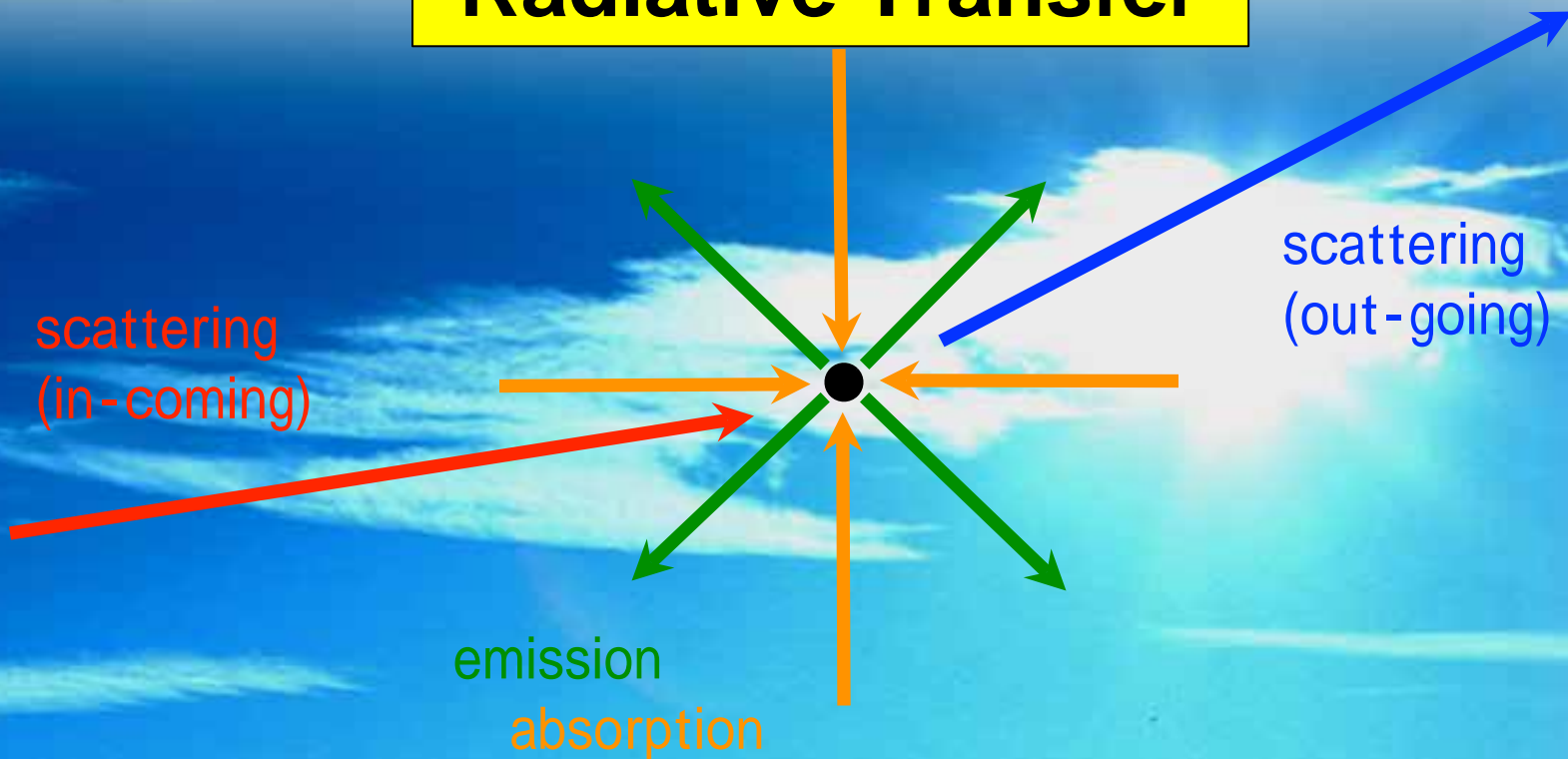
- Ø No exposure, non-invasive, no side effects like CT or MRI, applicable to new-born babies and infants
- Ø Bed side inspection, low cost, low restraint, high time resolution, automatic and remote diagnosis, monitoring, screening
- Ø Diagnosis of cancers in thyroid, mammary gland, surface tissue, blood flow, ventricles, subarachnoid, etc.



Imaging of thyroid gland



Radiative Transfer



Time-dependent Radiative Transfer Equation

(Photon Boltzmann equation in phase space of 3D space, 2D direction, and 1D frequency.)

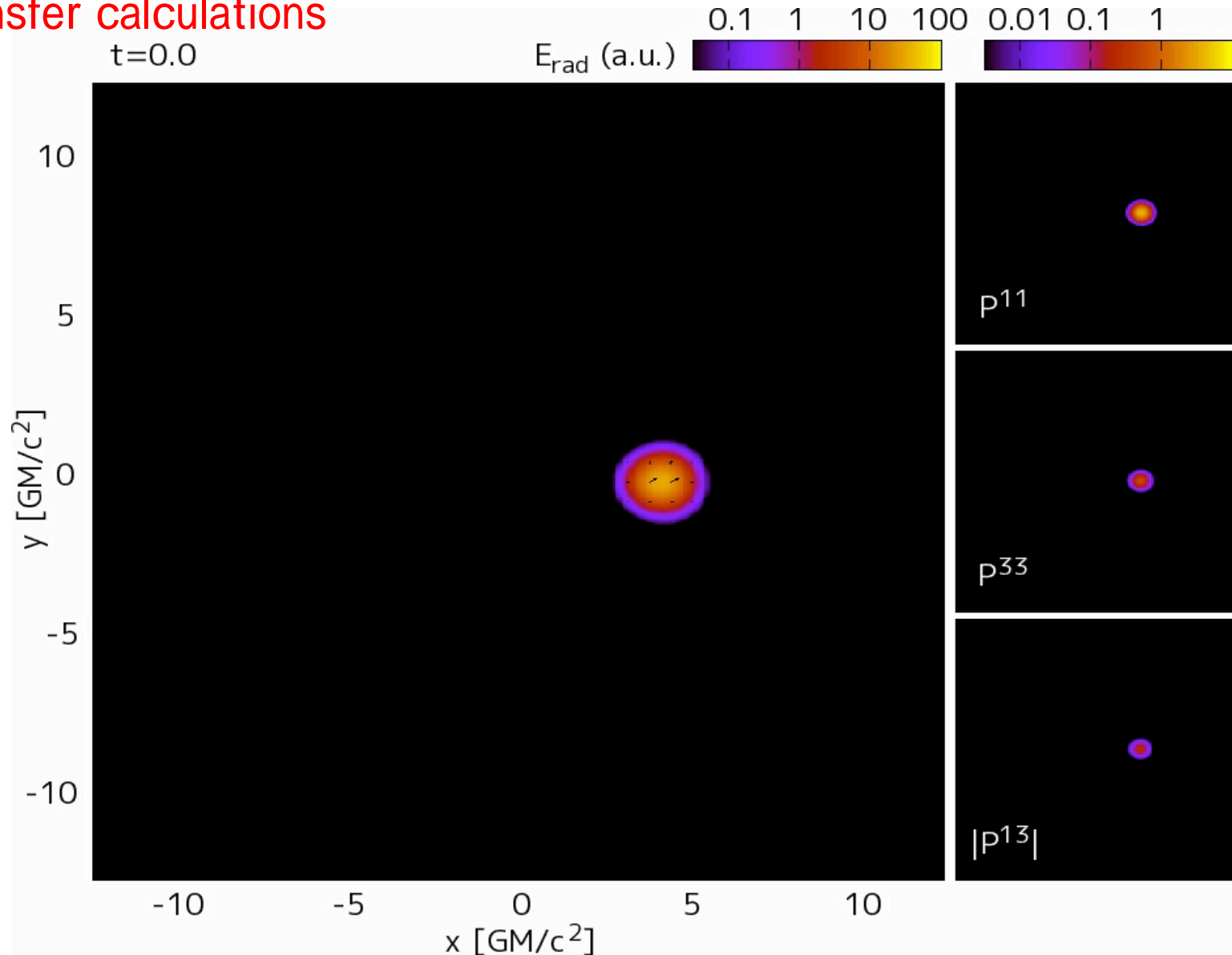
$$\frac{1}{c} \frac{\partial I_\nu(\mathbf{n})}{\partial t} + \mathbf{n} \cdot \nabla I_\nu(\mathbf{n}) = \frac{\eta_\nu}{4\pi} - \kappa_\nu I_\nu(\mathbf{n}) - \sigma_\nu I_\nu(\mathbf{n}) + \sigma_\nu \int \phi(\mathbf{n}; \mathbf{n}') I_\nu(\mathbf{n}') d\Omega'$$

emission
absorption

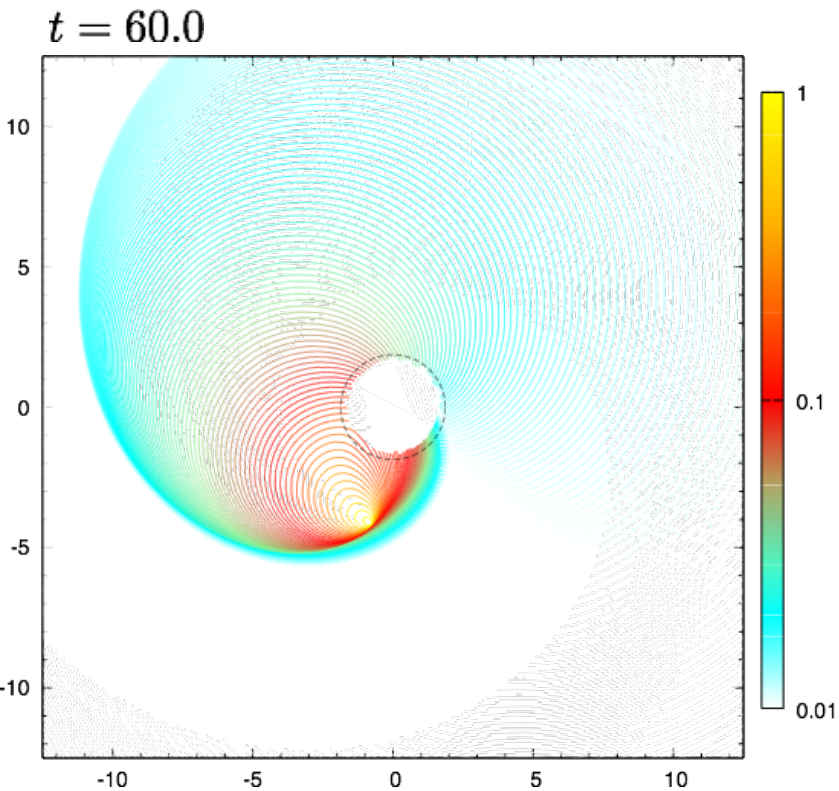
scattering (out-going)
scattering (in-coming)

Dynamical Test 2 by *ARTIST* : Photon wave front from a rotating hot spot

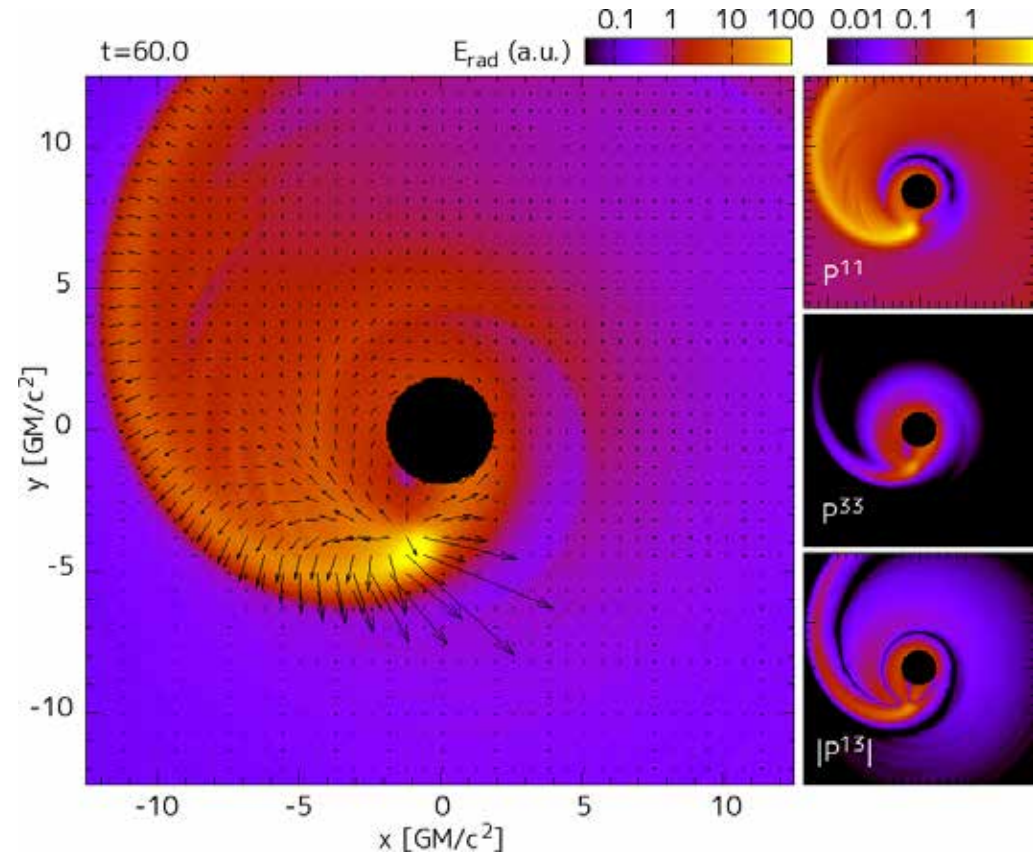
Transfer calculations



Ray-tracing calculations by *MASTER*



Transfer calculations by *ARTIST*



$$N_i(r)=90, N_j(f)=256$$

$$\text{Geodesics}=4608$$

Accurate Algorithm of Image Reconstruction

Biomedical Optics

Diffuse Optical Tomography (DOT) , Photoacoustic Imaging (PAI), Fluorescence Tomography

Direct Problem Analysis

Numerical solution of radiative transfer with absorption, scattering, and reflection.

Inverse Problem Analysis

Mathematical Analysis of non-linear optimization problem.