

STATUS OF THE STRUCTURAL BIOLOGY AT THE PHOTON FACTORY



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Shohei Watanabe

Kazutaka Demura

Tomoya Oota

Takayuki Kubota

Shinji Naito

Rei Tanabe

Kenichi Kawasaki

Kotaro Kuroya



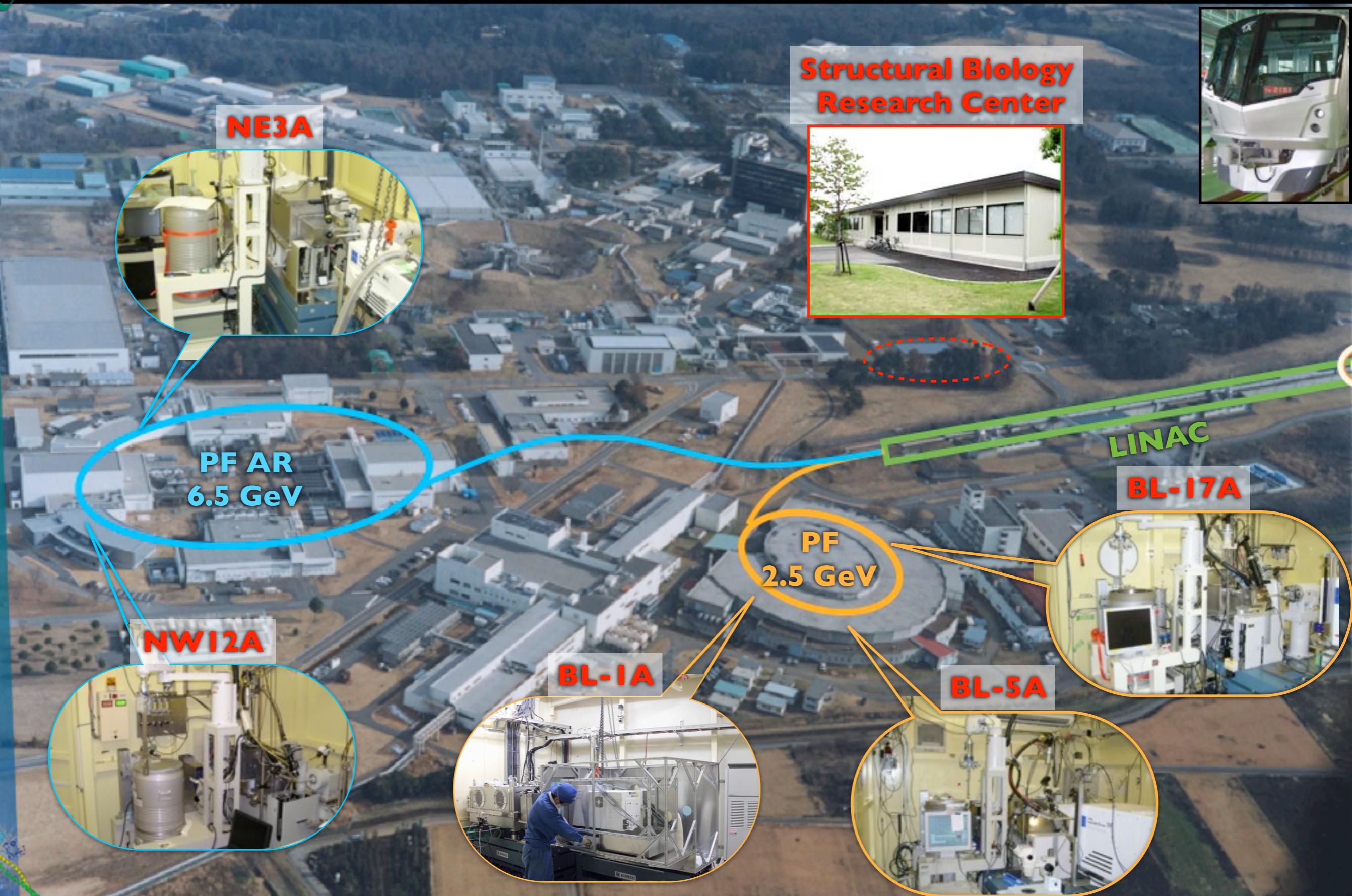


OUTLINE

- **Where are we?**
- **What are we doing?**
 - ~ **introduction on structural biology**
 - ~ **R&D : application to structural biology**
 - ~ **synchrotron science**
 - ... **AR-NE3A: high-throughput beamline**
 - ... **BL-1A: low energy beamline**
- **Structural biology applied to influenza virus and human neuraminidases: involvement R&D and synchrotron science**
- **New applications**
 - ~ **R & D : UV-based centering and phasing**
 - ~ **synchrotron science : high-pressure studies**
- **Conclusions**
- **Acknowledgments**



PHOTON FACTORY - SBRC

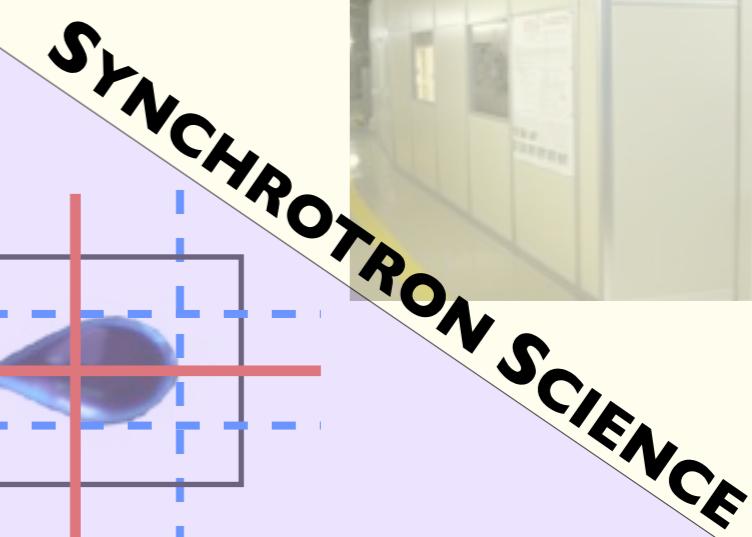
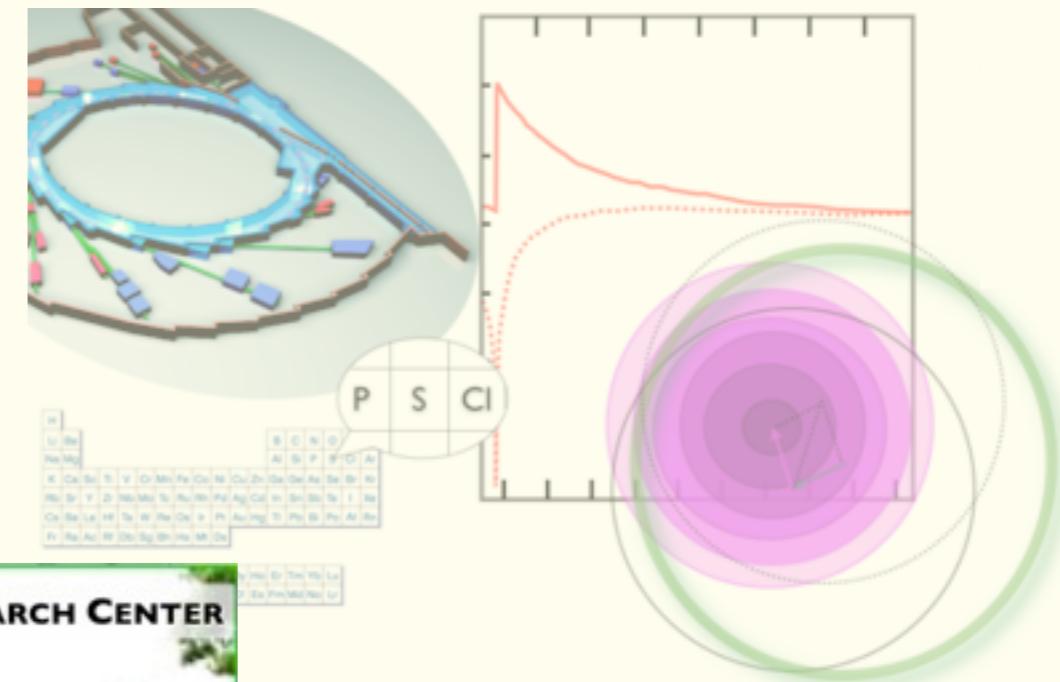
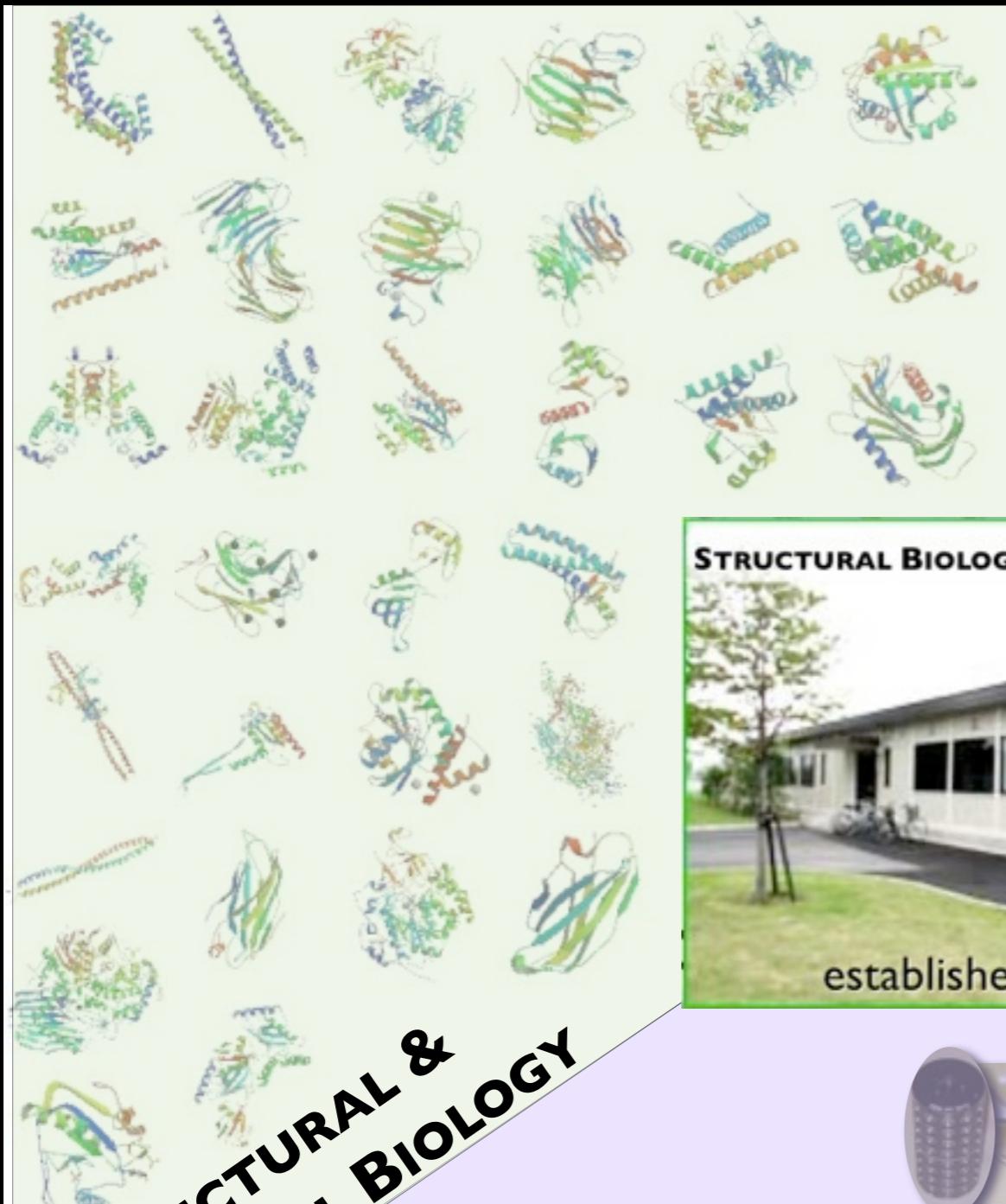


• Where are we?

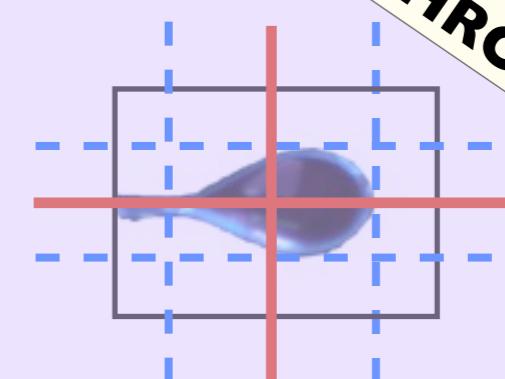
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RESEARCH AND SUPPORT ACTIVITIES

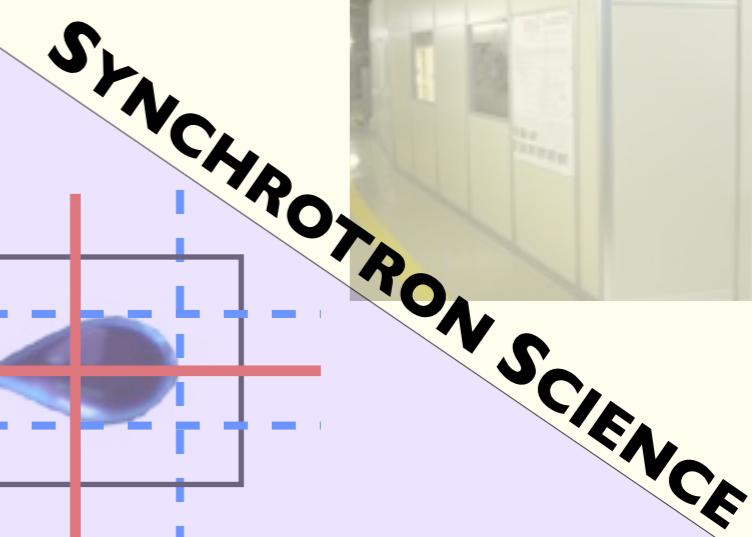
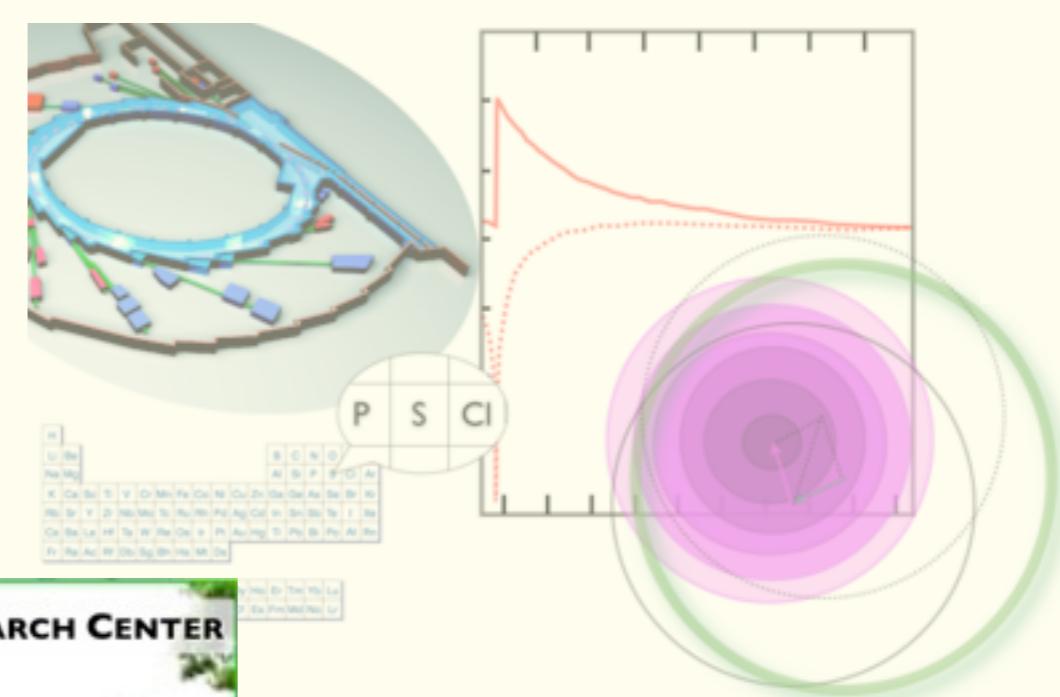
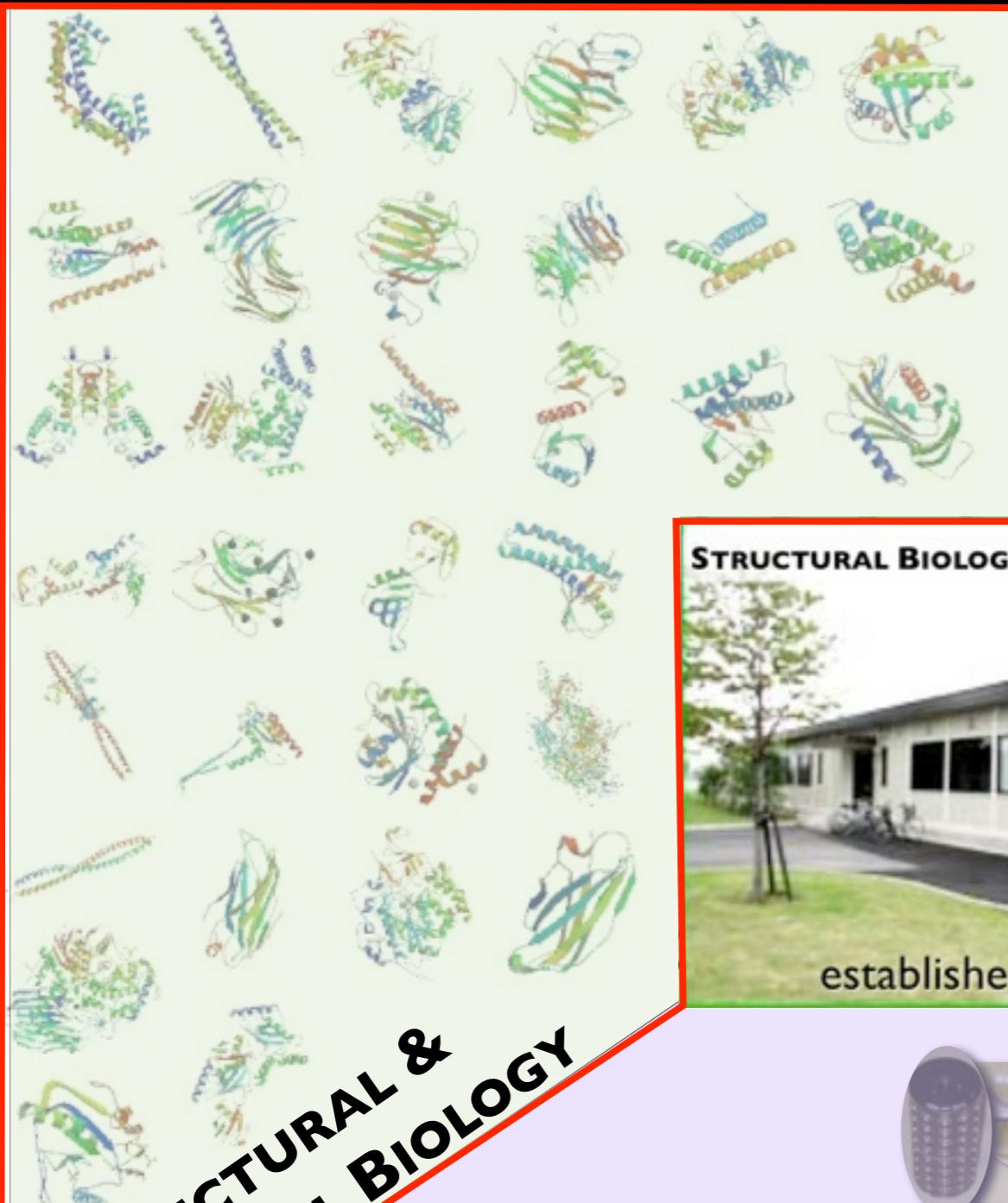


R&D



• What are we doing?

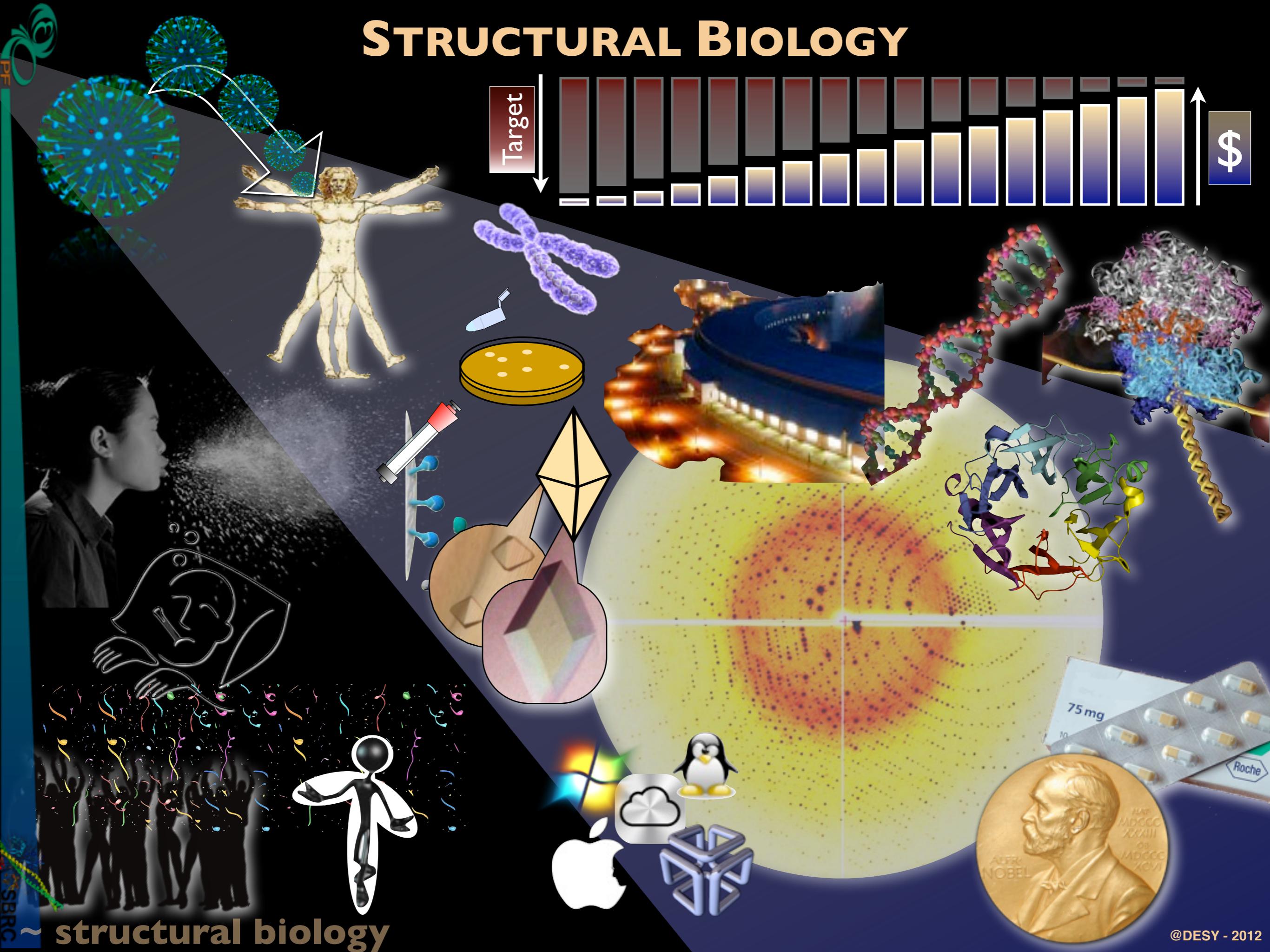
RESEARCH AND SUPPORT ACTIVITIES



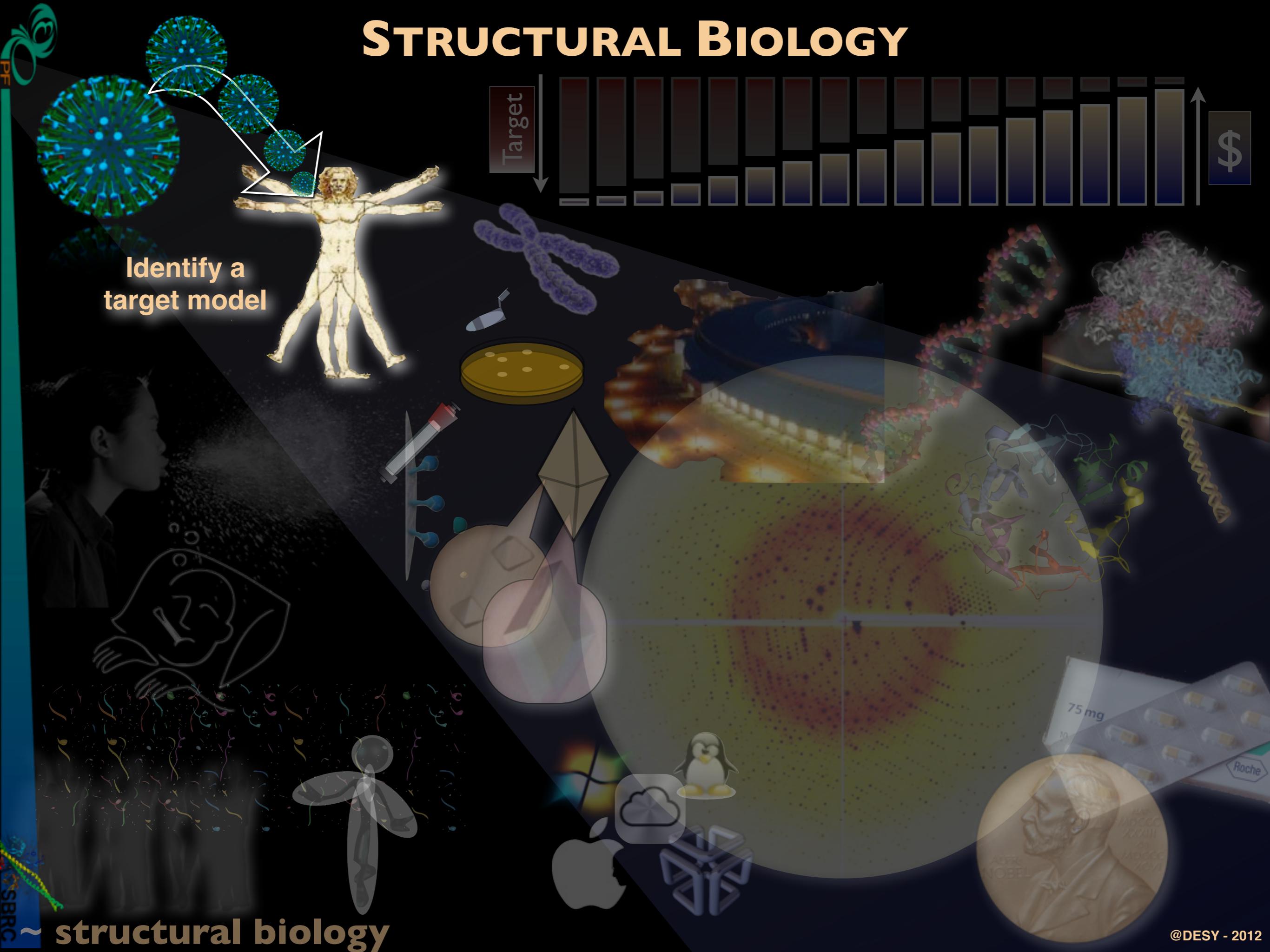
R&D

• What are we doing?

STRUCTURAL BIOLOGY



STRUCTURAL BIOLOGY



BIOLOGICAL MODEL: INFLUENZA

Oxford American Dictionaries:

“...a highly contagious viral infection of the respiratory passages causing fever, severe aching, and catarrh, and often occurring in epidemics...”



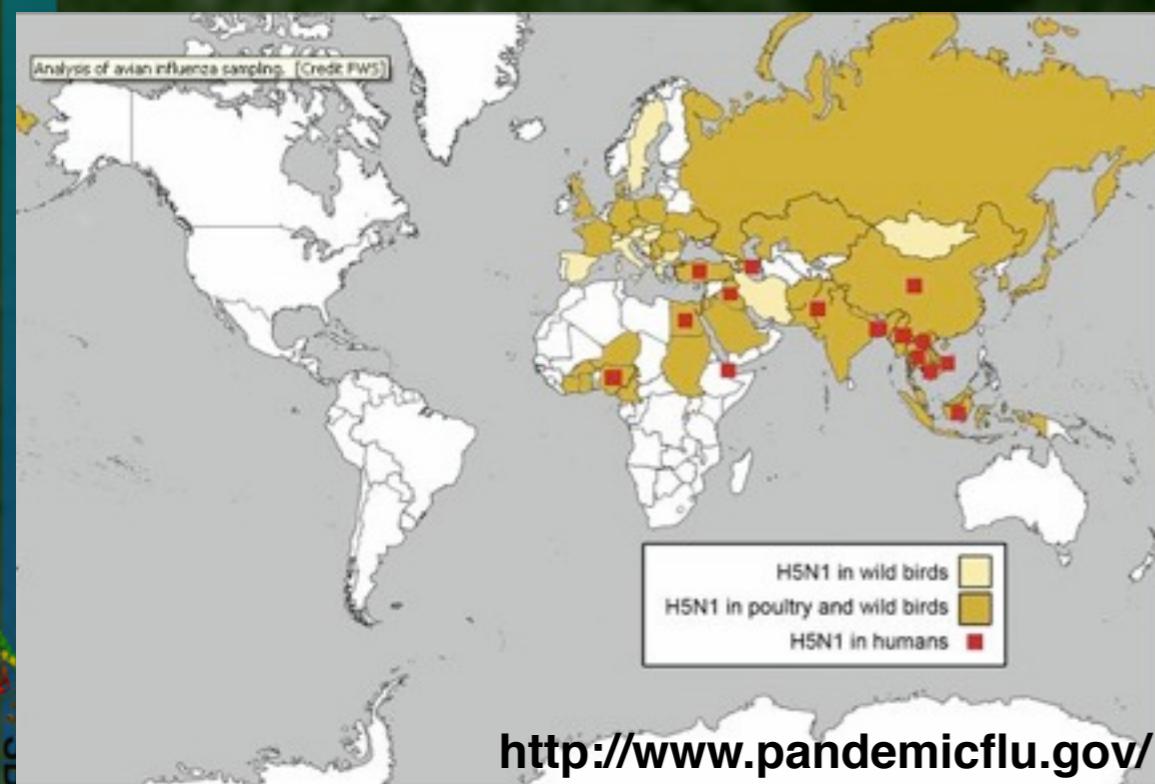
abnormal blue comb



swelling of the wattles



congestion of hocks and shanks



“Avian (or bird) flu is caused by influenza viruses that occur naturally among wild birds. The H5N1 variant is deadly to domestic fowl and can be transmitted from birds to humans. There is no human immunity and no vaccine is available.”

As of 2010:

- 508 human cases
- 302 deaths

BIOLOGICAL MODEL: INFLUENZA

H5N1

As of 2010:

- 508 human cases
- 302 deaths



H1N1

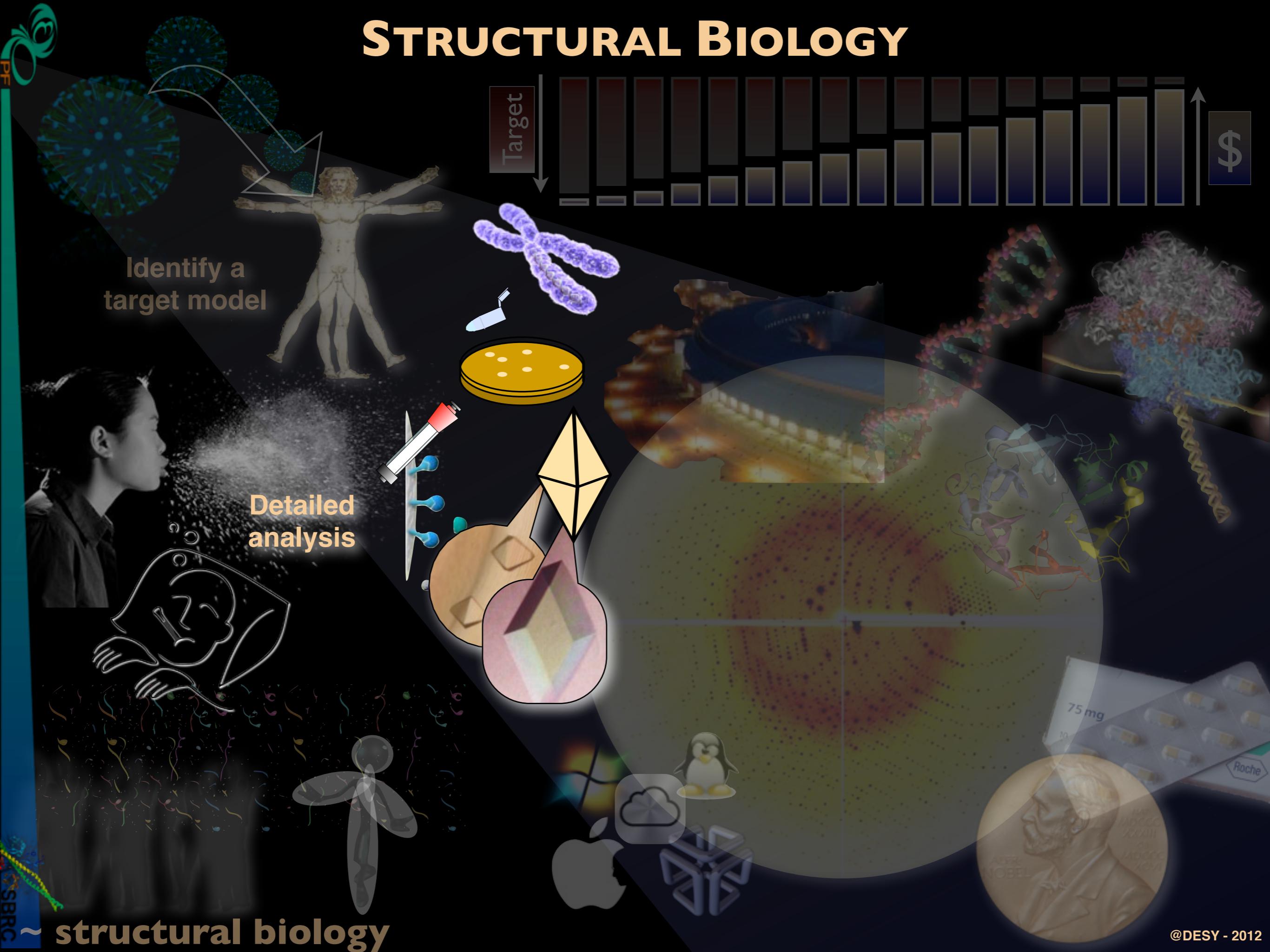
As of 2010:

- ~ 61 million human cases
- ~ 12 470 deaths



“H1N1 is a new influenza A virus that has never before circulated among humans. This virus is not related to previous or current human seasonal influenza viruses.”

STRUCTURAL BIOLOGY

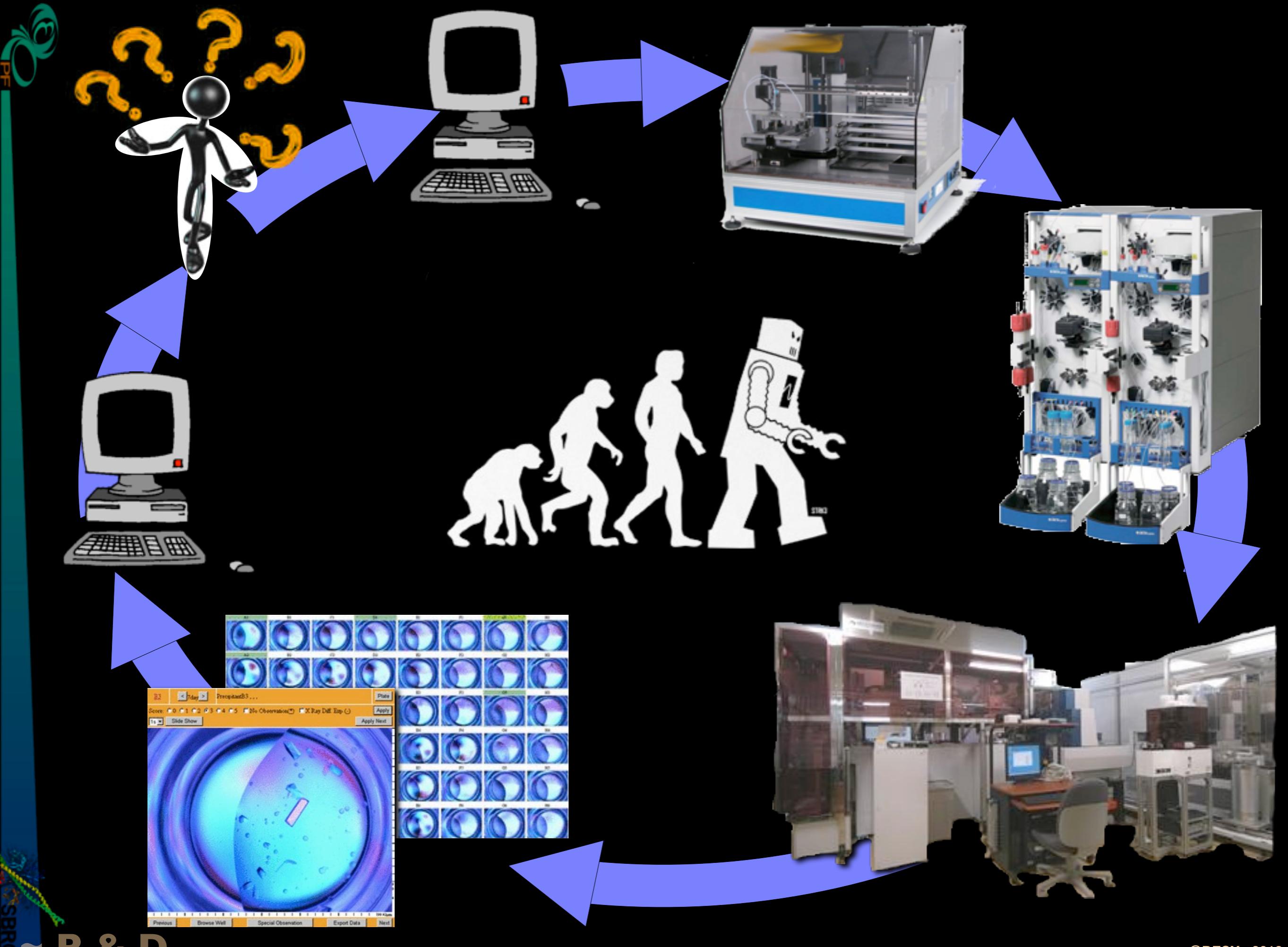


RESEARCH AND SUPPORT ACTIVITIES

The collage includes:

- A grid of 24 molecular structures (ribbons) in various colors.
- A detailed diagram of a protein-DNA complex.
- A graph showing a red curve and concentric circles with labels P, S, CI.
- A circular diagram with concentric rings and a central triangle.
- A photograph of a white building labeled "STRUCTURAL BIOLOGY RESEARCH CENTER" with the text "established in 2002" below it.
- A 3D surface plot of experimental data.
- A photograph of a laboratory interior with equipment and a sign reading "AR-NE3A astellas".
- A photograph of a laboratory bench with various pieces of equipment.
- A photograph of a robotic arm or similar laboratory equipment.
- A large blue arrow pointing diagonally from the bottom left towards the center, containing the text "STRUCTURAL & FUNCTIONAL BIOLOGY" and "R&D".
- A large blue arrow pointing diagonally from the bottom right towards the center, containing the text "SYNCHROTRON SCIENCE".

• What are we doing?

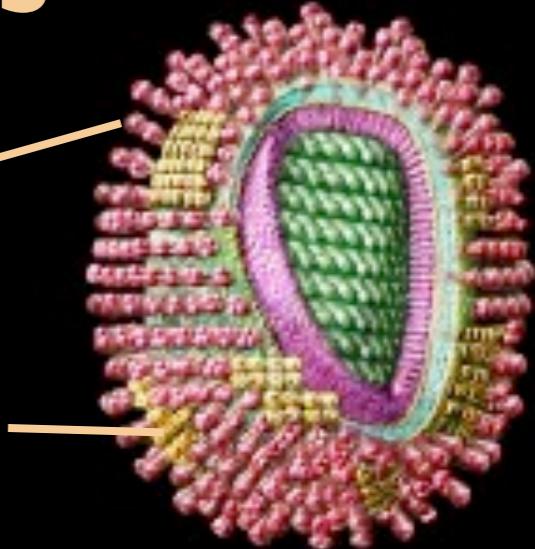


INFLUENZA VIRUS PROTEINS



Haemagglutinin (HA)

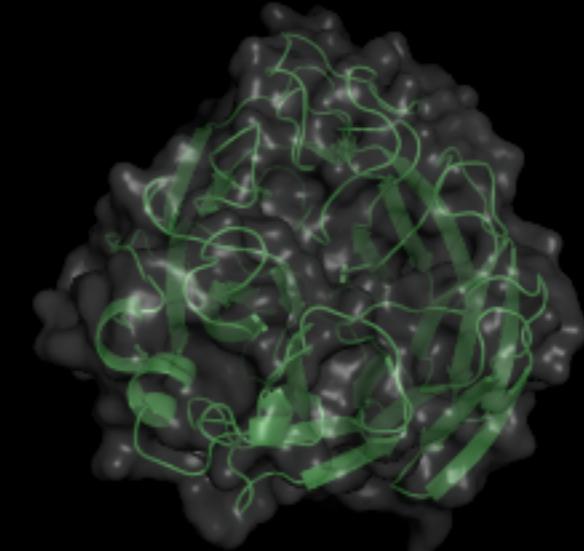
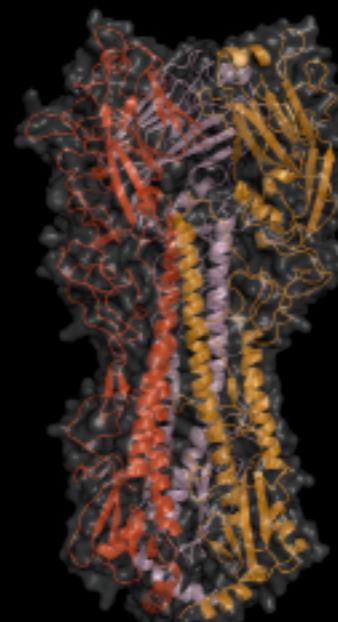
Neuraminidase (NA)



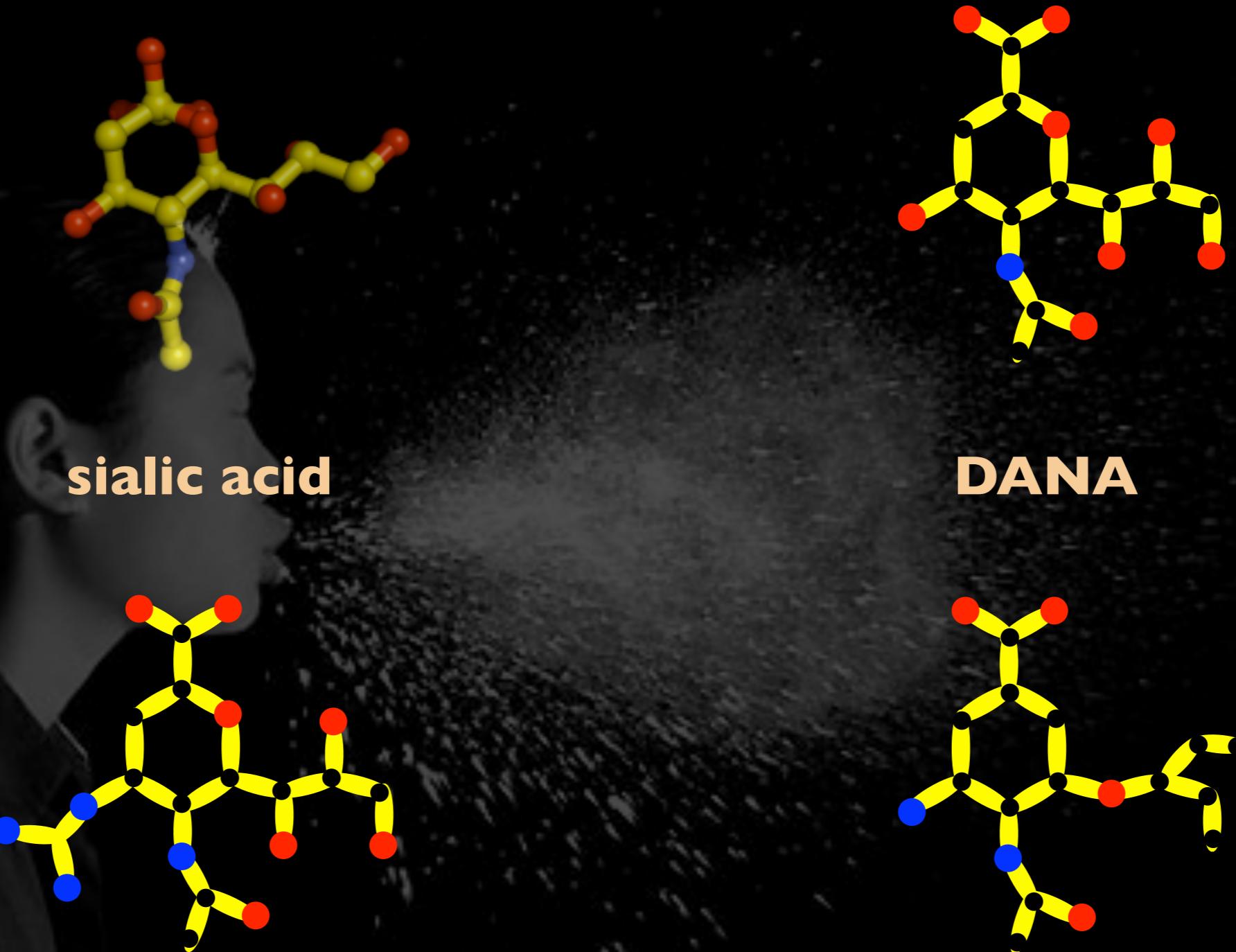
Influenza virus is an RNA type virus

Haemagglutinin (HA) & Neuraminidase (NA) are 2 surface proteins that help to parasite the host cell

- **HA recognizes sialic acid receptors at the plasma membrane; necessary for the virus entrance**
- **NA hydrolyzes the sialic acid of the receptor; necessary for newly formed virion's release and virus propagation**



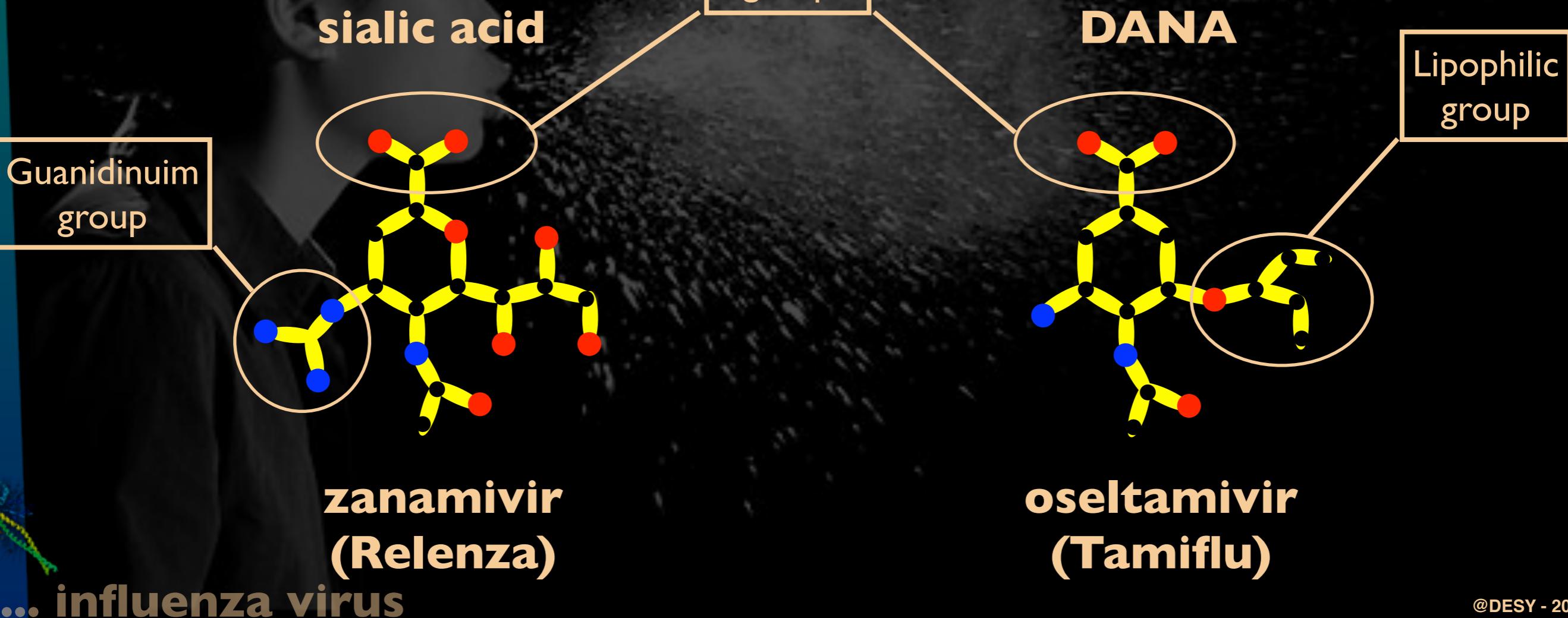
ANTI-VIRAL DRUGS FOR FLU TREATMENT



ANTI-VIRAL DRUGS FOR FLU TREATMENT

Recognition specificity:

- carboxyl head group
- guanidinium group
- lipophilic group

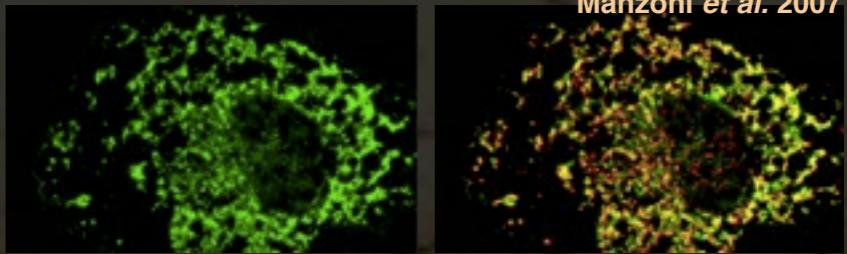




HUMAN NAs

- Neu1

- lysosomal membrane sialidase
- mutations associated with lysosomal storage diseases
(sialidosis; galactosialidosis)



- Neu2

- cytoplasmic sialidase
- low expression, mainly in skeletal muscle
- hydrolysis of GM₃; weak against GM₂ or GM₁
- induces proliferation and spontaneous differentiation of myoblast cells
- potential role in early steps of neuronal differentiation

- Neu3

- plasma membrane sialidase
- implicated in cell signaling and insulin signaling



- Neu4

- mitochondrial? lysosomal sialidase?
- implicated in apoptosis?

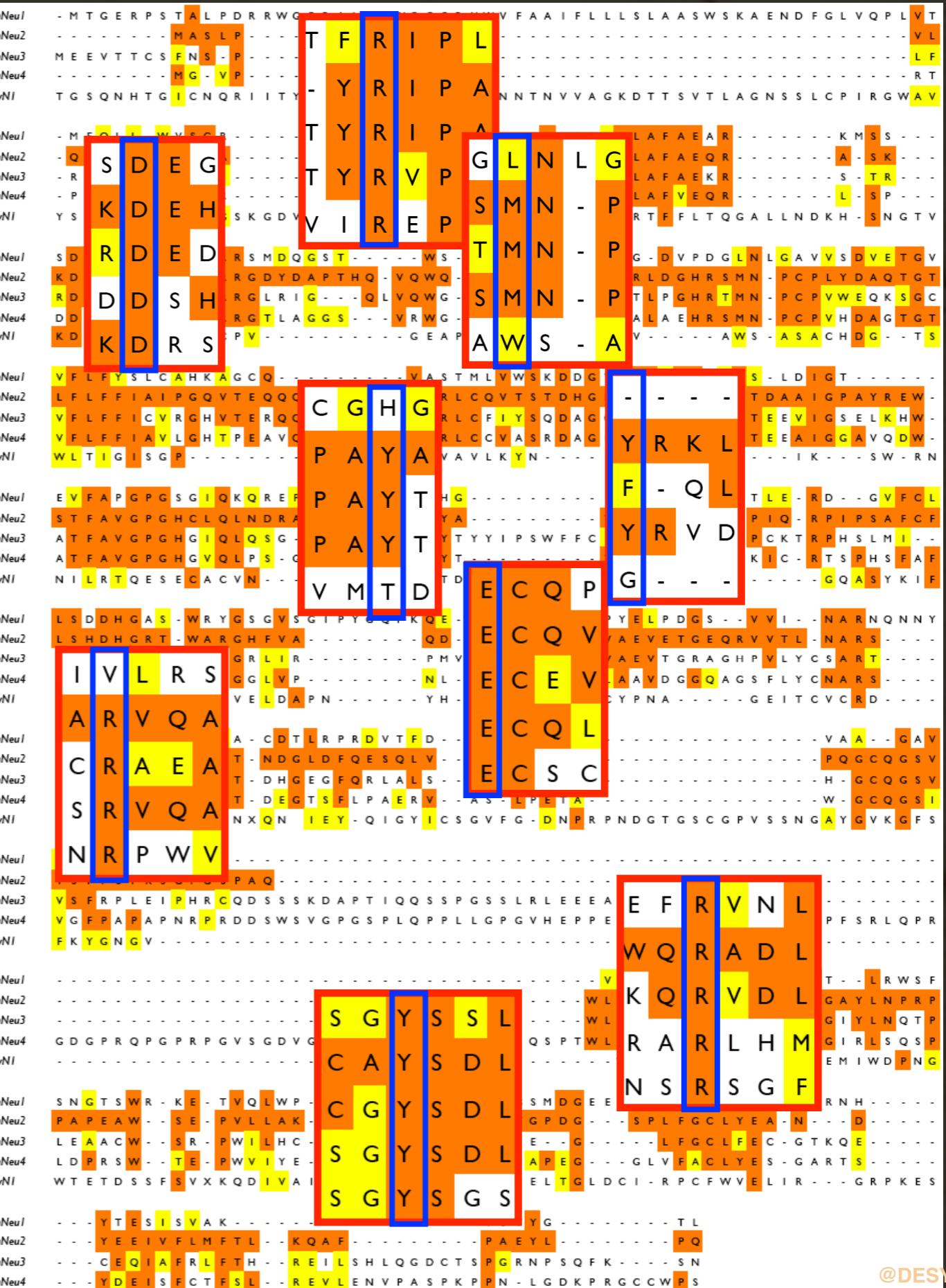


HUMAN NAS

- Human sialidases share a high sequence identity when comparing to the influenza virus neuraminidase

- Neu2 compared with:
 - Neu1.....23 %
 - Neu3.....38 %
 - Neu4.....34 %
 - influ NA...17 %

- Conserved residues for sialic acid coordination and enzymatic activity



... influenza virus

EFFECTS OF ANTI-FLU DRUGS IN HUMAN?

**WED., NOV.
15TH, 2006**

Tamiflu will carry caution on new label

Roche Laboratories Inc.:
“ ...dose dependent increase in the incidence rates of a variety of minor skeletal abnormalities...”

GlaxoSmithKline:

“ ...increase in the incidence rates of a variety of minor skeleton alterations...”

um, hallucinations and other unusual psychiatric behavior in children treated with the drug. Most were Japanese children.

The Food and Drug Administration said a relationship between the drug and the behavior had not been established and that the

spark the abnormal behavior flagged in the updated label.

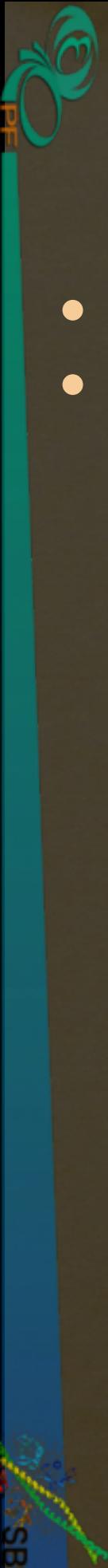
The surprise change came three days before an FDA panel of outside experts was to discuss whether to recommend that the agency add the precautionary language to the Tamiflu label.

In documents

flu and users' reported bizarre behavior. REUTERS PHOTO

of bird flu. The drug doesn't prevent flu but can reduce the length and severity of its symptoms.

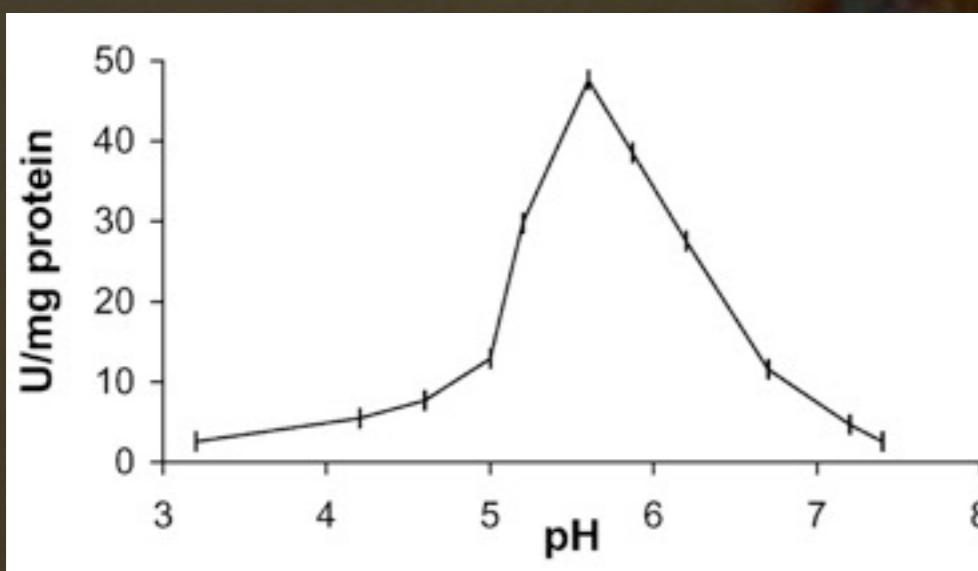
Previously, Roche has cited studies from the United States and Canada that



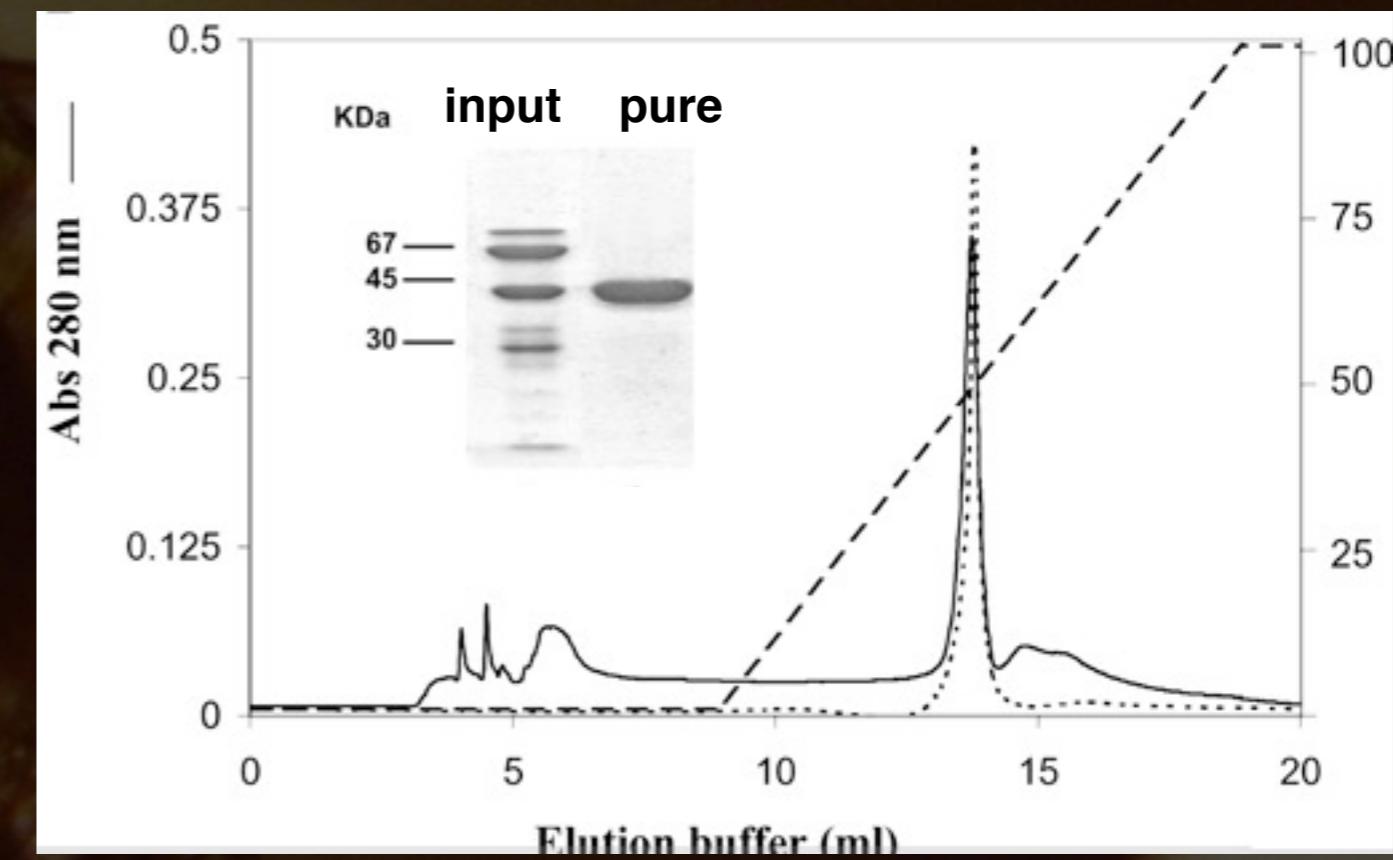
NEU2 SAMPLE PREPARATION

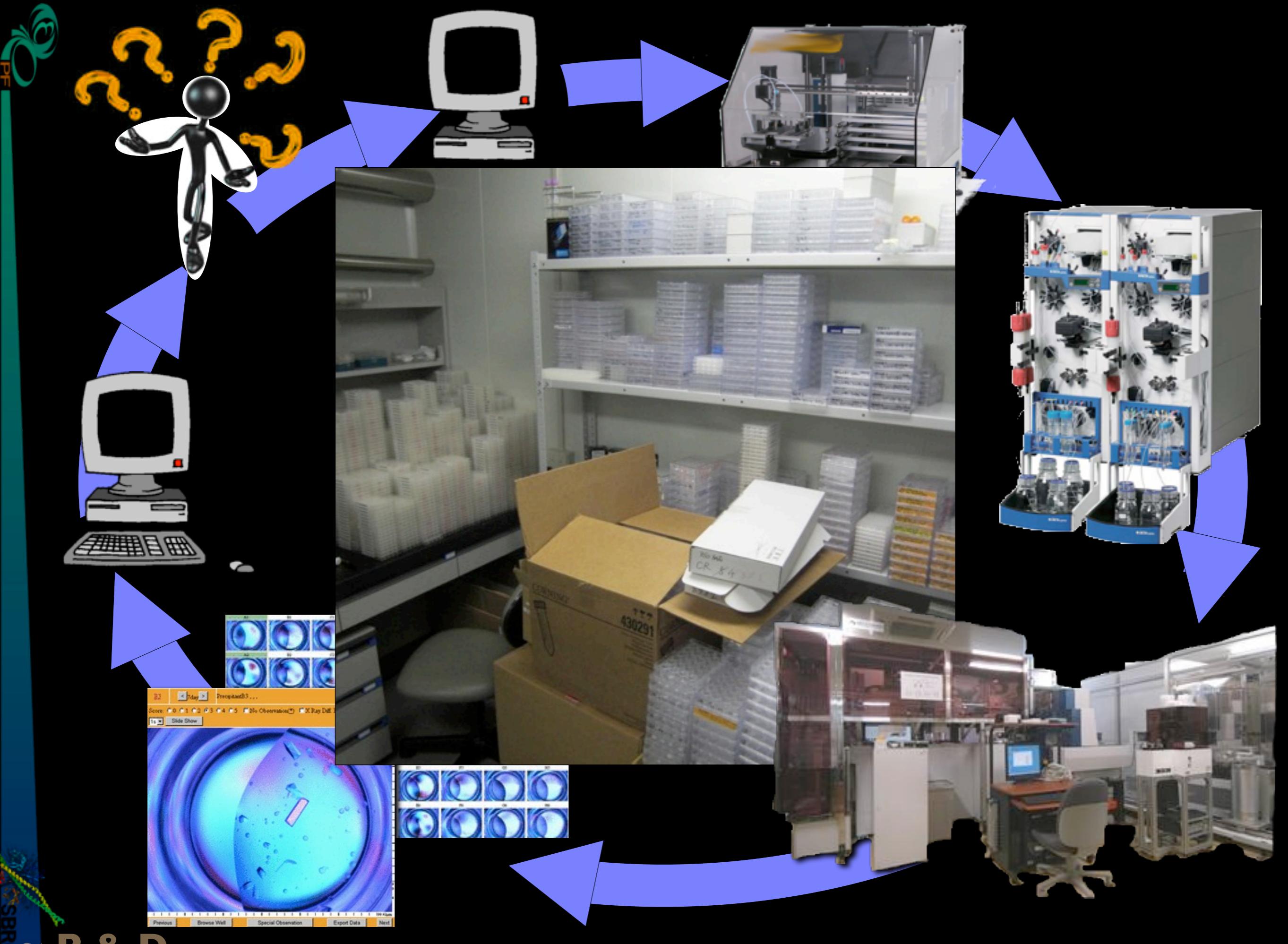
- Recombinant expression (GST-tagged protein)
- Protein purification (affinity, ion exchange chromatography)

- > monodisperse sample
- > highly pure and intact
- > ready for crystallization
- > active protein



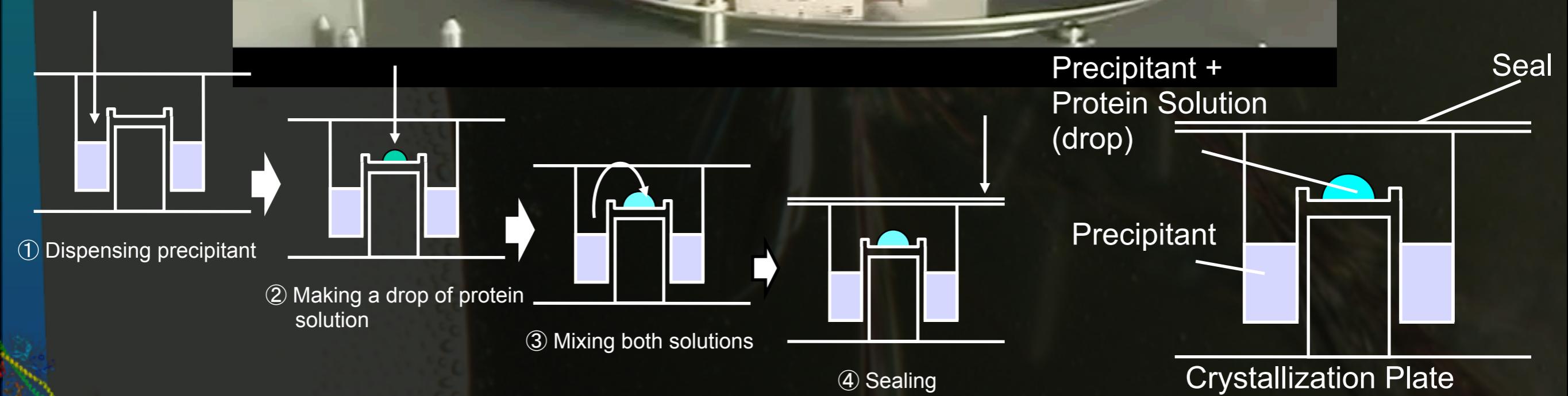
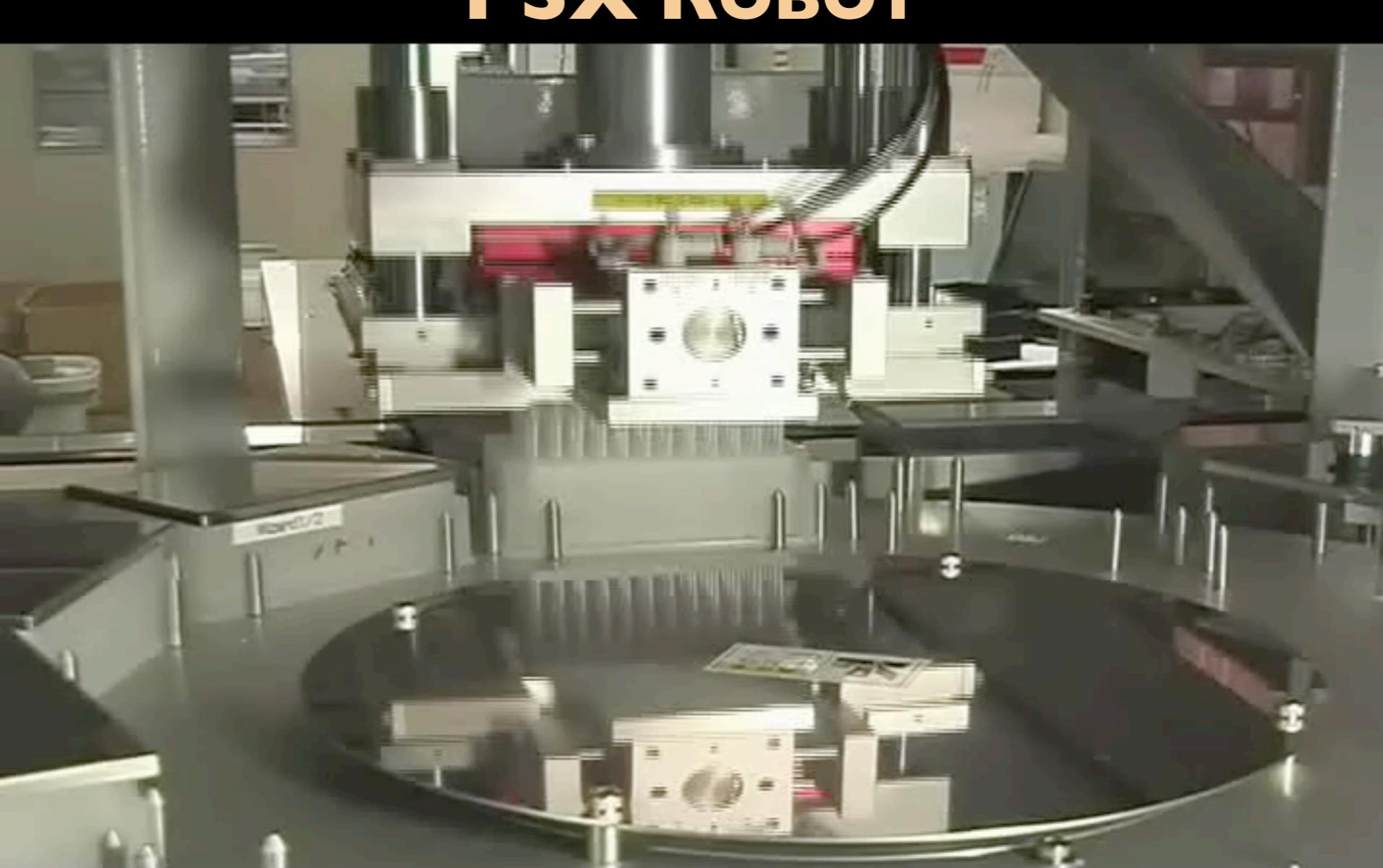
activity test at different pH



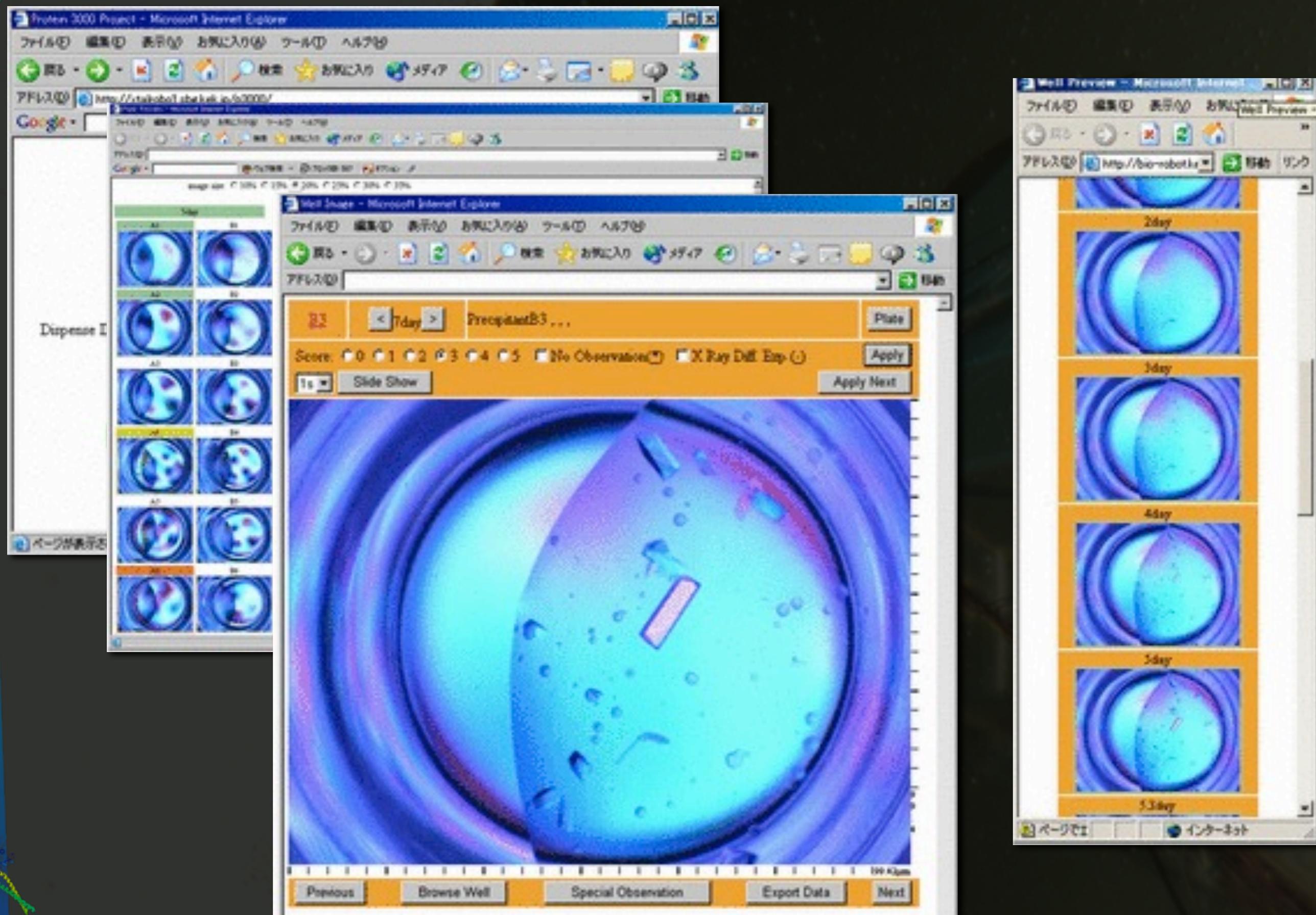




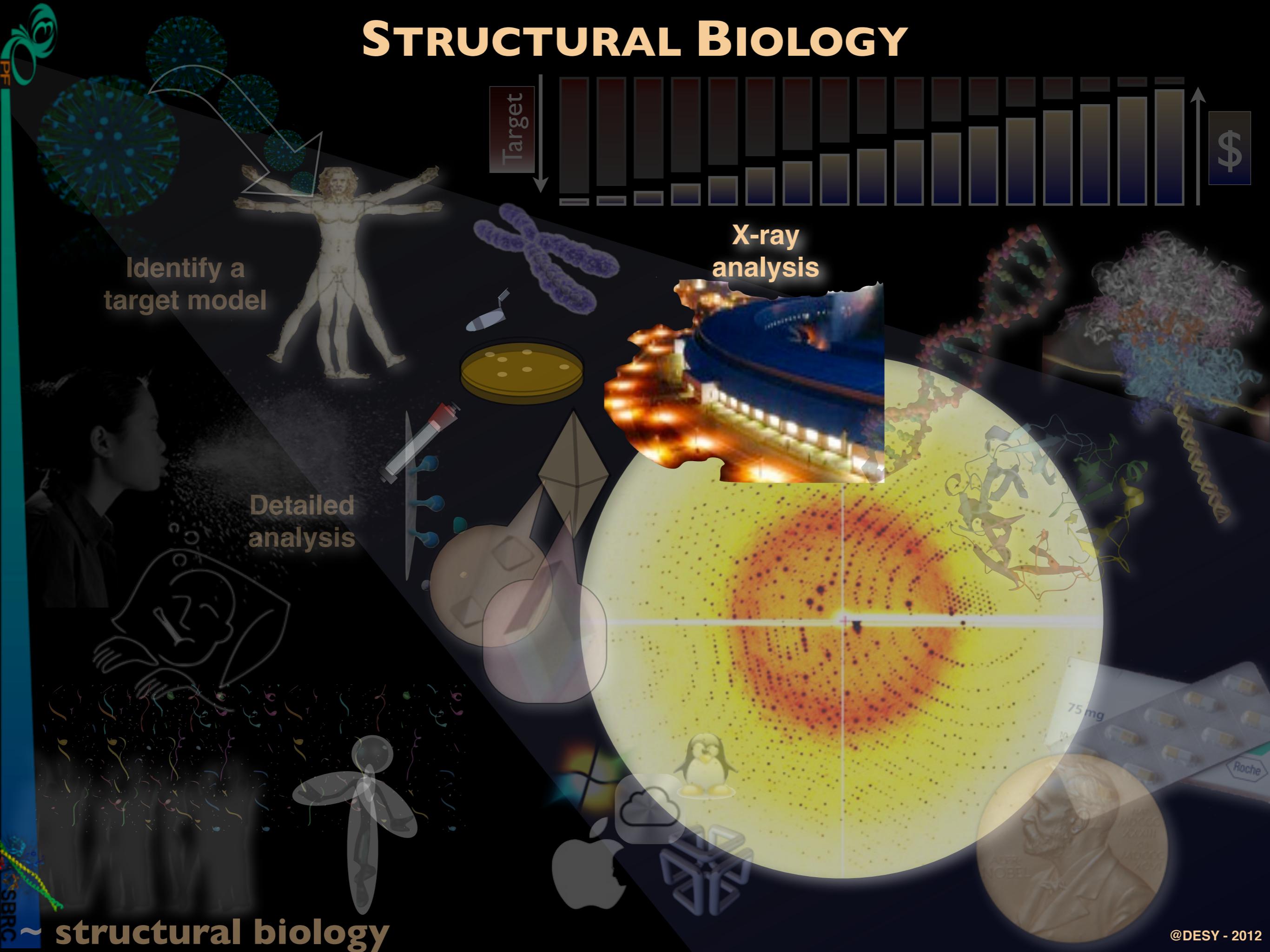
PSX ROBOT



OBSERVATION SYSTEM



STRUCTURAL BIOLOGY



RESEARCH AND SUPPORT ACTIVITIES

The collage illustrates various research and support activities:

- STRUCTURAL & FUNCTIONAL BIOLOGY:** A grid of protein structures (ribbons) and a photograph of the Structural Biology Research Center building, established in 2002.
- R&D:** Laboratory equipment including a robotic pipetting station and a circular array plate.
- SYNCHROTRON SCIENCE:** A diagram of a synchrotron beamline, a plot of experimental data, and a photograph of the ARNE3A beamline at DESY.
- Support Activities:** A graph showing energy loss fine-structure (EXELFS) data, a periodic table of elements, and a molecular model.

Diagonal text labels: **STRUCTURAL & FUNCTIONAL BIOLOGY** and **SYNCHROTRON SCIENCE**.

• What are we doing?

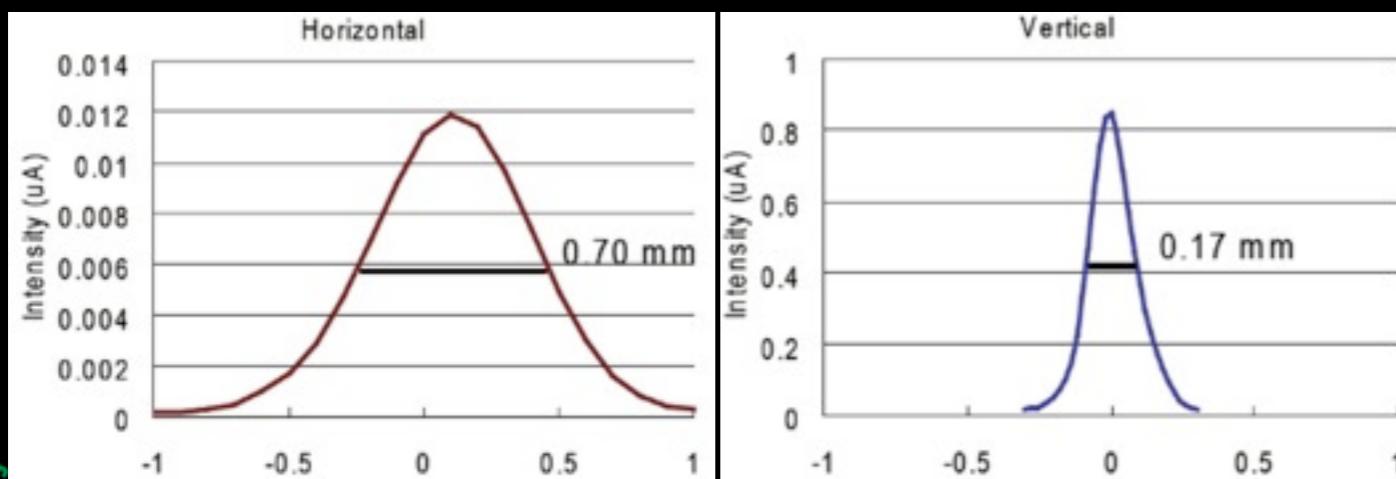
RESEARCH AND SUPPORT ACTIVITIES

	BL-1A 2010 Operational	BL-5A 2004 Operational PF Top-up	BL-17A 2006 Operational	AR-NW12A 2003 Operational PF AR	AR-NE3A 2009 Operational Twice / day (8:30. 20:30)
Starting year	2010	2004	2006	2003	2009
Status	Operational	Operational	Operational	Operational	Operational
Synchrotron Ring					
Injection					
X-ray source	Short-gap undulator	Multipole wiggler	Short-gap undulator	Undulator	Undulator
Wavelength (Å)	0.9 - 1.1 / 2.7 - 3.0	0.7 - 1.9	0.9 - 2.1	0.7 - 1.9	0.7 - 1.9
Energy resolution ($\Delta E / E$)	2.5×10^{-4}	2.5×10^{-4}	2.5×10^{-4}	2.5×10^{-4}	2.5×10^{-4}
Photon flux (photons / sec, @ 1.0 Å)	2×10^{10}	2×10^{11}	6.6×10^9	2.0×10^{11}	8.0×10^{11}
Slit size for flux measurement (μm^2)	10 x 10	200 x 200	20 x 20	200 x 200	200 x 200
Detector	Quantum 270	Quantum 315r	Quantum 210r	Quantum 210r	Quantum 270
Type	CCD	CCD	CCD	CCD	CCD
Active area (mm²)	270 x 270	315 x 315	210 x 210	210 x 210	270 x 270
Pixel size (μm^2)	64.8 x 64.8	102 x 102	51 x 51	102 x 102	64.8 x 64.8
Pixel number	4168 x 4168	3072 x 3072	4096 x 4096	2048 x 2048	4168 x 4168
Frame data size (MByte)	34	19	32	8.0×10^{11}	34
Readout time (sec)	1.1	0.3	1	0.3	1.1
Typical exposure time (sec / deg)	Exp. time + 2.7	Exp. time + 1.6	Exp. time + 2.7	Exp. time + 1.6	Exp. time + 2.7
Data collection time per frame (sec)	11	14	24	14	11
Typical data collection time (min, 180 frames) (using the IEEE1394 interface)	6	6 (24)	6	6	6
Camera distance (mm)	40 - 500	60-950	40-700	60-950	60-500
Detector vector offset (mm)	0 - 100	0 - 150	0-100	0-100	0-150
Sample changer			PAM		
Software image processing			HKL2000, Mosflm, XDS...		



RESEARCH AND SUPPORT ACTIVITIES

Astellas Beamline : Pharmaceuticals Beamline

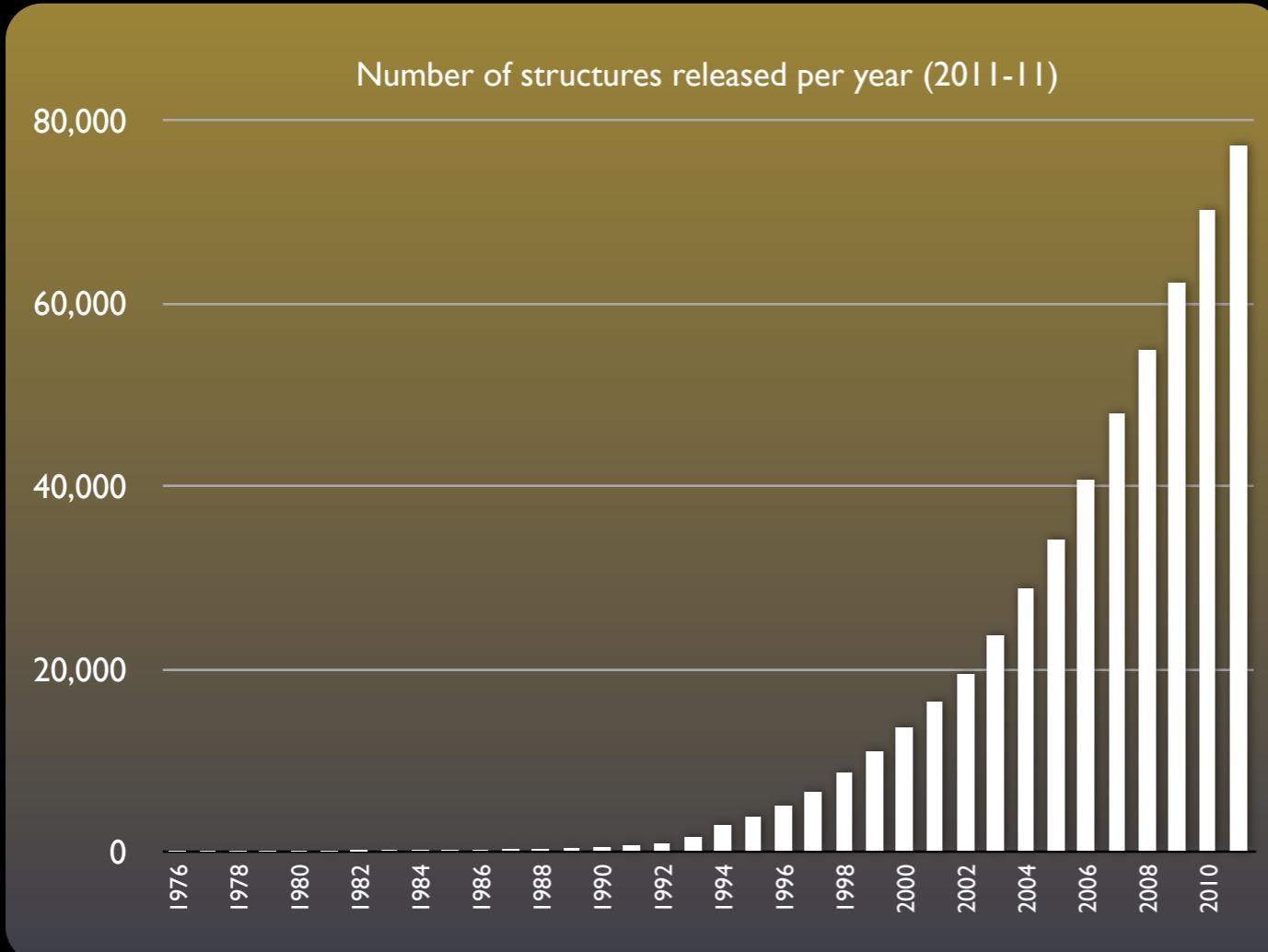


~ synchrotron science

Starting year	AR-NE3A
Status	2009
Synchrotron Ring	Operational
Injection	PF AR
X-ray source	Twice / day
Wavelength (Å)	Undulator
Energy resolution ($\Delta E / E$)	0.7 - 1.9
Photon flux (photons / sec, @ 1.0 Å)	2.5×10^{-4}
	8.0×10^{11}
Slit size for flux measurement (μm^2)	200 x 200
Detector	Quantum 270
Type	CCD
Active area (mm²)	270 x 270
Pixel size (μm²)	64.8 x 64.8
Pixel number	4168 x 4168
Frame data size (MByte)	34
Readout time (sec)	1.1
Typical exposure time (sec / deg)	Exp. time + 2.7
Data collection time per frame (sec)	11
Typical data collection time (min, 180 frames) (using the IEEE 1394 interface)	6
Camera distance (mm)	60-500
Detector vector offset (mm)	0-150
Sample changer	PAM
Software image processing	HKL2000 etc.

RESEARCH AND SUPPORT ACTIVITIES

Astellas Beamline : Pharmaceuticals Beamline



Starting year

Status

Synchrotron Ring

Injection

X-ray source

Wavelength (Å)

Energy resolution ($\Delta E / E$)

Photon flux (photons / sec, @ 1.0 Å)

AR-NE3A

2009

Operational

PF AR

Twice / day

Undulator

0.7 - 1.9

2.5×10^{-4}

8.0×10^{11}

200 x 200

Quantum 270

CCD

270 x 270

64.8 x 64.8

4168 x 4168

34

1.1

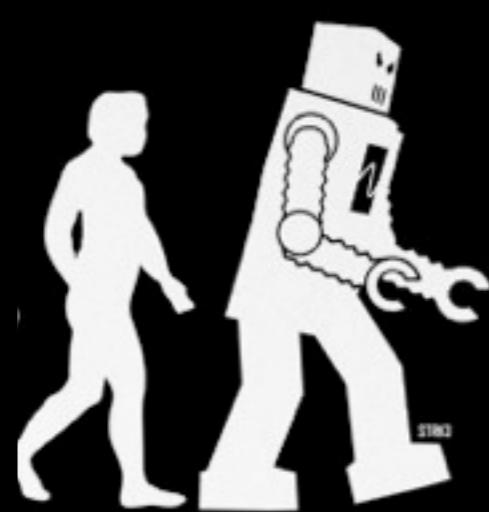
Exp. time + 2.7

11

6

60-500

0-150



Slit size for flux measurement (μm^2)

Detector

Type

Active area (mm^2)

Pixel size (μm^2)

Pixel number

Frame data size (MByte)

Readout time (sec)

Typical exposure time (sec / deg)

Data collection time per frame (sec)

**Typical data collection time (min, 180 frames)
(using the IEEE 1394 interface)**

Camera distance (mm)

Detector vector offset (mm)

Sample changer

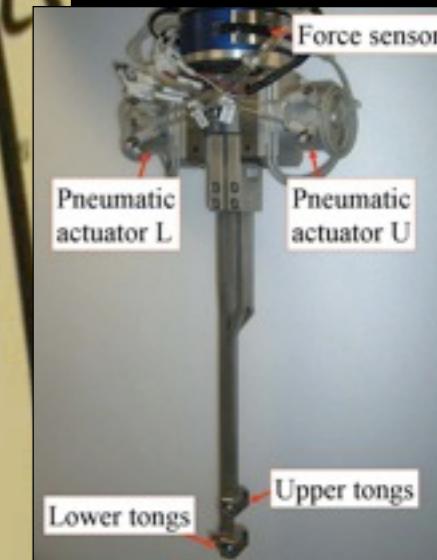
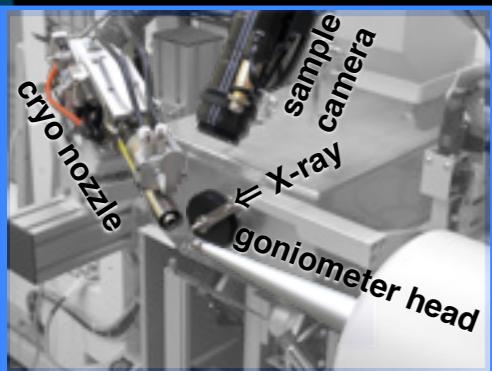
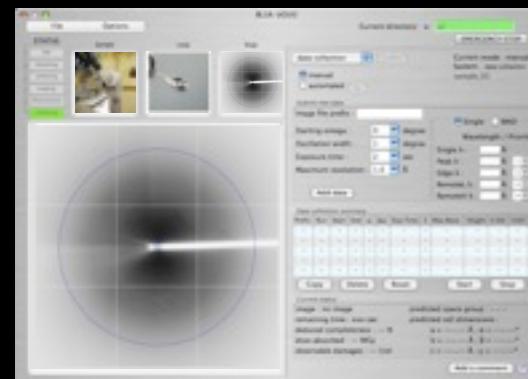
PAM

Robots to the rescue !

synchrotron science

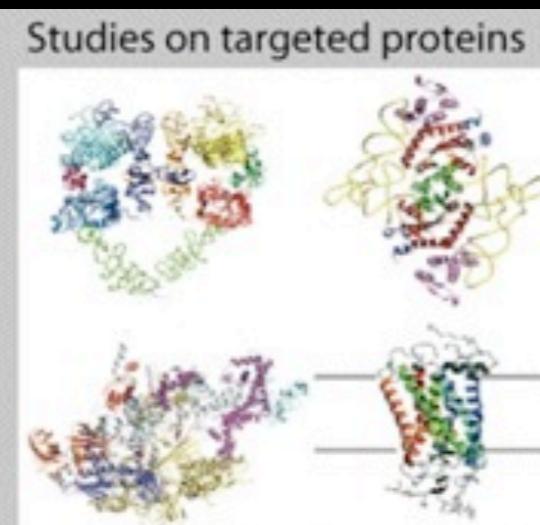
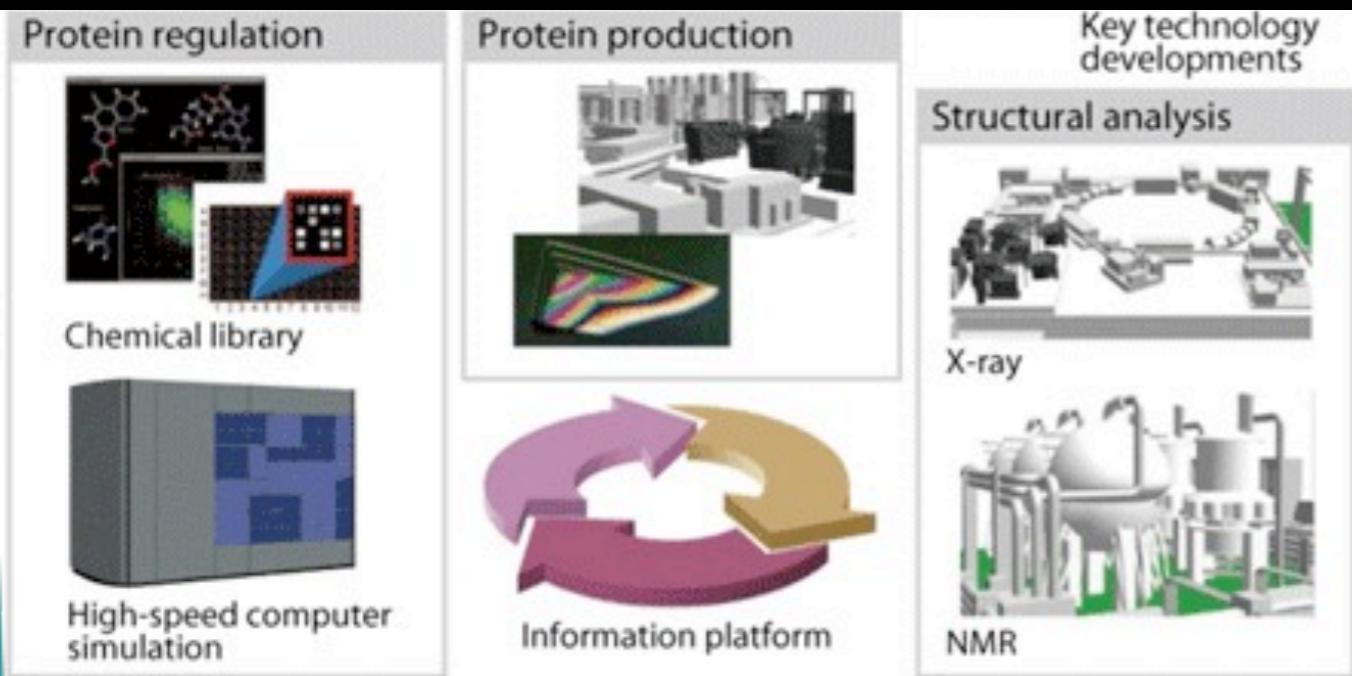
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PF AUTOMATED MOUNTING SYSTEM



RESEARCH AND SUPPORT ACTIVITIES

Low Energy Beamline

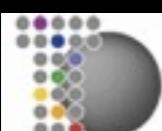


Contribution to medical and pharmaceutical sciences

Applications to food production and bioremediation, etc

Investigations of fundamental biological phenomena

Return the research results to the society



TPRP

Targeted Proteins Research Program

MEXT



Starting year

Status

Synchrotron Ring

Injection

X-ray source

Wavelength (Å)

Energy resolution ($\Delta E / E$)

Photon flux (photons / sec, @ 1.0 Å)

Slit size for flux measurement (μm^2)

Detector

Type

Active area (mm²)

Pixel size (μm^2)

Pixel number

Frame data size (MByte)

Readout time (sec)

Typical exposure time (sec / deg)

Data collection time per frame (sec)

**Typical data collection time (min, 180 frames)
(using the IEEE1394 interface)**

Camera distance (mm)

Detector vector offset (mm)

Sample changer

Software image processing

BL-1A

2010

Operational

PF

Top-up

Short-gap undulator

0.9 - 1.1 / 2.7 - 3.0

2.5×10^{-4}

2×10^{10}

2×10^9 (@2.7 Å)

10×10

Quantum 270

CCD

270×270

64.8×64.8

4168×4168

34

1.1

Exp. time + 2.7

11

6

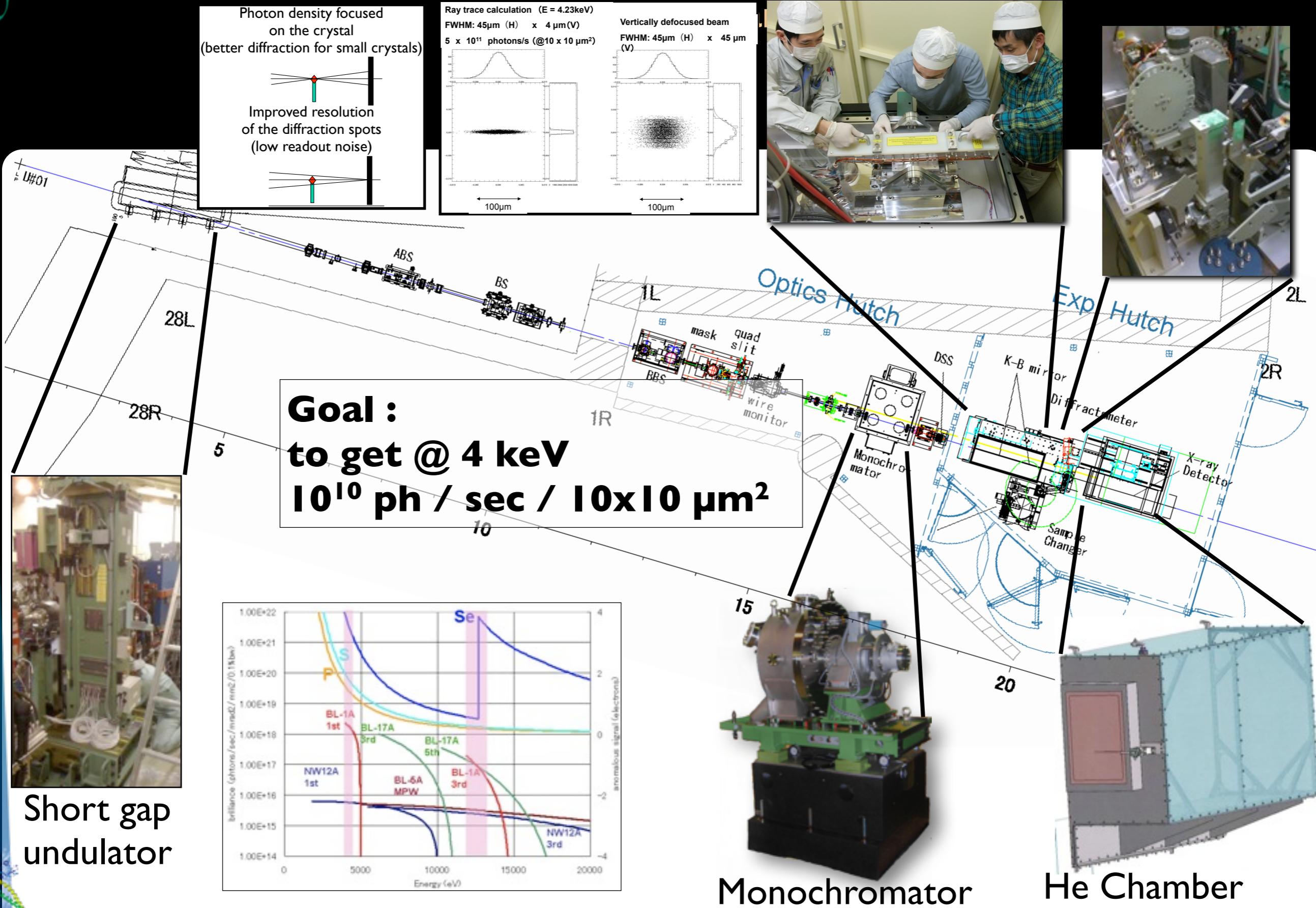
40 - 500

0 - 100

PAM

HKL2000 etc.

RESEARCH AND SUPPORT K/B Mirrors'ITIES





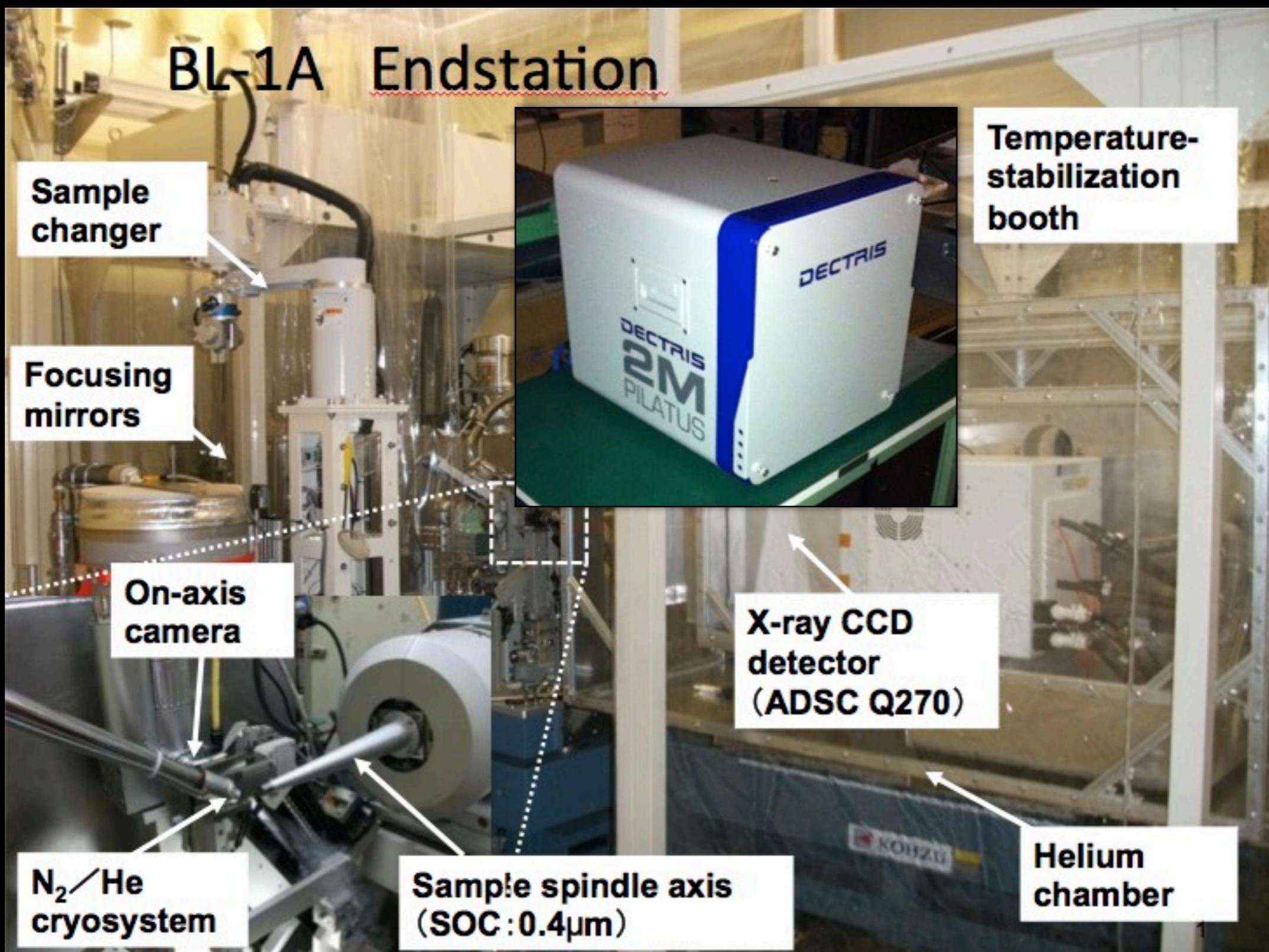
Energy (eV)

Monochromator

He Chamber

RESEARCH AND SUPPORT ACTIVITIES

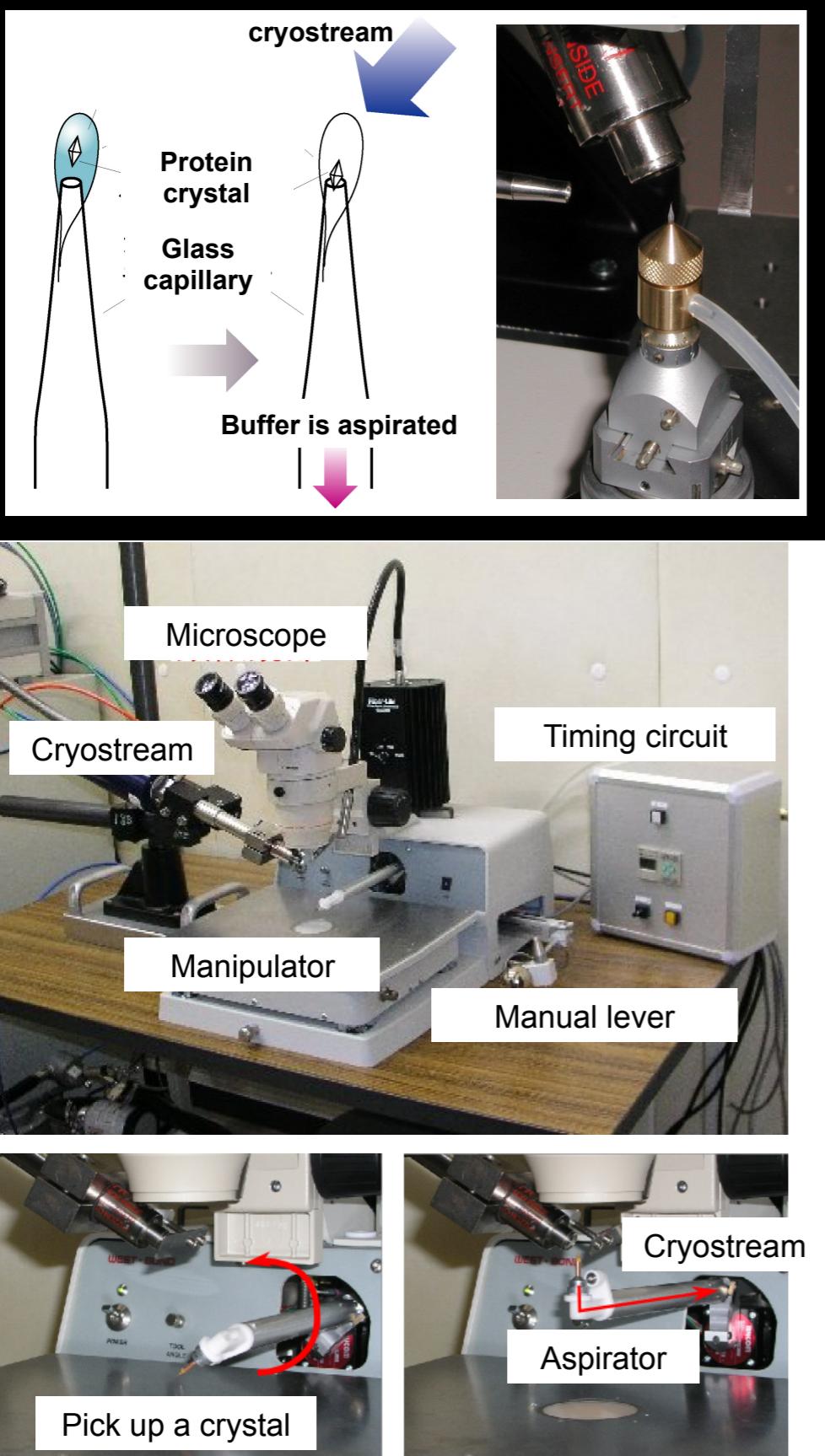
Low Energy Beamline



BL-1A	2010
Operational	PF
Top-up	Short-gap undulator
0.9 - 1.1 / 2.7 - 3.0	2.5×10^{-4}
	2×10^{10}
	2×10^9 (@2.7 Å)
	10 x 10
Quantum 270	CCD
	270 x 270
	64.8 x 64.8
	4168 x 4168
	34
	1.1
Exp. time +	2.7
	11
	6
	40 - 500
	0 - 100
	PAM
	HKL2000 etc.

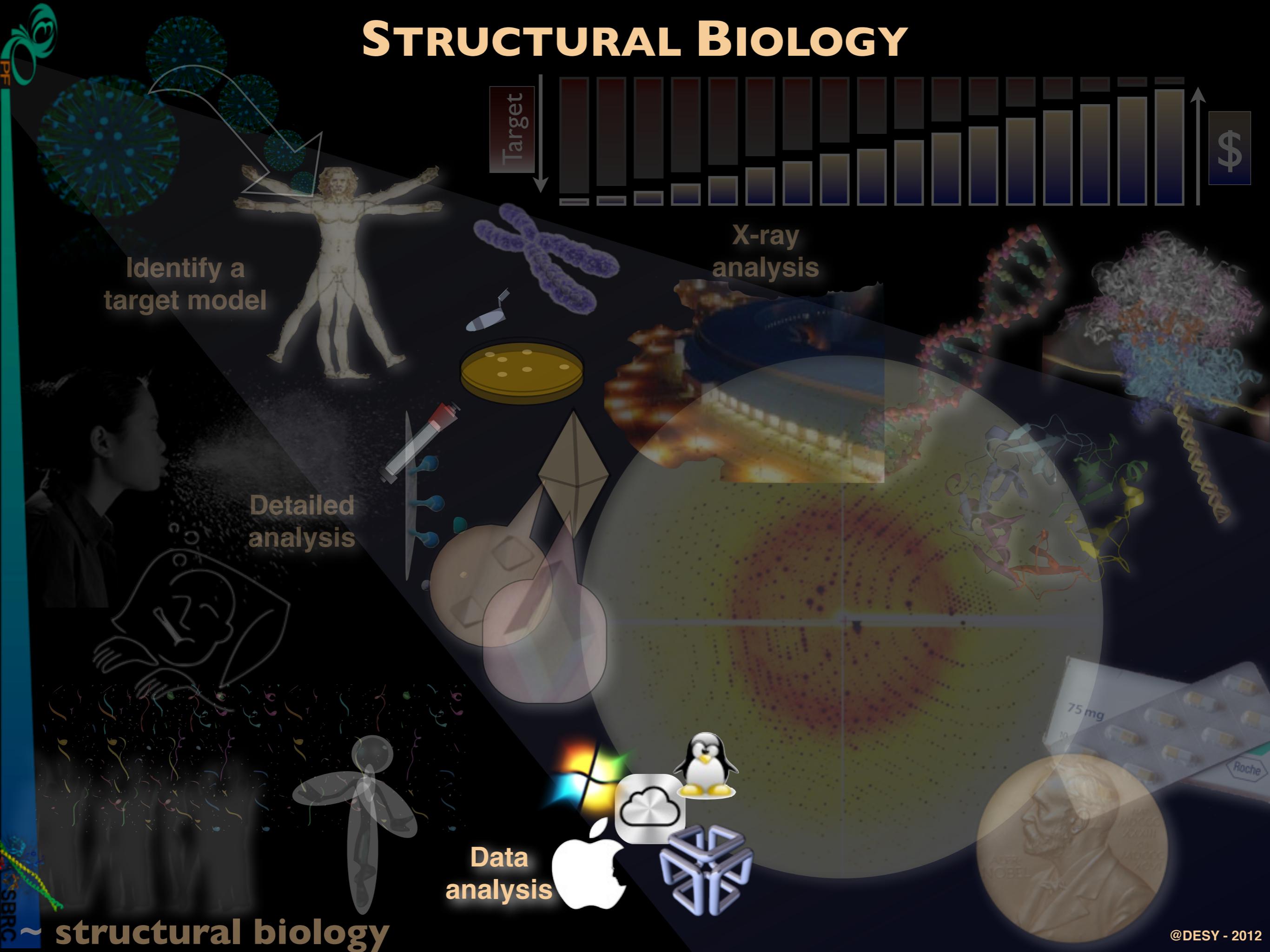
RESEARCH AND SUPPORT ACTIVITIES

Low Energy Beamline

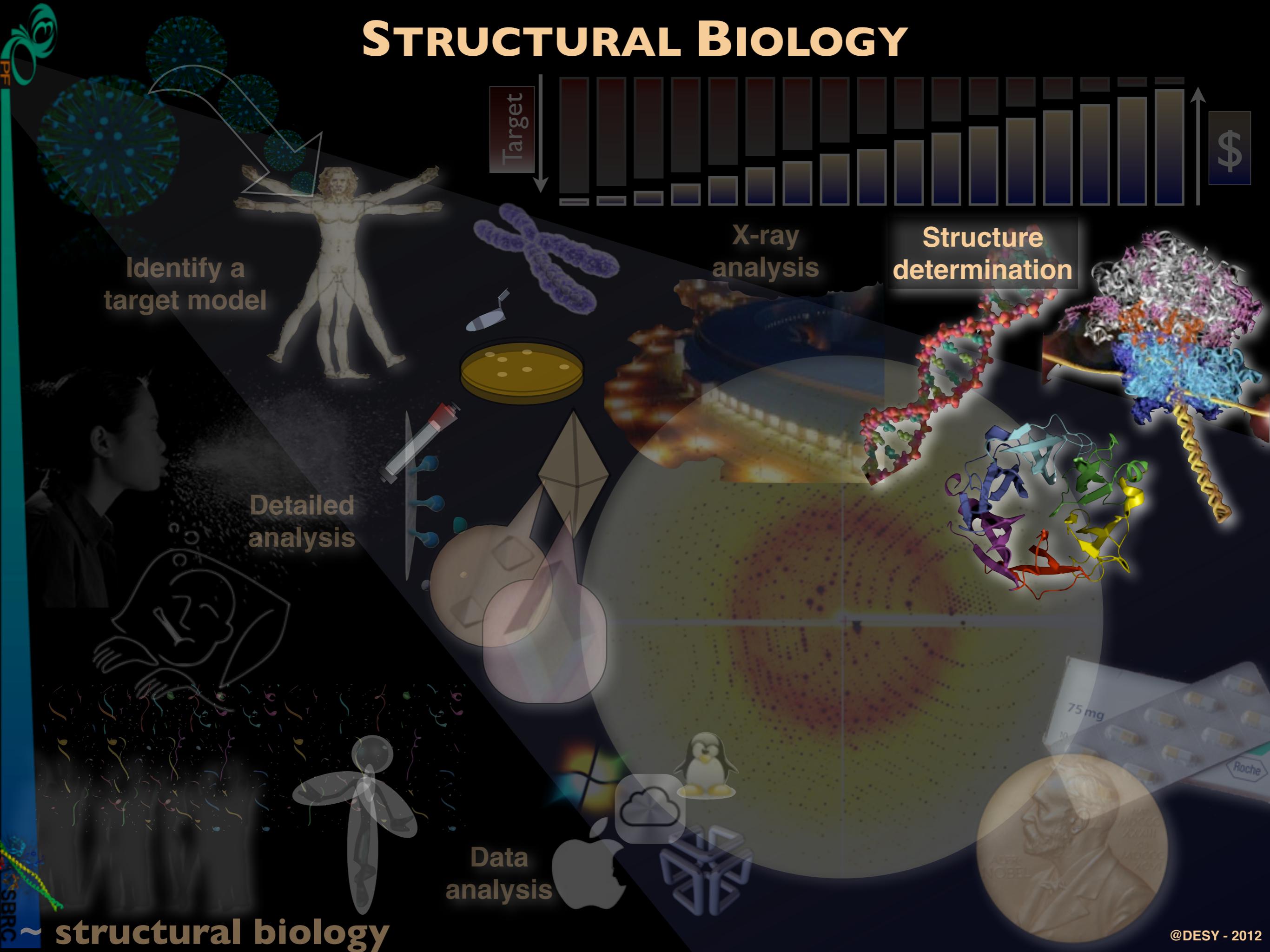


Starting year	BL-1A
Status	2010
Synchrotron Ring	Operational
Injection	PF
X-ray source	Top-up
Wavelength (Å)	Short-gap undulator
Energy resolution ($\Delta E / E$)	0.9 - 1.1 / 2.7 - 3.0
Photon flux (photons / sec, @ 1.0 Å)	2.5×10^{-4}
	2×10^{10}
	2×10^9 (@2.7 Å)
Slit size for flux measurement (μm^2)	10×10
Detector	Quantum 270
Type	CCD
Active area (mm²)	270 x 270
Pixel size (μm²)	64.8 x 64.8
Pixel number	4168 x 4168
Frame data size (MByte)	34
Readout time (sec)	1.1
Typical exposure time (sec / deg)	Exp. time + 2.7
Data collection time per frame (sec)	11
Typical data collection time (min, 180 frames) (using the IEEE1394 interface)	6
Camera distance (mm)	40 - 500
Detector vector offset (mm)	0 - 100
Sample changer	PAM
Software image processing	HKL2000 etc.

STRUCTURAL BIOLOGY



STRUCTURAL BIOLOGY



Identify a target model

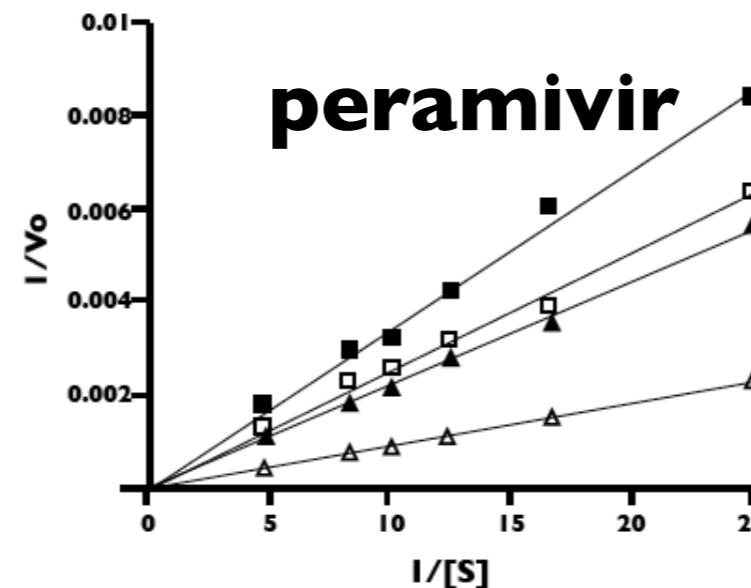
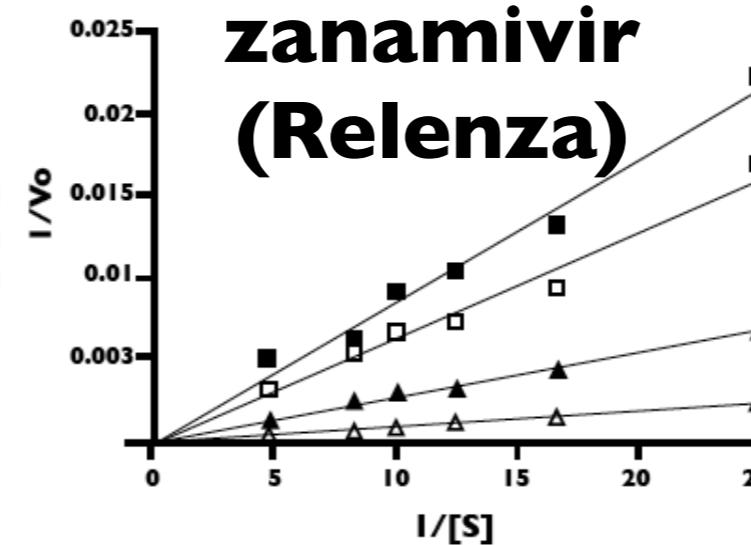
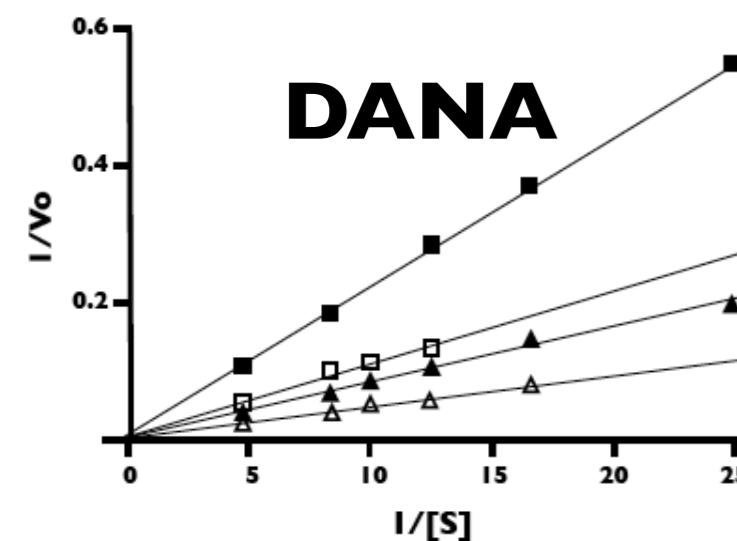
X-ray analysis

Structure determination

Detailed analysis

Data analysis

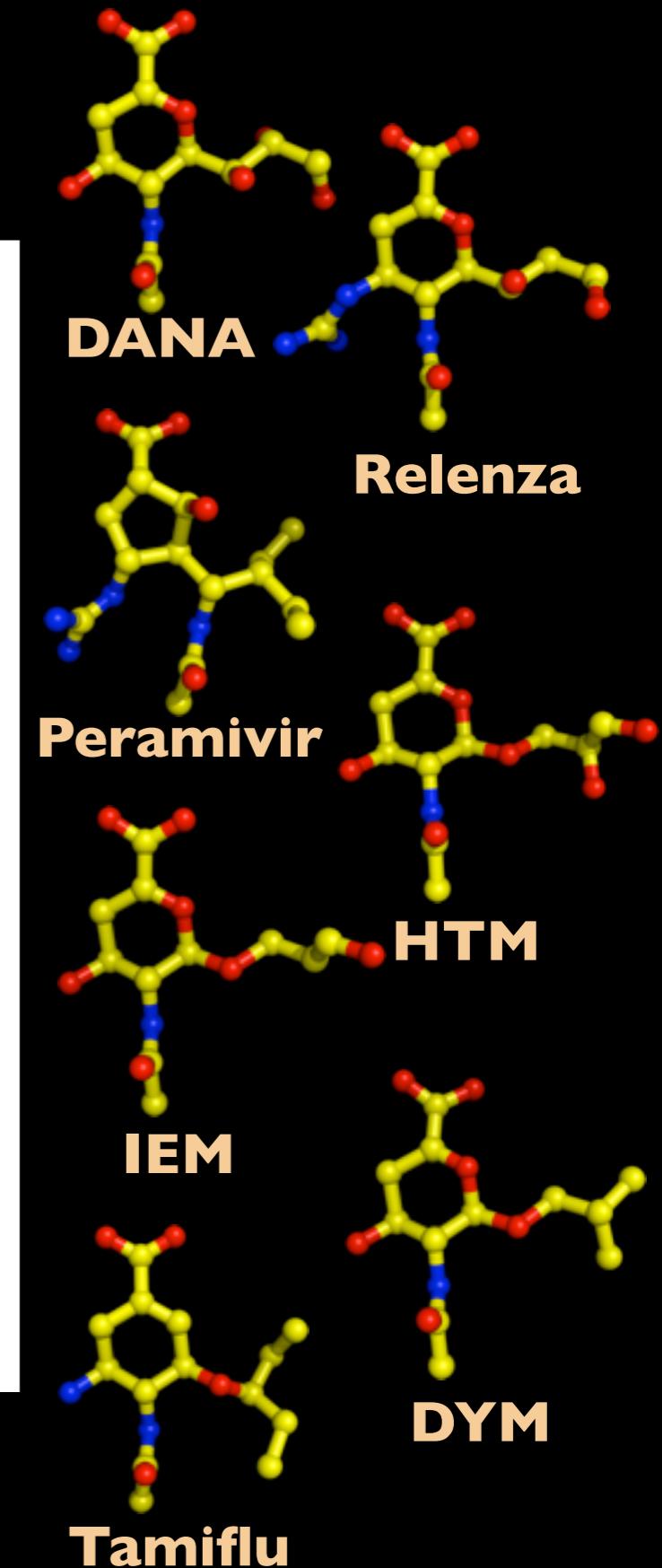
NEU2 INHIBITION BY NIs

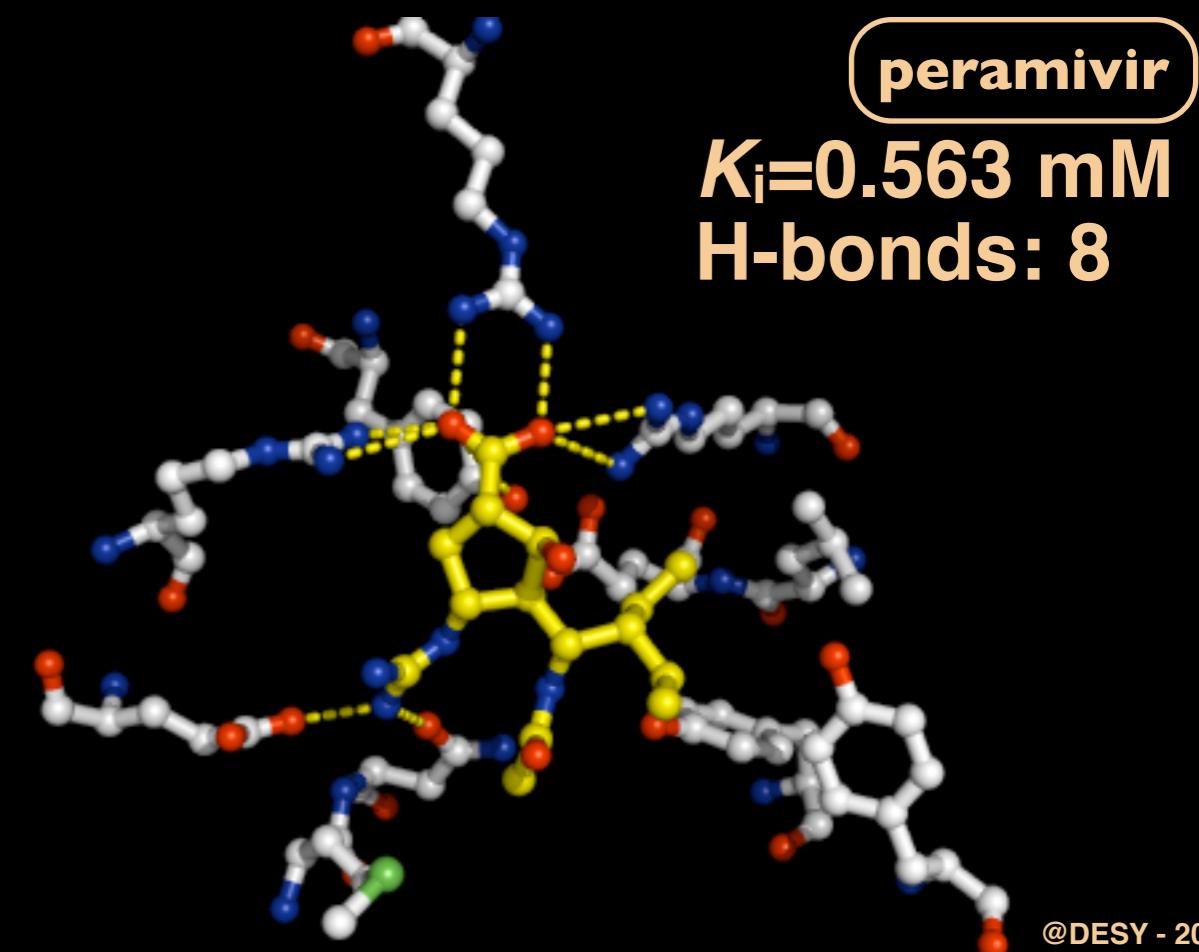
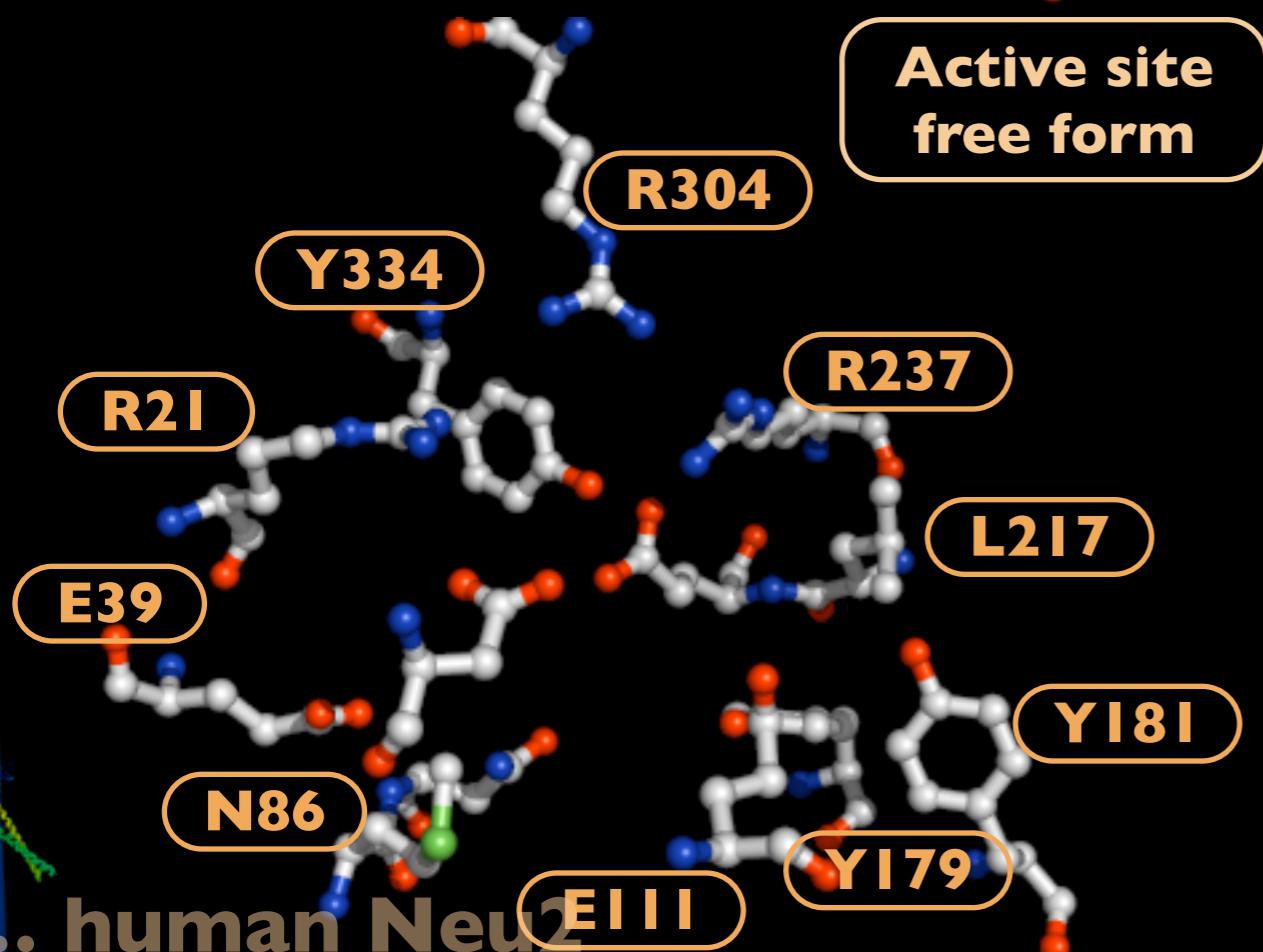
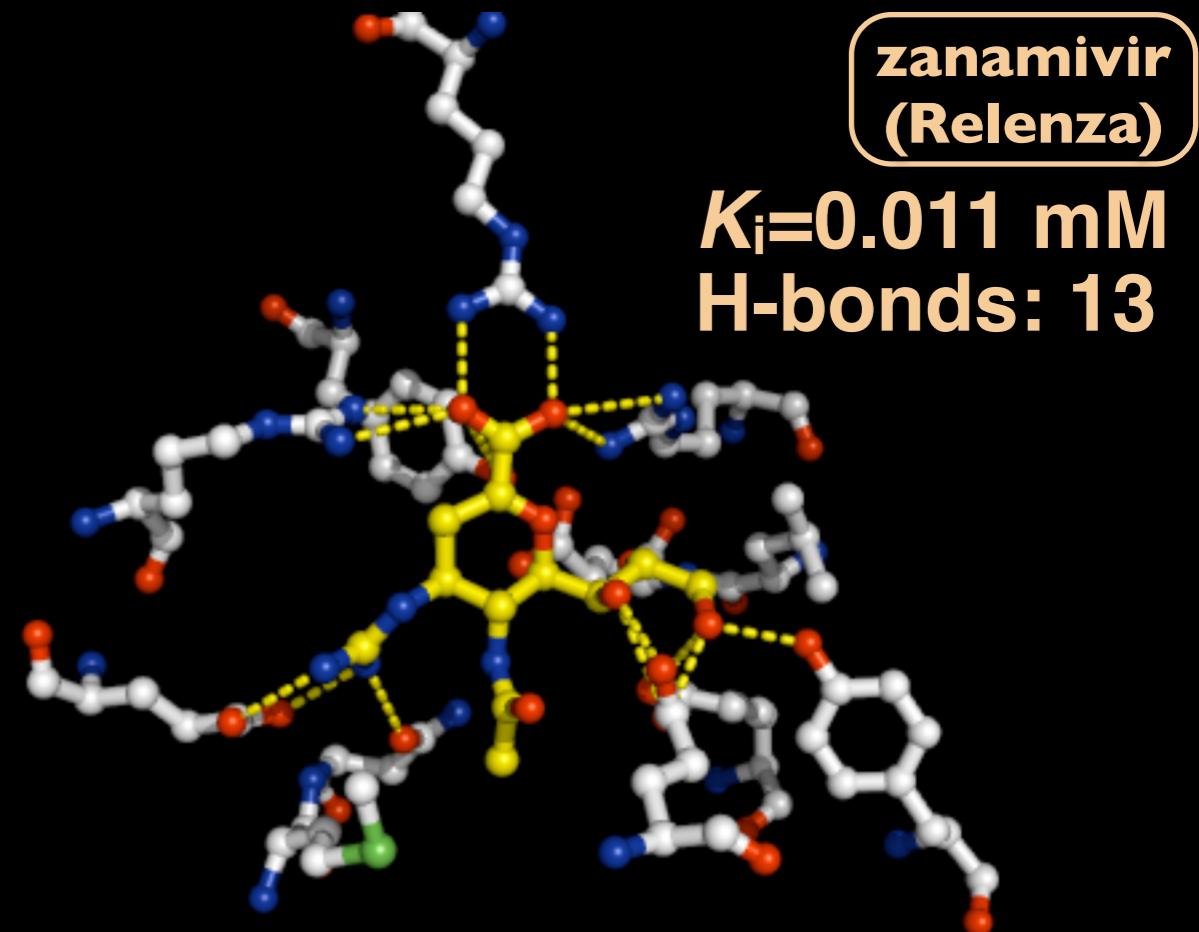
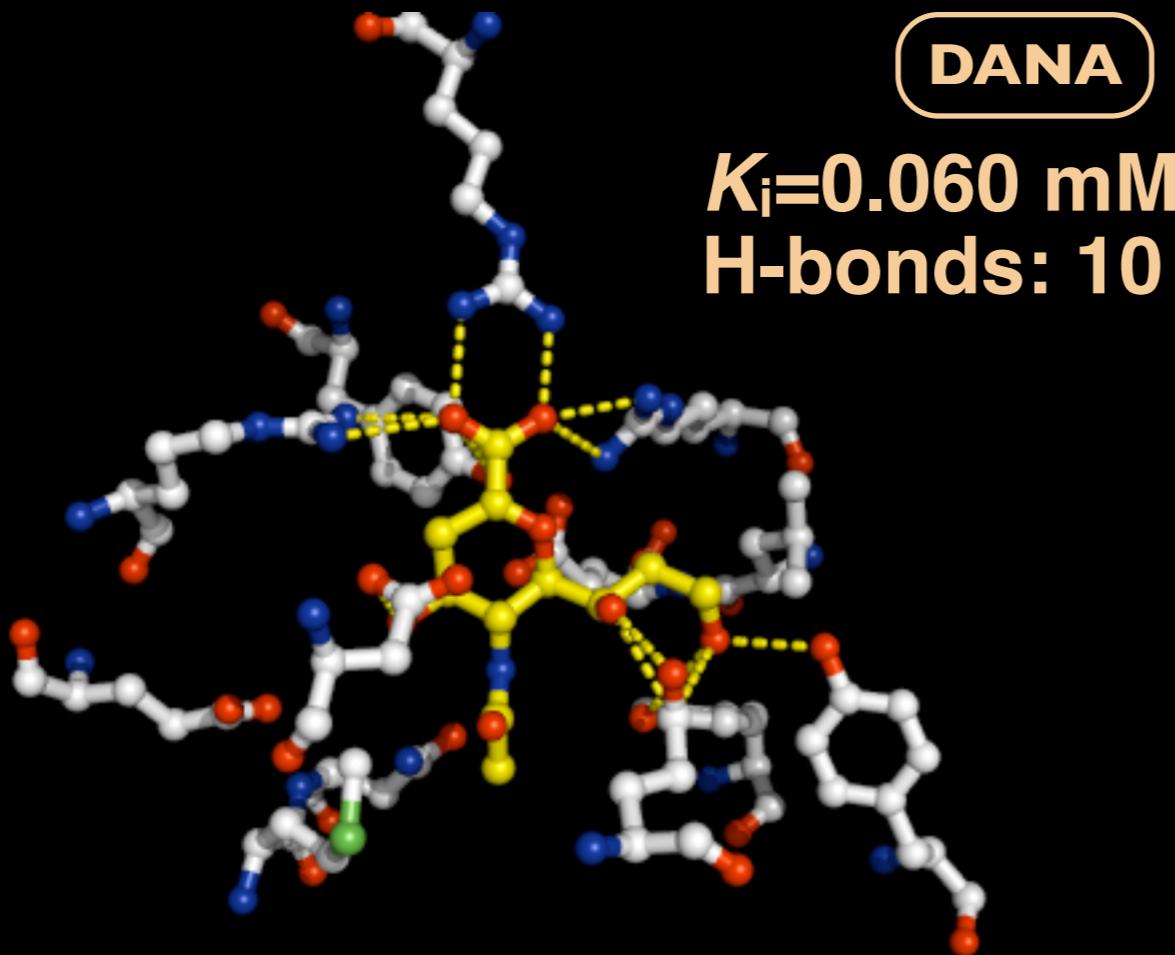


DANA: $K_i=0.060 \text{ mM}$

zanamivir (Relenza):
 $K_i=0.011 \text{ mM}$

peramivir: $K_i=0.563 \text{ mM}$





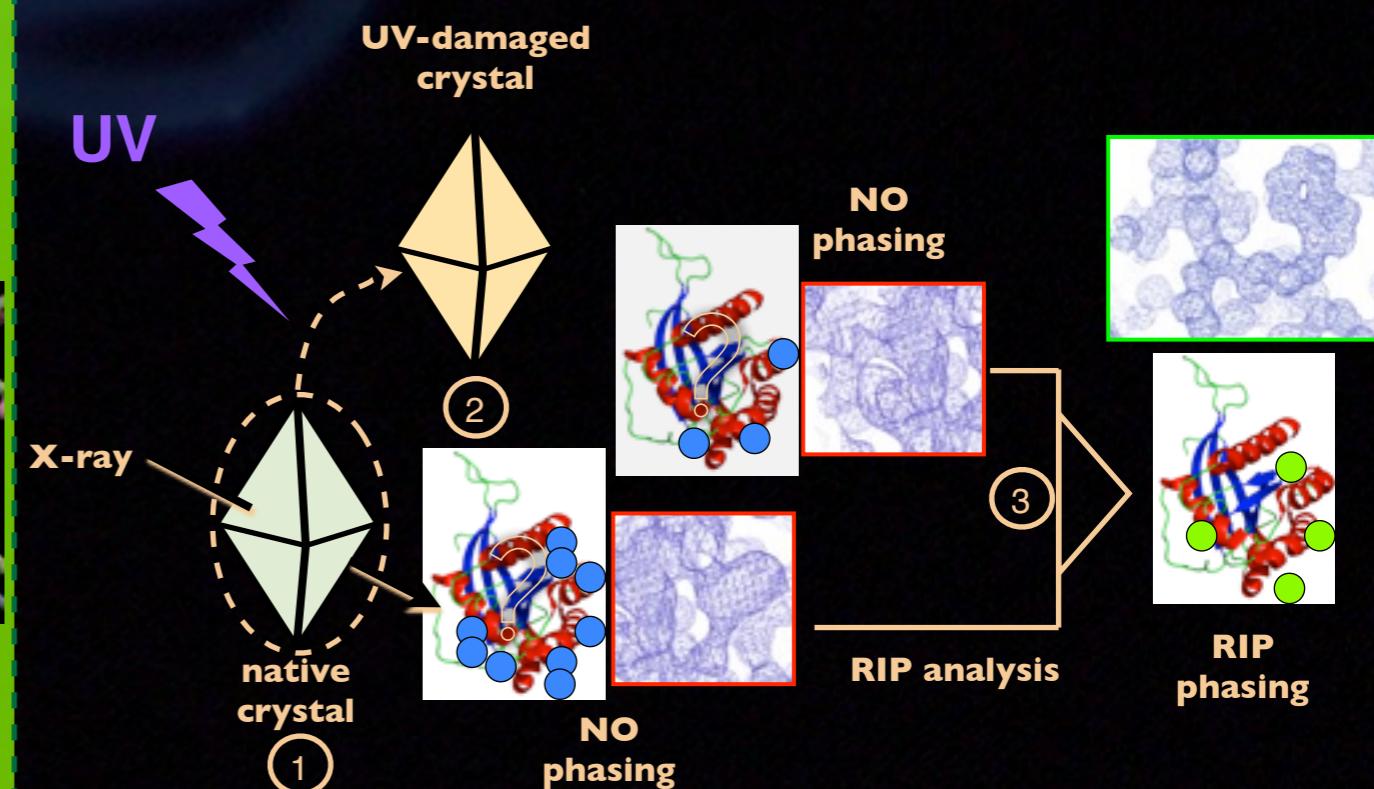
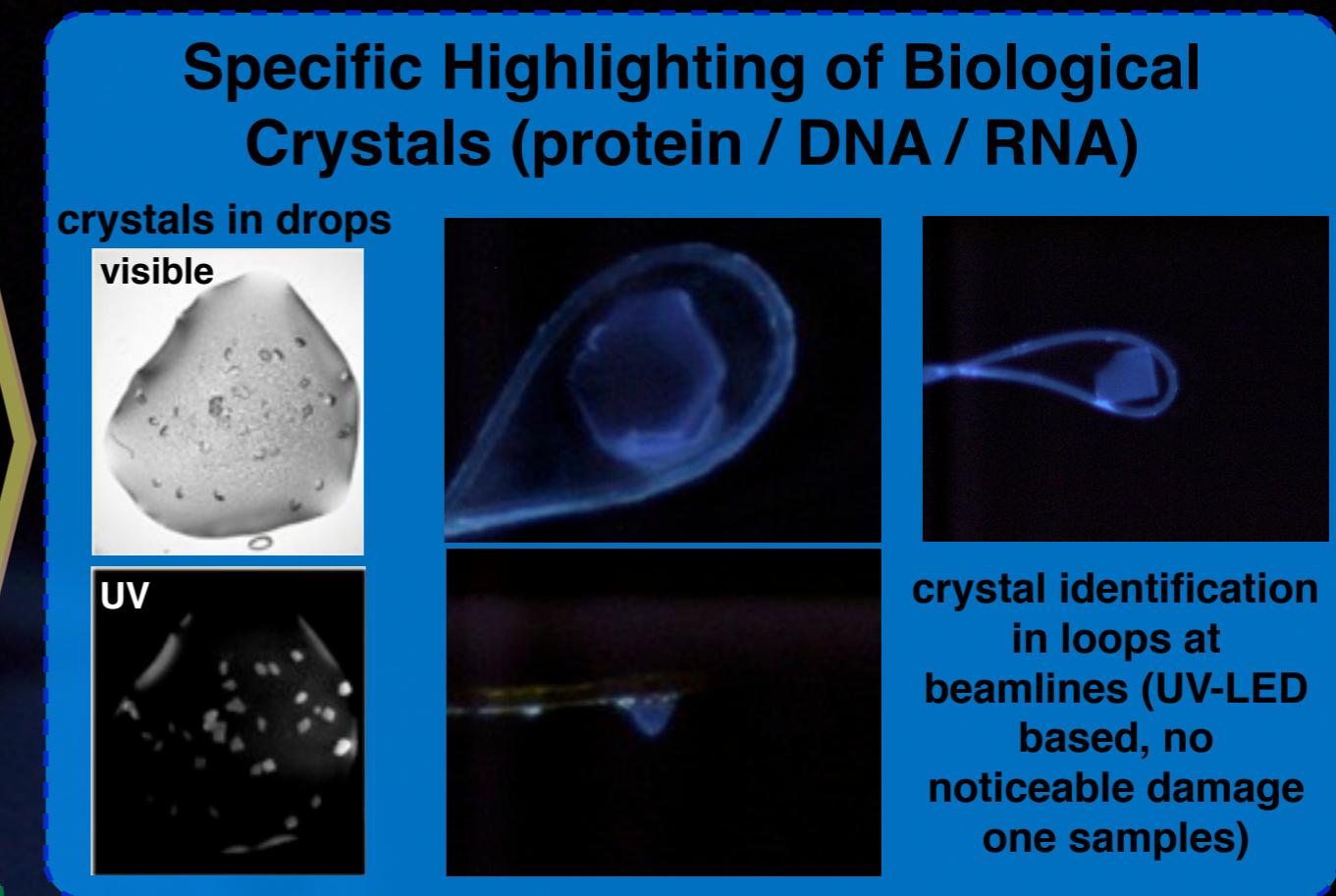
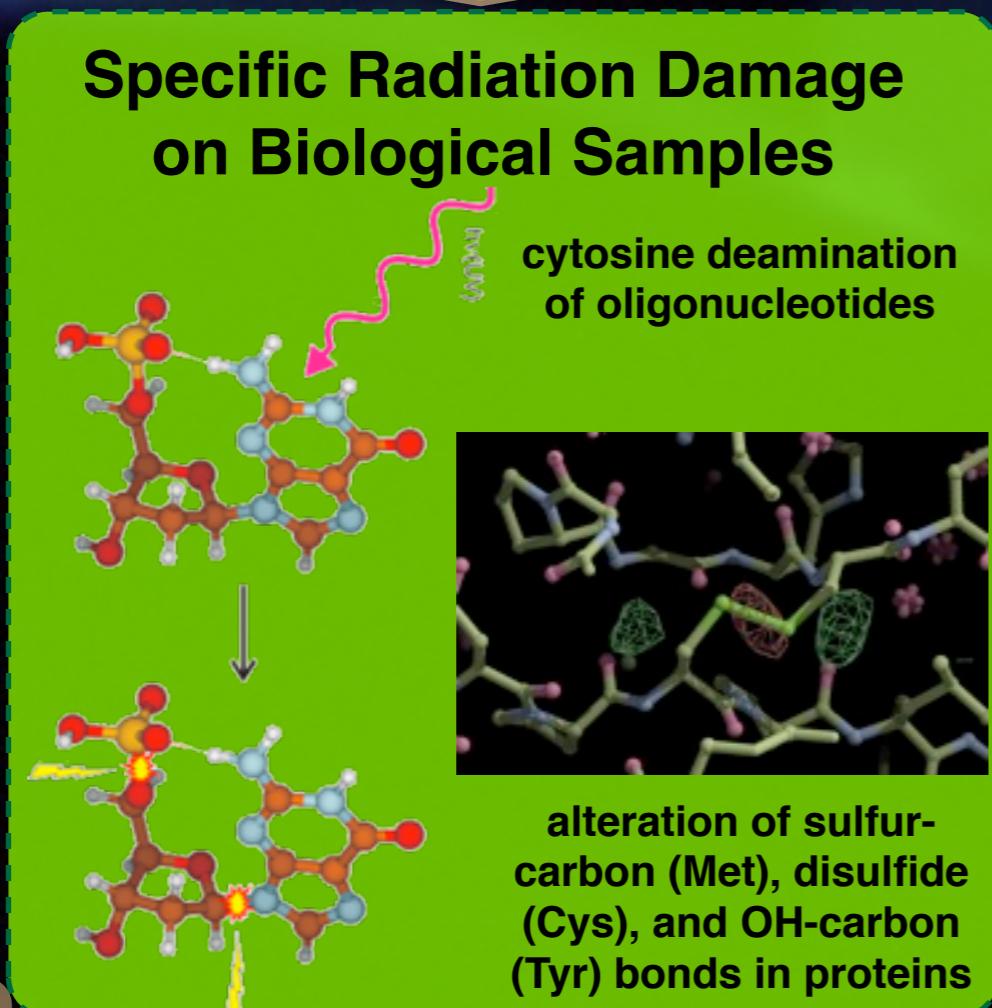
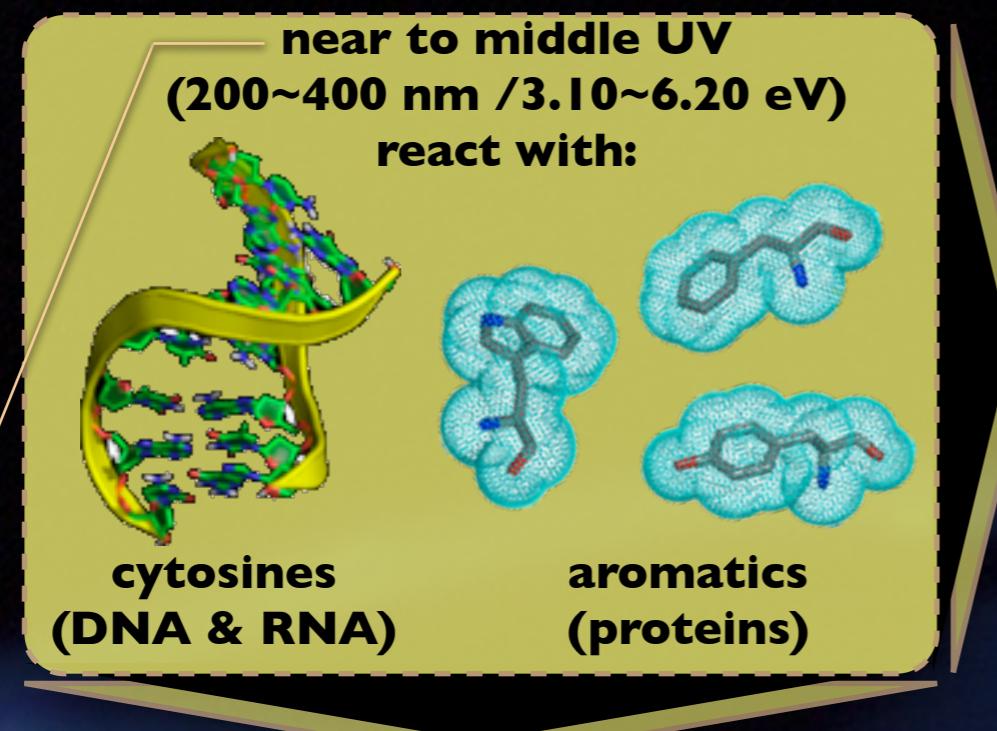
RESEARCH AND SUPPORT ACTIVITIES

The collage includes:

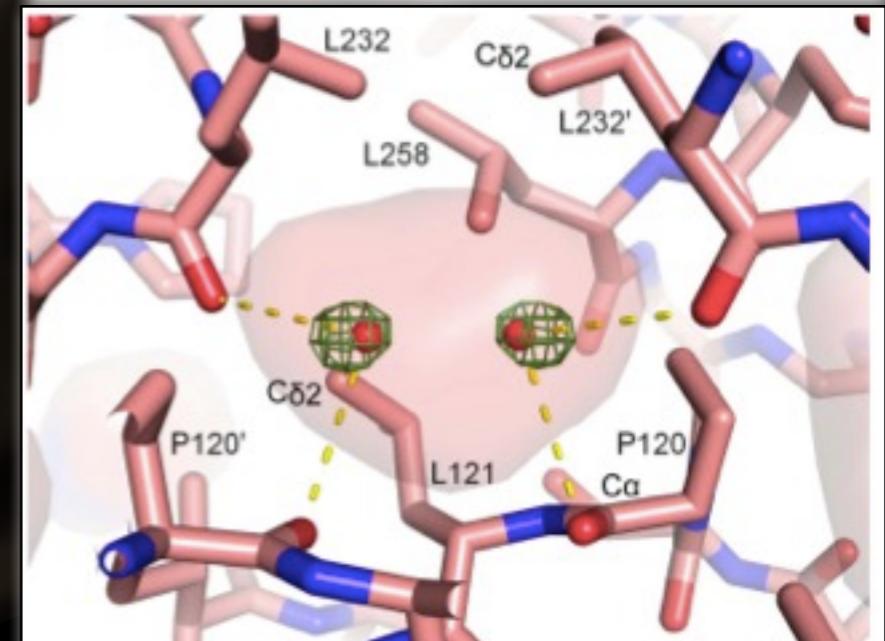
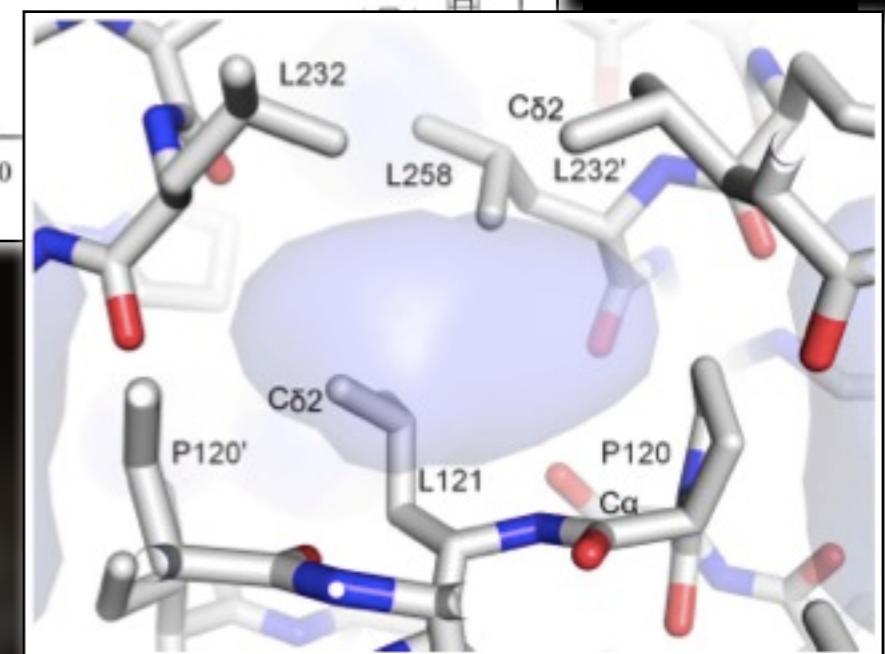
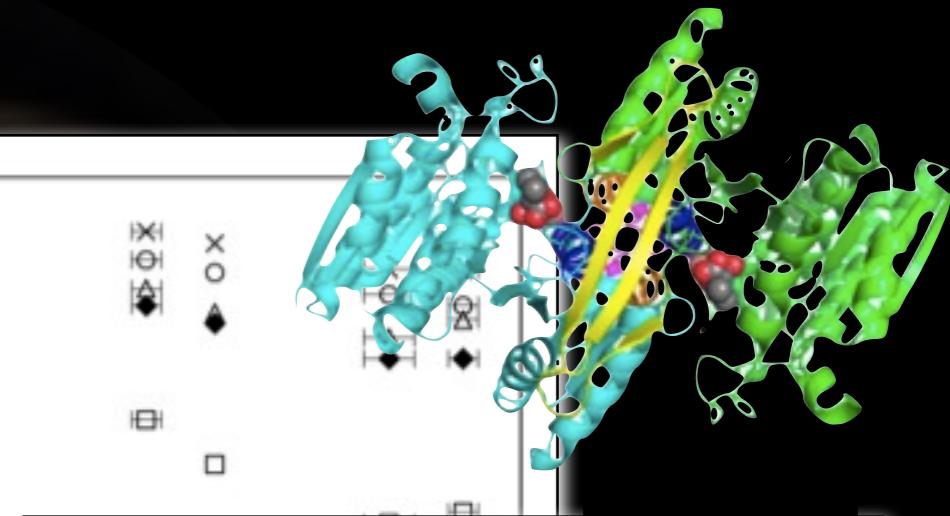
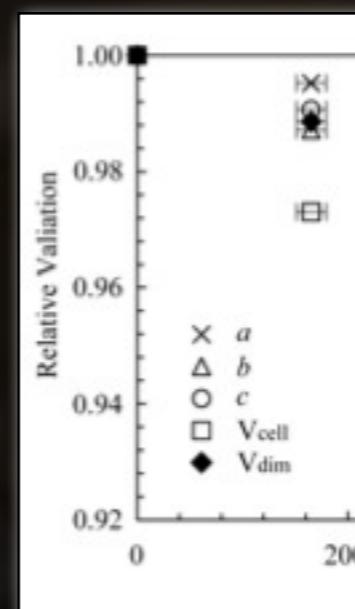
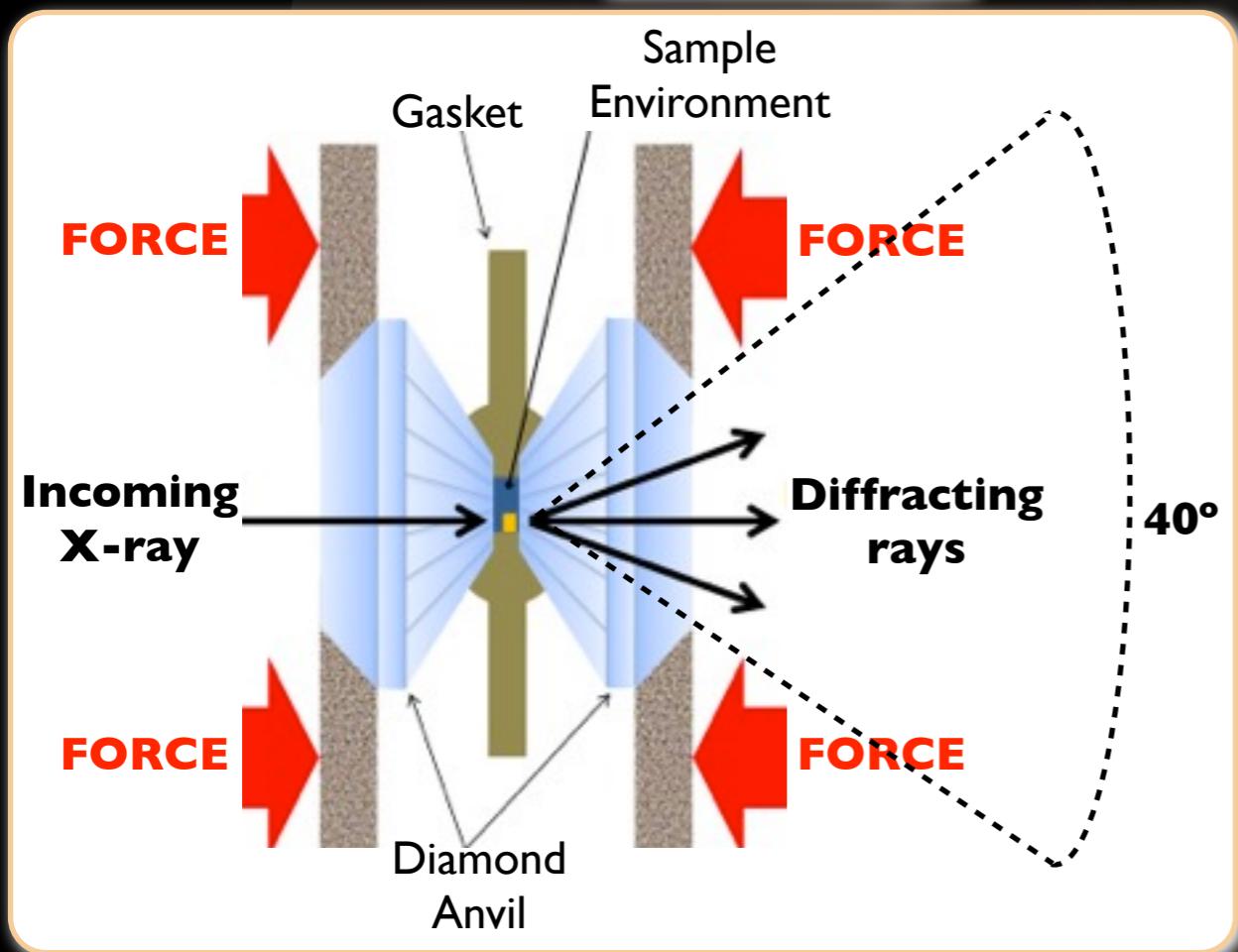
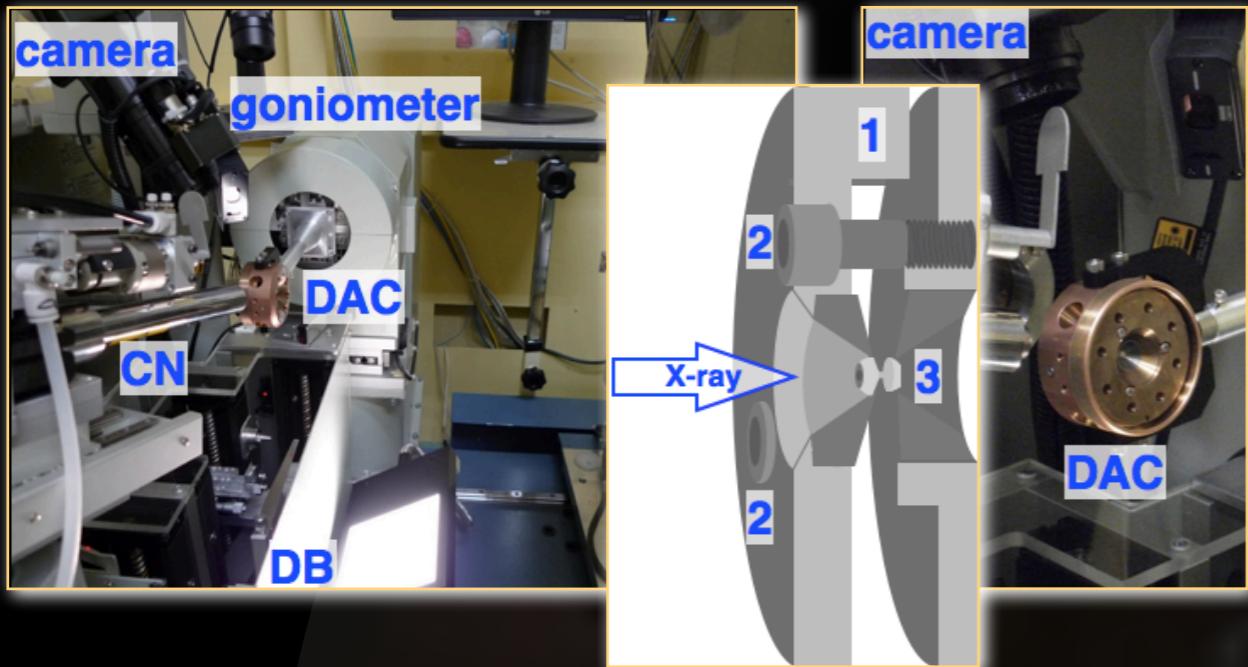
- A grid of 24 molecular structures (ribbons) representing various proteins or complexes.
- A detailed diagram of a protein structure with side chains labeled.
- A graph showing a red curve and a green circle with concentric rings.
- A circular diagram with labels P, S, C, I and a periodic table.
- A photograph of a white building labeled "STRUCTURAL BIOLOGY RESEARCH CENTER" with the text "established in 2002" below it.
- A 3D surface plot of experimental data.
- A photograph of a laboratory area with a sign reading "AR-NE3A astellas".
- A photograph of laboratory equipment, including two circular devices (possibly X-ray sources or detectors) and a complex machine.
- A graphic of a synchrotron beamline with a central beam and surrounding dashed lines.
- A large diagonal banner at the bottom left reads "STRUCTURAL & FUNCTIONAL BIOLOGY" in black text on a light blue background.
- A large diagonal banner at the bottom right reads "SYNCHROTRON SCIENCE" in black text on a light blue background.

• What are we doing?

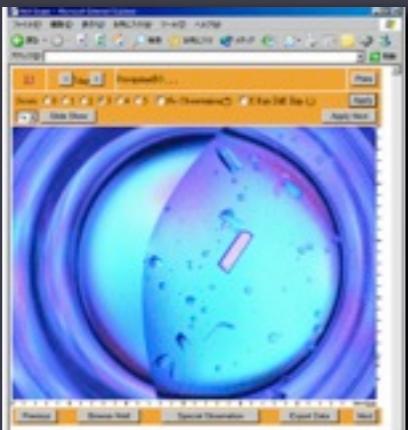
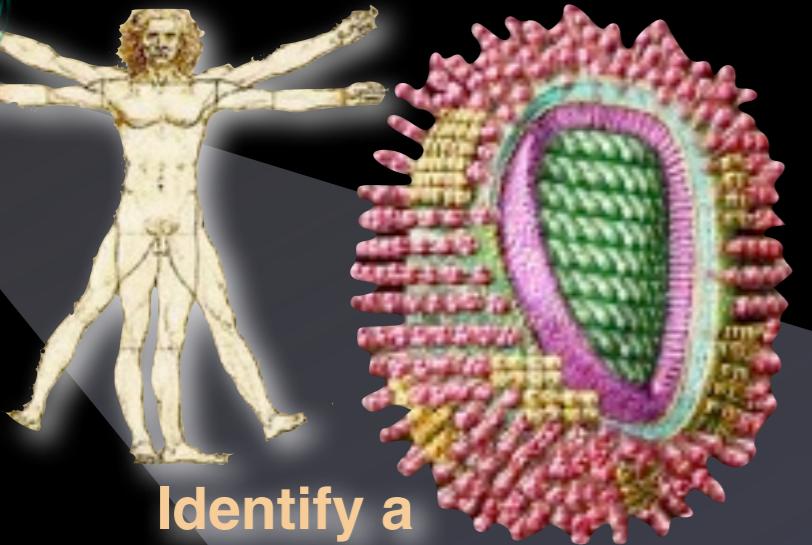
UV FOR CENTERING & PHASING



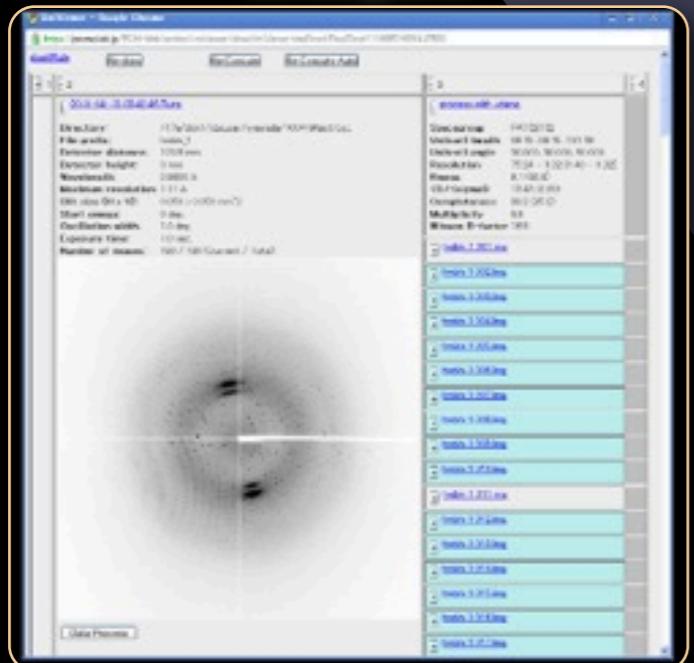
HIGH-PRESSURE SET-UP



CONCLUSIONS



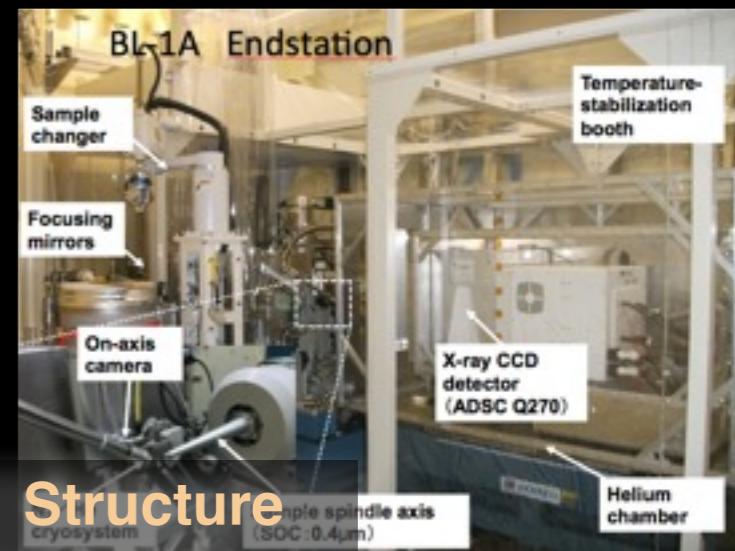
Detailed analysis



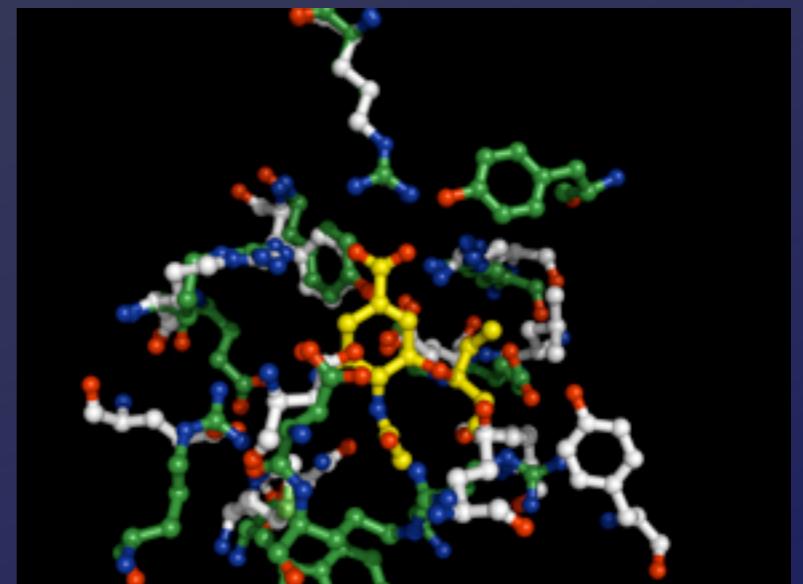
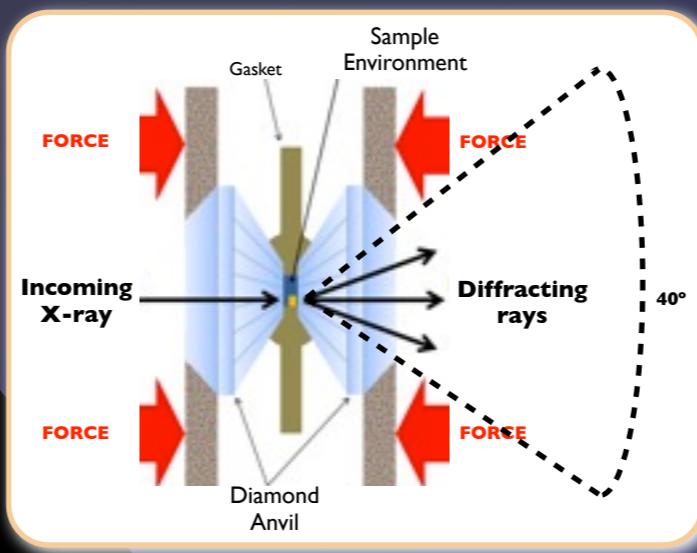
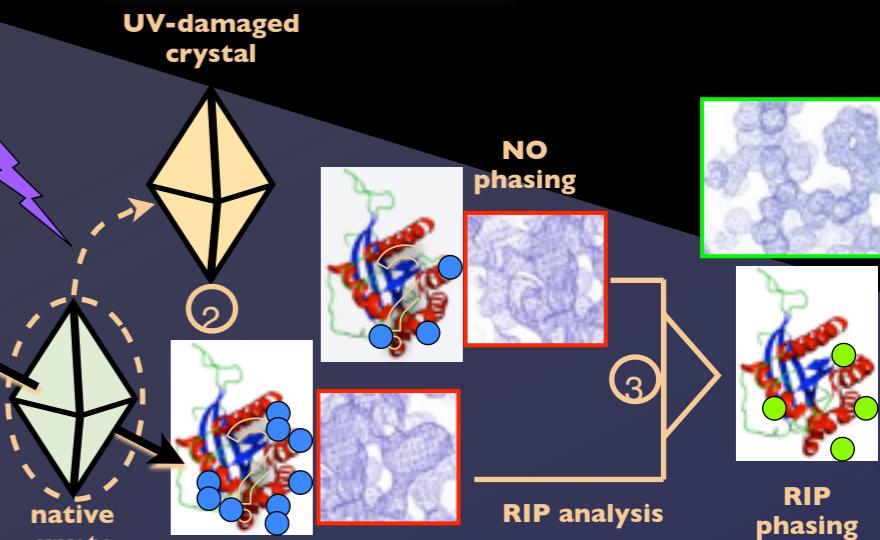
Data analysis



X-ray analysis



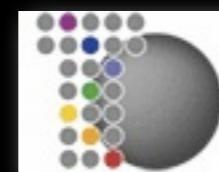
Structure determination



ACKNOWLEDGMENTS



KEK - PF



TPRP

Targeted Proteins Research Program

MEXT



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**... and more than
40 different collaborations worldwide
(SACLA XFEL, SPring-8, AS...)**



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