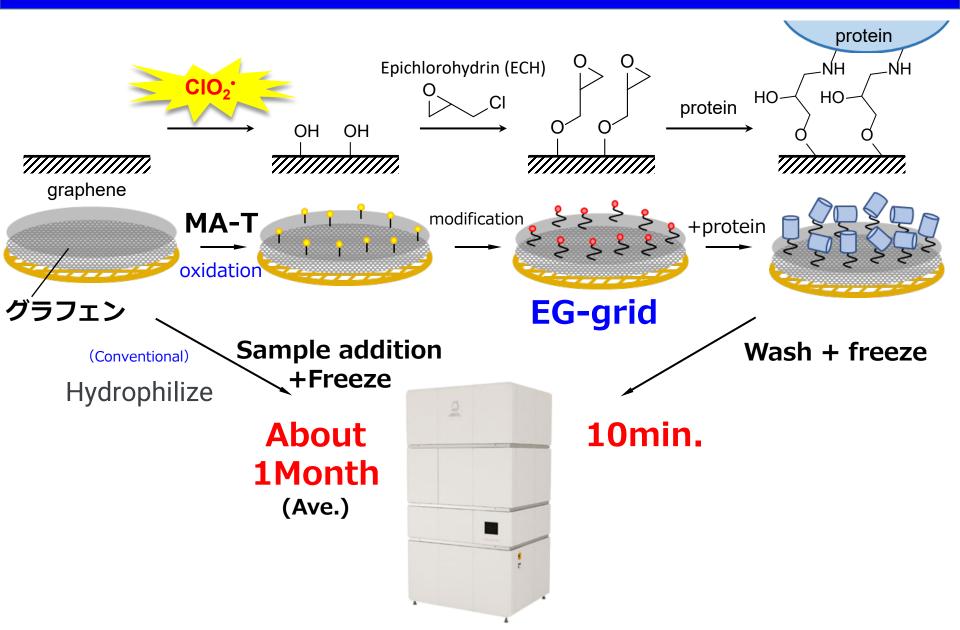
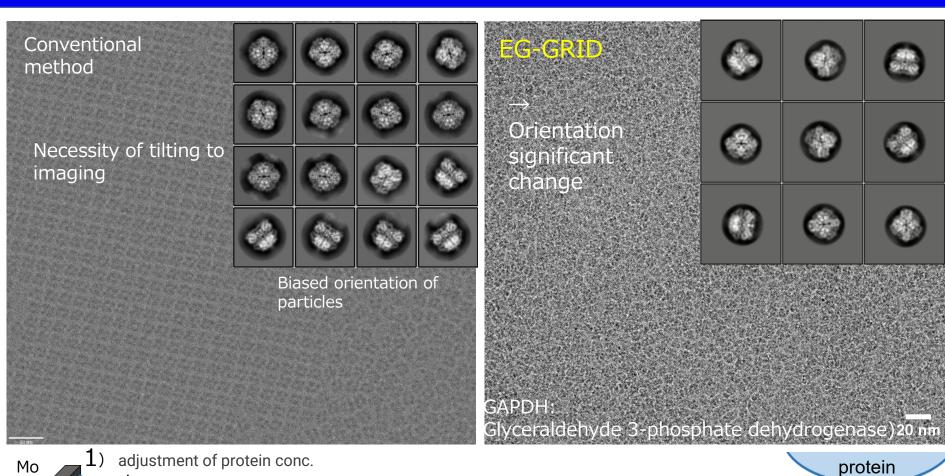
Introduction of EG-GRID for Cryo-EM

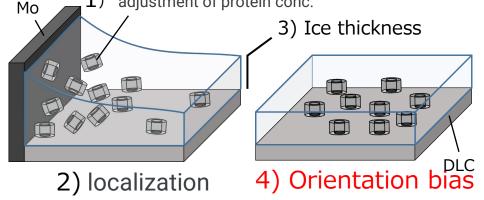
School of Pharmaceutical Sciences Osaka University Prof. Tsuyoshi INOUE

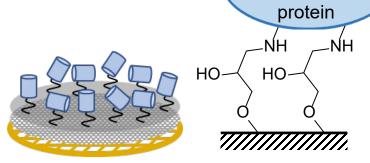
Dramatically accelerated pre-analysis preparation: 1 month to 10 minutes



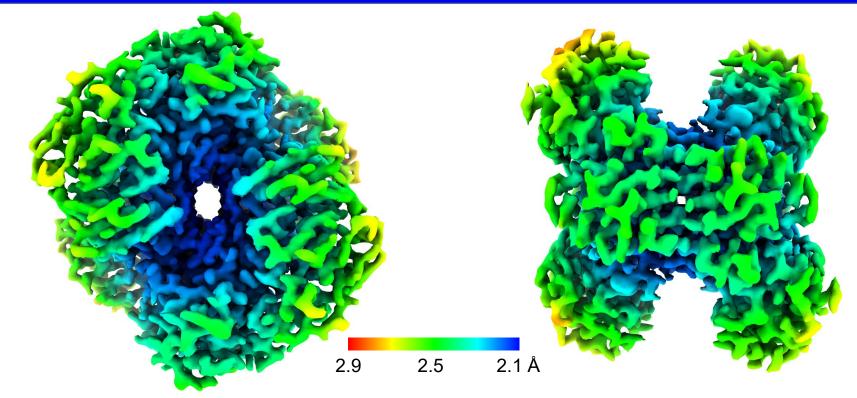
Fujita, J. et al., Sci. Rep., 13, 2279 (2023)







Example: Structural analysis of Protein-B (Cryo-EM)



	conventional	EG-GRID
number of images	4,415	241
Effective number of particles	813,828	88,731
Effective number of particles per image	184.3	368.2
resolution(Å)	2.3	2.15

Reach higher resolution with less than 1/20th of the number of images

significant efficiency (2 hours shooting time)

Sci. Rep. 2023

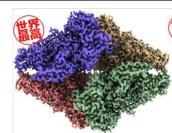
We have many achievements



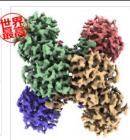
GroEL 1.99 Å



1.29 Å (最高1.22Å)



Apoferritin β-galactosidase 1.81 Å



GAPDH 2.16 Å



V1-ATPase 3.03 Å



Aldolase 2.28 Å

mBio 投稿中

Sci Rep.

2022

SARS-CoV-2 spike proteins

VHH complex: 3種

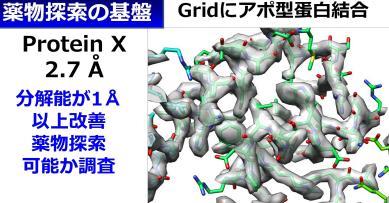
Fab complex: 3種

2種

mutant: 2種

Protein X 2.7 Å 分解能が1点

以上改善 薬物探索 可能か調査



Accelerating antibody drug development

Analysis completed within 3 days! **BA.4/5**

 $IC_{50} = 0.29 \, \mu g/ml$

BQ1.1

 $IC_{50} = 4.20 \, \mu g/ml$

XBB

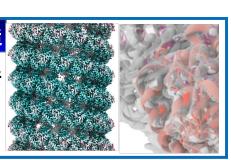
 $IC_{50} = 0.45 \, \mu g/ml$

新規抗菌標的構造

フィラメントで解析

2.7 Å

(1回のグリド調製)



Nat. Comm., in press

Looking for joint research with companies. Try EG-GRID with your protein!

- Perfect confidentiality. (highly confidential compounds? No ploblem.)
- · No cryo-EM? You can use Osaka University's cryo-EM.
- Only 5ul (0.1~1 mg/ml) protein is required

Many companies have already started joint research. First, check the performance through trial joint research.