

Structure Determination

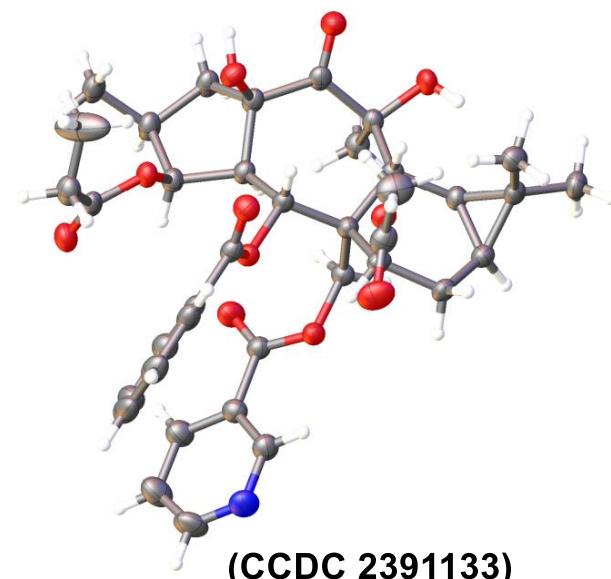
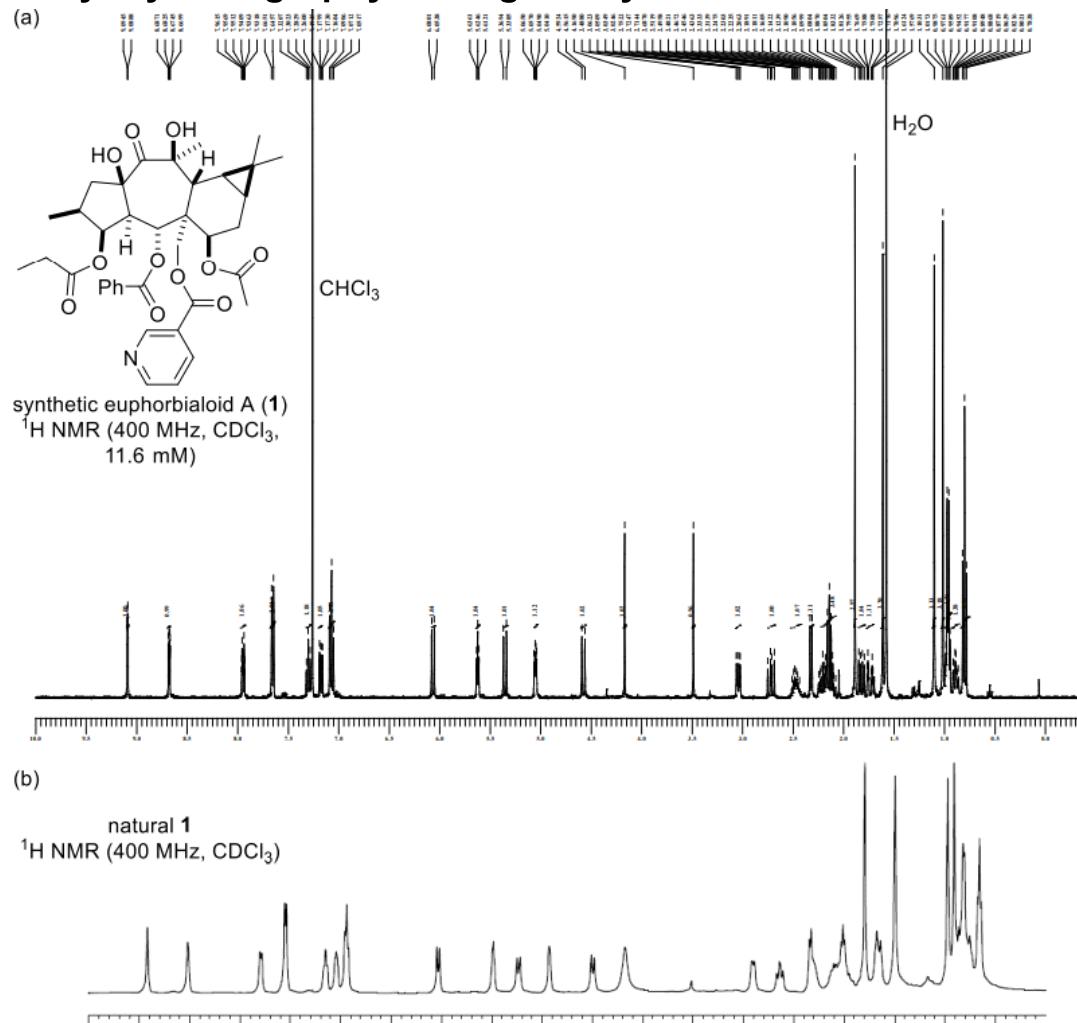
Literature Seminar
2024. 12. 27

Kyohei Takaoka

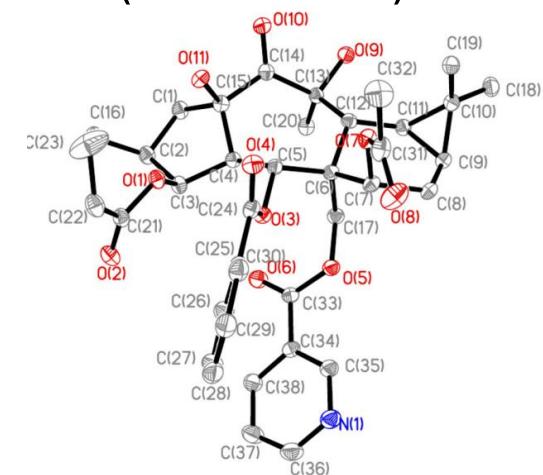
- 1. Introduction**
- 2. Strategy for Crystal Analysis**

X-Ray Crystallography as a Powerful Tool for Structure Determination

X-ray crystallography unambiguously determines the structure.

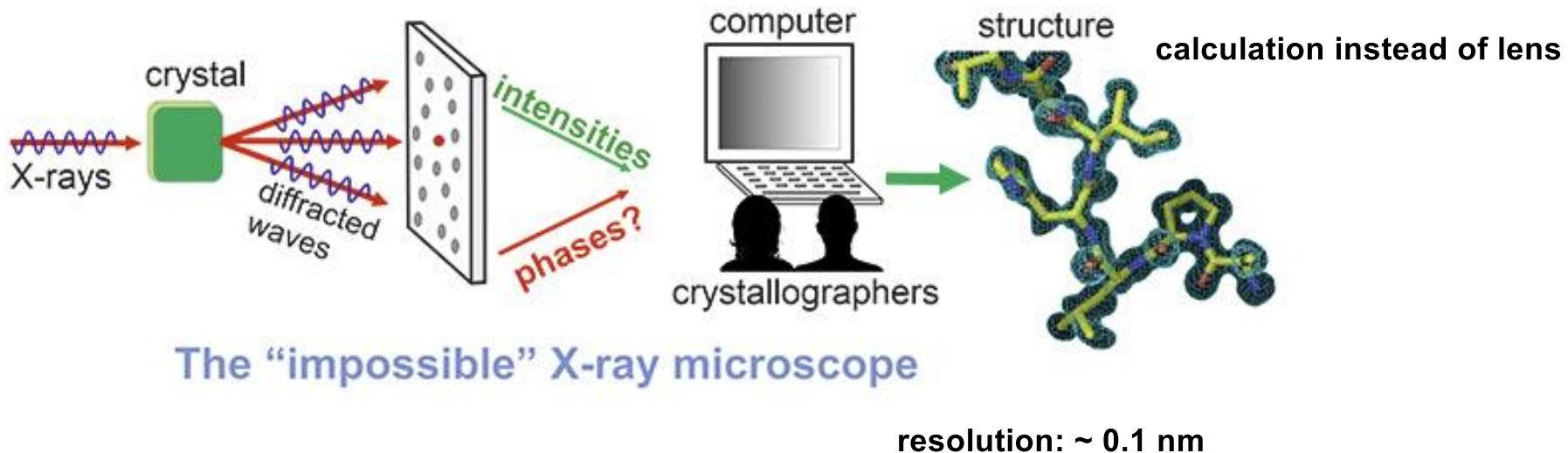
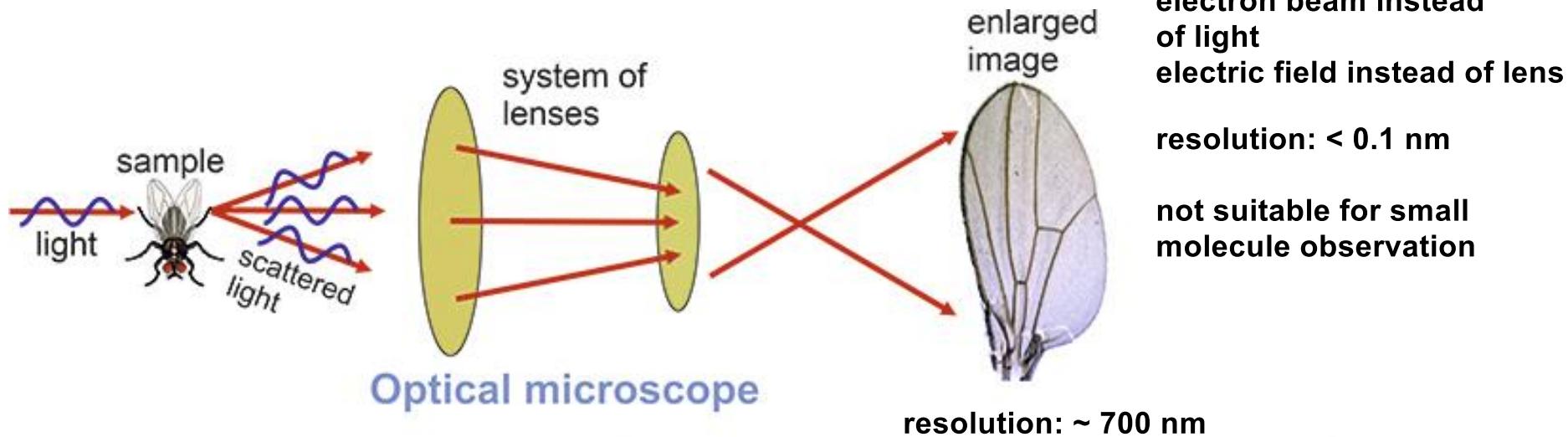


(CCDC 2391133)



originally determined by
X-ray crystallography

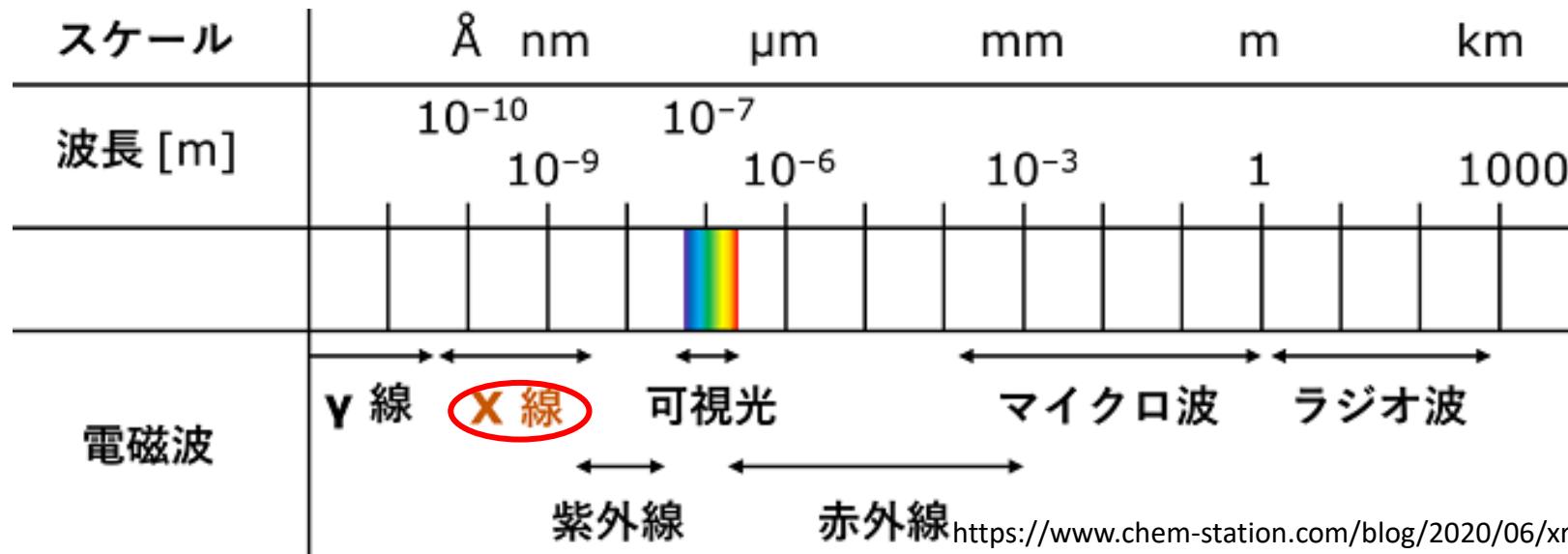
Observing Substances in Atom Scale



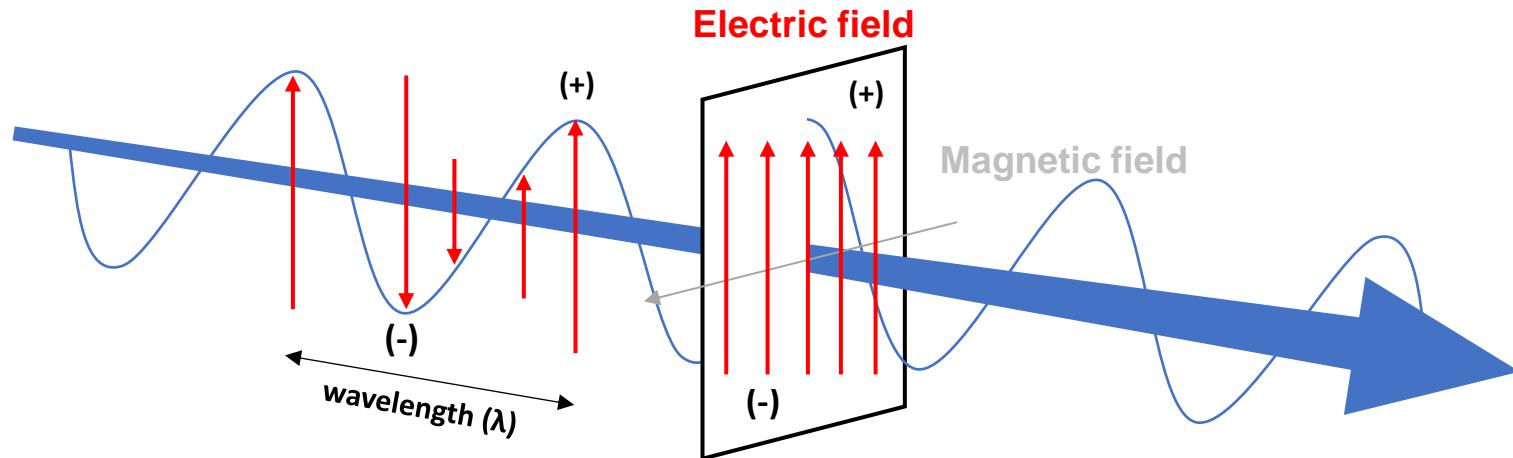
What is X-Ray?

X-ray: A form of high-energy electromagnetic radiation

Wavelength (λ) \sim 10 pm to 10 nm

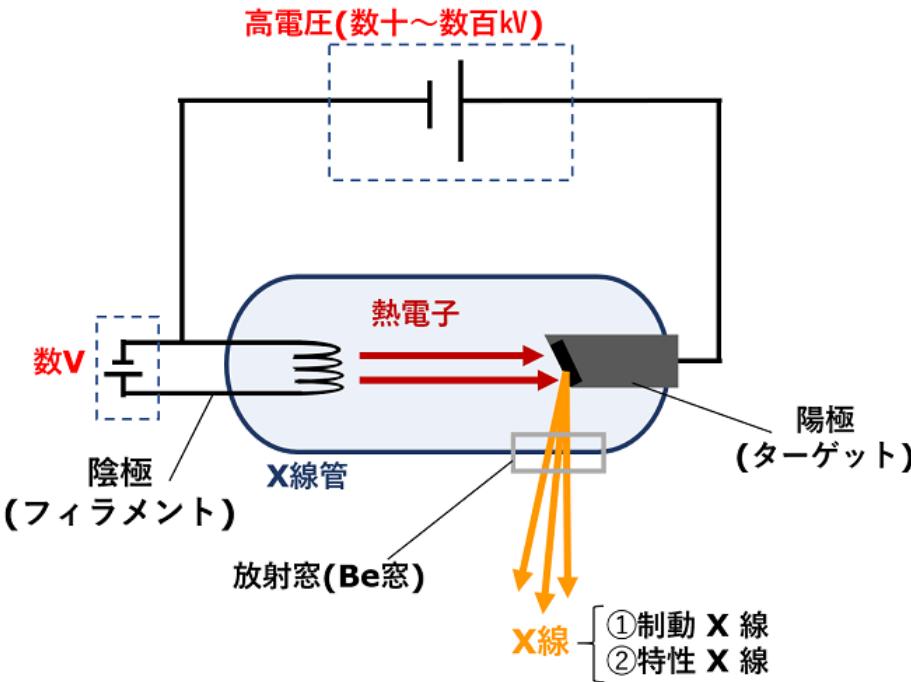


Electromagnetic waves can be regarded as the progression of the wavefront of an electric magnetic field



X-Ray Generation

Example of X-ray tube:

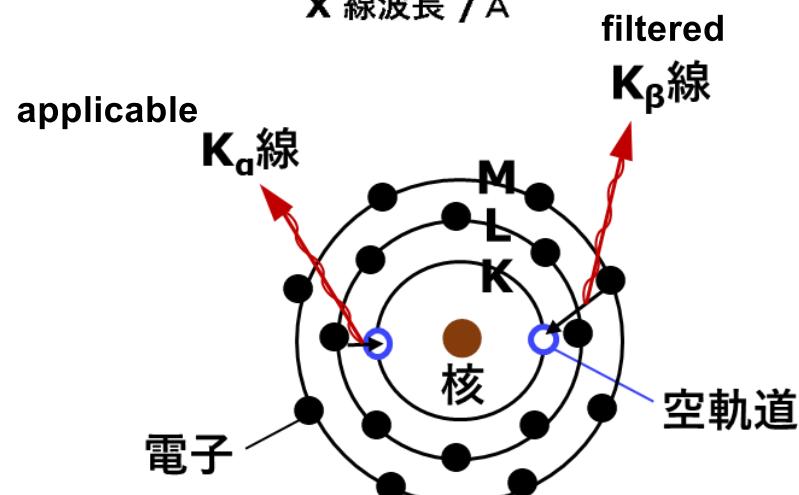
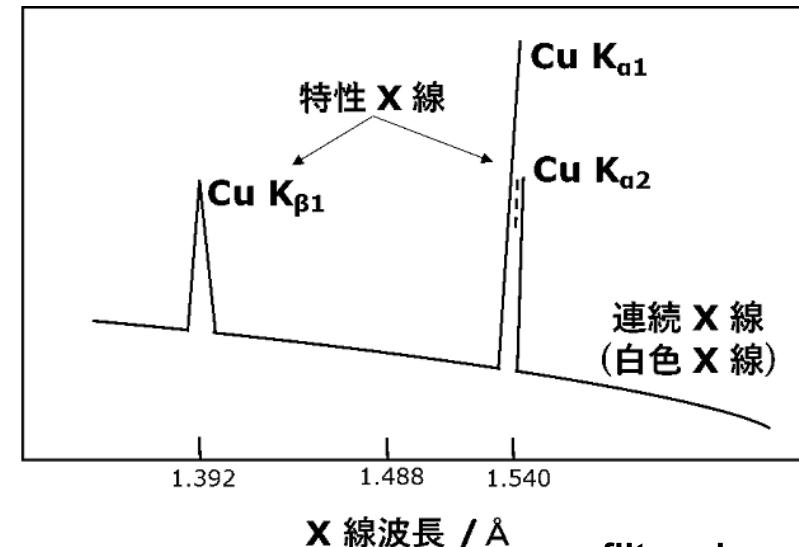


2 types of X-ray are emitted:

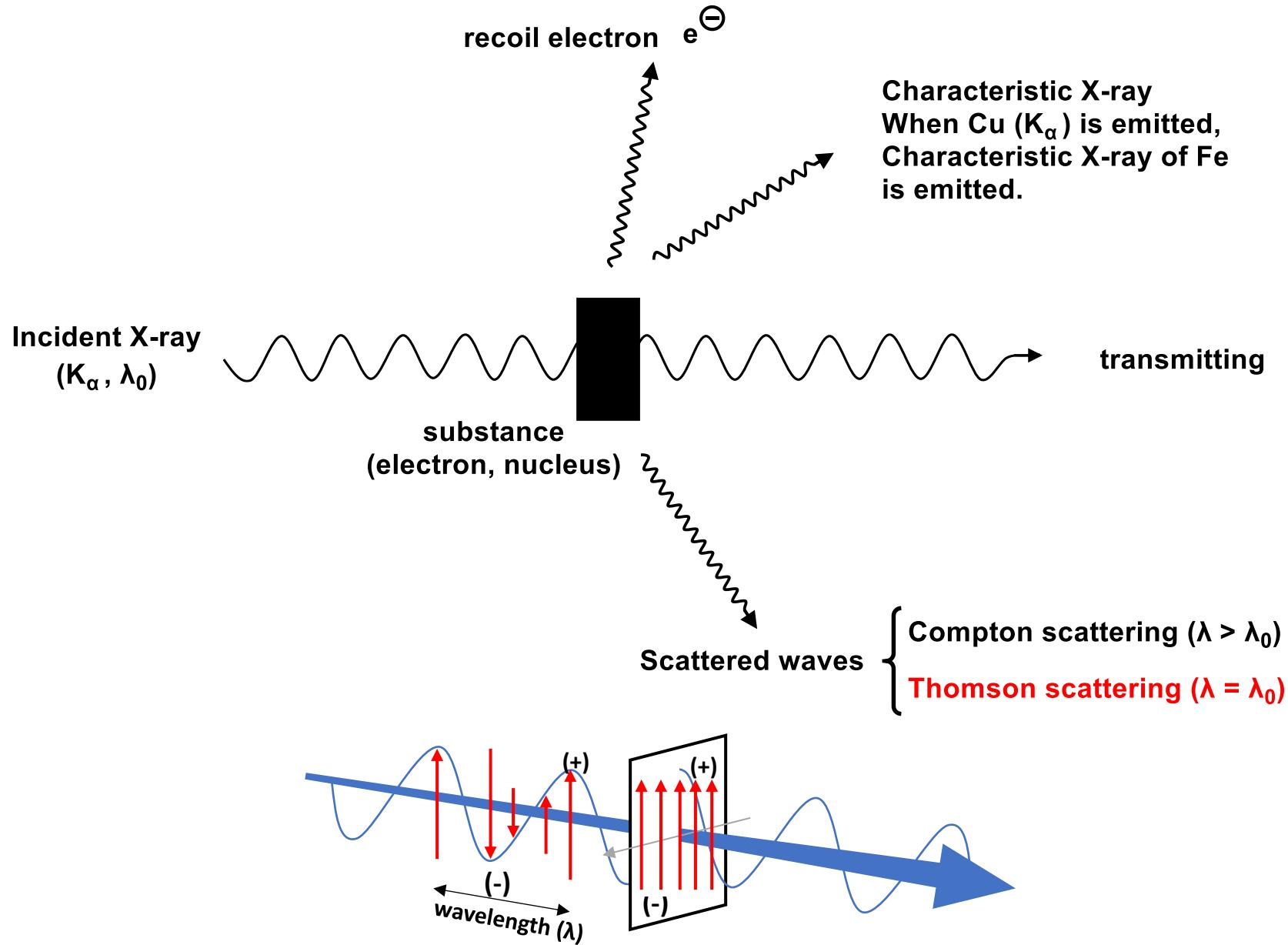
① Continuous X-ray (制動X線、連続X線)
generated when electrons are decelerated
as they pass near the nuclei of target atoms
not used in X-ray crystallography = filtered

② Characteristic X-ray (used in X-ray crystallography, 特性X線)
generated when high-energy electrons collide with
inner-shell electrons of target atoms, causing them to be ejected.
used in X-ray crystallography (K_{α}) e.g. Mo: 0.711 Å, Cu: 1.542 Å

X-ray spectrum generated from Cu tube

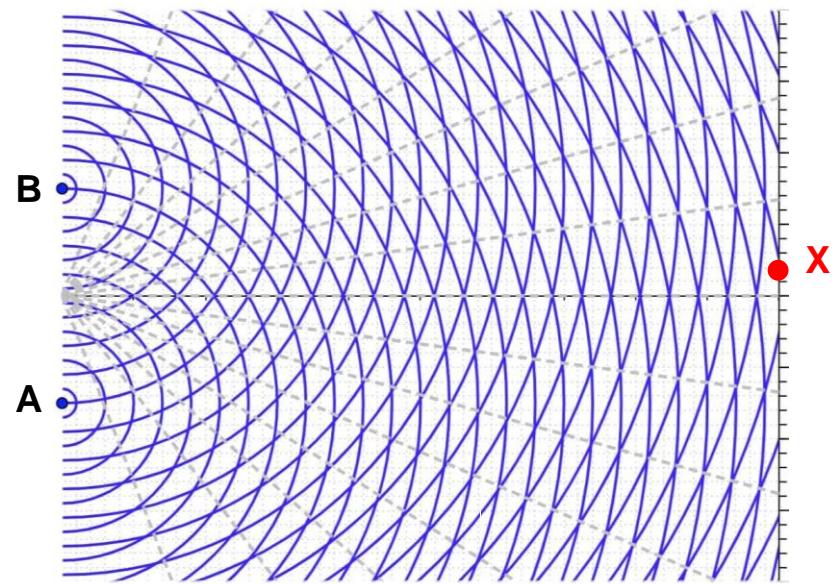


Interaction between X-ray and Substance



Wave Interaction

Wave interaction will be observed when waves have same frequency



Composite waves E can be described as

Supposing waves from A and B are same and phase difference between A and B at point X is $2\pi\delta$, Wave E_A and E_B can be described as

$$E_A = A_0 e^{2\pi i(vt - \frac{x}{\lambda})}$$

$$E_B = A_0 e^{2\pi i(vt - \frac{x}{\lambda} + \delta)}$$

A_0 : amplitude of wave A and B

x: distance between point X and A

v: wave frequency

λ : wavelength

t: time

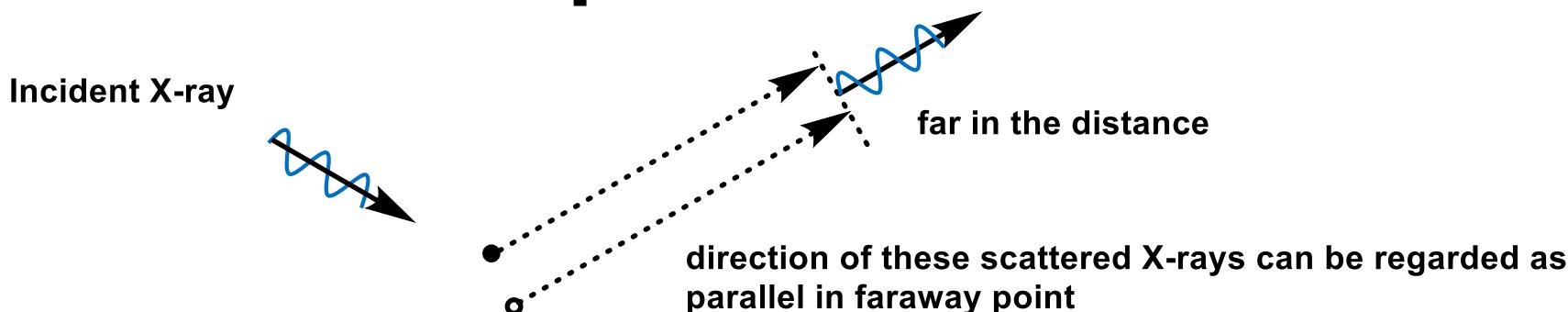
$$E = E_A + E_B = A_0 e^{2\pi i\left(vt - \frac{x}{\lambda}\right)} \underline{\left(1 + e^{2\pi\delta i}\right)}$$

determined by underlined formula

Wavelength of Thomson scattered X-ray from each electron is the same.
Therefore, these X-ray must be interacted.

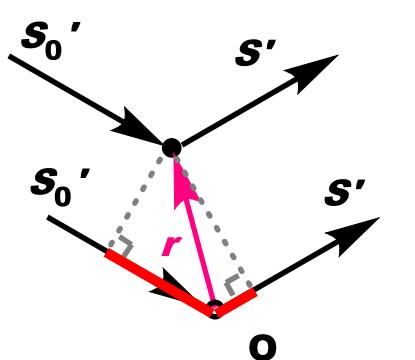
When considering composite wave, we only need to consider phase difference of each wave.

Phase Difference can be Expressed as Simple Formula

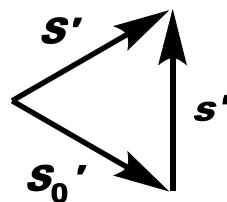


How can we express path difference and phase difference?

- Path difference can be described as **below red lines**.



\mathbf{S}_0' : vector of incident X-ray, $|\mathbf{S}_0'| = 1$
 \mathbf{S}' : vector of scattered X-ray, $|\mathbf{S}'| = 1$



$$|\mathbf{S}'| = |\mathbf{S}'_0| = 1$$

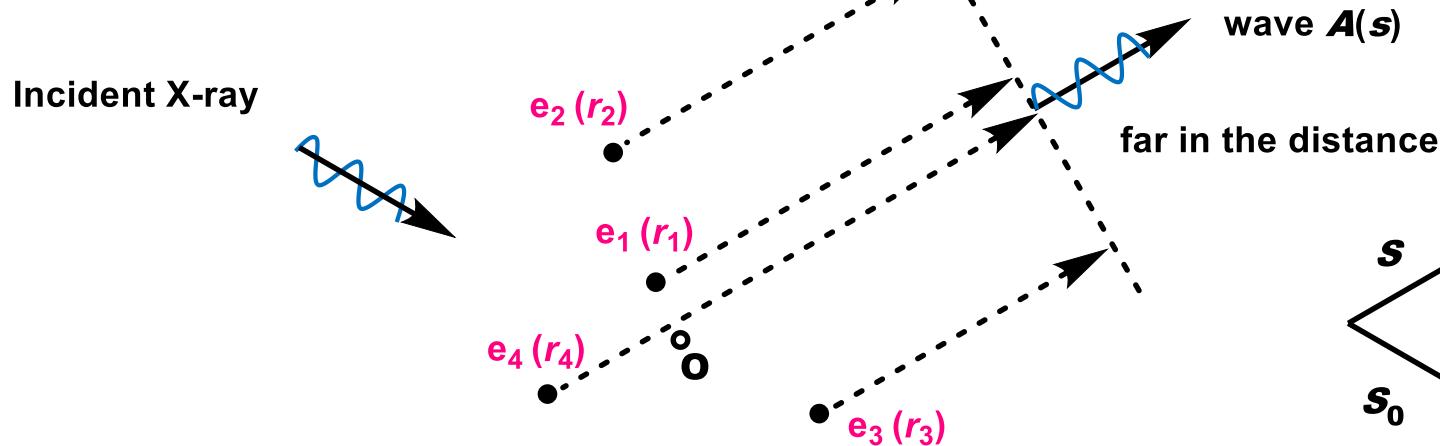
$$\text{path difference} = |r| \times |\mathbf{S}'| \times \cos(\text{angle between } \mathbf{r} \text{ and } \mathbf{S}') - |r| \times |\mathbf{S}'_0| \times \cos(\text{angle between } \mathbf{r} \text{ and } \mathbf{S}'_0) = \mathbf{r} \cdot \mathbf{S}' - \mathbf{r} \cdot \mathbf{S}'_0 = \mathbf{r} \cdot (\mathbf{S}' - \mathbf{S}'_0)$$

When \mathbf{S}_0 and \mathbf{S} are defined as $\mathbf{S}_0 = \mathbf{S}_0'/\lambda$, $\mathbf{S} = \mathbf{S}/\lambda$, phase difference parameter $\delta(r)$ is described as:

$$\delta(r) = \frac{\text{pathdifference}}{\lambda} = \frac{\mathbf{r} \cdot (\mathbf{S}' - \mathbf{S}'_0)}{\lambda} = \mathbf{r} \cdot \frac{(\mathbf{S}' - \mathbf{S}'_0)}{\lambda} = \mathbf{r} \cdot (\mathbf{S} - \mathbf{S}_0) = \mathbf{r} \cdot \mathbf{s} \quad (s = \mathbf{S} - \mathbf{S}_0)$$

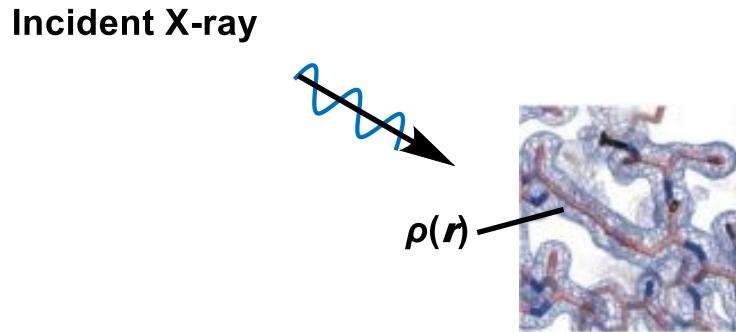
Relationship between Wave and Electron Density

Wave interaction among various electrons:



$$A(s) = A_e e^{2\pi i(r_1 \cdot s)} + A_e e^{2\pi i(r_2 \cdot s)} + \dots = \sum_{n=1,2,\dots} A_e e^{2\pi i(r_n \cdot s)}$$

Wave interaction in single molecule

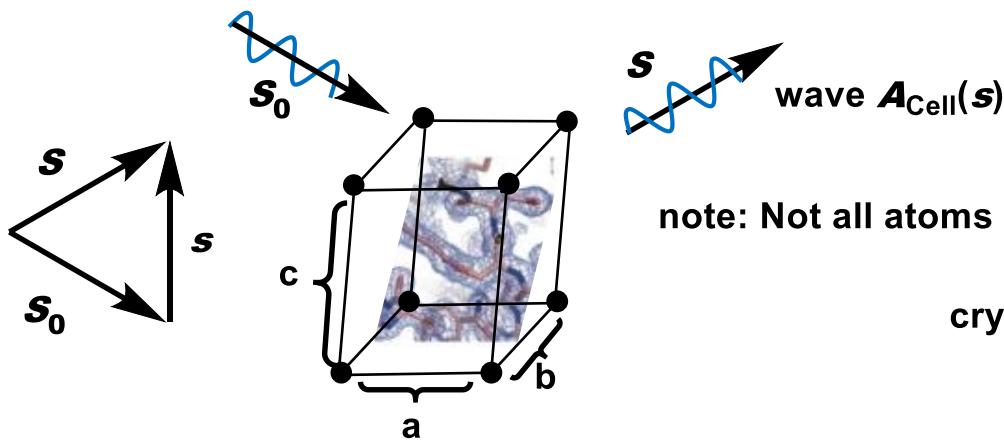


$$A(s) = \int \rho(r) e^{2\pi i(r \cdot s)} dv_r$$

* A_e is omitted
from Fourier integral theorem...

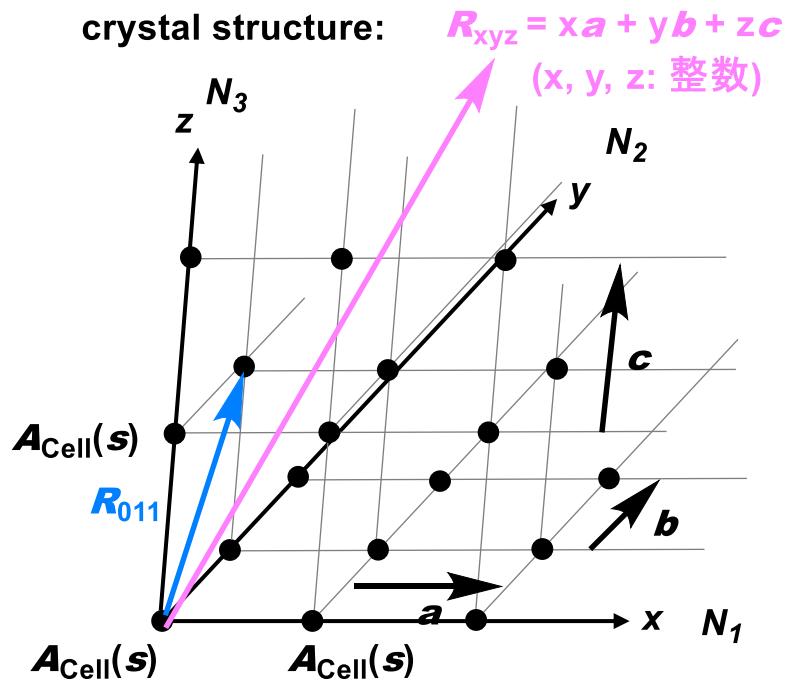
$$\rho(r) = \int A(s) e^{-2\pi i(r \cdot s)} dv_s$$

Mechanism of X-Ray Diffraction (1)



Composite waves of crystal $\mathbf{A}_{\text{crystal}}(\mathbf{s})$ is regarded as the sum of each unit cell's composite waves $\mathbf{A}_{\text{Cell}}(\mathbf{s})$.

$$\begin{aligned}\mathbf{A}_{\text{crystal}}(\mathbf{s}) &= \mathbf{A}_{\text{Cell}}(\mathbf{s})e^{2\pi i[(\mathbf{a}+\mathbf{b}+\mathbf{c}) \cdot \mathbf{s}]} + \mathbf{A}_{\text{Cell}}(\mathbf{s})e^{2\pi i[(2\mathbf{a}+\mathbf{b}+\mathbf{c}) \cdot \mathbf{s}]} + \dots \\ &= \sum_{x=-\frac{N_1-1}{2}}^{\frac{N_1-1}{2}} \sum_{y=-\frac{N_2-1}{2}}^{\frac{N_2-1}{2}} \sum_{z=-\frac{N_3-1}{2}}^{\frac{N_3-1}{2}} \mathbf{A}_{\text{Cell}}(\mathbf{s})e^{2\pi i[(x\mathbf{a}+y\mathbf{b}+z\mathbf{c}) \cdot \mathbf{s}]} \\ &= \mathbf{A}_{\text{Cell}}(\mathbf{s}) \left\{ \frac{\sin N_1 \pi (\mathbf{a} \cdot \mathbf{s})}{\sin \pi (\mathbf{a} \cdot \mathbf{s})} \cdot \frac{\sin N_2 \pi (\mathbf{b} \cdot \mathbf{s})}{\sin \pi (\mathbf{b} \cdot \mathbf{s})} \cdot \frac{\sin N_3 \pi (\mathbf{c} \cdot \mathbf{s})}{\sin \pi (\mathbf{c} \cdot \mathbf{s})} \right\}\end{aligned}$$



Mechanism of X-Ray Diffraction (2)

X-ray intensity is proportional to the square of the amplitude.

Composite wave $\mathbf{A}(\mathbf{s})$ is complex number, so X-ray intensity $I_{\text{crystal}}(\mathbf{s})$ is expressed as

$$I_{\text{crystal}}(\mathbf{s}) \propto A_{\text{crystal}}(\mathbf{s}) \cdot A_{\text{crystal}}^*(\mathbf{s}) = |A_{\text{Cell}}(\mathbf{s})|^2 \left\{ \frac{\sin^2 N_1 \pi (\mathbf{a} \cdot \mathbf{s})}{\sin^2 \pi (\mathbf{a} \cdot \mathbf{s})} \cdot \frac{\sin^2 N_2 \pi (\mathbf{b} \cdot \mathbf{s})}{\sin^2 \pi (\mathbf{b} \cdot \mathbf{s})} \cdot \frac{\sin^2 N_3 \pi (\mathbf{c} \cdot \mathbf{s})}{\sin^2 \pi (\mathbf{c} \cdot \mathbf{s})} \right\}$$

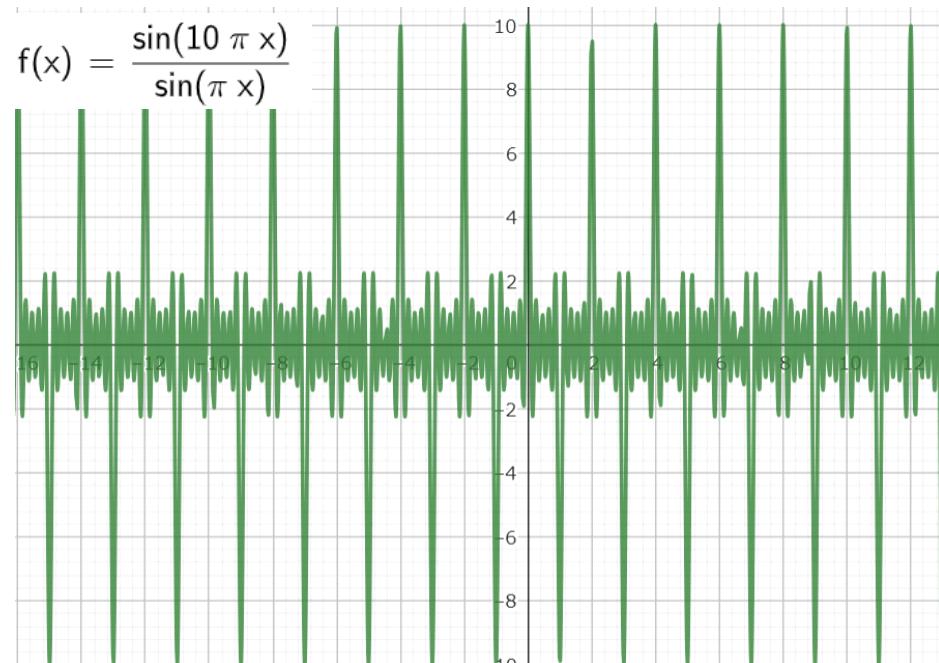
According to the graph of right,
 $I_{\text{crystal}}(\mathbf{s})$ becomes strong when
the following rules are satisfied.

$$(\mathbf{a} \cdot \mathbf{s}) = h, \quad (\mathbf{b} \cdot \mathbf{s}) = k, \quad (\mathbf{c} \cdot \mathbf{s}) = l$$

h, k, l : integer (整数) *Laue equation

Otherwise, $I_{\text{crystal}}(\mathbf{s})$ would be neglected

→ Scattered X-rays are observed only at certain points.
X-ray diffraction!



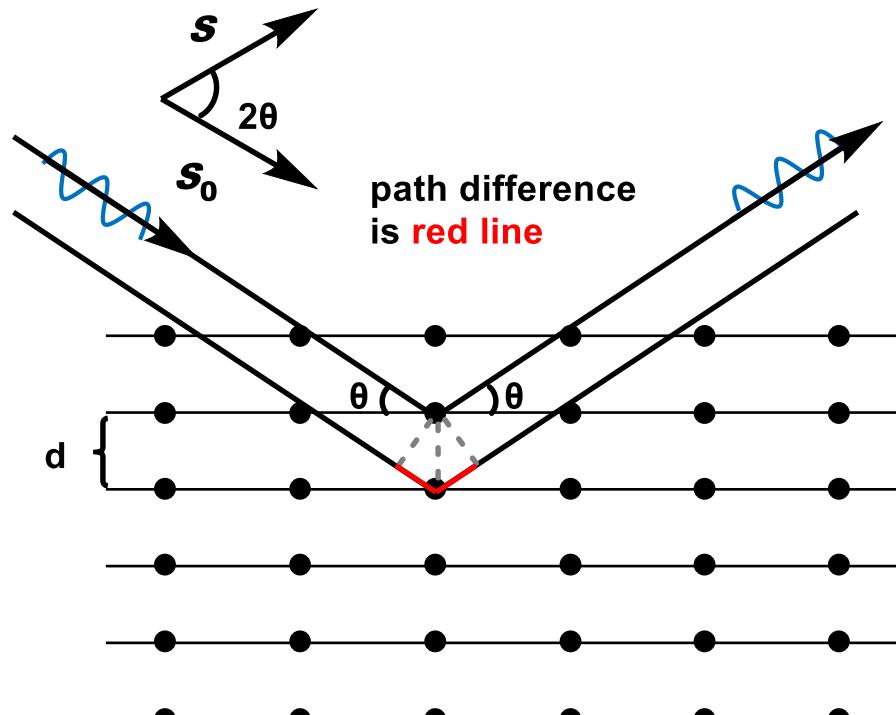
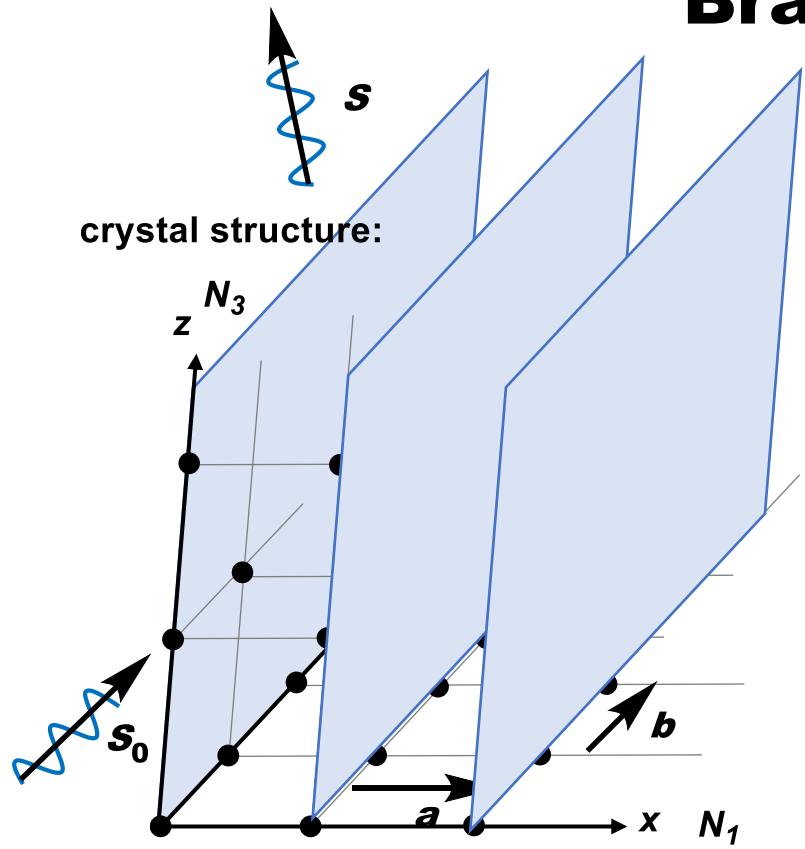
$$\mathbf{r} = x\mathbf{a} + y\mathbf{b} + z\mathbf{c}$$

$$\mathbf{r} \cdot \mathbf{s} = (x\mathbf{a} + y\mathbf{b} + z\mathbf{c}) \cdot \mathbf{s} = x(\mathbf{a} \cdot \mathbf{s}) + y(\mathbf{b} \cdot \mathbf{s}) + z(\mathbf{c} \cdot \mathbf{s}) = hx + ky + lz$$

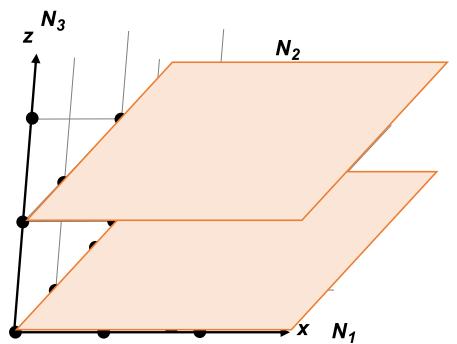
$$A_{\text{Cell}}(\mathbf{s}) = \int \rho(\mathbf{r}) e^{2\pi i(\mathbf{r} \cdot \mathbf{s})} d\mathbf{v}_r = \iiint \rho(x, y, z) e^{2\pi i(hx + ky + lz)} V dx dy dz = \underline{F(h, k, l)}$$

structure factor

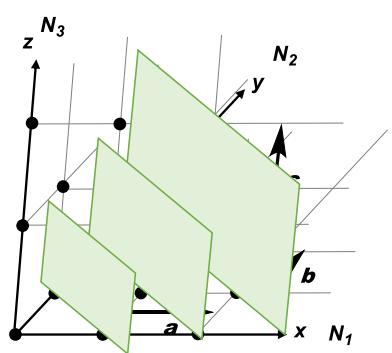
Bragg's law



crystal structure:



crystal structure:

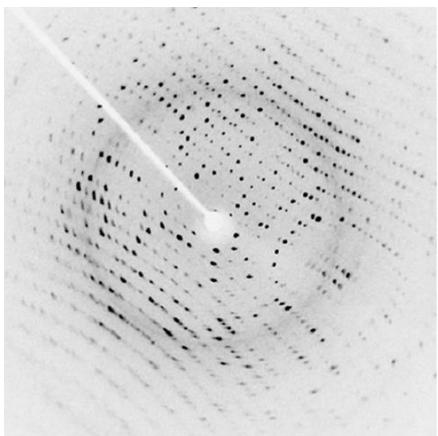


$$2ds\sin\theta = n\lambda \quad (n = 1, 2, 3, \dots) \quad \text{Bragg's law}$$

Such face can be drawn in various way.
→ Many spots are observed in X-ray diffraction

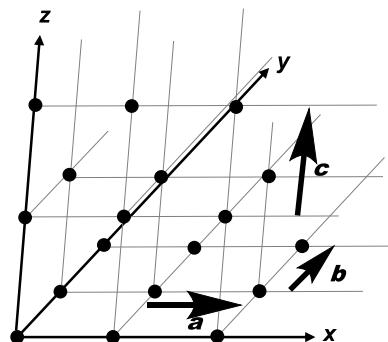
Introduction of Reciprocal Lattice

diffracted images:



1. define unit cell

crystal lattice:



We know the direction of incident and scattered X-rays, therefore we can determine vector H from each diffraction spot.
→ Reciprocal lattice can be decided.
= original (crystal lattice) can be decided.

definition of reciprocal lattice vector (\mathbf{a}^* , \mathbf{b}^* , \mathbf{c}^*):

$$\mathbf{a}^* \cdot \mathbf{a} = \mathbf{b}^* \cdot \mathbf{b} = \mathbf{c}^* \cdot \mathbf{c} = 1$$

$$\mathbf{a}^* \cdot \mathbf{b} = \mathbf{a}^* \cdot \mathbf{c} = \mathbf{b}^* \cdot \mathbf{a} = \mathbf{b}^* \cdot \mathbf{c} = \mathbf{c}^* \cdot \mathbf{a} = \mathbf{c}^* \cdot \mathbf{b} = 0$$

note: The reciprocal of the reciprocal lattice is the original lattice.

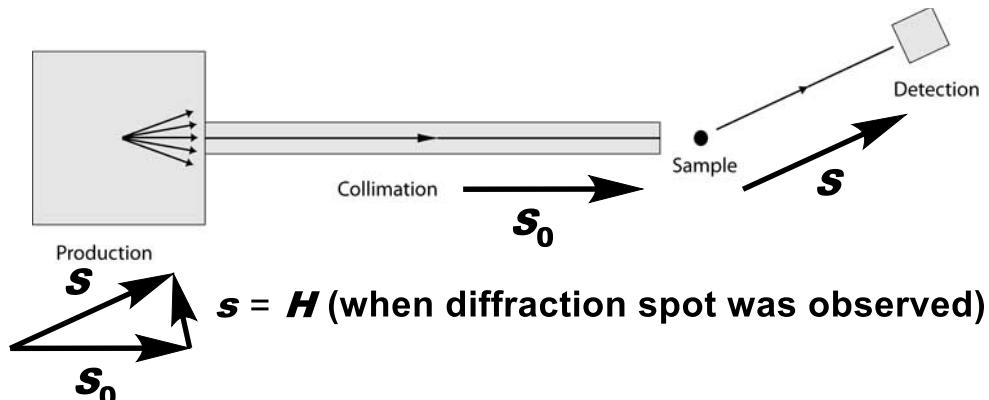
Defining \mathbf{H} as the vector toward the reciprocal grid point, the following relationships hold

$$\mathbf{H} = h\mathbf{a}^* + k\mathbf{b}^* + l\mathbf{c}^* \quad (h, k, l: \text{整数})$$

$$\mathbf{a} \cdot \mathbf{H} = \mathbf{a} \cdot (h\mathbf{a}^* + k\mathbf{b}^* + l\mathbf{c}^*) = h\mathbf{a} \cdot \mathbf{a}^* + k\mathbf{a} \cdot \mathbf{b}^* + l\mathbf{a} \cdot \mathbf{c}^* = h$$

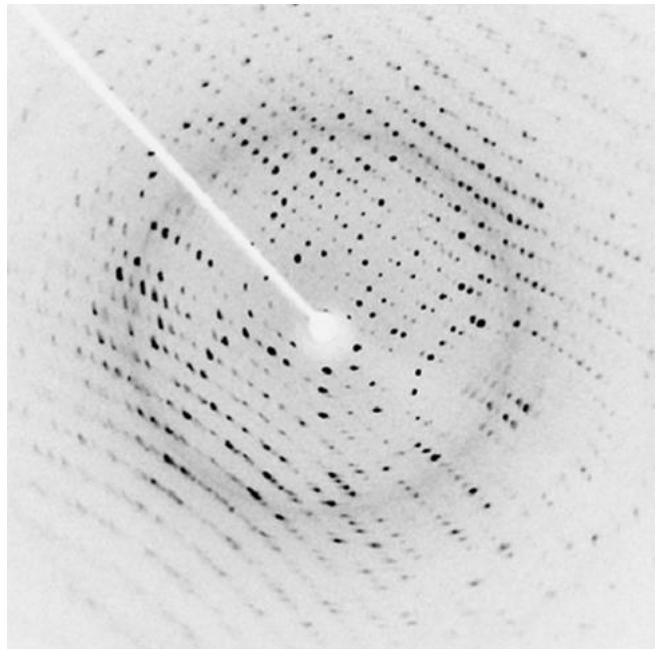
$$\mathbf{b} \cdot \mathbf{H} = k, \mathbf{c} \cdot \mathbf{H} = l$$

→ \mathbf{H} satisfies Laue equation: $(\mathbf{a} \cdot \mathbf{s}) = h$, $(\mathbf{b} \cdot \mathbf{s}) = k$, $(\mathbf{c} \cdot \mathbf{s}) = l$



$s = \mathbf{H}$ (when diffraction spot was observed)

Phase Problem in X-ray Analysis



X-ray diffraction images

$$F(h, k, l) = \iiint \rho(x, y, z) e^{2\pi i(hx+ky+lz)} V dx dy dz$$



Fourier integral theorem...

$$\begin{aligned}\rho(x, y, z) &= \frac{1}{V} \sum_{-\infty}^{\infty} \sum_{-\infty}^{\infty} \sum_{-\infty}^{\infty} F(h, k, l) e^{-2\pi i(hx+ky+lz)} \\ &= \frac{1}{V} \sum_{-\infty}^{\infty} \sum_{-\infty}^{\infty} \sum_{-\infty}^{\infty} |F(h, k, l)| e^{-i\{2\pi(hx+ky+lz)-\Phi(h, k, l)\}}\end{aligned}$$

available from
observable
intensity

unavailable
from diffraction
images!

"Phase problem"

- Direct (*ab initio*) method: Developed by Hauptman and Karle
Solve phase problem by using mathematical processing
Commonly used in recent X-ray analysis

In determination of absolute configuration of the structure,
the effect of anomalous scattering (異常散乱) is utilized.
Anomalous scattering was mainly observed on heavy atoms,
so introduction of heavy atom is recommended.



Hauptman



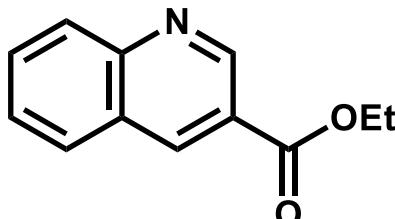
Karle

The Novel Prize in
Chemistry 1985

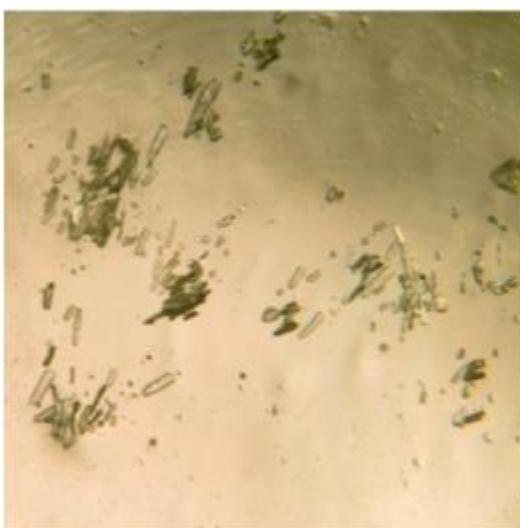
Limitation of X-ray Crystallography

Limitation:

Crystal with proper size and good quality is required for X-ray analysis



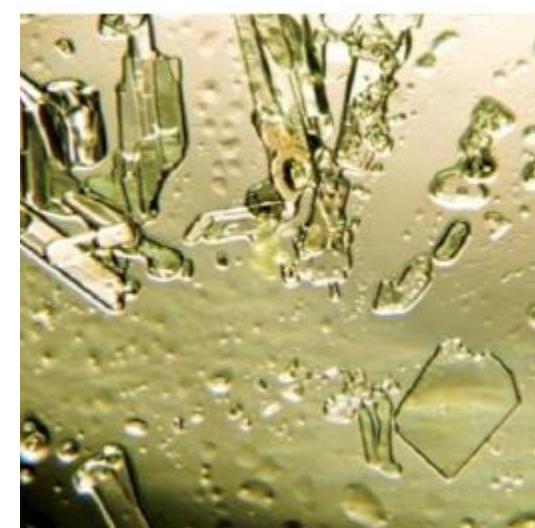
1 mm



Too small for X-ray analysis



Too thin for X-ray analysis



Proper crystal

3 approaches will be explained:

- Micro ED
- Crystalline Sponge
- Structure modification for better crystallinity

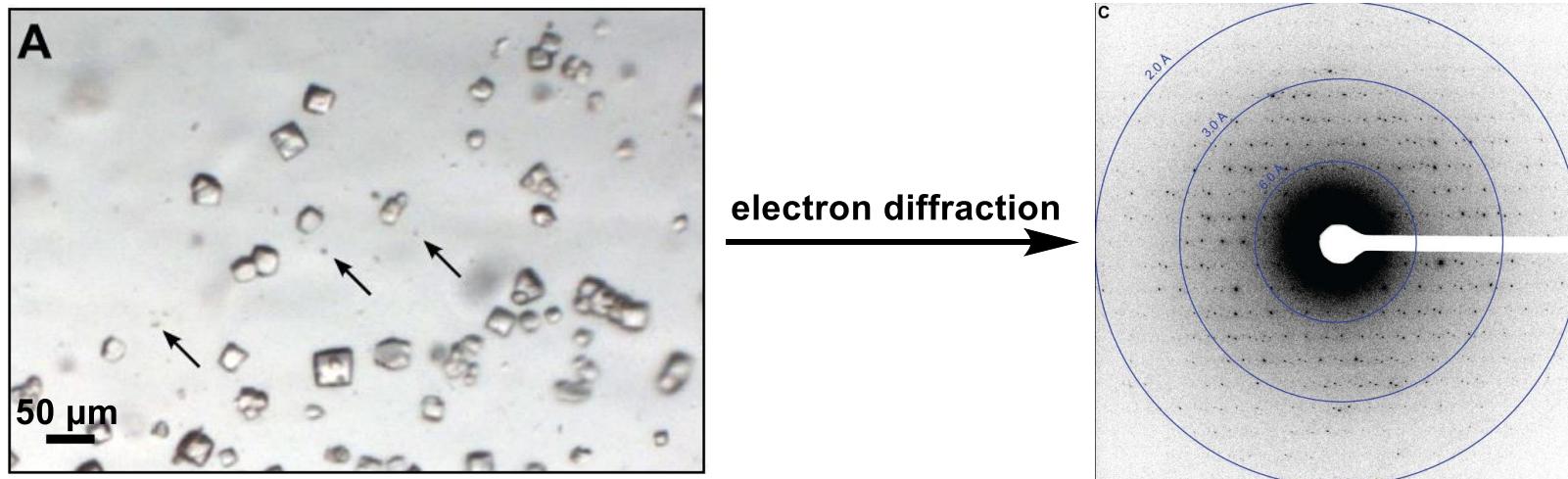
1. Introduction

2. Strategy for Crystal Analysis

Micro ED

micro ED: Use electron beam (interacts with nucleus and electrons)

Interactions are stronger than X-ray (interacts with only valence electron),
so stronger diffraction patterns can be obtained.
→ Small amount of molecule is needed for sufficient analysis



Usually, electron beam is too strong for small molecules, so Gonen's group reduced the electron dose to 1/200 compared with usual method.

After obtaining the diffraction pattern, the operation is the same as for X-ray crystallography.

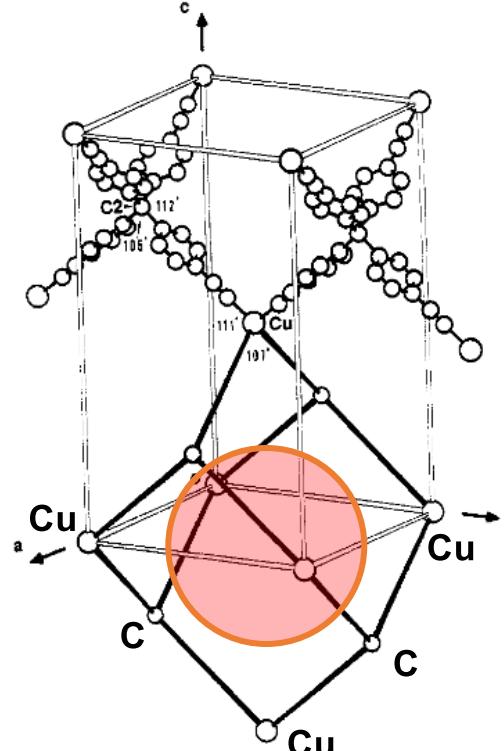
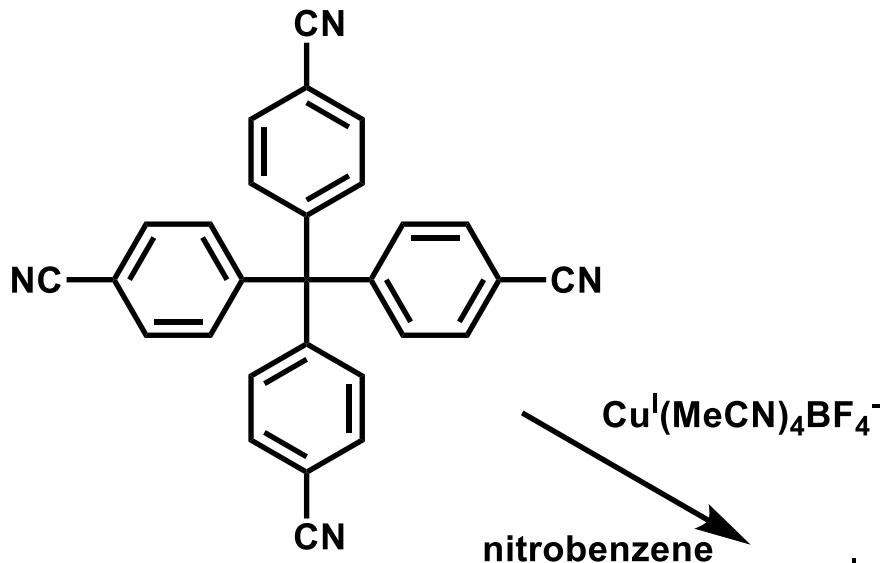
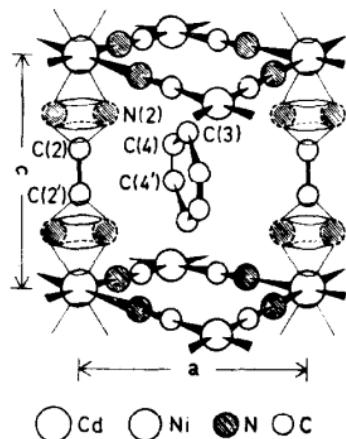
See also 240420_LS_Manaka_Matsumoto

1. Jones, C. G.; Martynowycz, M. W.; Hattne, J.; Fulton, T. J.; Stoltz, B. M.; Rodriguez, J. A.; Nelson, H. M.; Gonen, T. *ACS Cent. Sci.* **2018**, 4, 1587.
2. Shi, D.; Nannenga, B. L.; Iadanza, M. G.; Gonen, T. *eLife*, **2013**, 2, e01345.

Crystalline Sponge: Background

crystalline sponge method: Absorb the compounds to crystalline sponge

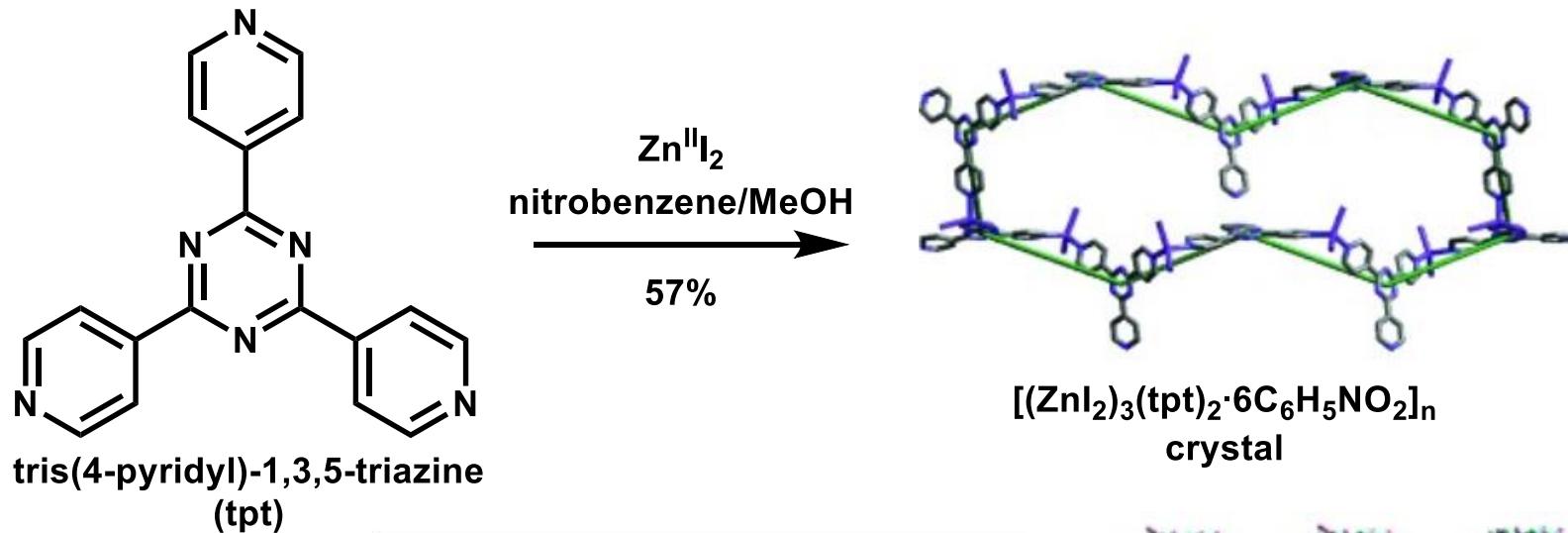
→ Crystallization is not necessary. Oil compounds can be applied to X-ray analysis in this method.



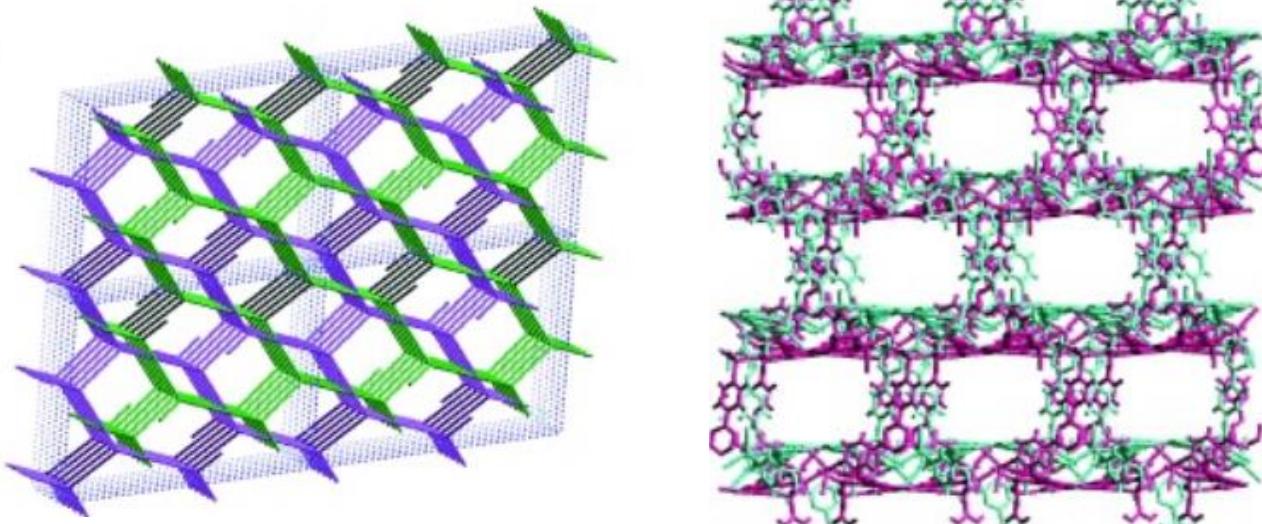
1. Miyoshi, T.; Iwamoto, T.; Sasaki, Y. *Inorg. Chim. Acta*. **1972**, 6, 59.
2. Hoskins, B. F.; Robson, R. *J. Am. Chem. Soc.* **1990**, 112, 1546.

MOF with Flexibility and Robstness (1)

Interpenetrated network MOF developed by Fujita's group:

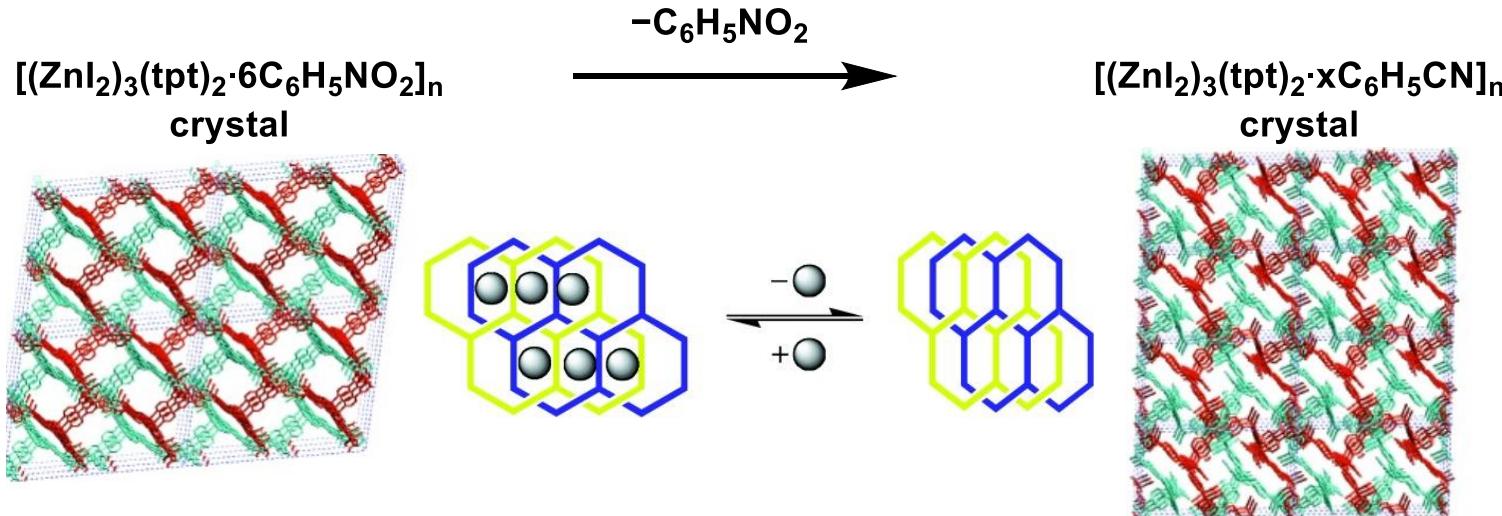


They found that two types of networks (green and purple) are interpenetrated in crystal structure



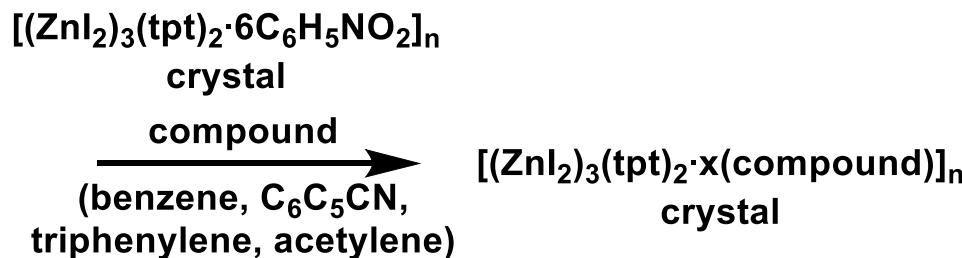
MOF with Flexibility and Robstness (2)

- Compression change of the networks "without destroying the crystalline nature "

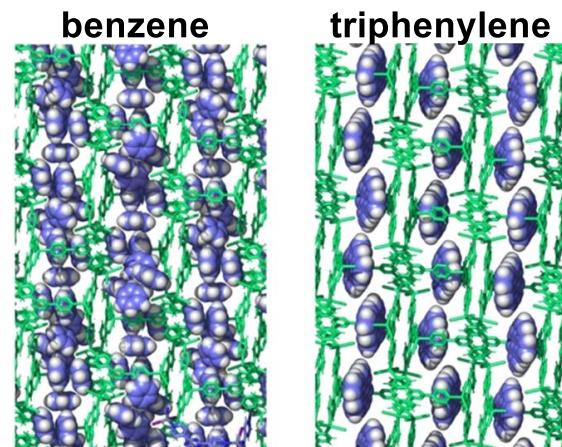


The guest molecule is released from the crystal and unit-cell parameters changed.

- Single-crystal-to-single-crystal (SCSC) guest exchange



"Substituted by prolonged soaking in a saturated solution of the guest"

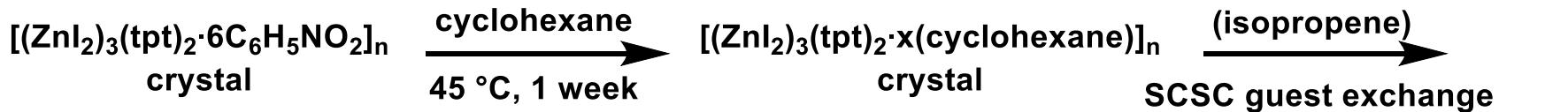


1. Zigon, N.; Duplan, V.; Wada, N.; Fujita, M. *Angew. Chem. Int. Ed.* **2021**, *60*, 25204.
2. Biradha, K.; Fujita, M. *Angew. Chem. Int. Ed.* **2002**, *41*, 3392.
3. Ohmori, O.; Kawano, M.; Fujita, M. *J. Am. Chem. Soc.* **2004**, *126*, 16292.

Crystalline Sponge

Crystalline sponge: Developed by Fujita's group

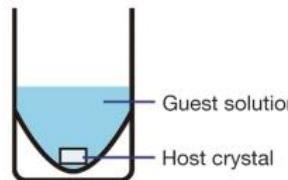
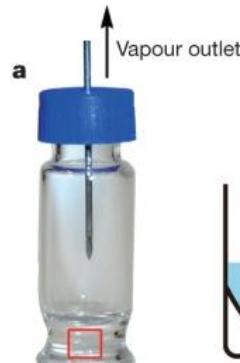
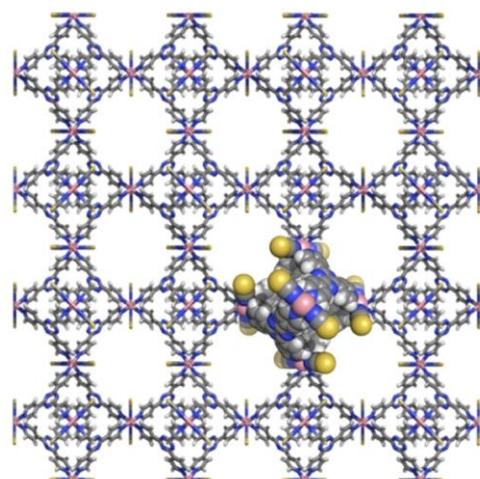
Enabled X-ray analysis of non-crystal compound at nano gram scale



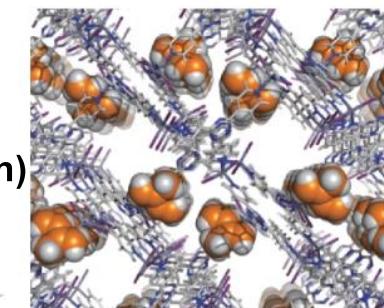
Simple operation

"Any guest that shows stronger host-guest interactions than hydrocarbons can, in principle, be absorbed."

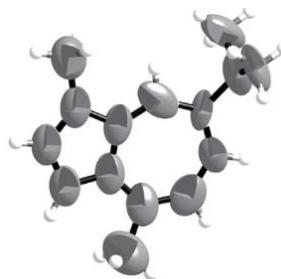
Other crystalline sponge in the paper:



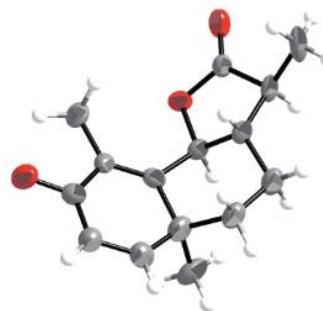
(cyclohexane/ CH_2Cl_2 solution)



Other examples:



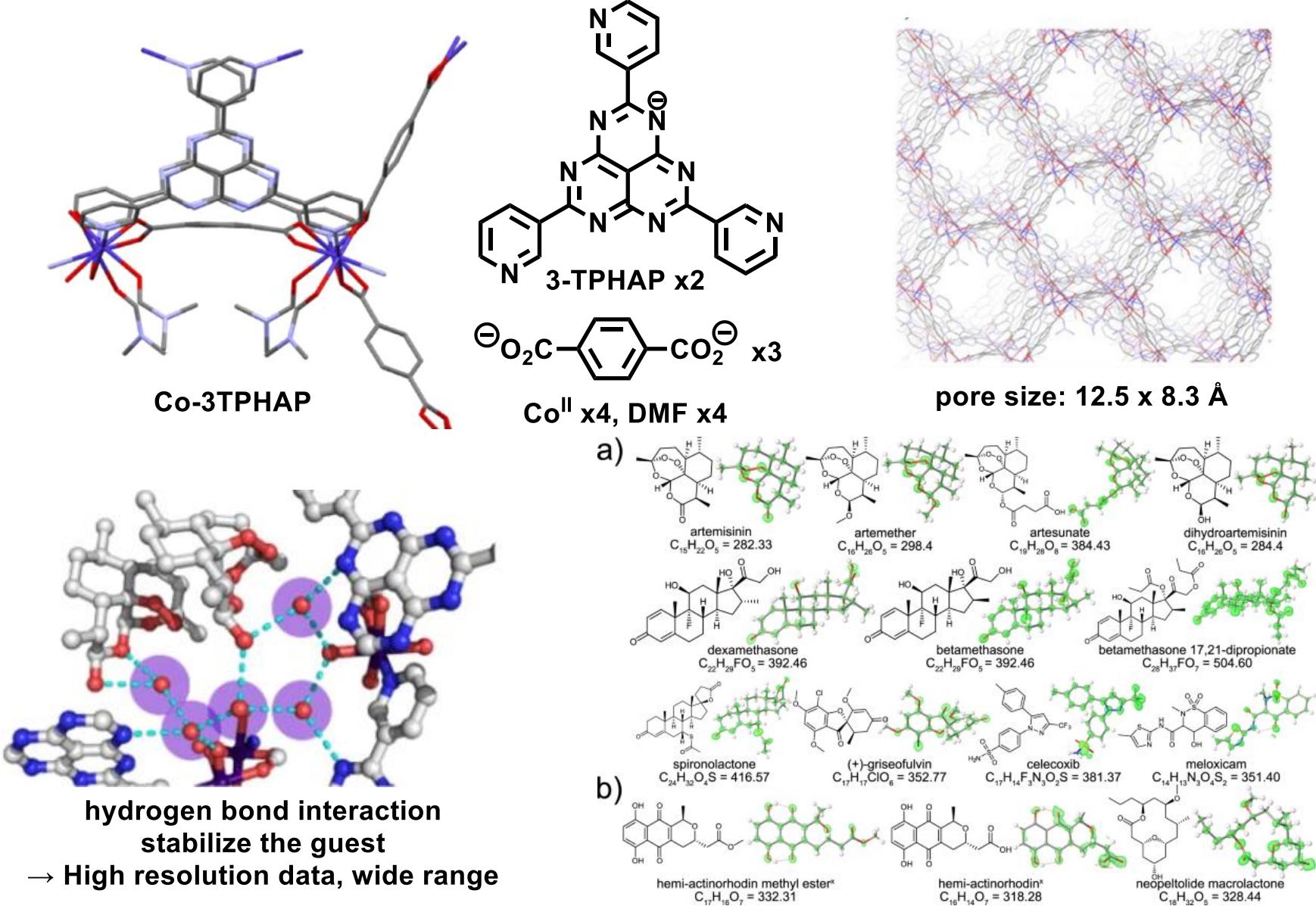
guaiazulene
(from 80 ng)



Santonin (from 15 μg)
absolute configuration
unambiguously determined

1. Inokuma, Y.; Yoshida, S.; Ariyoshi, J.; Arai, T.; Hitora, Y.; Takada, K.; Matsunaga, S.; Rissanen, K.; Fujita, M. *Nature*, **2013**, *495*, 461.
2. Zigon, N.; Duplan, V.; Wada, N.; Fujita, M. *Angew. Chem. Int. Ed.* **2021**, *60*, 25204.

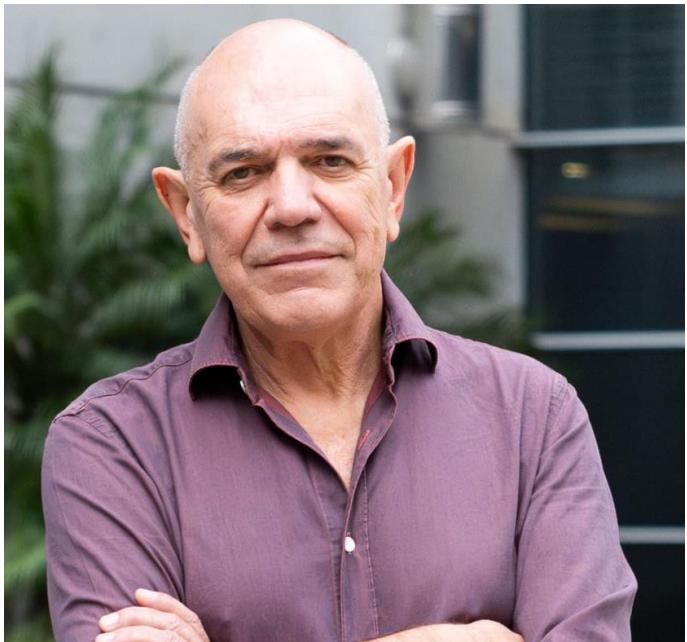
Crystalline Sponge: Recent Progress



1. Wada, Y.; Usov, P. M.; Mukaida, M.; Ohmori, K.; Ando, Y.; Fuwa, H.; Ohtsu, H.; Kawano, M. *Nat. Comm.* **2024**, 15, 81.

Prof. Antonio M. Echarvarren

Prof. Antonio M. Echarvarren



Career:

- 1982:** Ph. D. @The Universidad Autónoma de Madrid
(Prof. F. Fariña)
- 1982-1984:** Postdoctoral Fellow @Boston College
(Prof. T. R. Kelly)
- 1984-1986:** Assistant professor
@The Universidad Autónoma de Madrid
- 1986-1988:** NATO-fellow @The Colorado State University
(Prof. J. K. Stille)
- 1988-1992:** Institute of Organic Chemistry of Spanish National Council (CSIC) researcher
- 1992-2009:** Professor @The Universidad Autónoma de Madrid
- 2004-:** Research Professor @CSIC
- 2004-:** Group Leader @Institute of Chemical Research of Catalonia (ICIQ)

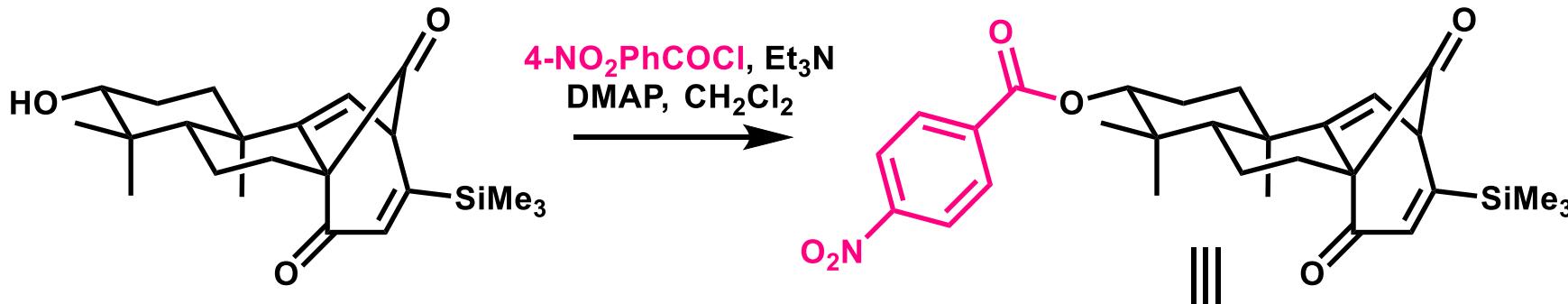
Research Interest:

C-H activation, Total synthesis (Au catalyzed)

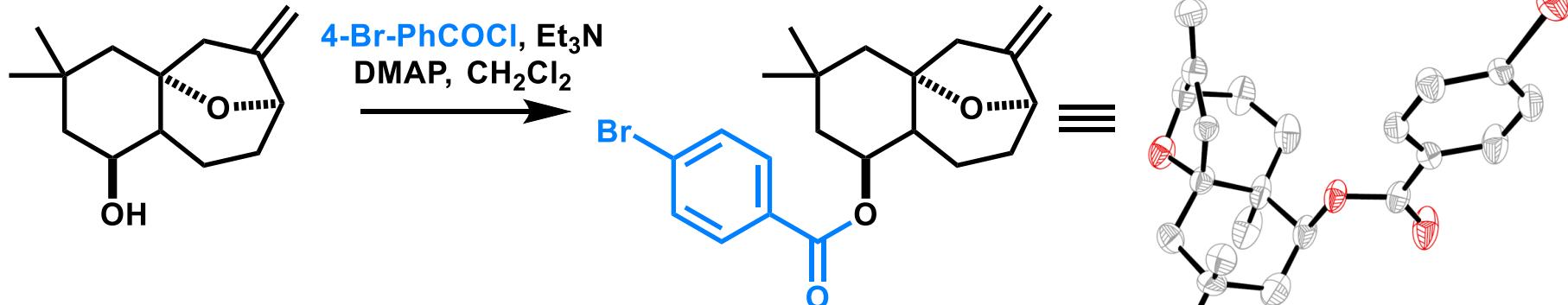
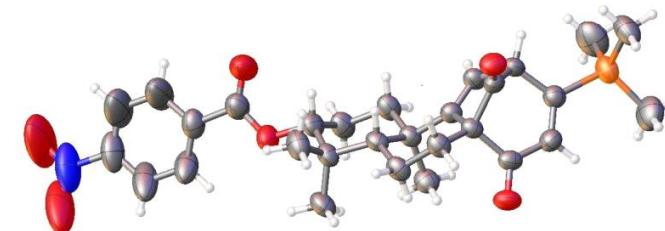
Structure Modification for Better Crystallinity

Acylation of alcohol was sometimes conducted for better crystallinity of substrates.

Intermolecular interactions (hydrogen bond, electrostatic interactions) increase the crystallinity

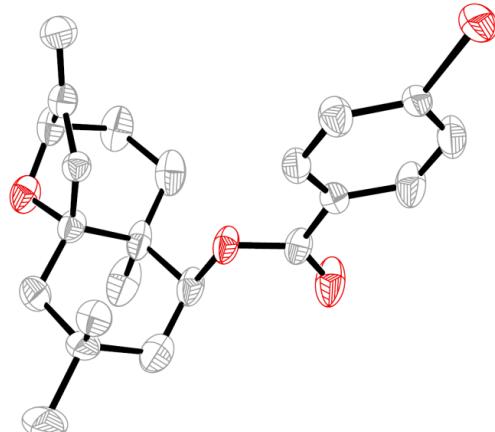


Not suitable for absolute configuration determination.



Suitable for absolute configuration determination

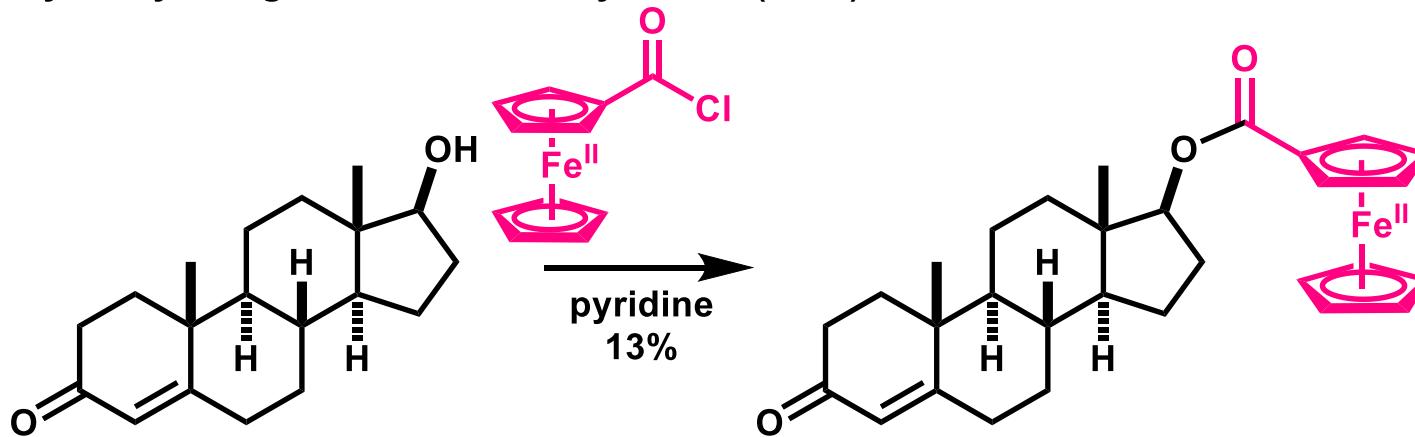
Less crystallinity compared with *p*-nitrobenzoyl



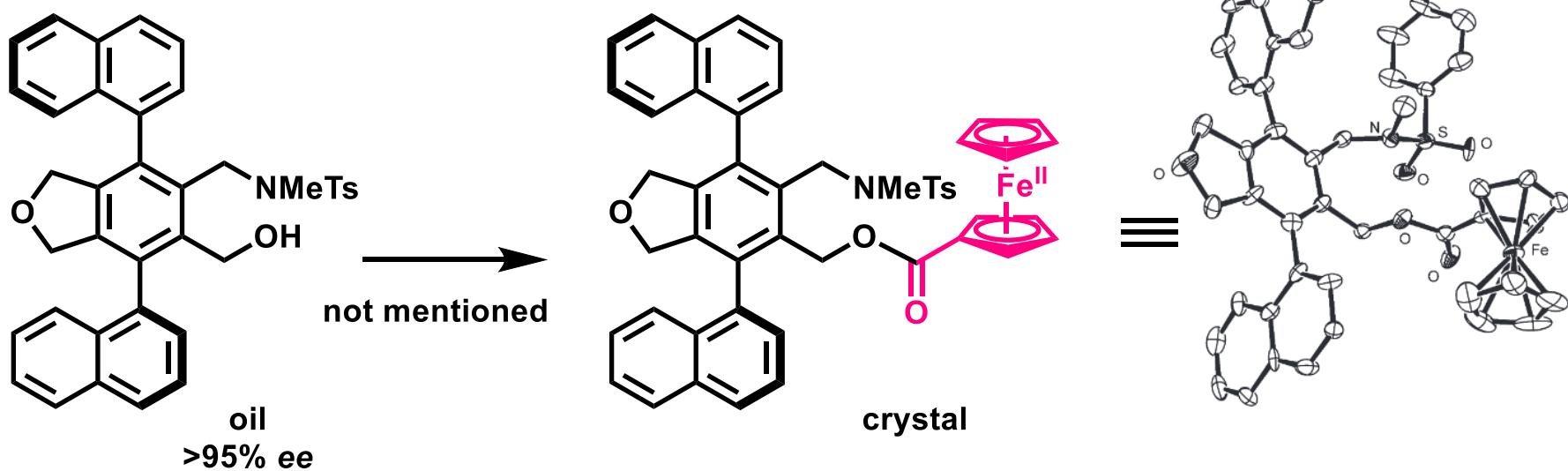
1. Song, L.; Zhu, G.; Liu, Y.; Liu, B.; Qin, S. *J. Am. Chem. Soc.* **2015**, *137*, 13706.
2. Qin, X.-Q.; Wu, Q.-L.; Han, F.-S. *Org. Lett.* **2021**, *21*, 8570.

Prior Work Using Ferrocene Carboxylic Acid

Early study using ferrocene carboxylic acid (1980)

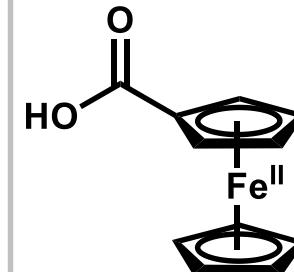
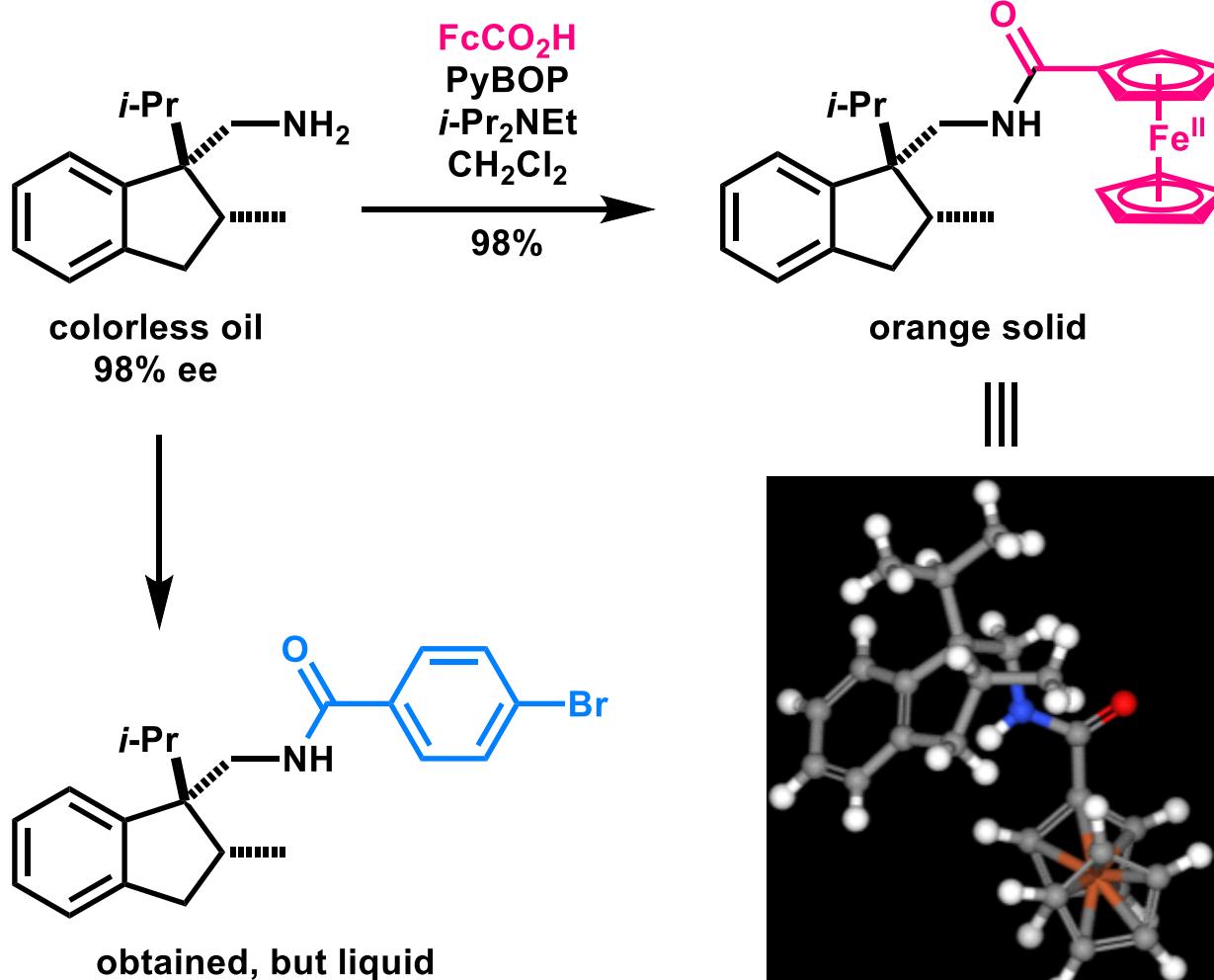


Previous studies of ferrocenyl esters as crystallizing reagents



1. Hoffman, K.; Rießelmann, B.; Wenzel, M. *Liebigs Ann. Chem.* **1980**, 1980, 1181.
2. Shibata, T.; Arai, Y.; Takami, K.; Tsuchikama, K.; Fujimoto, T.; Takebayashi, S.; Takagi, K.; *Adv. Synth. Catal.* **2006**, 348, 2475.

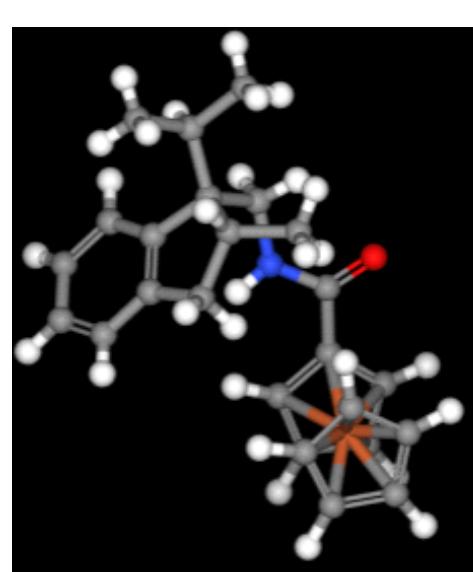
Proof of Concept



FcCO₂H
7,300 yen/ 1 g

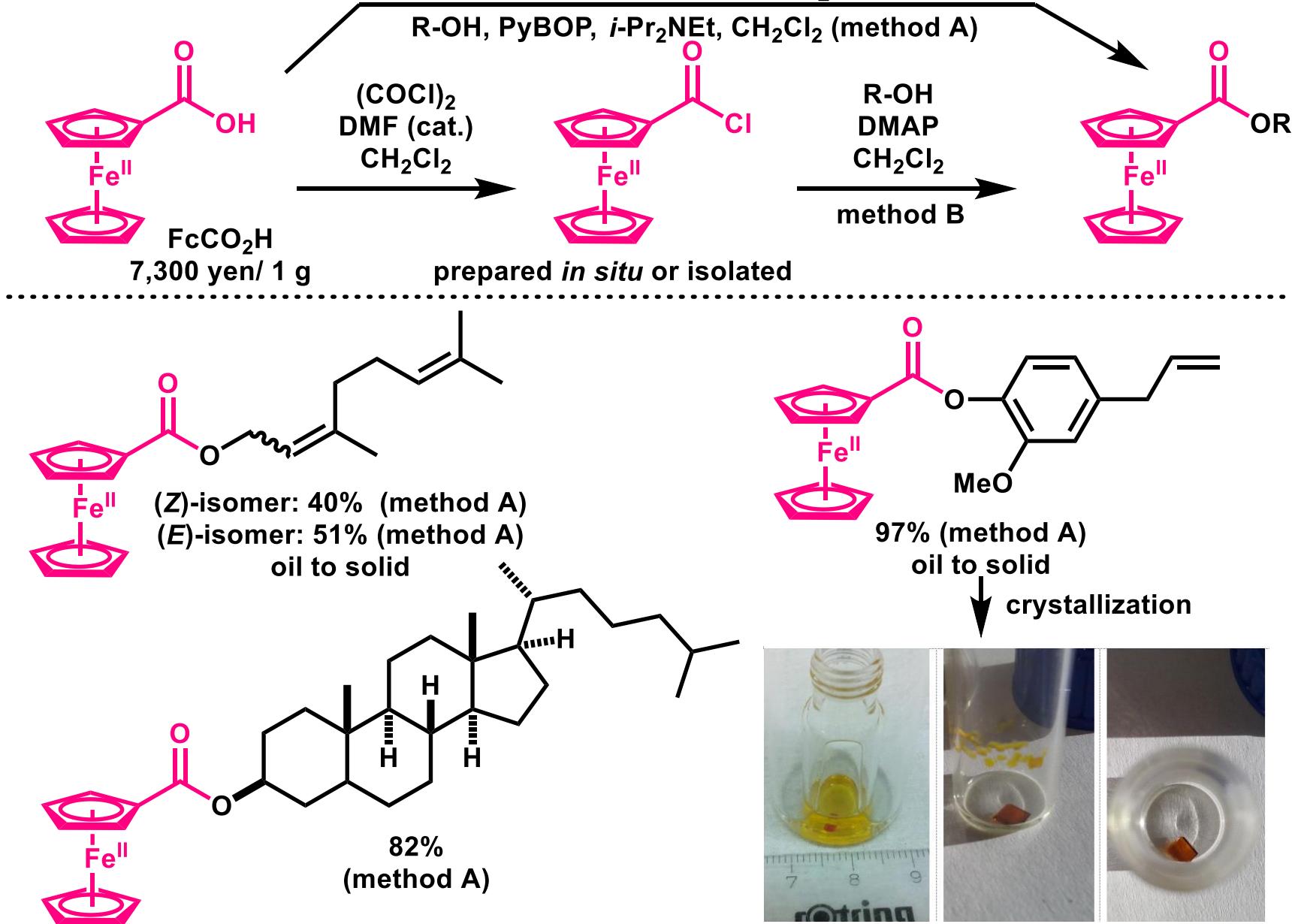
Characteristic of FcCO₂H:

- Highly crystalline
- Contains heavy atom
- Stable
- Orange color
- Relatively inexpensive



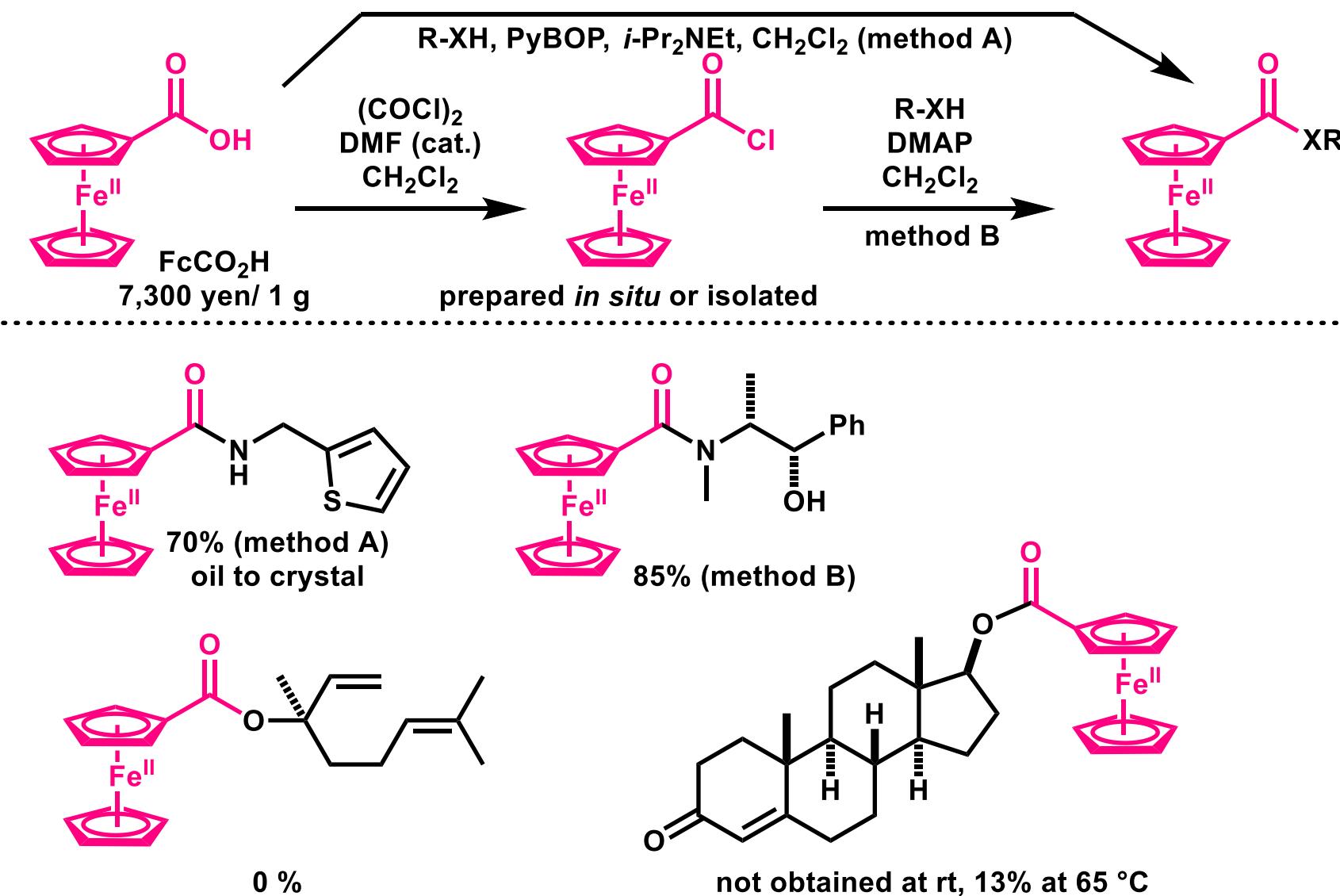
absolute configuration
unambiguously determined

Substrate Scope



1. Holstein, P. M.; Holstein, J. J.; Escudero-Adan, E. C.; Baudoin, O.; Echavarren, A. M. *Tetrahedron Asymmetry*, 2017, 28, 1321.

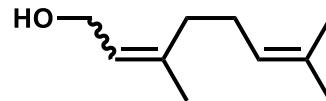
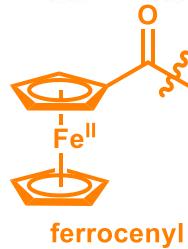
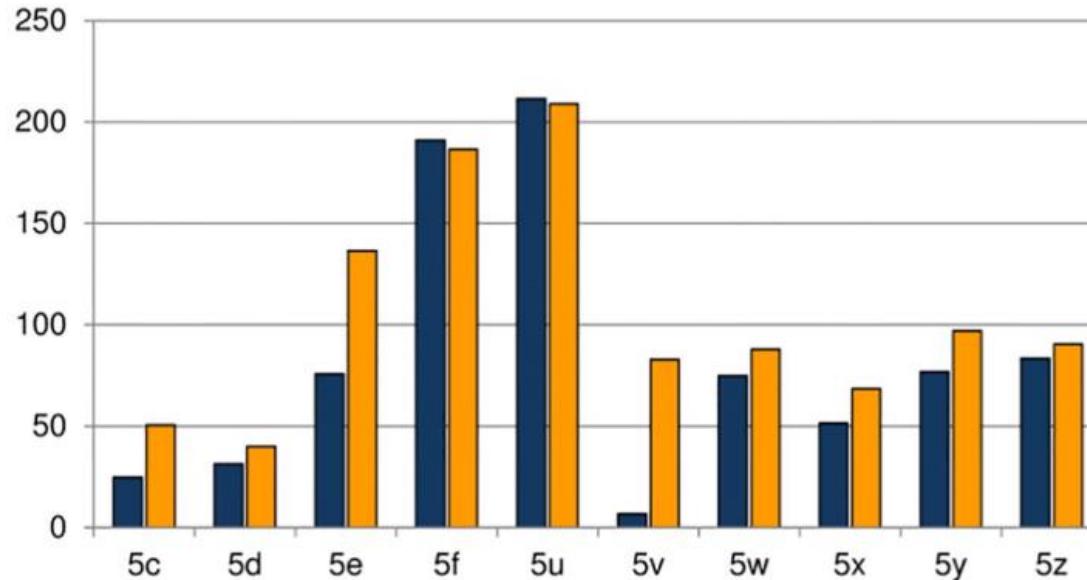
Substrate Scope and Limitation



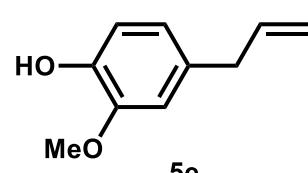
1. Hoffman, K.; Rießelmann, B.; Wenzel, M. *Liebigs Ann. Chem.* **1980**, 1980, 1181.
2. Holstein, P. M.; Holstein, J. J.; Escudero-Adan, E. C.; Baudoin, O.; Echavarren, A. M. *Tetrahedron Asymmetry*, **2017**, 28, 1321.

Comparison of the Melting Point

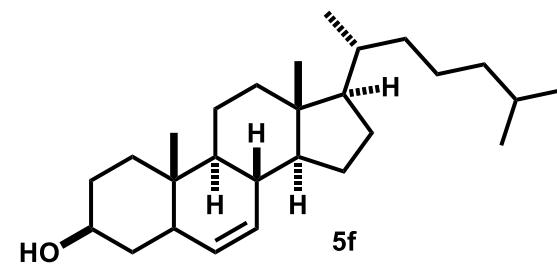
m.p. in °C



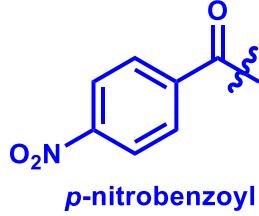
5c: (Z)-isomer
5d: (E)-isomer



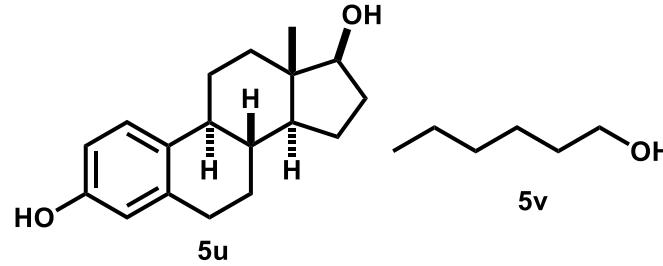
5e



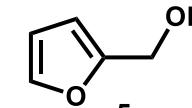
5f



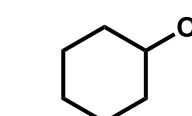
p-nitrobenzoyl



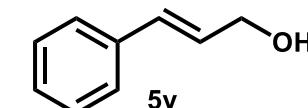
5u



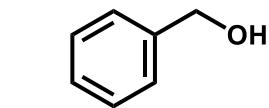
5w



5v



5y



5z

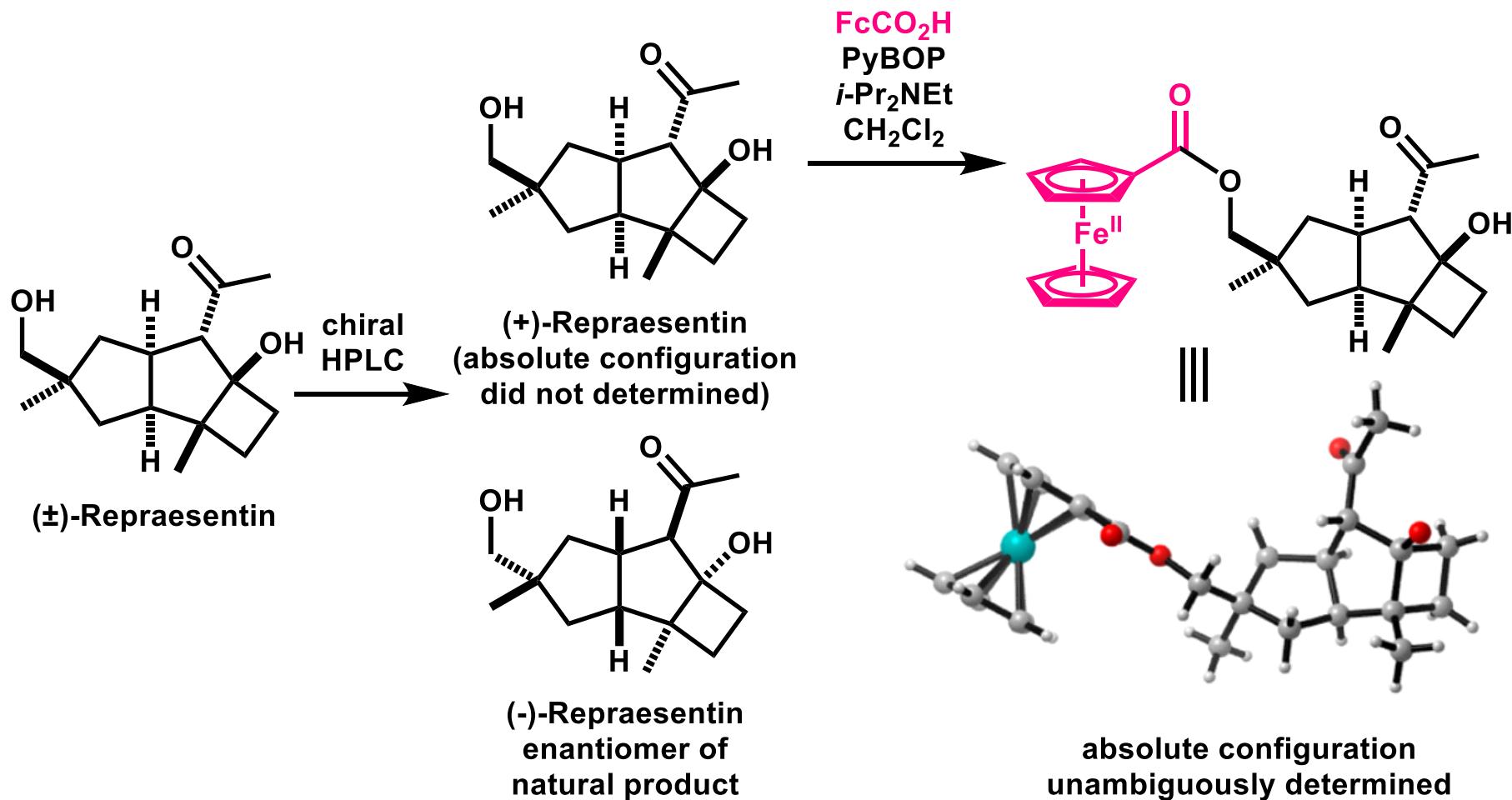
1. Holstein, P. M.; Holstein, J. J.; Escudero-Adan, E. C.; Baudoin, O.; Echavarren, A. M. *Tetrahedron Asymmetry*, 2017, 28, 1321.

blue: *p*-nitrobenzoyl derivatives
orange: ferrocenyl derivatives

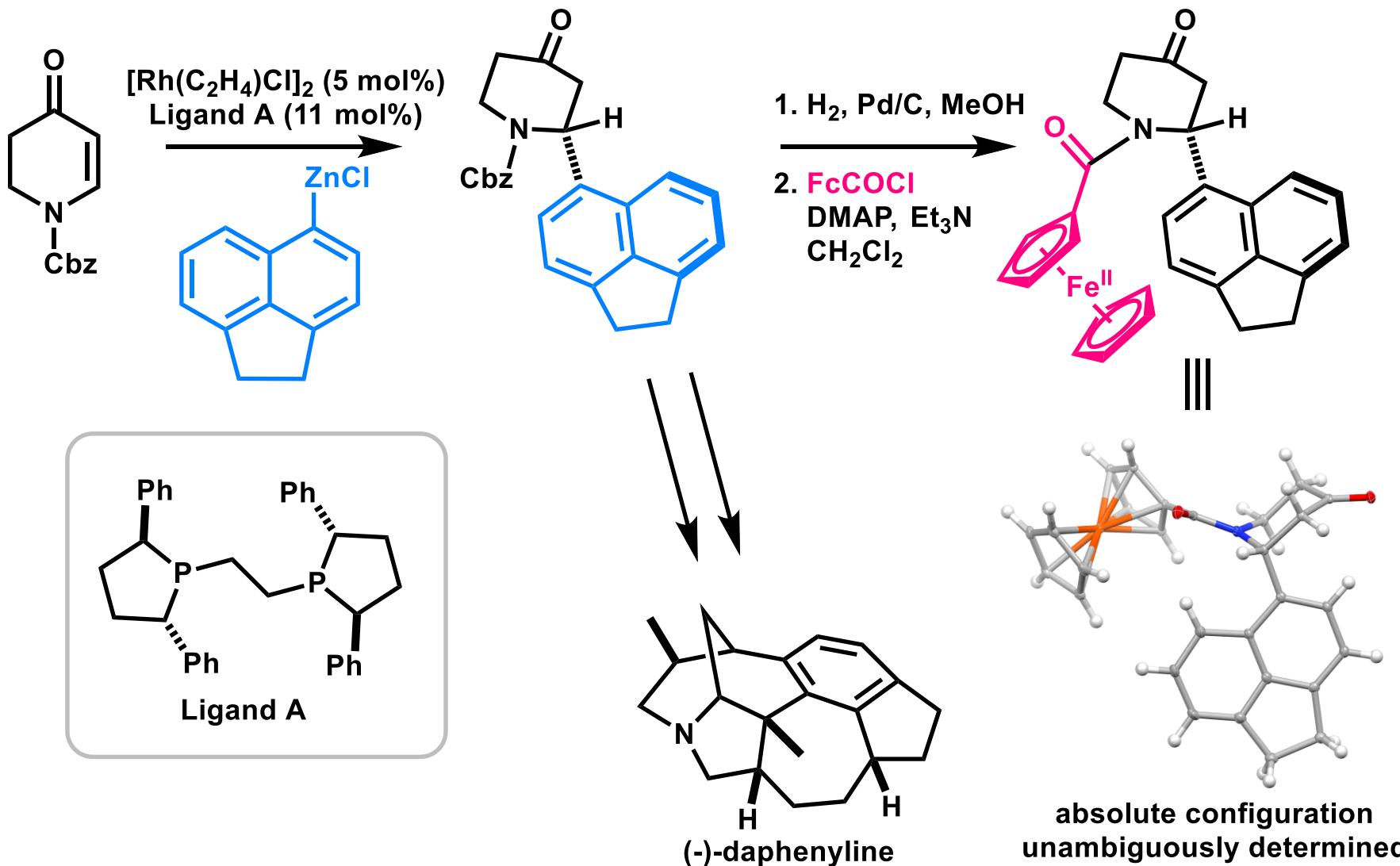
similar melting point was observed.

Indicating that ferrocenyl derivative is as good crystallinity as *p*-nitrobenzoyl derivative.

Application in Total synthesis (1)



Application in Total Synthesis (2)



Summary

Method	NMR	X-ray crystallography	Micro ED	Crystalline sponge
Range	◎ (almost all)	✗ (crystal with good size, quality)	△ (crystal with good quality)	potentially ○ (size of molecule)
Certainty of the structure	○	◎	◎	Potentially ○
Comment	~ mg order is required for data collection	Structure modification is sometimes effective	~μg scale would be possible	Further development will be expected