

Ion Mobility-Mass Spectrometry (IM-MS)

Literature Seminar #3 (2014.9.27)

Kiyomichi SHINODA

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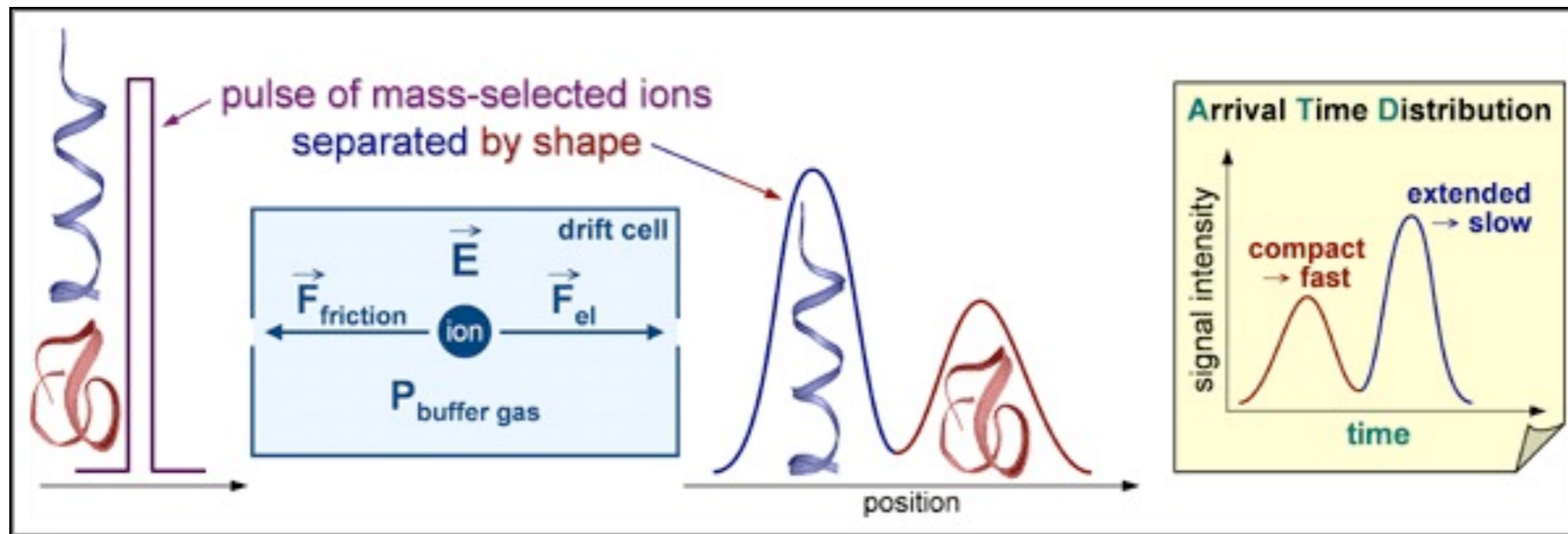
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What is Ion Mobility?

- Ion mobility achieves separation of compounds by their **size** and **shape**.

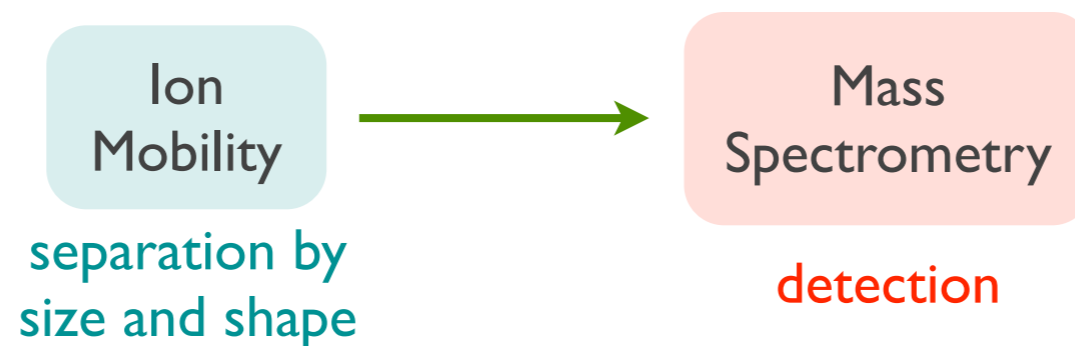


- E = uniform electric field [均一電場]
- F_{friction} = force of friction (caused by collisions of ions with the buffer gas)
- F_{el} = force of elimination
- $P_{\text{buffer gas}}$ = pressure of buffer gas

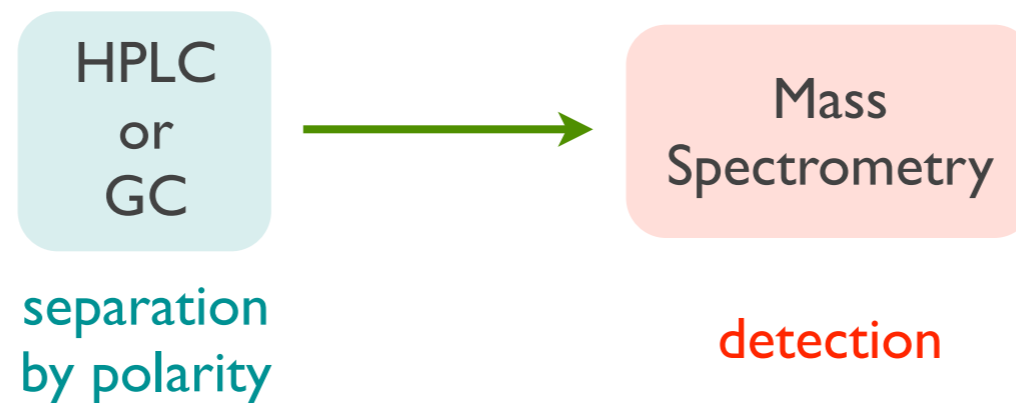
This figure is cited from Bowers group web page.
 (http://bowers.chem.ucsb.edu/theory_analysis/ion-mobility/index.shtml)

What is Ion Mobility-Mass Spectrometry (IM-MS)?

- conceptual diagram of IM-MS

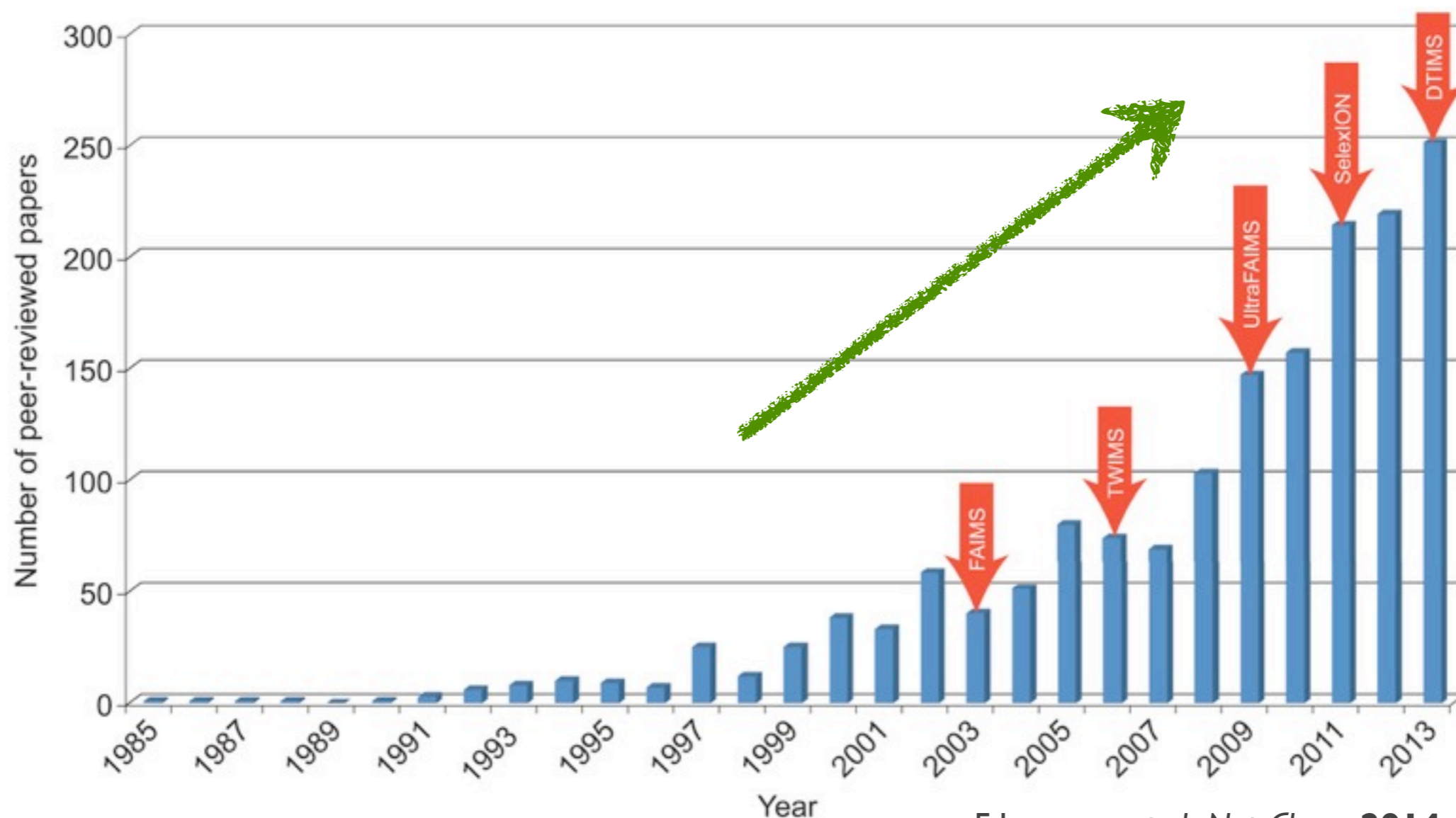


cf.) How about LC-MS or GC-MS??



Recent Explosion in Research Using IM-MS

- number of peer-reviewed papers published annually combining IM and MS



F. Lanucara et al., *Nat. Chem.* **2014**, 6, 281.

- Especially, the application toward biomolecules is the current hot topic.

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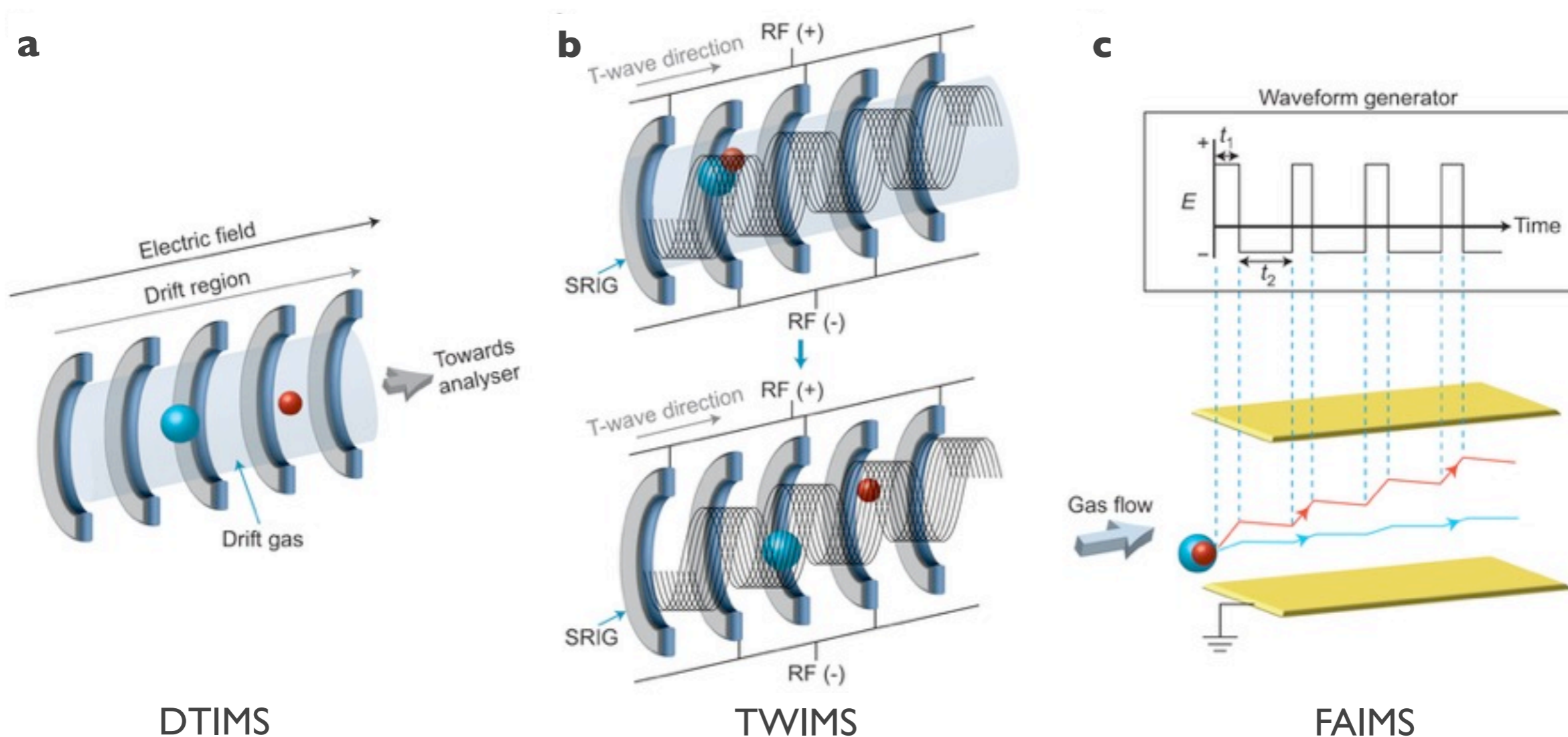
§3.4 Mechanism Analysis of Aggregation Inhibitor

Three Types of Ion Mobility Spectrometry

a: Drift-Time Ion Mobility Spectrometry (DTIMS)

b: Traveling-Wave Ion Mobility Spectrometry (TWIMS)

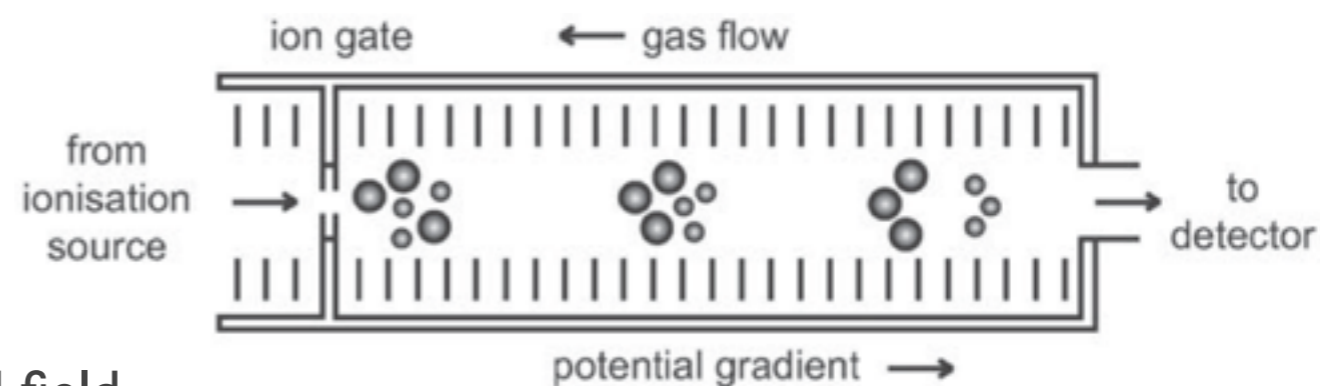
c: Field-Asymmetric Ion Mobility Spectrometry (FAIMS)



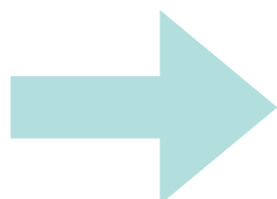
DTIMS (I): System Outline

- Pulse of ions is introduced into a **drift cell**.

- Static uniform electric field is applied.
- filled with drift gas (typically helium)
- Ions travel in the direction of the applied field in uniform motion.



C. Uetrecht *et al.*, *Chem. Soc. Rev.* **2010**, 39, 1633.



Ions collide with gas and are separated based on its size and shape.
(Small ions travel faster than big ions.)

- The time taken for an ion to drift through the cell (= drift time) is related to its rotationally averaged cross-sectional area (= **collision cross-section (CCS)**).



index of ion size and shape

DTIMS (2): How to Calculate CCS?

- Mason-Schamp equation: provide the **relationship between ion mobility and CCS**

$$K = \frac{v_d}{E} = \frac{3}{16} \frac{q}{N} \left(\frac{1}{m} + \frac{1}{M} \right)^{\frac{1}{2}} \left(\frac{2\pi}{k_B T} \right)^{\frac{1}{2}} \frac{1}{\Omega}$$

$$\left(\text{in the case } \frac{E}{N} \leq 2 \times 10^{-17} \text{ (V} \cdot \text{cm}^2) \right)$$

K : measured mobility at 273.15 K, 101325 Pa (イオン移動度)

v_d : drift velocity (ドリフト速度)

E : electric field (電場)

q : charge of the analyze ion (分析対象イオンの価数)

N : density of the drift gas (ドリフトガスの密度)

m : mass of the analyte ion (分析対象イオンの質量)

M : mass of the drift gas (ドリフトガス分子の質量)

k_B : the Boltzmann constant (ボルツマン定数)

T : gas temperature (ガスの温度)

Ω : collision cross section (CCS) (衝突(散乱)断面積)

F. Lanucara et al., *Nat. Chem.* **2014**, 6, 281.

C. S. Creaser et al., *Analyst* **2004**, 129, 984.

T. Sugai, *J. Mass Spectrom. Soc. Jpn.* **2010**, 58, 47.

**As the recorded drift time of an ion can be easily converted to v_d ,
we can calculate CCS using this apparatus!!**

DTIMS (3): Advantages and Disadvantages

• advantages

- **ability to determine CCS:** direct determination of CCS from Mason-Schamp eq.
- **high resolving power:** An ion with CCS of 100 \AA^2 can theoretically be separated from an ion with $\pm 1 \text{ \AA}^2$ difference (= 1% uncertainty).

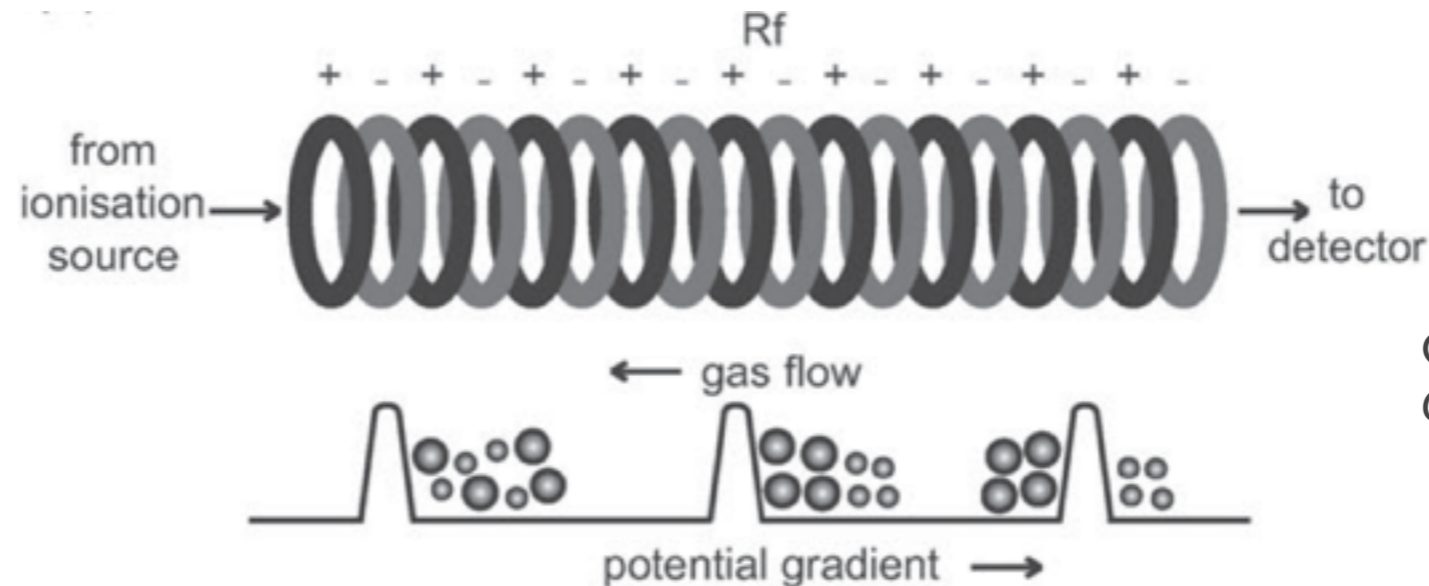
• disadvantages

- **low detection efficiency:** Ions are lost on several devices where entry or exit of ions into the drift cell occurs.
- **hugeness of apparatus:** Apparatus tend to become huge to provide high resolving power.

F. Lanucara *et al.*, *Nat. Chem.* **2014**, 6, 281.

TWIMS (I): System Outline

- developed by Waters Corporation (S. Pringle *et al.*, *Int. J. Mass Spectrom.* **2007**, 261, 1.)



C. Uetrecht *et al.*,
Chem. Soc. Rev. **2010**, 39, 1633.

- used for CCS determination:** the condition $\frac{E}{N} \leq 2 \times 10^{-17} \text{ (V} \cdot \text{cm}^2)$ is met.
 - Radio-frequency voltages (高周波電圧) of opposite phases are applied to adjacent electrodes and this voltages consist a sine curve potential barrier which confine the ions.
 - Direct current voltage (直流電圧) is applied to each electrode sequentially providing “**traveling waves**” and this “wave” propels ions from cell entrance to the exit.
- Higher mobility ions are carried by the wave, whereas lower mobility ions are trapped by the wave, thus taking longer to move through the drift cell.

TWIMS (2): Advantages and Disadvantages

• advantages

- **high detection efficiency:** RF voltages confine ions and prevent their diffusion.
- **ability to determine CCS:** Although drift time calibration with analytes of similar physical and chemical features with known CCS is needed, it is possible.

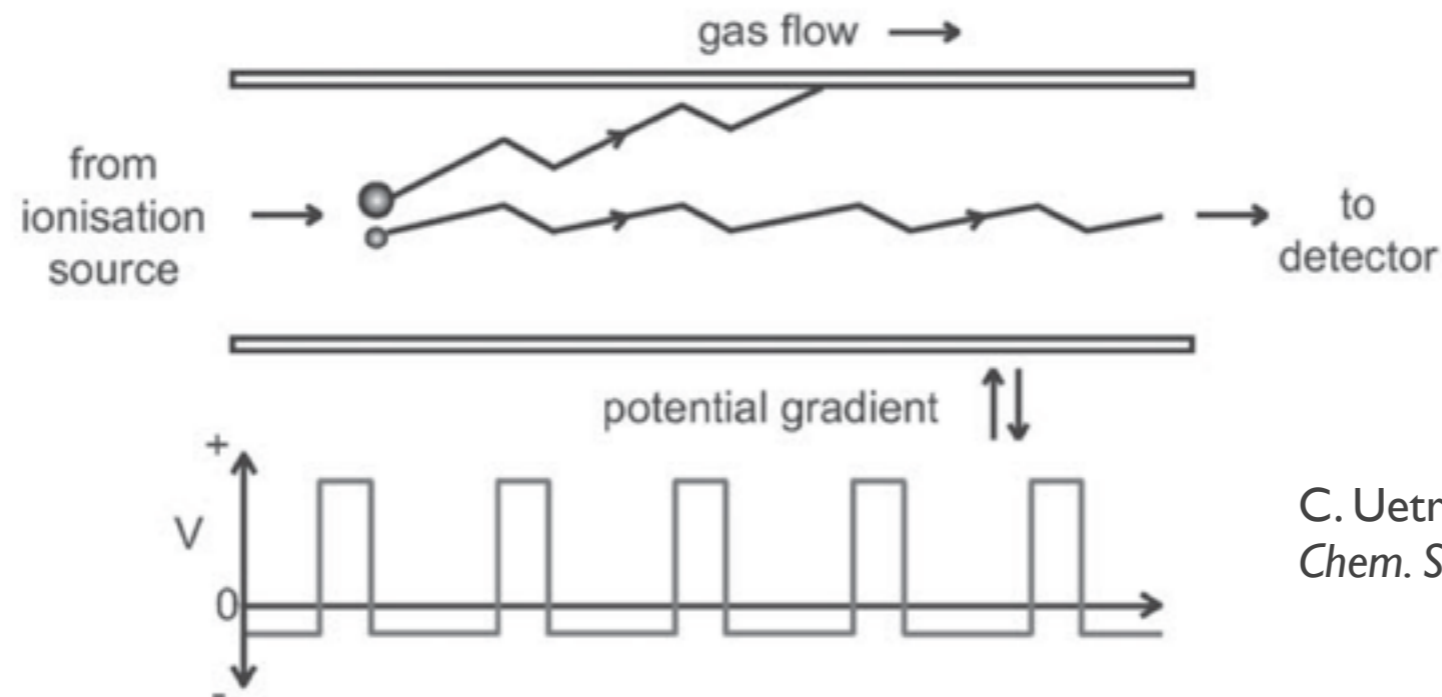
• disadvantages

- **relatively low resolving power:** An ion with CCS of 900 \AA^2 is theoretically separated from an ion with $\pm 20 \text{ \AA}^2$ difference at most.
- **limitation of CCS determination:** When system is complex and calibration is difficult, CCS determination is no longer carried out.

F. Lanucara *et al.*, *Nat. Chem.* **2014**, 6, 281.

FAIMS (I): System Outline

- constructed of two electrodes, across which an electric field is established



C. Uetrecht *et al.*,
Chem. Soc. Rev. **2010**, 39, 1633.

- As $\frac{E}{N}$ is greater than 2×10^{-17} , **determination of CCS is no longer possible.**
- Alternating asymmetric waveform makes ions oscillating and moving toward one electrode.
- To protect ions of interest from contact with electrode and following neutralization, **the compensation voltage** (補償電圧) is applied.
- Thus, this apparatus **operate as a mobility filter** to achieve increased selectivity and peak capacity before MS analysis.

FAIMS (2): Advantages and Disadvantages

- **advantages**

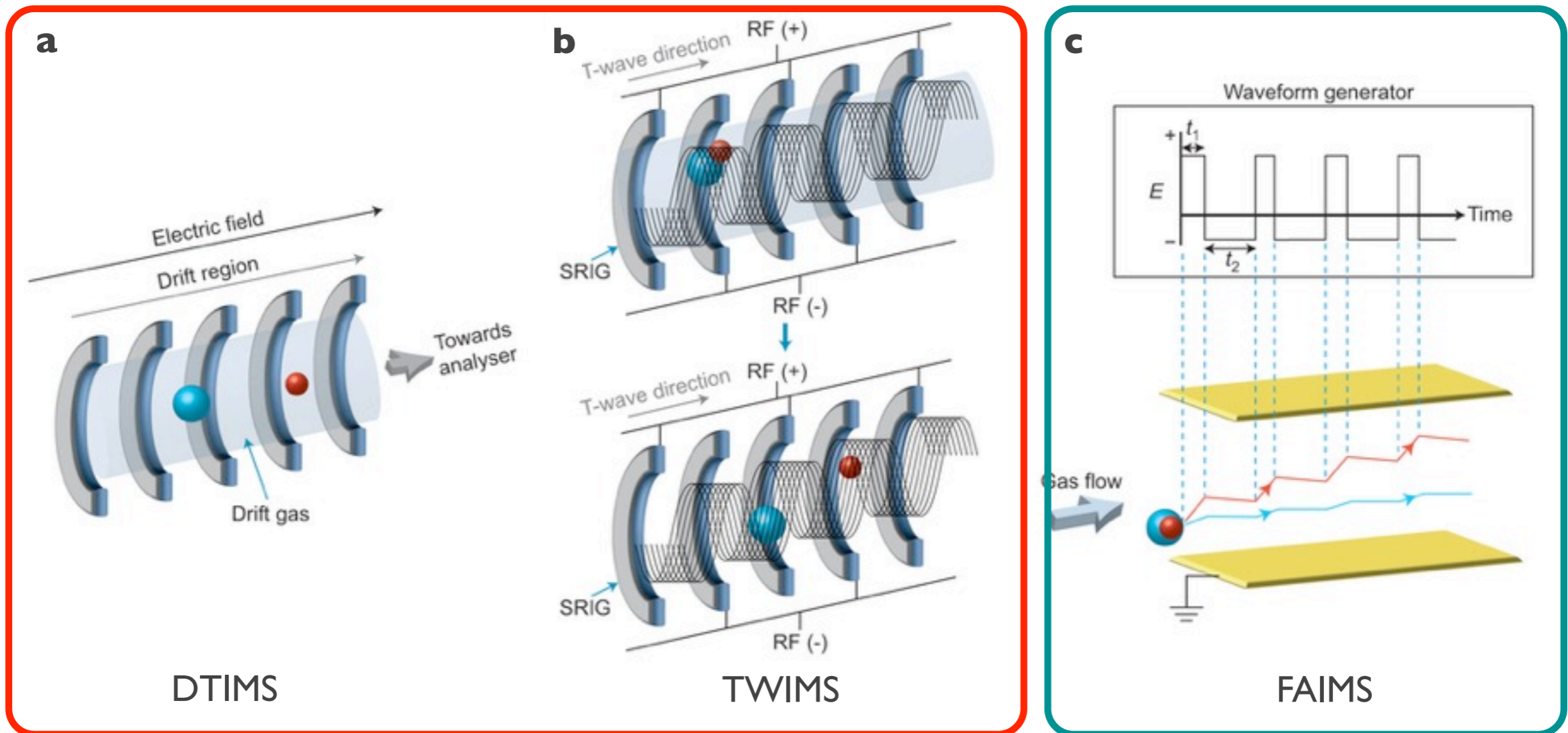
- **orthogonality toward MS:** As there's no correlation with CCS, FAIMS can be a great filter of undesired ions.
- **High resolving power** can be achieved by appropriate selection of buffer gas.

- **disadvantages**

- **CCS can't be determined.**
- **limitation of application:** limited only to post-ionization separation so far

F. Lanucara *et al.*, *Nat. Chem.* **2014**, 6, 281.

Short Summary



**Structural
Analysis**

**Post-Ionization
Separation**

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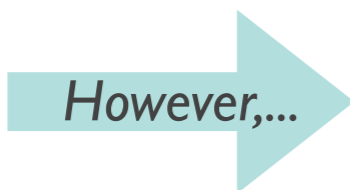
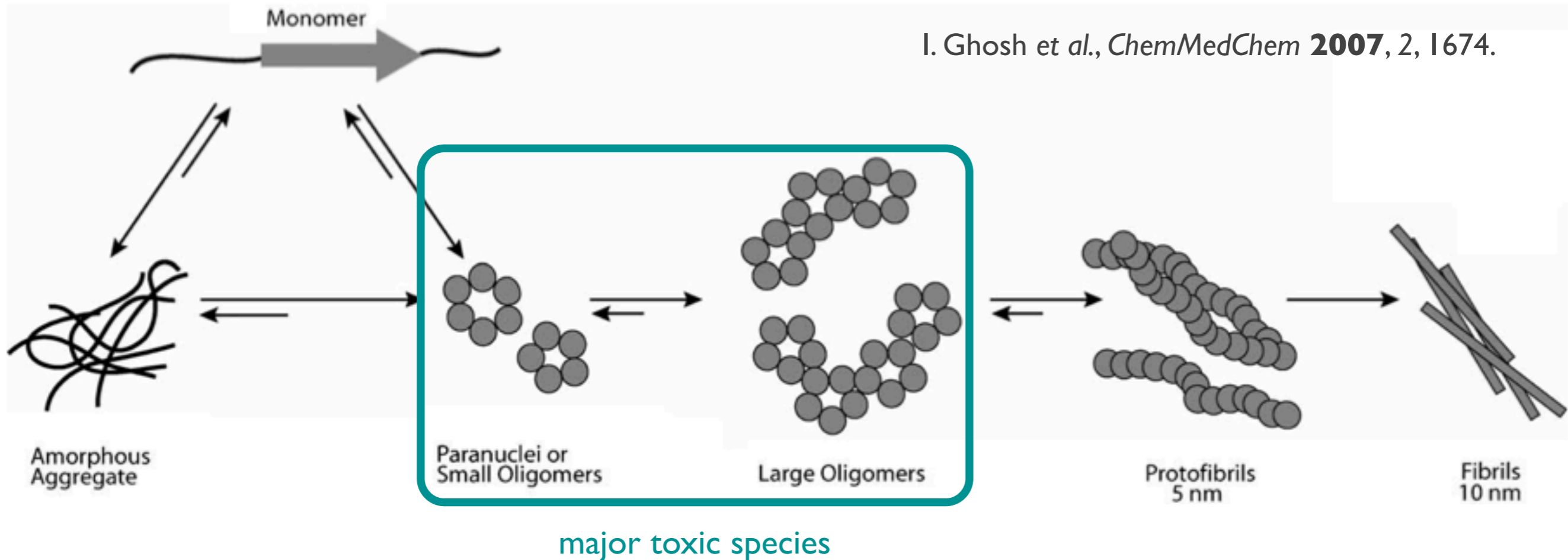
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Current Problem in A β Study



oligomer structure hasn't been studied sufficiently.

∴ A β changes its structure dynamically, so that well-established structural study (e.g. NMR or X-ray crystal analysis) which needs pure oligomer can't be conducted smoothly.

Advantages and Disadvantages of IM-MS Application

• advantages

- **quite small sample requirement:** In contrast to NMR or X-ray analysis which require milligrams of sample, IM-MS requires micrograms sample at most.
- **ability to detect intermediates:** Very short analysis time (millisecond time scale) enables the detection of short-living species (e.g. oligomers).
- **no purification requirement:** In contrast to NMR or X-ray, crude sample can be used.
- **ability to determine stoichiometry of complex:** Stoichiometry of complex such as A β oligomer and A β -inhibitor complex can be determined.

• disadvantages

- **requirement of MD simulation for precise structural information:** To obtain detailed structural information such as secondary structure or atomic level information, time-consuming and often challenging MD simulation is necessary.
- **low resolution power:** Compared to NMR or X-ray which give structural information at atomic level, 1% uncertainty in CCS is too big.

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Bowers' Home-Made IM-MS

- Prof. Michael T. Bowers

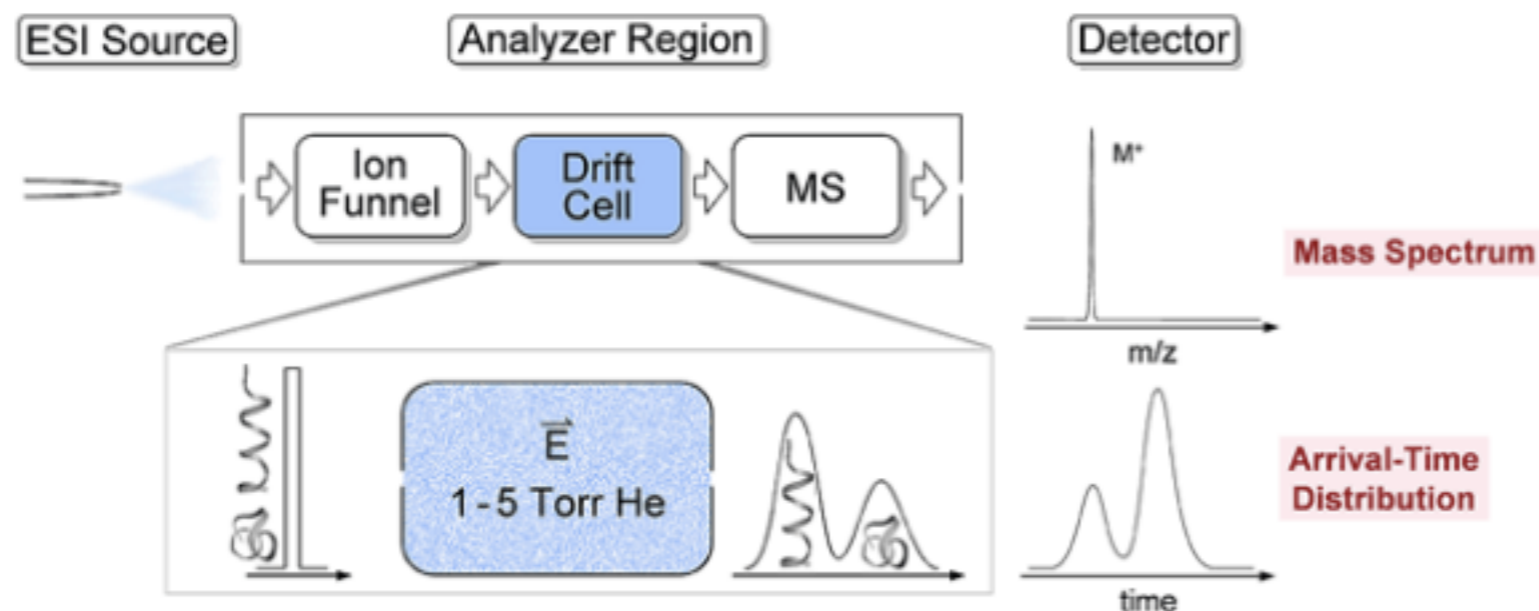


1962: obtained his B. S. from the Gonzaga University

1966: obtained his Ph. D. from the University of Illinois

1968: joined the University of California Santa Barbara faculty

- classified to **DTIMS** → **Direct estimation of CCS is possible!**

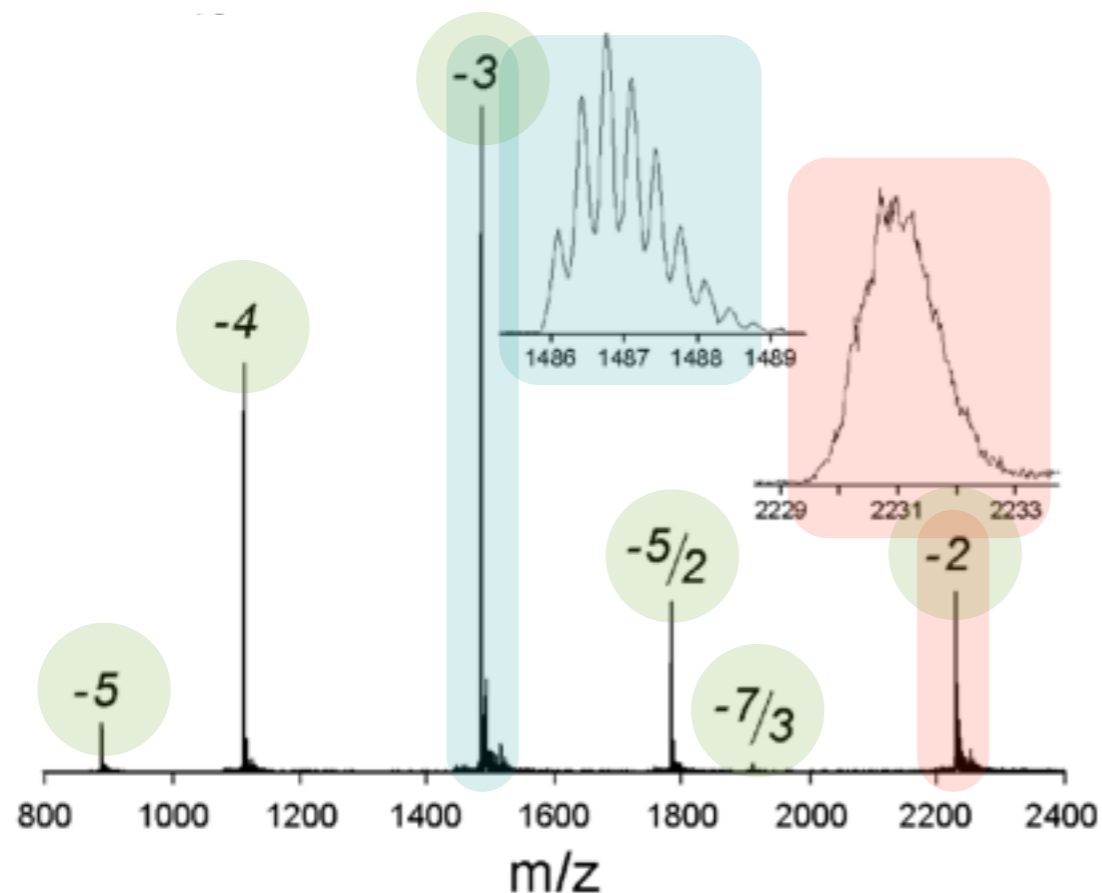


This figure is cited from Bowers group web page and is modified.
(http://bowers.chem.ucsb.edu/theory_analysis/ion-mobility/index.shtml)

Mass Spectrum of [Pro¹⁹]Aβ42

- [Pro¹⁹]Aβ42 (blue: negative charged side chain, orange: positive charged side chain)

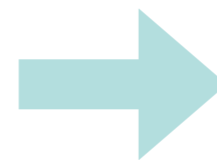
DAEFR HDSGY EVHHQ KLVPF AEDVG SNKGA IIGLM VGGV IA



(● = z/n (z = charge, n = oligomer order)
 Ex.) $z/n = -2 = -2/1, -4/2, -6/3, \dots$

several species in one peak

- **¹³C isotope distributions** is also shown: show the dominant species of the peak of interest.
- “ $z/n = -3$ ”: Peaks are separated by 0.33 ($1 \div 3$) amu.



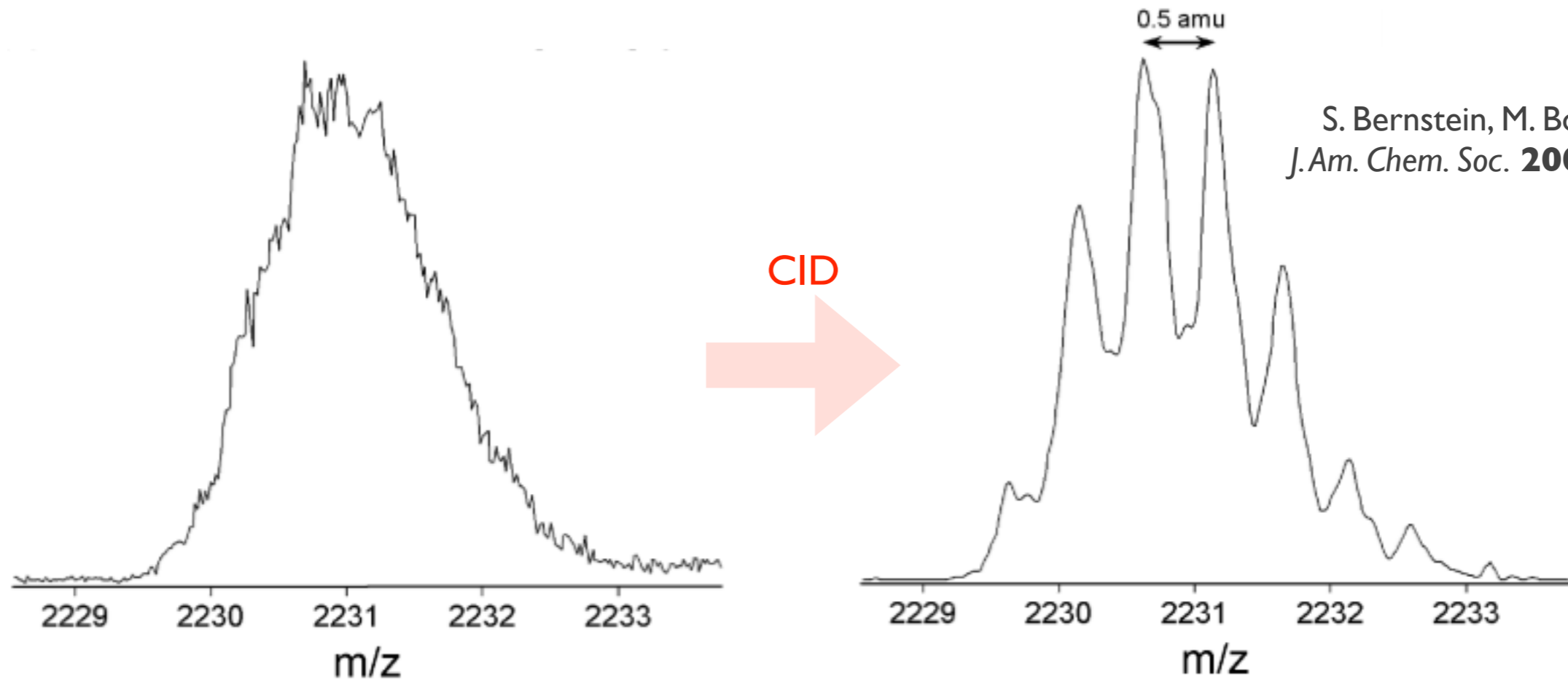
monomer-dominated

(In the dimer-dominated case, peak will be separated by $1 \div 6 = 0.17$ amu.)

- “ $z/n = -2$ ”: Peak separation is ambiguous...

Collision-Induced Dissociation (CID)

- The “ $z/n=-2$ ” charge state peak was selected by the quadrupole and subjected to CID.



- Peaks are separated by 0.5 (= 1/2) amu → indicating the monomer generation
 - Small peaks are observed at 0.25 (1÷4) amu, indicating some undissociated dimer remains.
- In summary, **components larger than dimer** construct the “ $z/n=-2$ ” charge state.

Arrival Time Distributions (ATD)

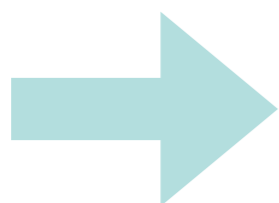
- In ATD, different compounds which have the same charge state (z/n) can be distinguished.
 - Ion mobility can separate ions by their size and shape.
 - In “ $z/n=-q$ ” peak of mass spectrum, the following species can be present:
Monomer^{-q}, Dimer^{-2q}, Trimer^{-3q}, ...
 - These species are different in size, so they can be separated using IM.



ATD shows how species with different size and shape are separated.

Assignment of Peaks in ATD (I)

- ATD of “ $z/n=-3$ ” charge state (injection energy = 40 mV)
 - From ^{13}C isotope distributions, this is a monomer-dominant peak.

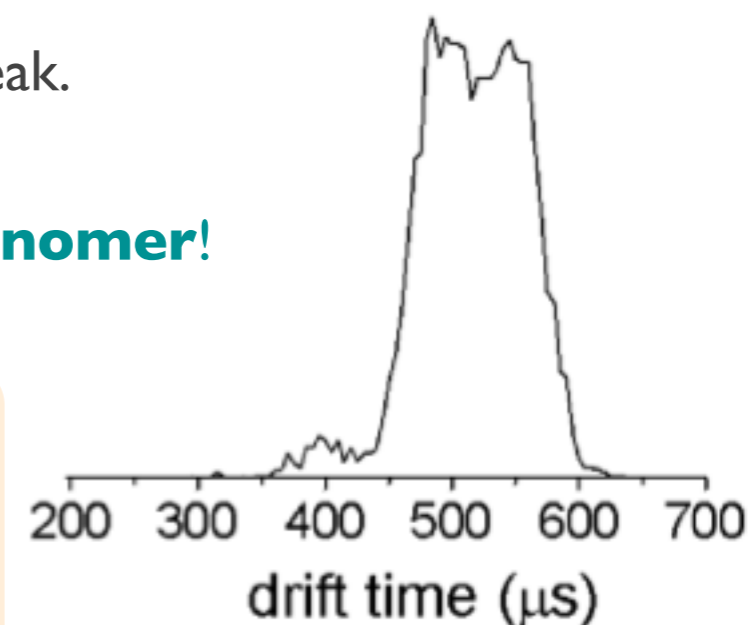


These two peaks should be predominantly **monomer!**

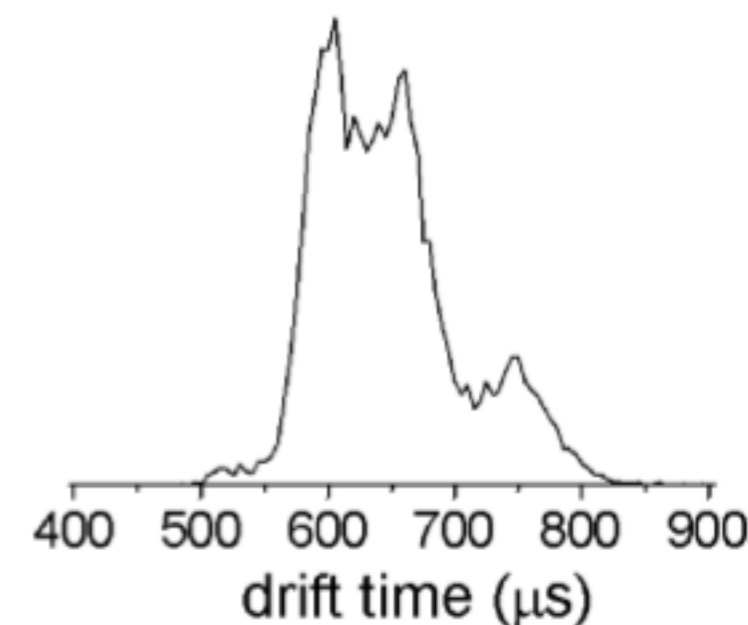
<caution>

Strictly speaking, to confirm these two peaks are monomer, we need to **calculate the CCS from drift times** and **compare them with the theoretical CCS obtained from modeling** (data not shown).

m/z 1487 (-3)



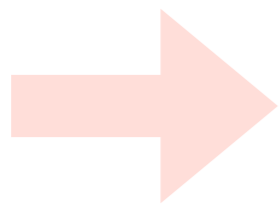
m/z 2231 (-2)



- ATD of “ $z/n=-2$ ” charge state (injection energy = 40 mV)
 - From ^{13}C isotope distributions, this is a mixture of oligomers and in fact, ATDs shows several peaks.

Assignment of Peaks in ATD (2)

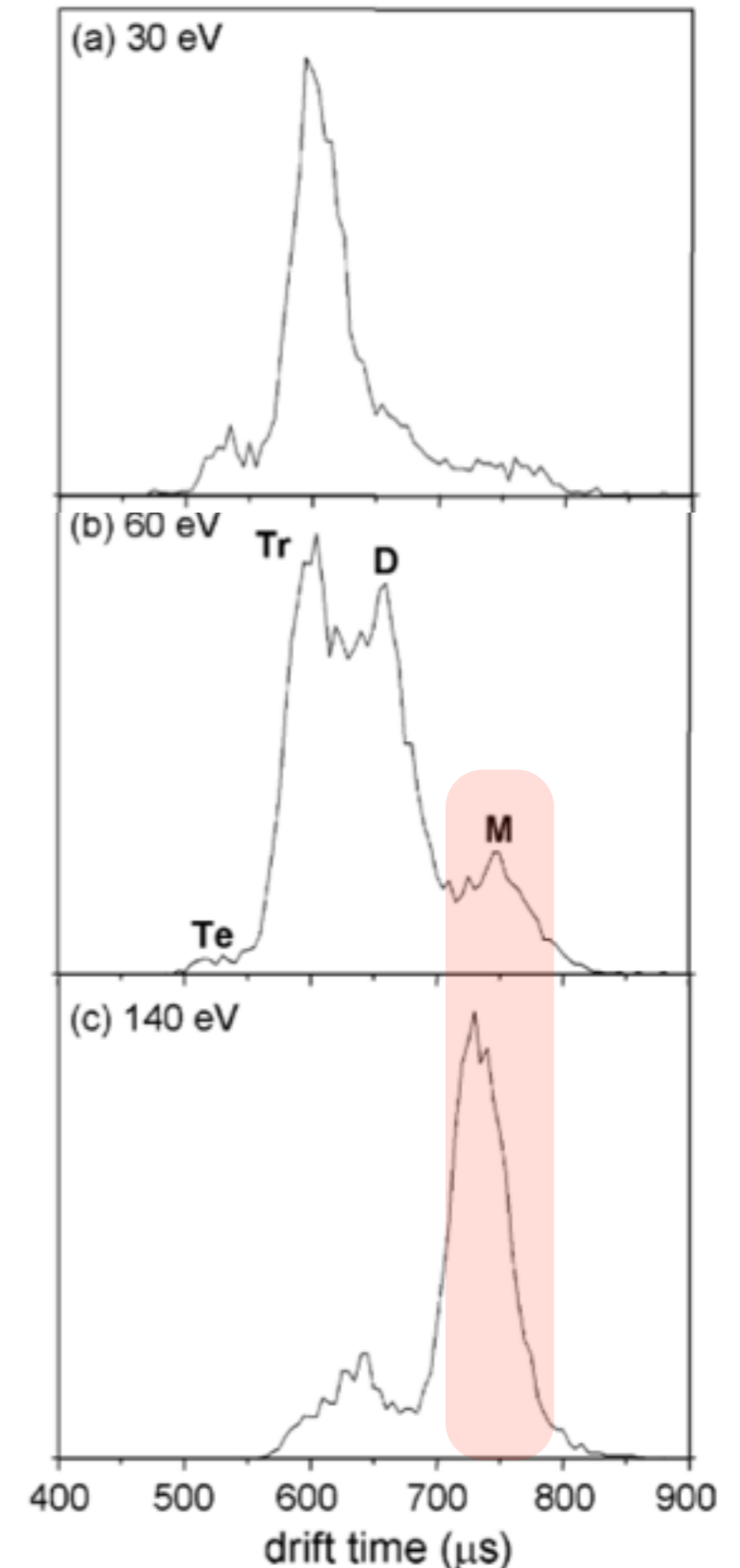
- injection energy dependence of the “ $z/n=-2$ ” charge state
 - As the injection energy increases, the peak at 740 μs becomes dominant and **no additional peaks were observed at longer drift times.**
 - Injection with high energy gives the ion higher energy, thus dissociation or conformation change occurs.



740 μs peak would be the **monomer**.

- Other peaks are assigned as **dimer**, **trimer** and **tetramer**.

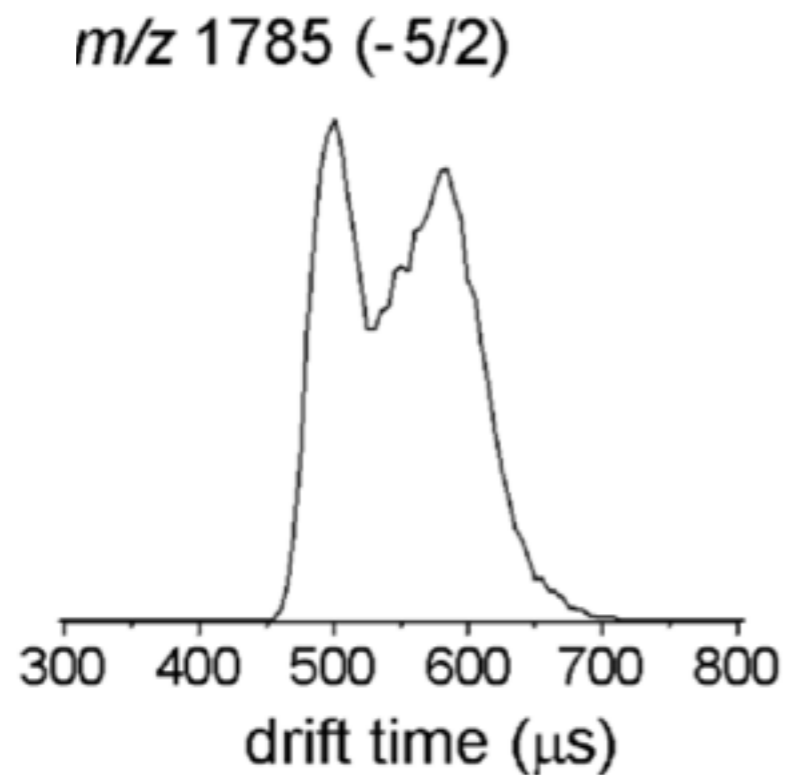
(This assignment is reasonable because CID-conducted ^{13}C isotope distributions shows the existence of dimer. Also, native ^{13}C isotope distribution clearly shows the existence of larger oligomer than dimer.)



S. Bernstein, M. Bowers *et al.*,
J. Am. Chem. Soc. **2005**, 127, 2075.

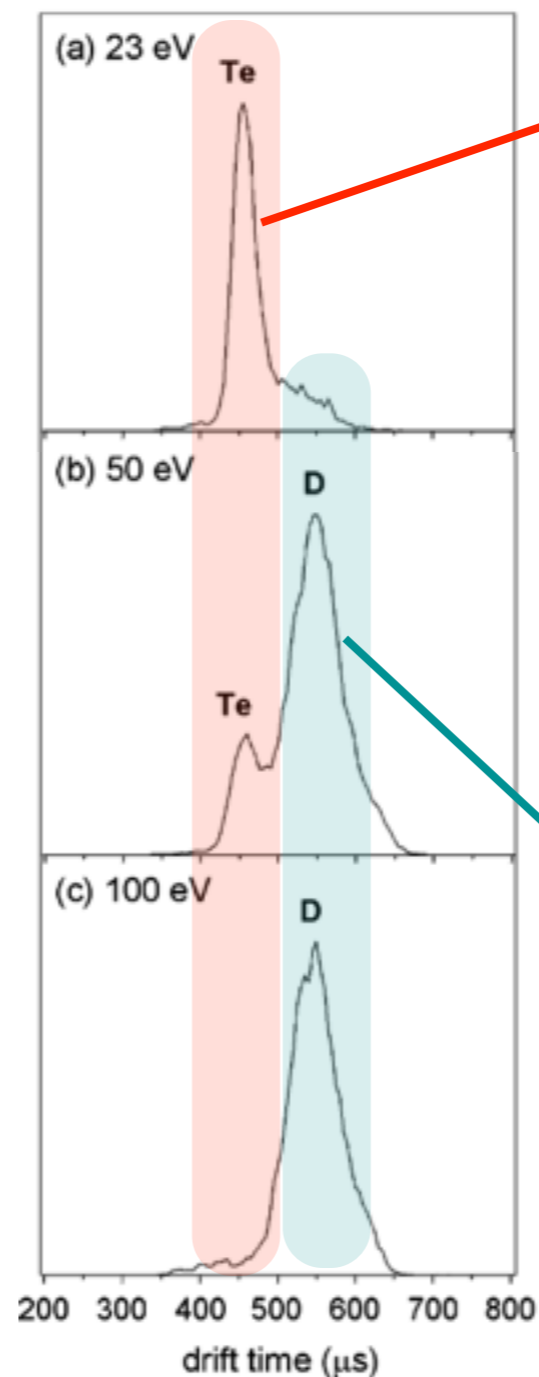
Assignment of Peaks in ATD (3)

- ATD of “ $z/n=-5/2$ ” charge state (injection energy = 40 mV)

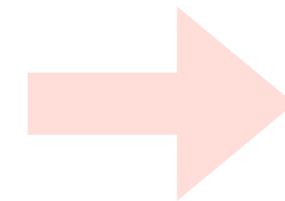


S. Bernstein, M. Bowers *et al.*,
J. Am. Chem. Soc. **2005**, 127, 2075.

- injection energy dependence

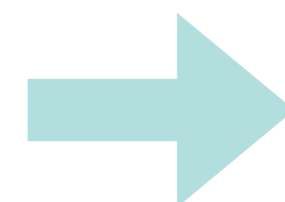


- Trimers can't be involved.
- bigger than dimer



tetramer

- Monomers can't be involved.
- No peak with longer drift time was observed.

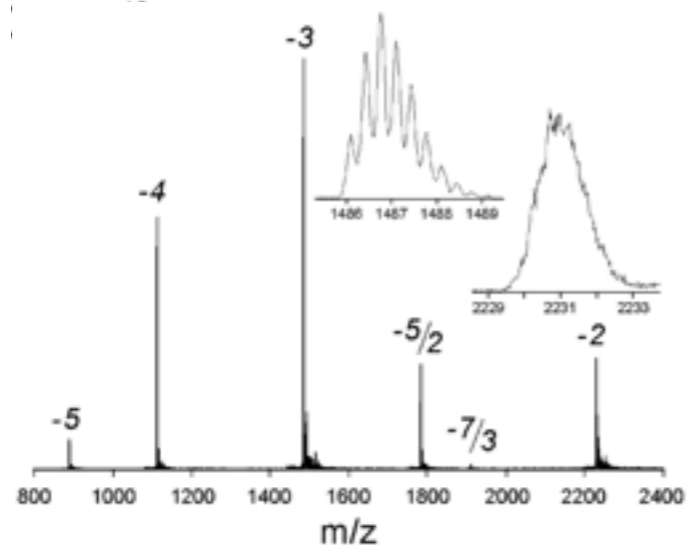


dimer

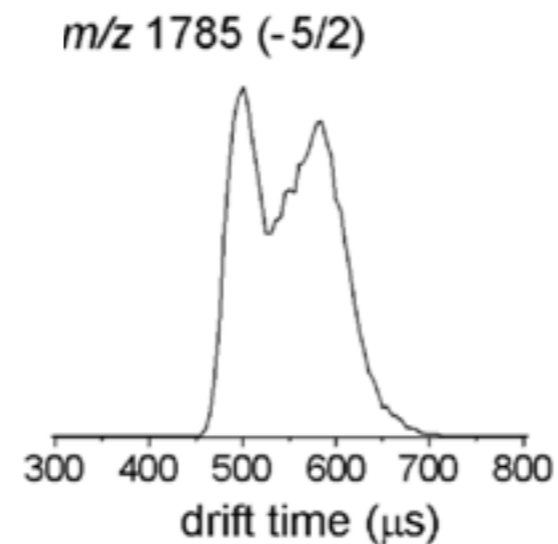
Short Summary

- flow chart of IM-MS analysis

1) analyze the **MS chart**



2) analyze the **ATD** of z/n which you are interested in



Structural Information

- ^{13}C isotope distributions
- Collision-Induced Dissociation

- injection energy dependence
- **CCS calculation and comparison of it with theoretical CCS** obtained from modeling.

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Subject of this Section

ARTICLES

PUBLISHED ONLINE: 14 JUNE 2009 | DOI: 10.1038/NCHEM.247

nature
chemistry

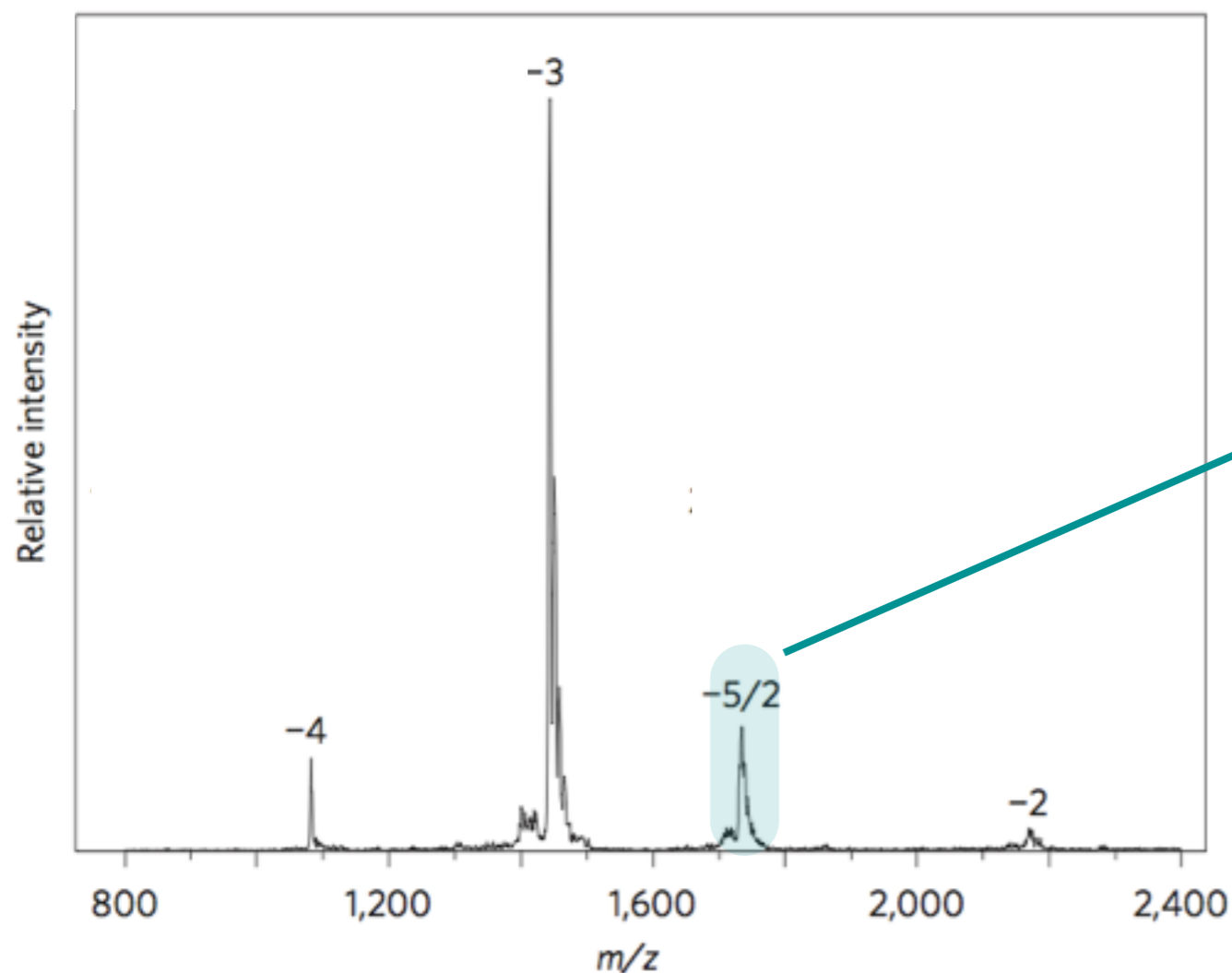
Amyloid- β protein oligomerization and the importance of tetramers and dodecamers in the aetiology of Alzheimer's disease

Summer L. Bernstein¹, Nicholas F. Dupuis¹, Noel D. Lazo², Thomas Wytttenbach¹, Margaret M. Condrón³, Gal Bitan³, David B. Teplow^{3,4}, Joan-Emma Shea¹, Brandon T. Ruotolo⁵, Carol V. Robinson⁵ and Michael T. Bowers^{1*}

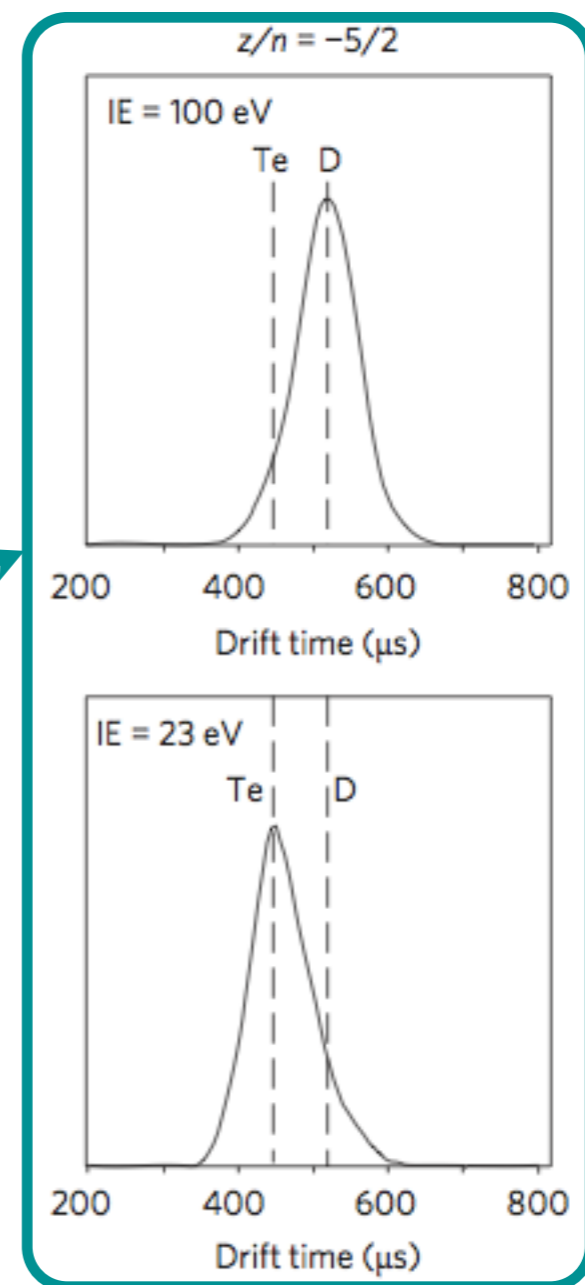
Nat. Chem. **2009**, *1*, 326.

IM-MS Analysis of A β 40

- mass spectrum and ATDs ($z/n = -5/2$)



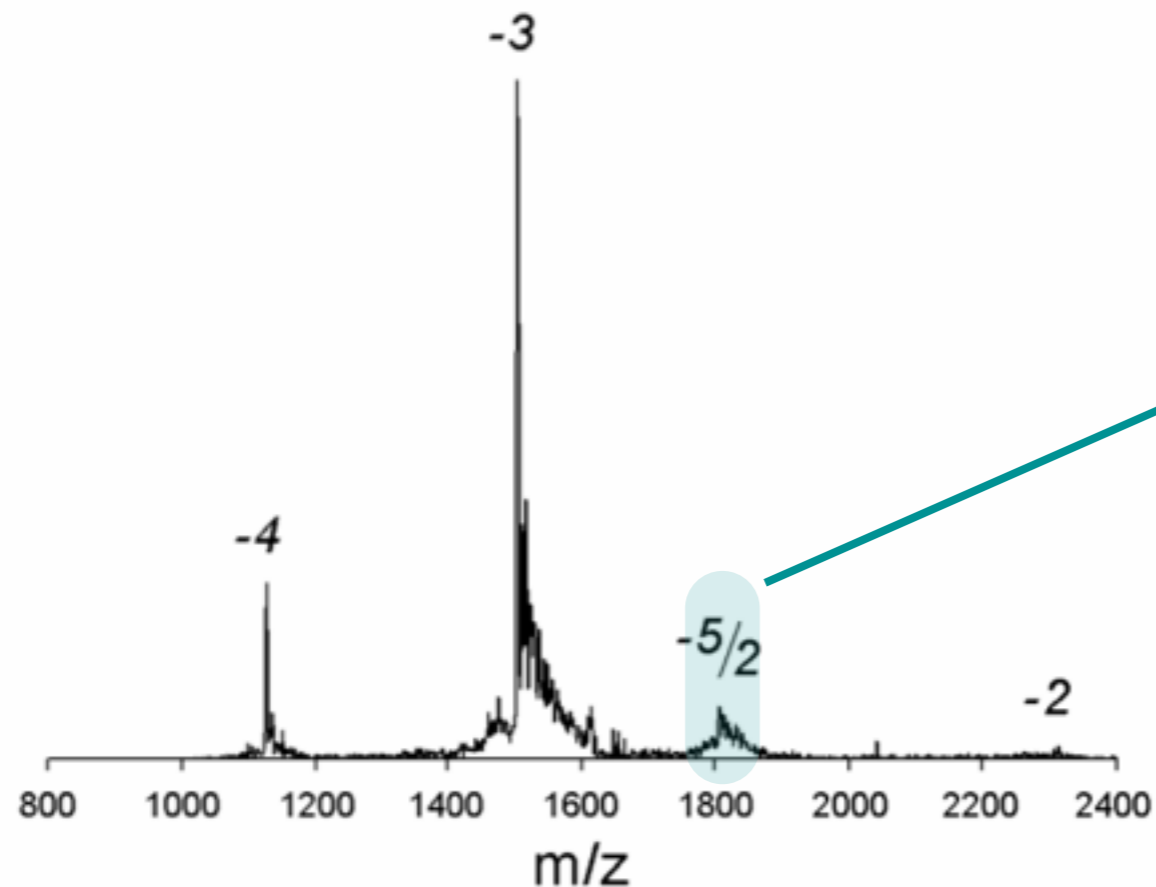
S. Bernstein, M. Bowers *et al.*, *Nat. Chem.* **2009**, *1*, 326.



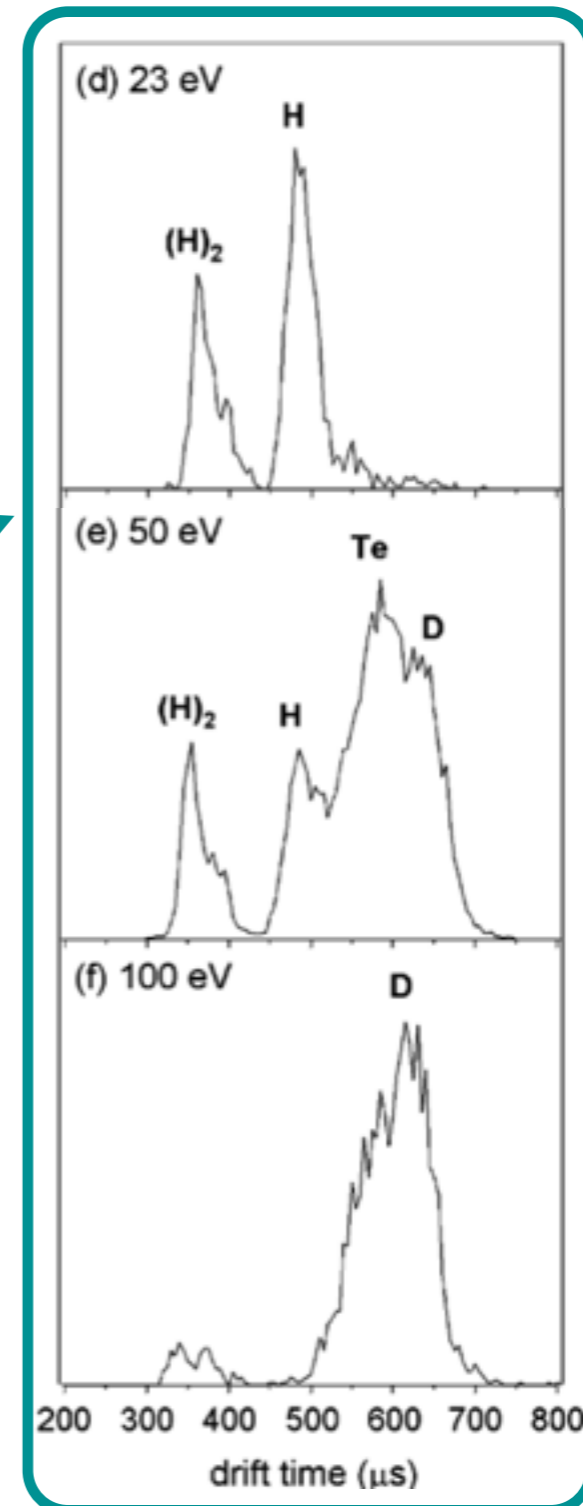
- As shown here, in “ $z/n=-5/2$ ” charge state, tetramer and dimer was observed.

IM-MS Analysis of A β 42

- mass spectrum and ATDs ($z/n = -5/2$)



- As shown here, in “ $z/n = -5/2$ ” charge state, dimer, tetramer, hexamer, and dodecamer (12-mer) was observed.



CCS Estimation of A β 42 Oligomer (I)

- Theoretical CCS for A β dimers were obtained as follows.

<assumption> A β monomer has a **spherical shape** (especially hard sphere)



Based on this, **theoretical CCS for the dimers were calculated** with the center-center distance of the two monomers as a variable parameter.





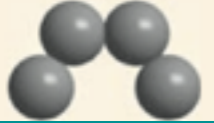


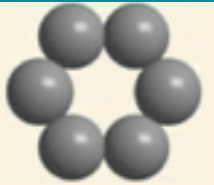

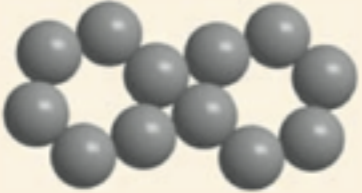
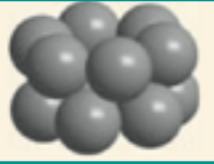
<**calibration**> Center-center distance was adjusted to give the experimental dimer CCS.



CCS of tetramer, hexamer, and dodecamer were calculated based on the theoretical dimer (**double hard sphere approximated dimer**).

CCS Estimation of A β 42 Oligomer (2)

- theoretical CCS and experimentally observed CCS for A β 42 oligomers

Oligomer	Structure [†]	Cross-section	
		Model	Experiment
Dimer*		1,256	1,256
Tetramer		2,358	
		2,277	2,332
		2,135	
Hexamer		3,450	
		3,100	2,898
		2,578	
Dodecamer		5,824	
		4,562	4,308

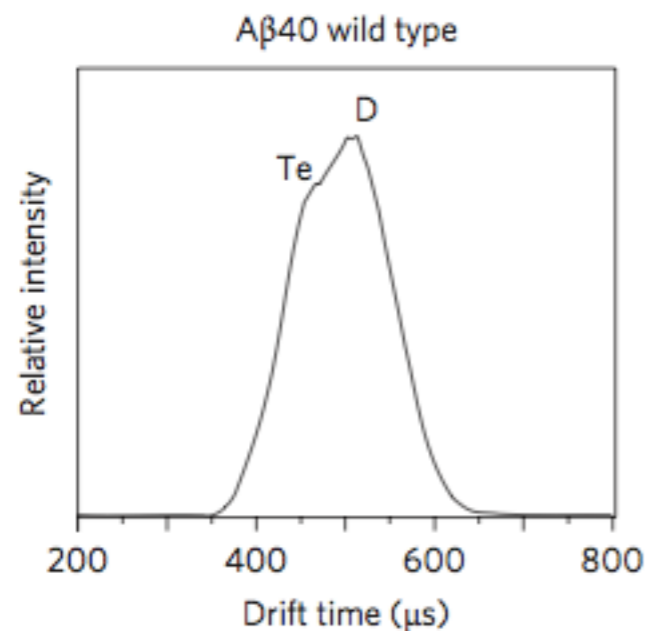
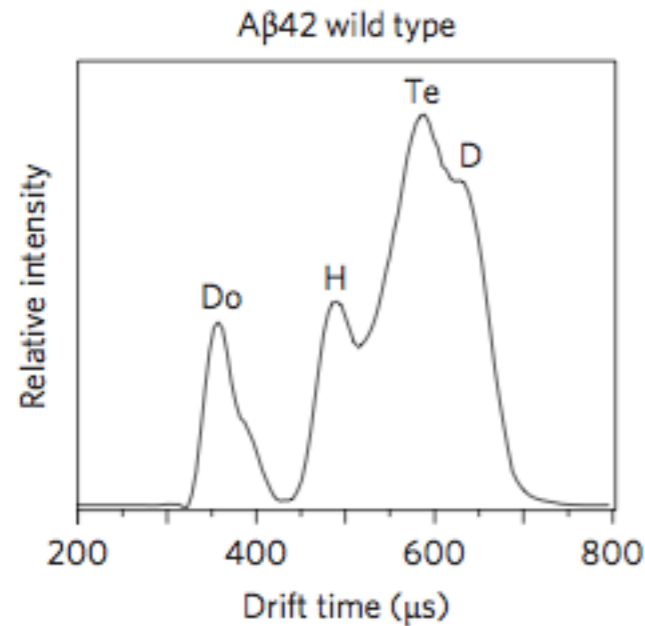
← Theoretical CCS was adjusted to observed CCS.

Model structure which had the **most close CCS value to experiment CCS** was successfully obtained.

*Model cross-section fit to experiment. [†]Limiting structures of high symmetry.

Comparison between A β 42 and A β 40 (I)

- ATDs of A β 42 and A β 40 for “z/n = -5/2” charge state



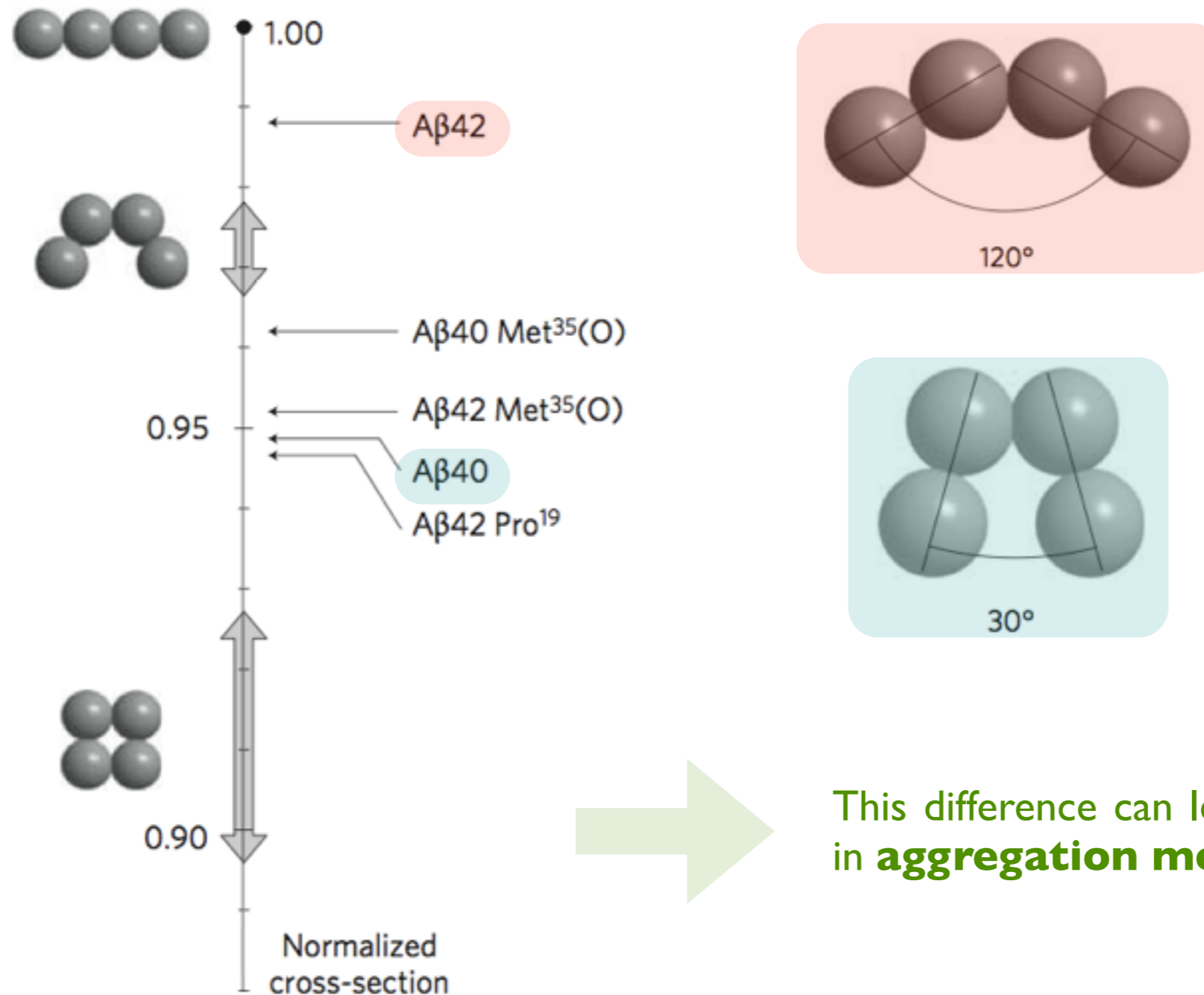
- In the case of A β 42, oligomers bigger than tetramer were observed.
- In contrast, in the case of A β 40, oligomers bigger than tetramer were not observed at all.



Tetramers can represent the difference between A β 40 and A β 42?

Comparison between A β 42 and A β 40 (2)

- comparison of tetramer conformation between A β 42 and A β 40

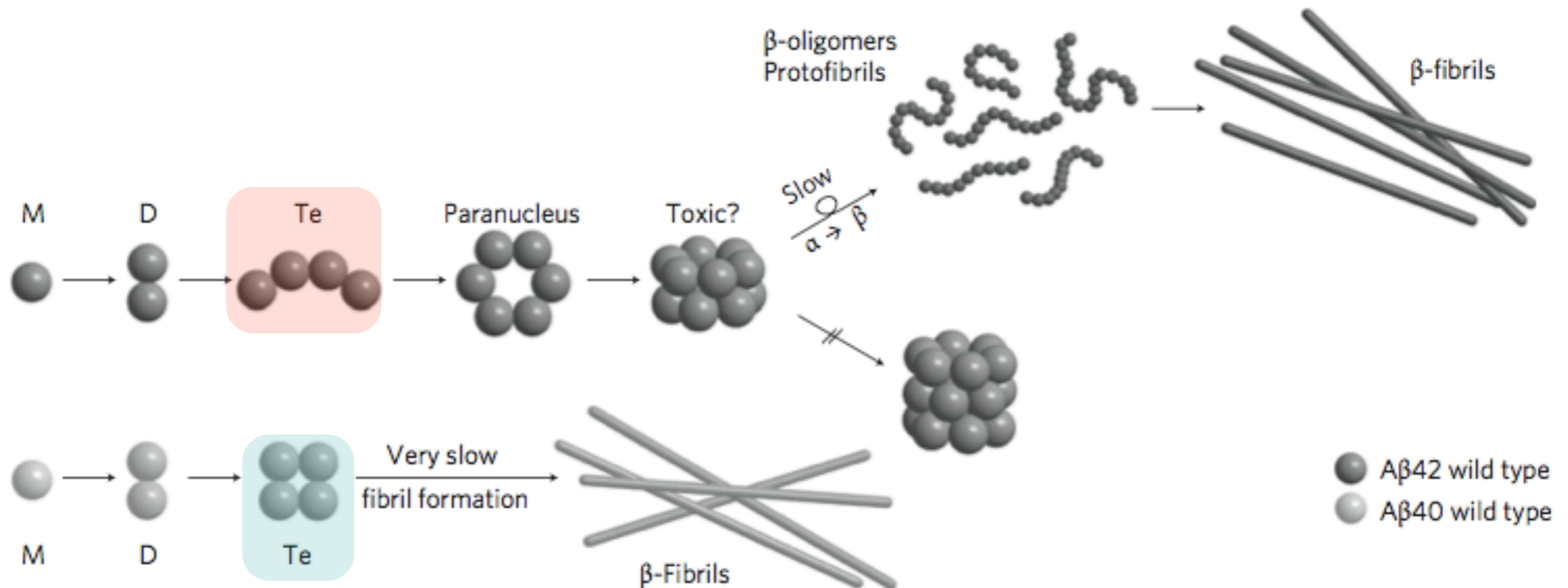


This difference can lead to the difference in **aggregation mechanism**.

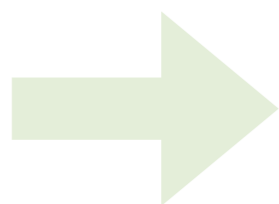
S. Bernstein, M. Bowers *et al.*, *Nat. Chem.* **2009**, *1*, 326.

Comparison between A β 42 and A β 40 (2)

- comparison of aggregation mechanism between A β 42 and A β 40



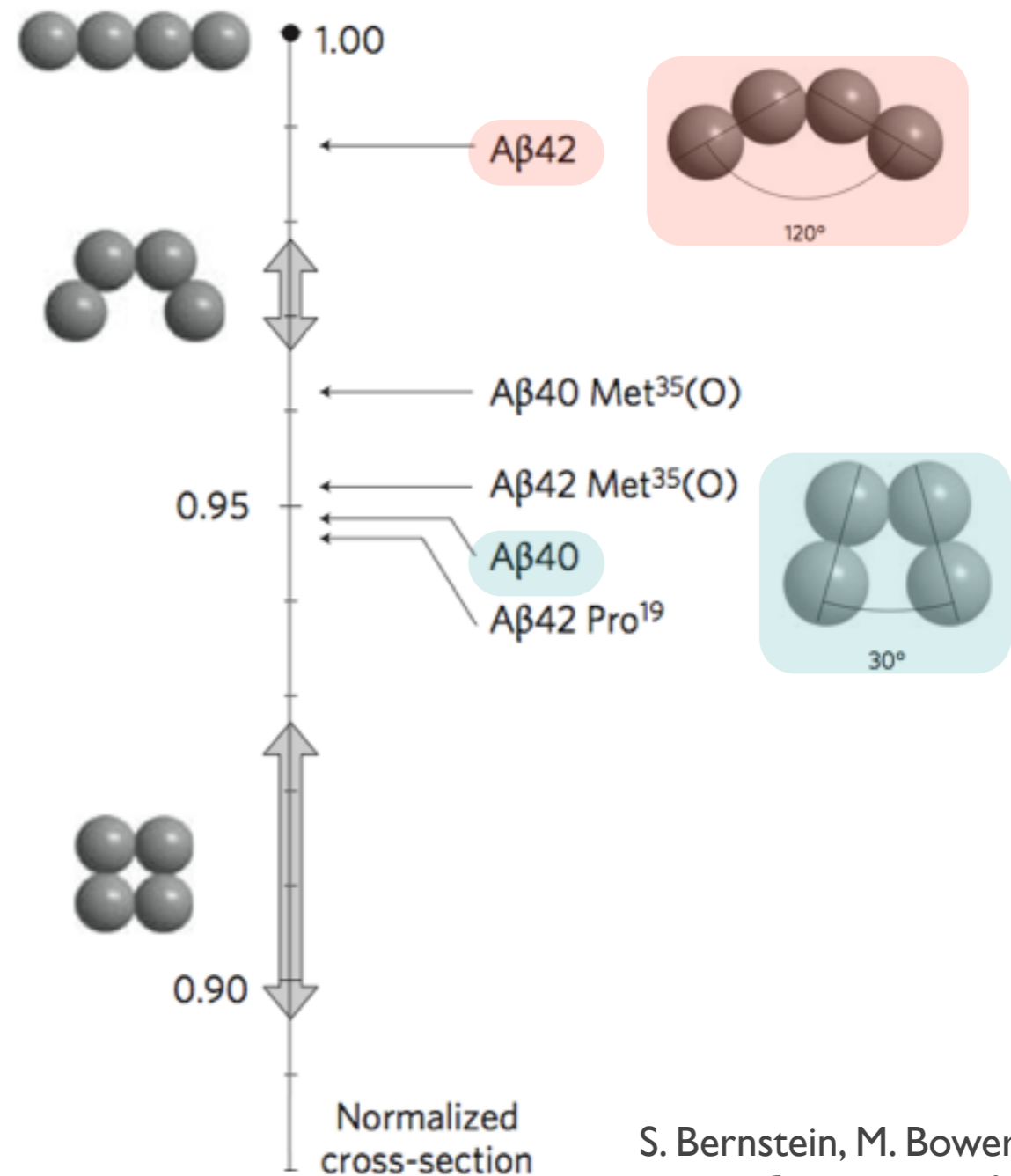
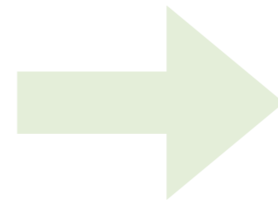
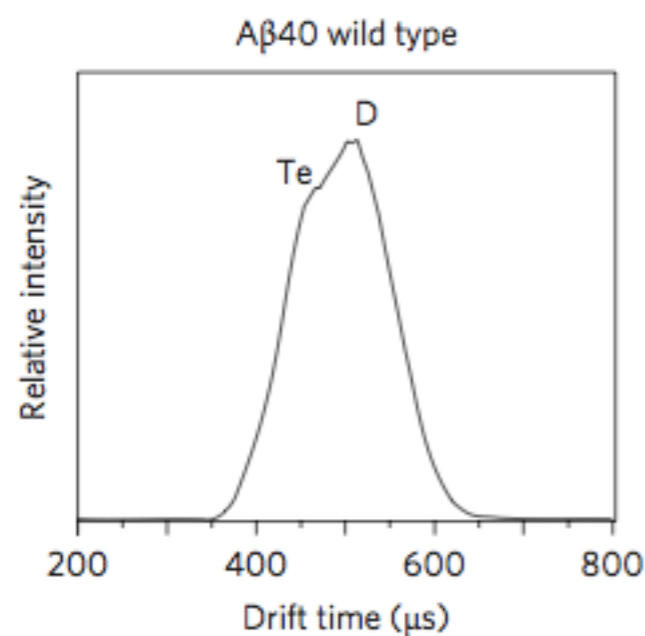
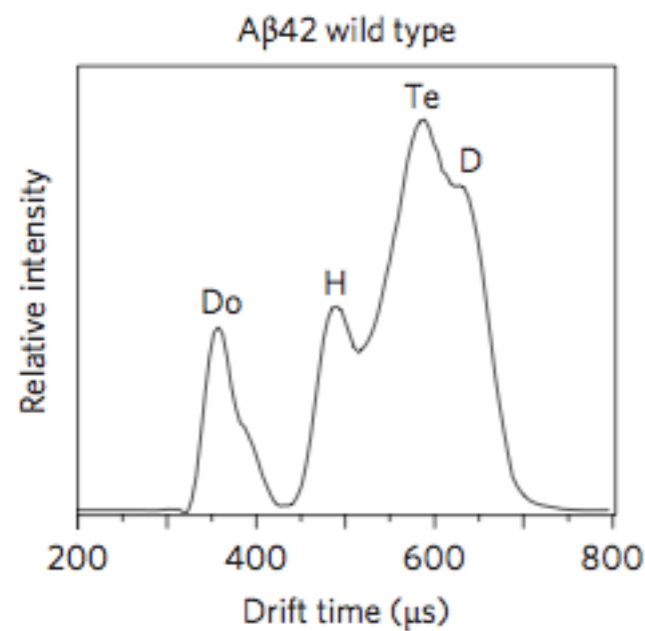
- A β 42 tetramer has a room of addition of another dimer to form hexamer or dodecamer. In contrast, A β 40 tetramer doesn't have a room of addition and doesn't form hexamer.



IM-MS could show the difference of aggregation mechanism between A β 42 and A β 40!

Short Summary

- Application of IM-MS achieves the mechanistic study of A β 42 and A β 40 aggregation based on the difference of tetramer conformation.



S. Bernstein, M. Bowers *et al.*,
Nat. Chem. **2009**, *1*, 326.

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§3.2 Obtainable Data and Their Explanation

§3.3 Oligomerization Analysis

§3.4 Mechanism Analysis of Aggregation Inhibitor

Subject of this Section



J | A | C | S
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Article

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Ion Mobility Spectrometry Reveals the Mechanism of Amyloid Formation of $A\beta(25-35)$ and Its Modulation by Inhibitors at the Molecular Level: Epigallocatechin Gallate and *Scyllo*-inositol

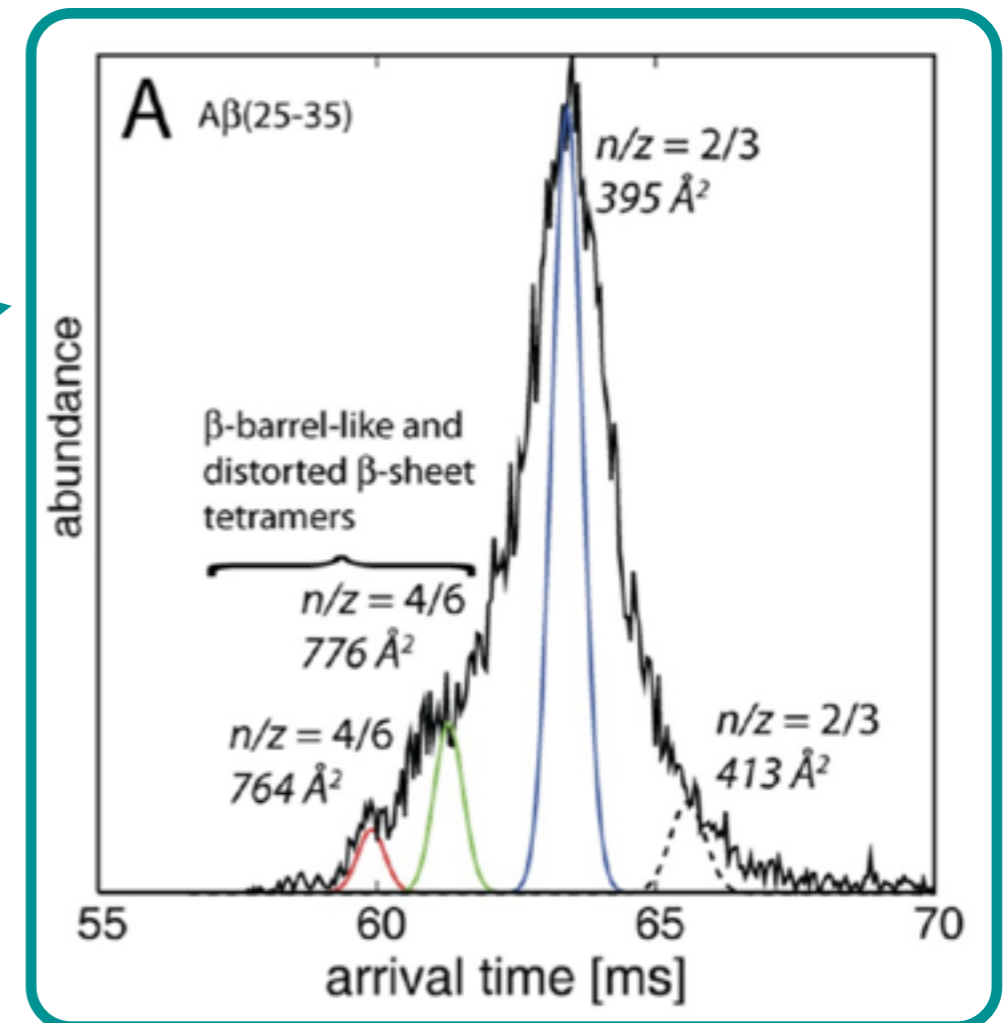
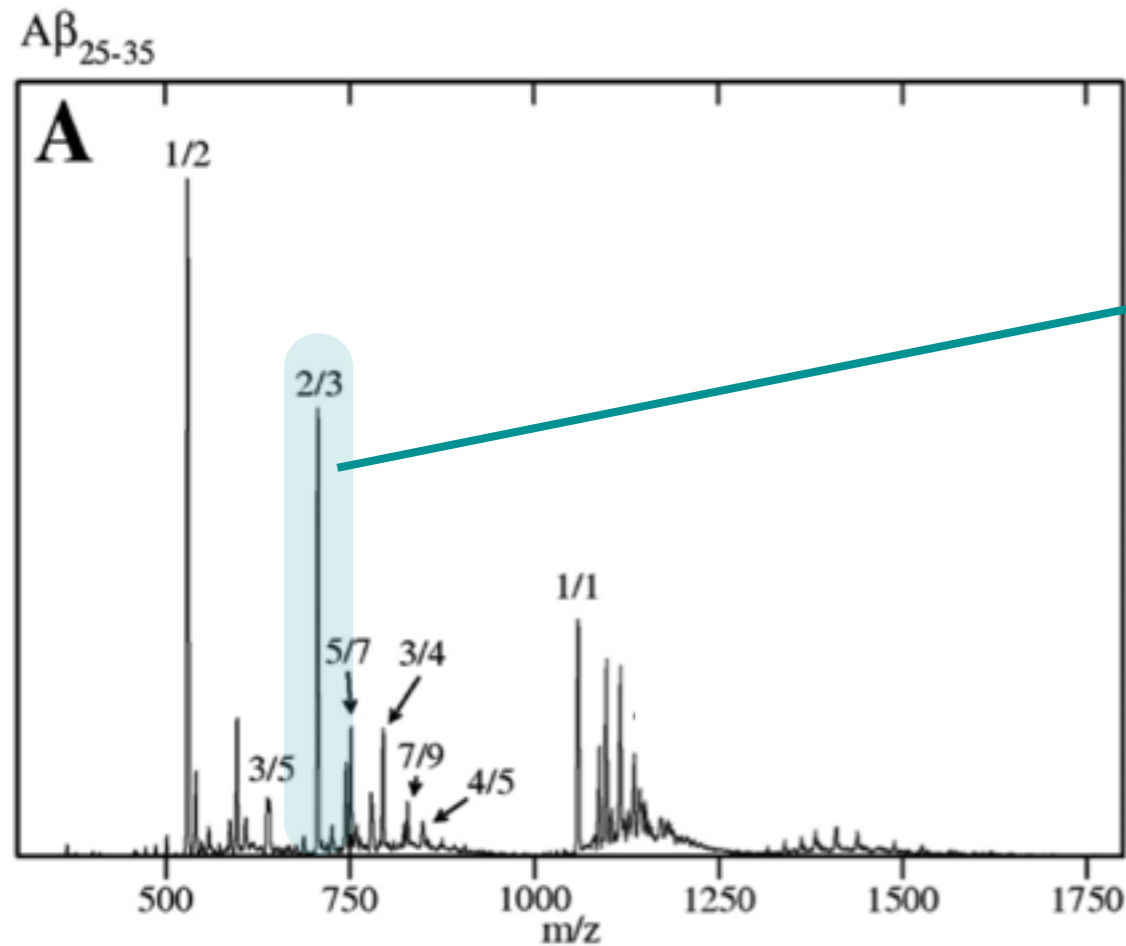
Christian Bleiholder,[†] Thanh D. Do,[†] Chun Wu, Nicholas J. Economou, Summer S. Bernstein, Steven K. Buratto, Joan-Emma Shea, and Michael T. Bowers*

Department of Chemistry and Biochemistry, University of California, Santa Barbara, California 93106-9510, United States

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IM-MS Analysis of A β (25-35)

- mass spectrum and ATD for $m/z = 707$ ($n/z = 2/3$)

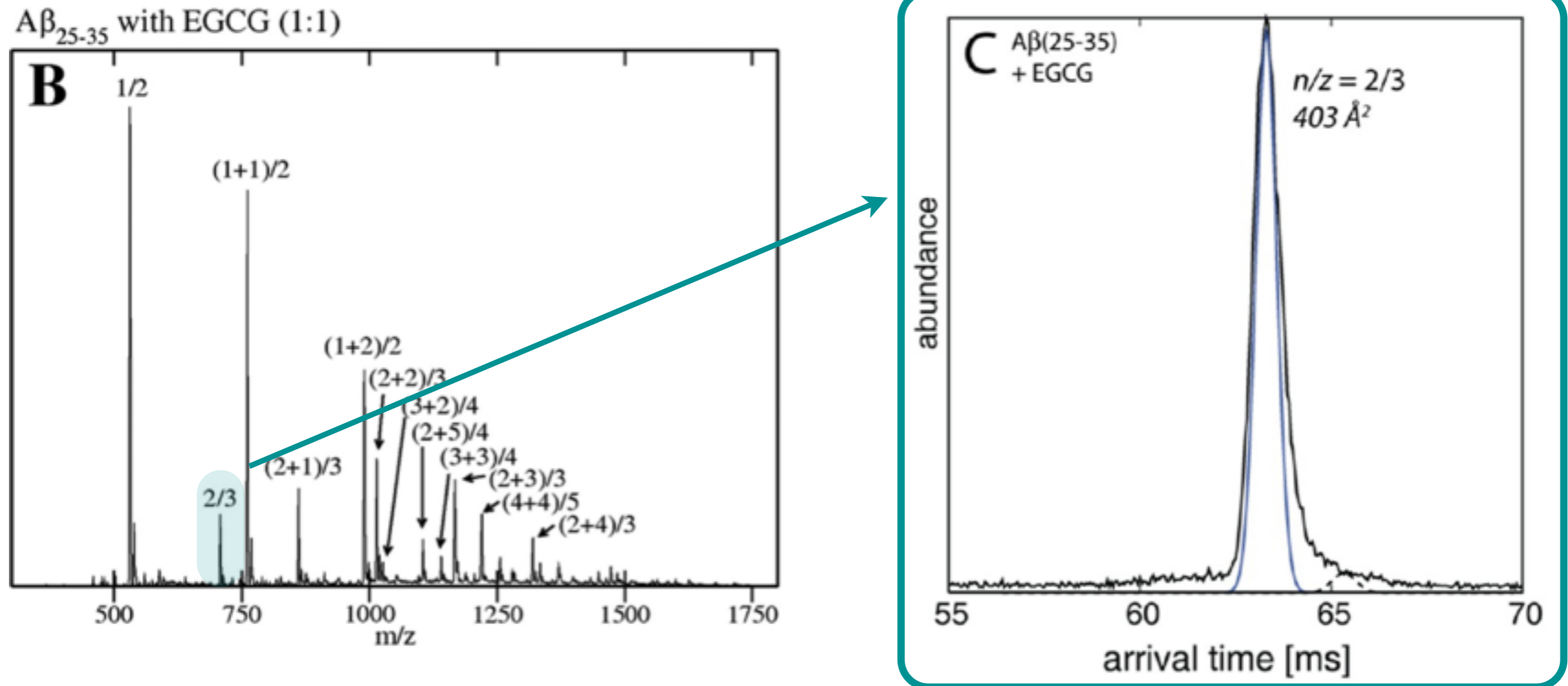


- Dimers and tetramers were observed.

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IM-MS Analysis of A β (25-35) with EGCG

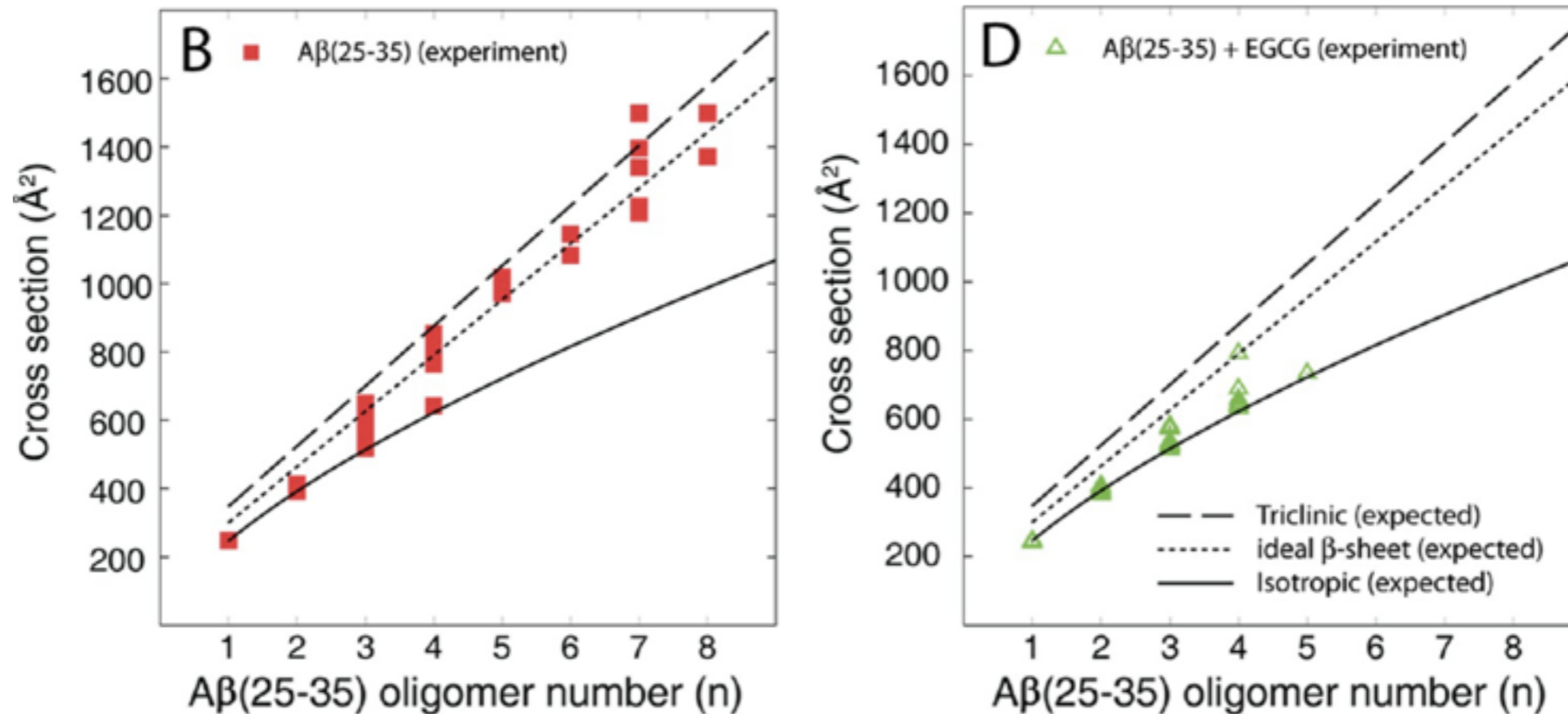
- mass spectrum and ATD for $m/z = 707$ ($n/z = 2/3$)



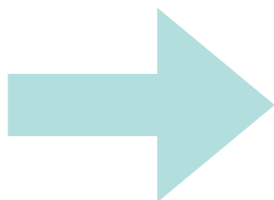
- Most peaks observed were hetero-oligomers of A β (25-35) and EGCG.
- **Tetramers were not observed.**

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Oligomer Growth Model (I)



- Correlation with linear line means that observed oligomer had a β -sheet structure.
 - triclinic: ideal out-of-register β -sheet (each monomer chain is shifted by one amino acid residue)
 - isotropic: ideal homolytic growth (oligomer grows equally in all spatial dimensions)

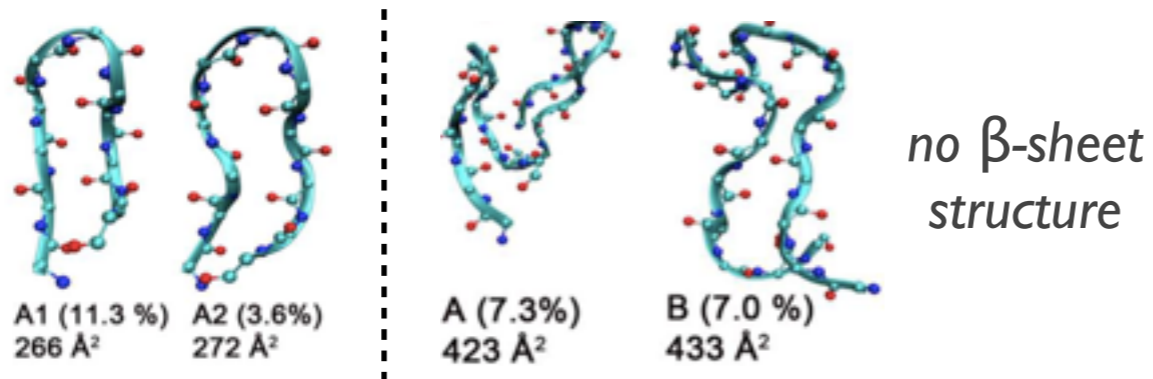


EGCG inhibited the β -sheet conformation appearance.

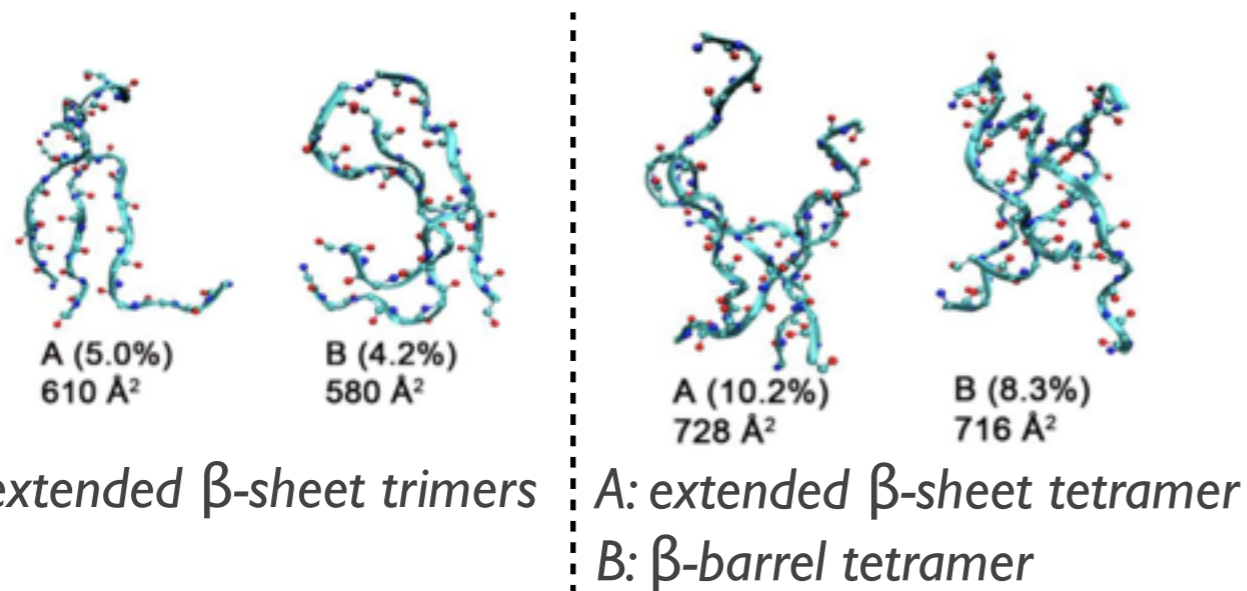
Oligomer Growth Model (2)

- As A β (25-35) oligomer grows, β -sheet tendency becomes dominant.

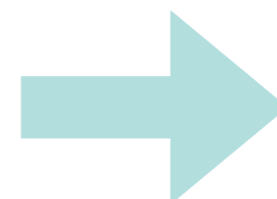
- monomer and dimer: isotropic



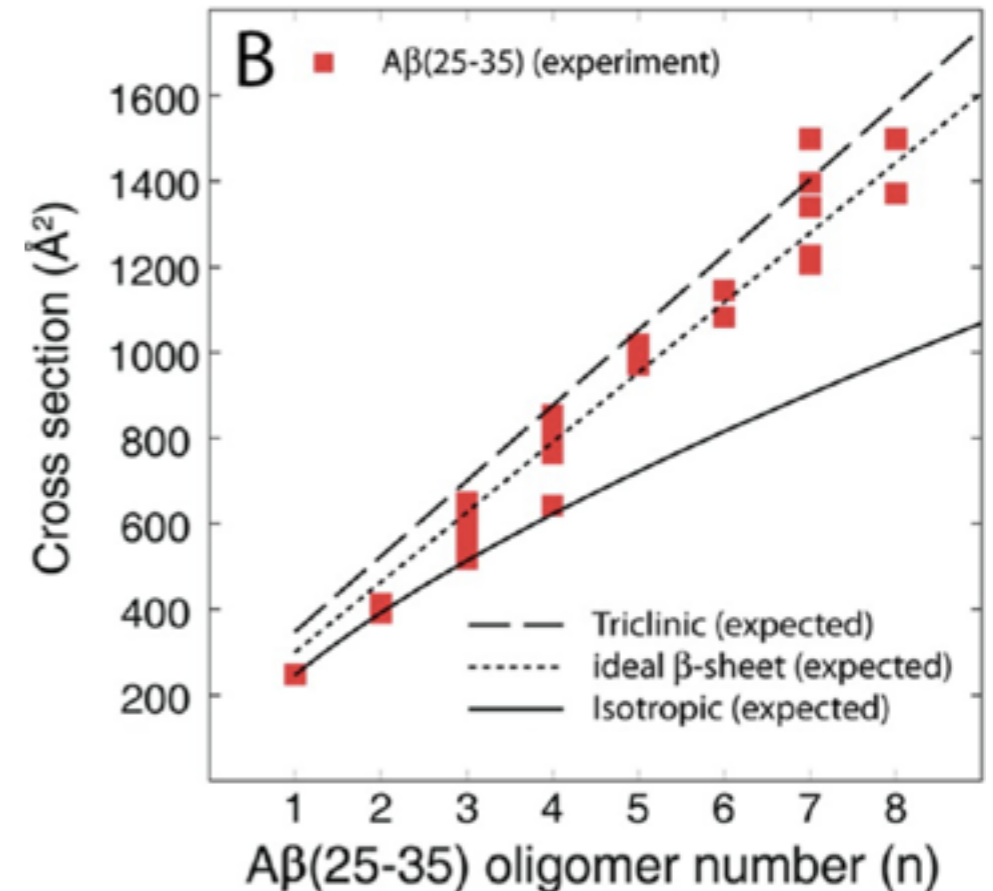
- trimer and tetramer: increasing β -sheet tendency



- bigger than pentamer: β -sheet



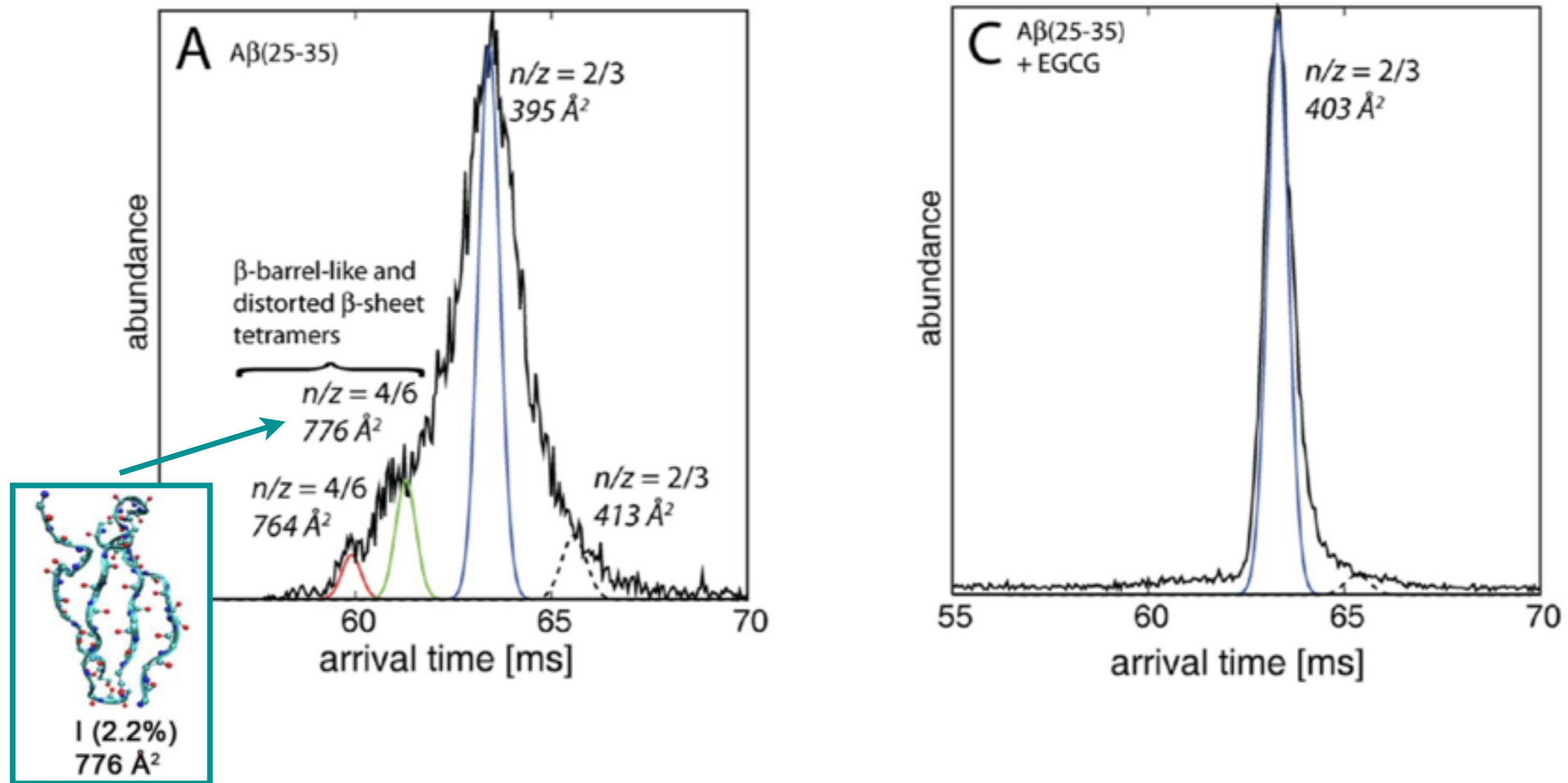
Trimer or tetramer supposed to be essential for the structural conversion to β -sheet.



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ATDs Comparison

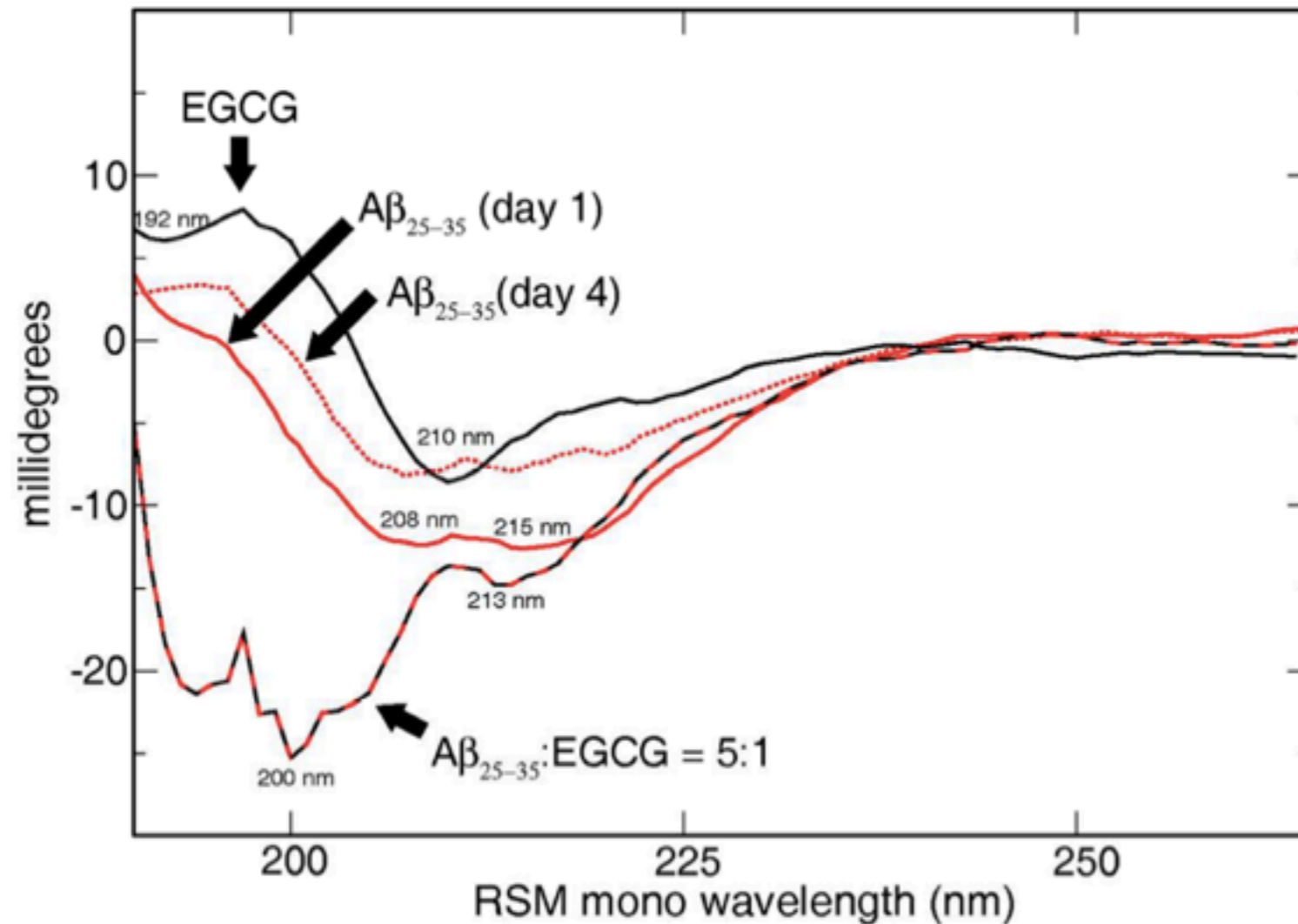
- ATDs for $m/z = 707$ ($n/z = 2/3$)



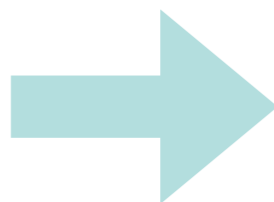
EGCG seemed to **inhibit the formation of β -barrel like tetramer or similar species.**

Circular Dichroism (CD)

- 0.5 mg/mL of A β (25-35) was incubated at r.t. with or without 10 mM EGCG (A β :EGCG=5:1).



- A β (25-35) spectrum indicated the presence of β -sheet structure.
- A β (25-35) + EGCG spectrum indicated the presence of random-coil structure.

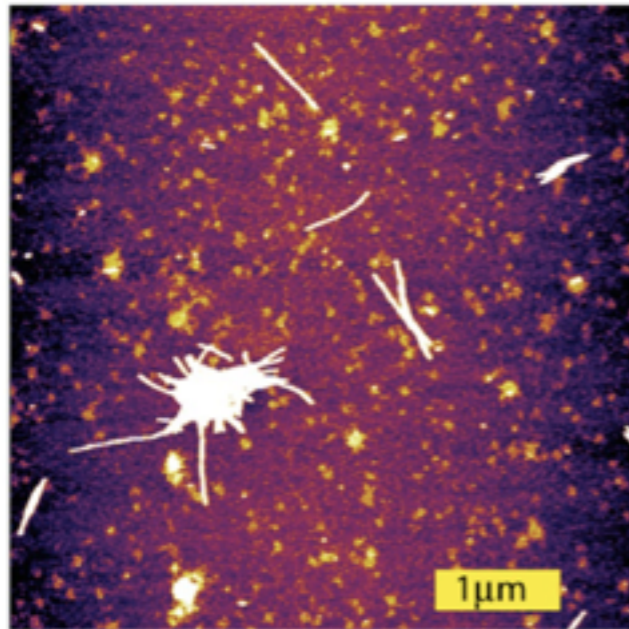


This result strongly support the previous IM-MS analysis.

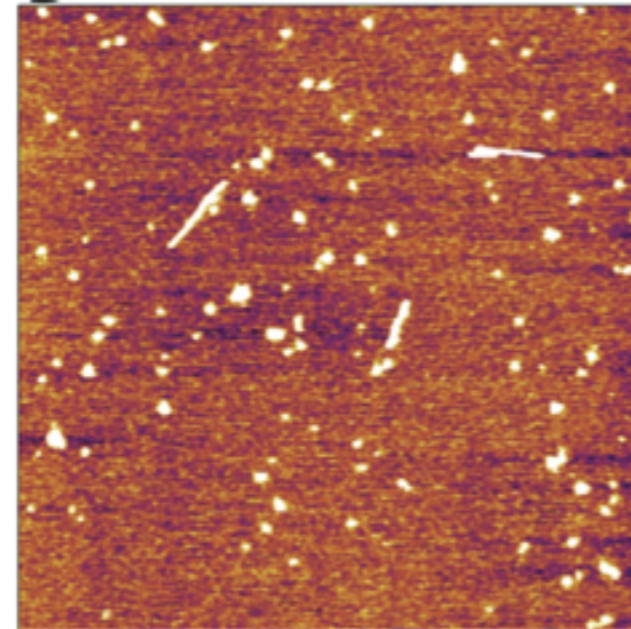
Atomic Force Microscopy (AFM)

- 200 μM of $\text{A}\beta(25-35)$ was incubated at r.t. with or without EGCG ($\text{A}\beta:\text{EGCG}=1:1$).

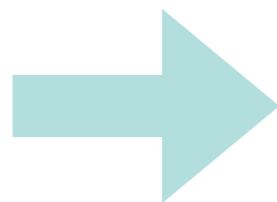
- without EGCG



- with EGCG



- Fibril formation was prevented using EGCG.

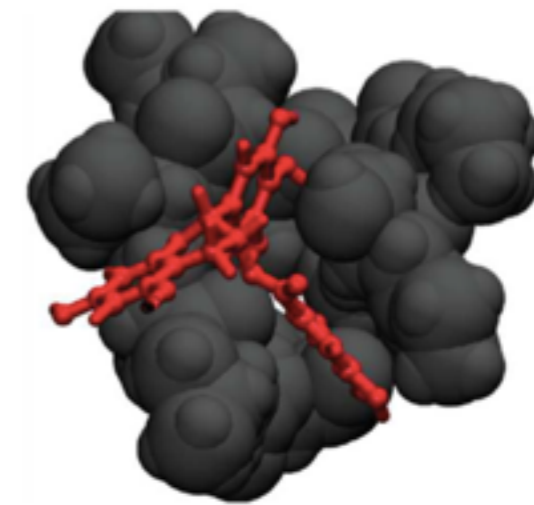
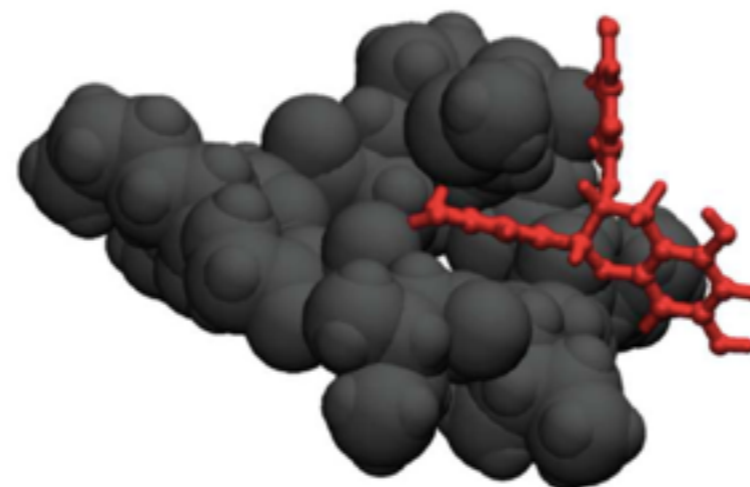
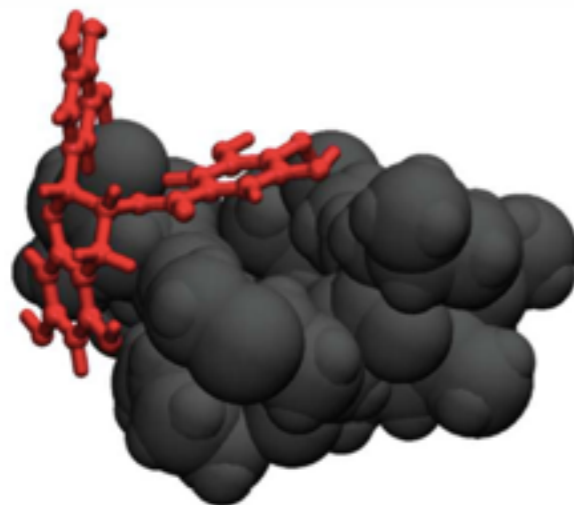
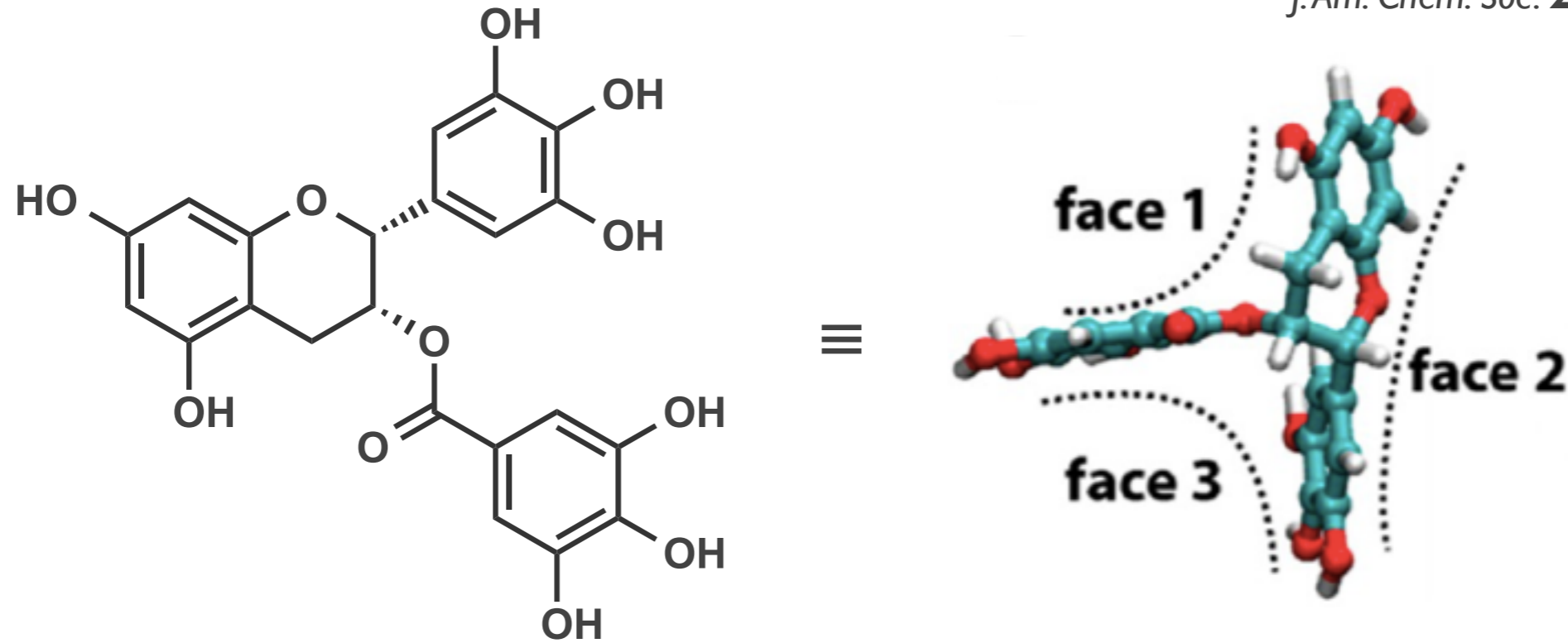


This result also support the previous IM-MS analysis.

Binding Simulation of EGCG (I)

- three binding sites of EGCG and binding simulation of each site

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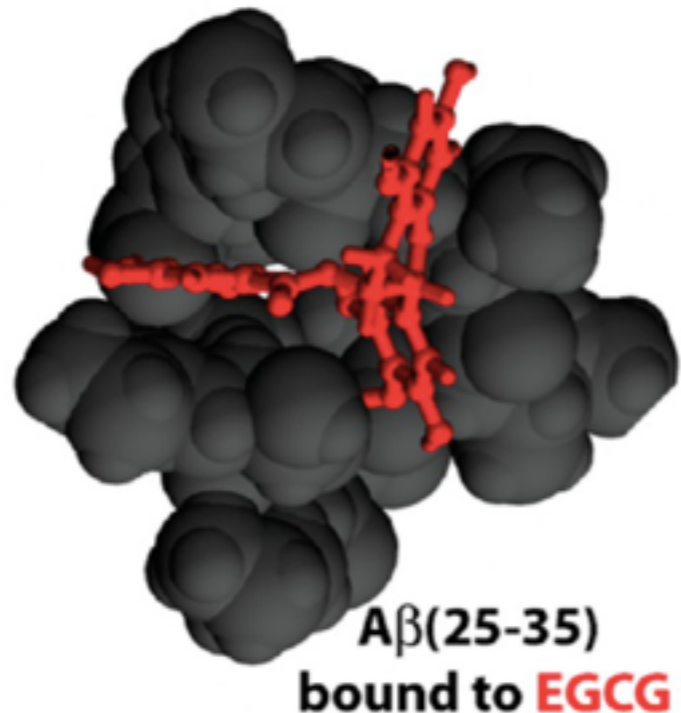
1

2

3

Binding Simulation of EGCG (2)

- A β (25-35) can bind to EGCG using all three binding sites.



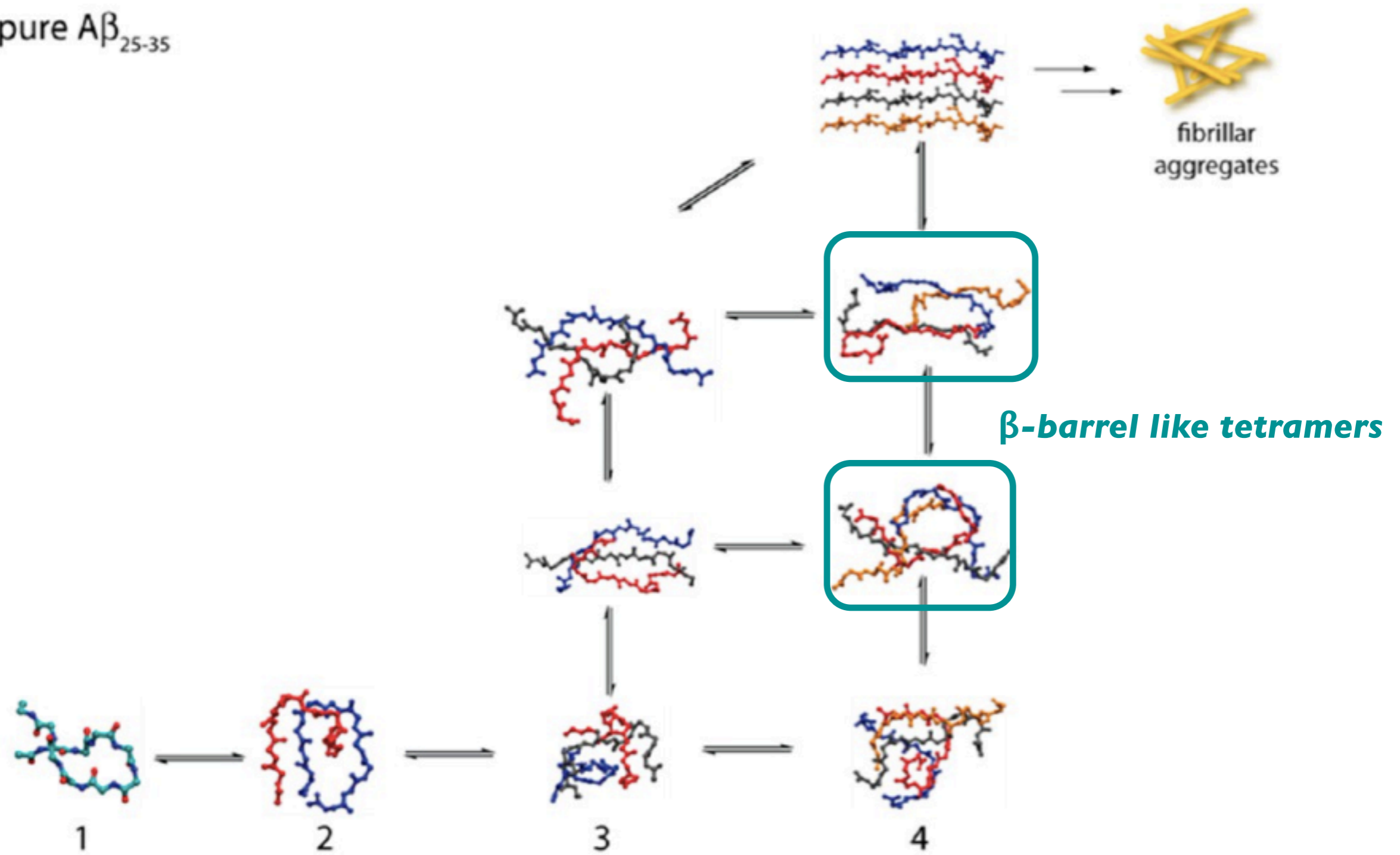
- This tridentate binding mode can only be accomplished when the bound A β chain takes on a unordered structure. (data was not shown in the paper...)

Formation of β -sheet which is observed in the aggregation of A β (25-35) **is prevented.**

This binding property might inhibit the generation of β -barrel like tetramer strongly and thus inhibit the formation of fibrils.

Short Summary: Aggregation Pathway of A β (25-35)

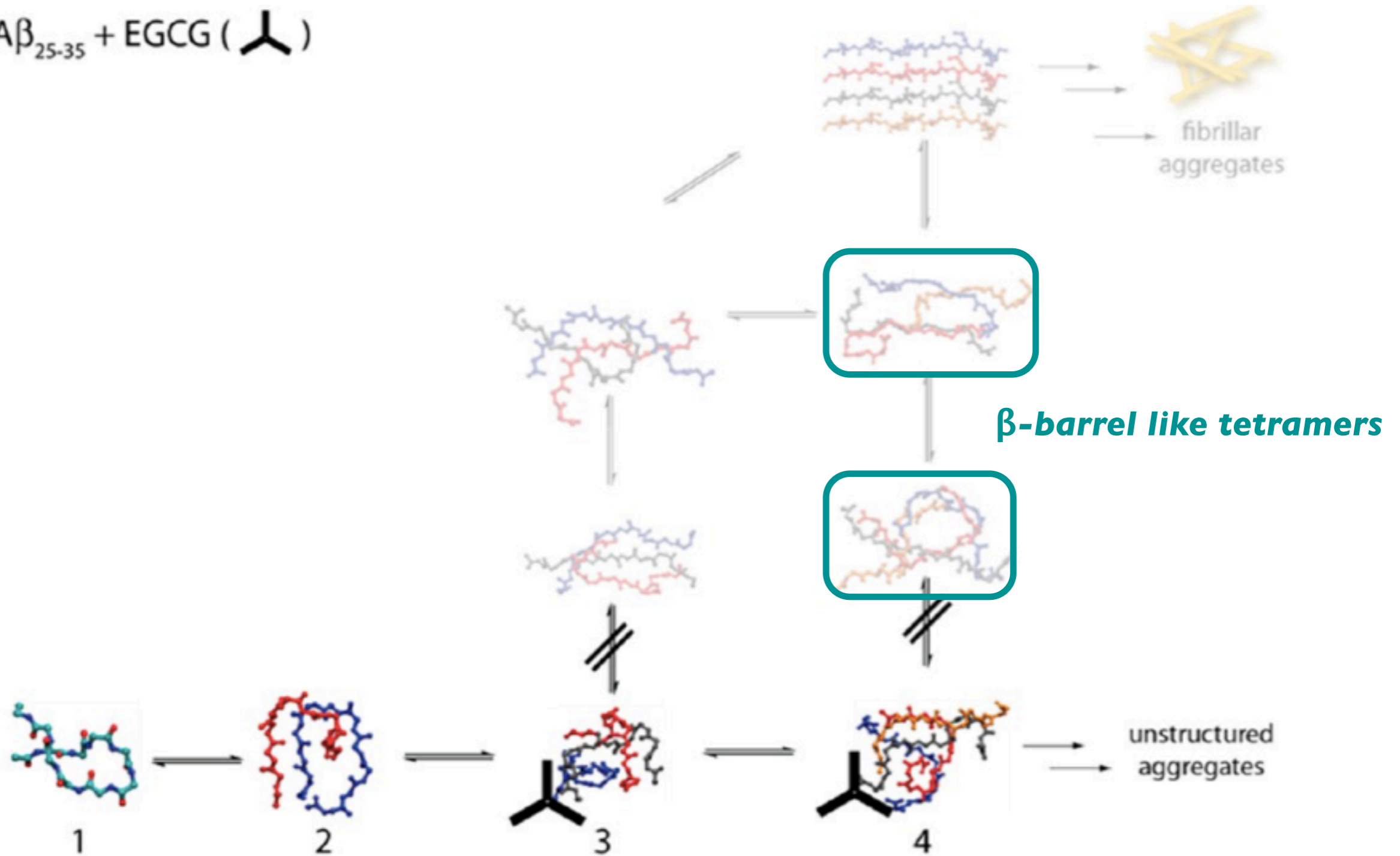
A) pure A β ₂₅₋₃₅



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Short Summary: Aggregate-Inhibiting Pathway of EGCG

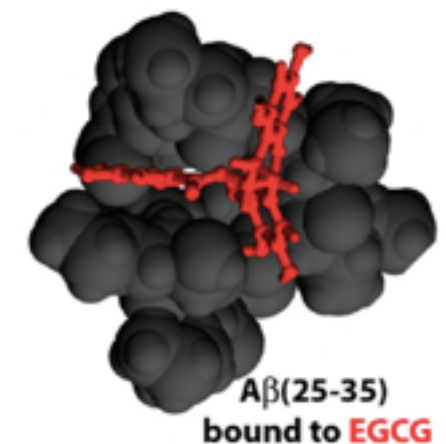
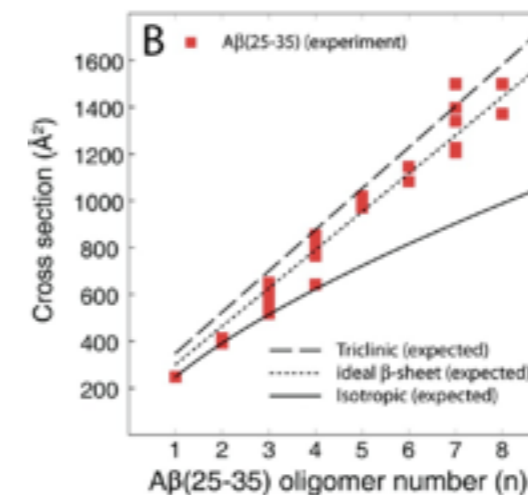
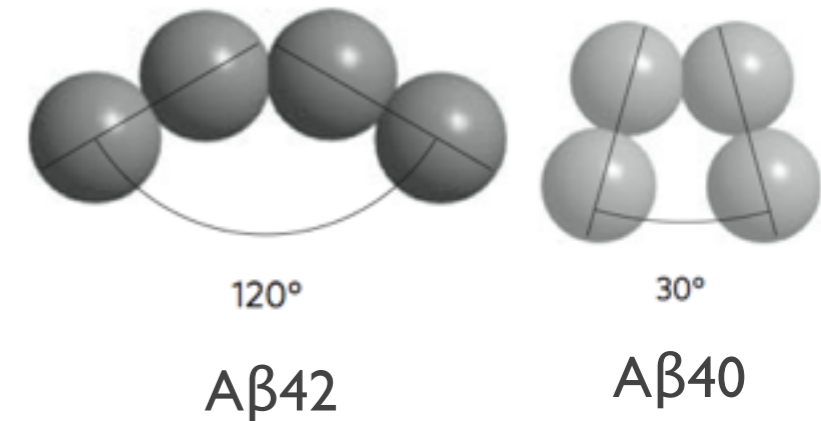
B) $A\beta_{25-35}$ + EGCG ()



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Summary

- application to A β 42 and A β 40 oligomerization analysis
 - Using appropriate approximation, structure difference between A β 42 tetramer and A β 40 tetramer was distinguished.
- application to mechanism analysis of aggregation inhibitor
 - Combining MD simulation and IM-MS analysis, advanced information of A β (25-35) aggregation and its inhibition by EGCG was obtained.



The most important ability of IM-MS to detect intermediates is successfully utilized!

Appendix

Derivation of Mason-Schamp Equation (I)

気相移動度測定においては、イオンの速度は $\frac{E}{N}$ に依存した一定の値をとる。特に、低電場極限(電場効果が熱運動に対し無視できるほど小さい)、すなわち

$$\frac{E}{N} \leq 2 \times 10^{-17} \text{ (V} \cdot \text{cm}^2) \quad (1)$$

が満たされているとき、イオンは等速運動するようになり、その速度は

$$v_d = KE \quad (2)$$

と書けることが知られている。

一方、イオンとガス分子との衝突時の運動量変化を解析すると、

$$v_d = qE\tau \left(\frac{1}{m} + \frac{1}{M} \right) \quad (3)$$

が得られる。ここで τ は、「イオンがバッファーガスに衝突する間の時間」であり、

$$\tau = \frac{1}{N \langle |\vec{v}_r| \rangle \Omega} \quad (4)$$

と書ける (分母 = 「単位時間でイオンが衝突するバッファーガス分子の数」)。

Derivation of Mason-Schamp Equation (2)

(4) 式を (3) 式に代入すると、下式が得られる。

$$v_d = \frac{qE}{N \langle |\vec{v}_r| \rangle \Omega} \left(\frac{1}{m} + \frac{1}{M} \right) \quad (5)$$

ただし $\langle |\vec{v}_r| \rangle$ は、「全イオンと全バッファーガスの相対速度の絶対値の平均値」である。

低電場極限においては、 $\langle |\vec{v}_r| \rangle$ に対し熱運動近似を施すことができ、(5) 式は

$$v_d = \frac{1}{\sqrt{3}} \frac{qE}{N} \left(\frac{1}{m} + \frac{1}{M} \right)^{\frac{1}{2}} \left(\frac{1}{k_B T} \right)^{\frac{1}{2}} \frac{1}{\Omega} \quad (6)$$

と書き直すことができる。近似せずより厳密に取り扱くと、

$$v_d = \frac{3}{16} \frac{qE}{N} \left(\frac{1}{m} + \frac{1}{M} \right)^{\frac{1}{2}} \left(\frac{2\pi}{k_B T} \right)^{\frac{1}{2}} \frac{1}{\Omega} \quad (7)$$

となるので、(2) 式とあわせて

$$K = \frac{v_d}{E} = \frac{3}{16} \frac{q}{N} \left(\frac{1}{m} + \frac{1}{M} \right)^{\frac{1}{2}} \left(\frac{2\pi}{k_B T} \right)^{\frac{1}{2}} \frac{1}{\Omega} \quad (8)$$

が得られる (Mason-Schamp equation)。

To Achieve High Resolving Power in DTIMS

半値幅分解能 (近接したピークを高さの半分の高さ位置で分離できること) は

$$R = \frac{1}{4} \sqrt{\frac{q}{k_B \ln 2}} \sqrt{\frac{V}{T}} \quad (9)$$

で表される。(9) 式は, 定数項を考えなければ

$$R \propto \sqrt{\frac{V}{T}}$$









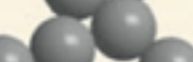

となり, 分解能は「電位とバッファーガス温度の比に依存する」といえる。

ゆえに, 高分解能を実現するためには, 電位を高く設定すればよいが, 高電位下では放電が起こる可能性がある。これを防ぐため, DTIMSではバッファーガス圧力が高く設定されている。なお, この圧力の高さは, 結果的に (8) 式成立の前提である

$$\frac{E}{N} \leq 2 \times 10^{-17} \text{ (V} \cdot \text{cm}^2) \quad (1)$$

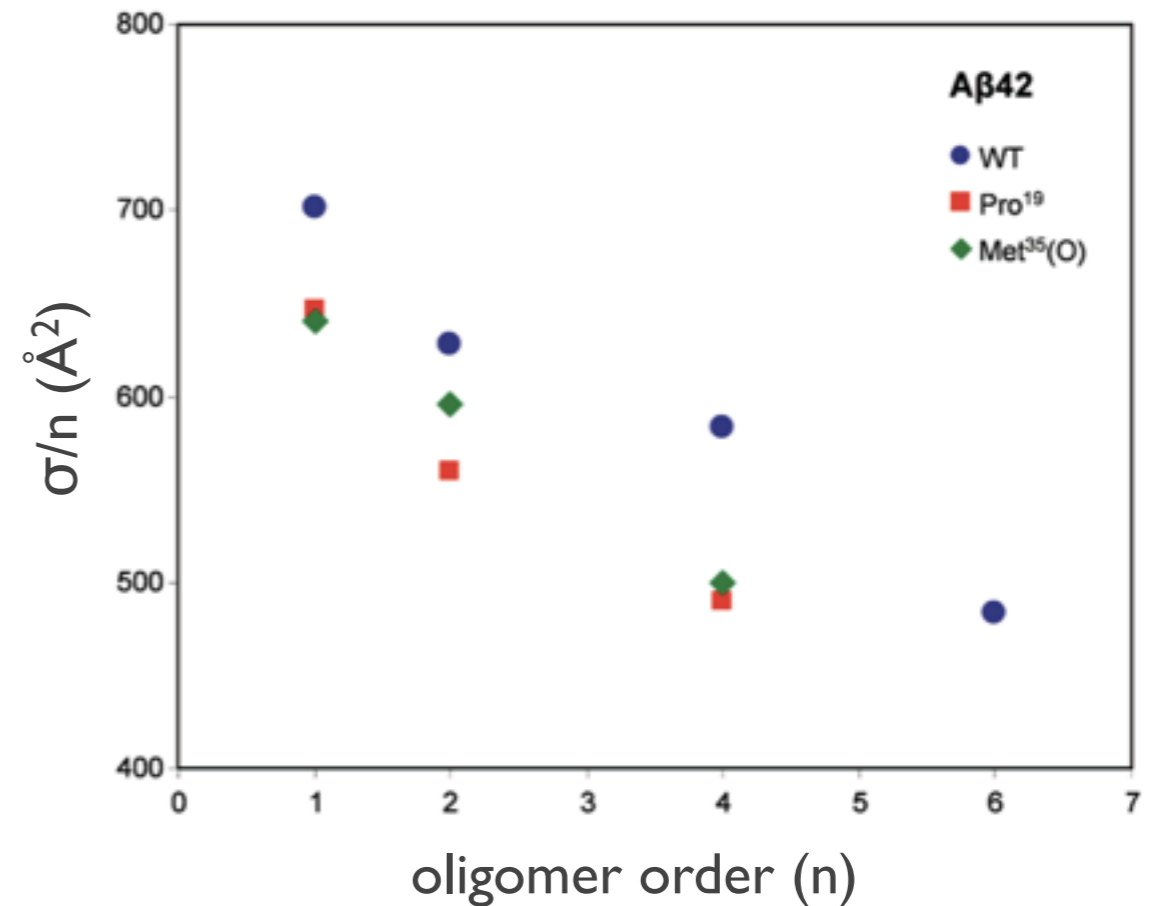
を満たすことにつながっている。また, $V = E \cdot \ell$ であるから, 「高電位かつ低電場」は, 装置を大きくすることによっても実現できる。

Gap Study between Model CCS and Exp. CCS

Oligomer	Structure [†]	Cross-section	
		Model	Experiment
Dimer*		1,256	1,256
Tetramer		2,358	
		2,277	2,332
		2,135	
Hexamer		3,450	
		3,100	2,898
		2,578	
Dodecamer		5,824	
		4,562	4,308
			

Model
v
Exp.

*Model cross-section fit to experiment. [†]Limiting structures of high symmetry.



- As n increases σ/n decreases.
∴ Oligomerization induces a certain amount of structural accommodation...?
- As model is based on dimer size information, it can overestimate CCS of larger oligomers. Thus, the inequality shown left is reasonable.