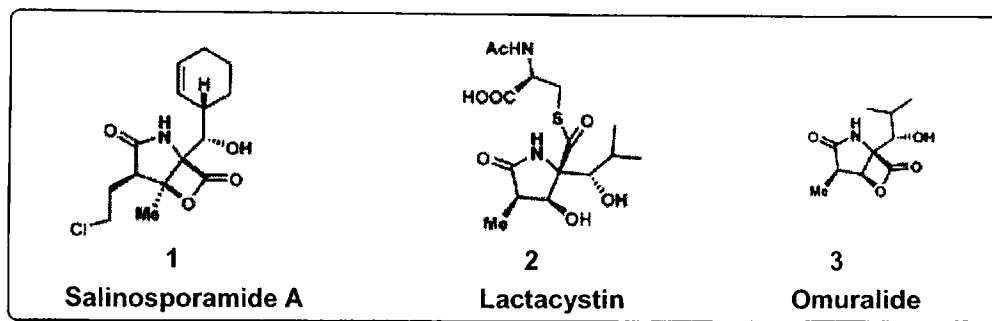


# Total Synthesis of Proteasome Inhibitor : Salinosporamide A and its Biological Aspects

Figure1:  
Proteasome inhibitor



	Salinosporamide A	Lactacystin	Omuralide
isolated from	bioactive product of a marine microorganism	Secondary metabolite derived from a Streptomyces in 1991	Key intermediate
Biological activity	Proteasome inhibitor IC <sub>50</sub> = 1.3 nM Antitumor against HCT-166 IC <sub>50</sub> = 11 ngML <sup>-1</sup>	Proteasome inhibitor IC <sub>50</sub> = 49 nM (Omuralide)	
First total synthesis	JACS.2004.126. 6230 (E. J. Corey) JSCS.2005.127. 8298 (Danishefsky, S. J.)	JACS.1992.114. 10677 (E. J. Corey) JACS.1993.115. 5302 (Omura-Smith) JACS.1994.116. 2139 etc...	TL.1993.34. 6977 (Key intermediate) (E. J. Corey)



Aaron Ciechanover



Avram Hershko



Irwin Rose

The Nobel Prize in Chemistry 2004  
" Ubiquitin-Proteasome system"

## Today's contents

<b>(A) Total Synthesis of Salinosporamide A.</b>	<b>(P.2~7)</b>
(A-1) Retro Synthesis	(P.2~3)
(A-2) Corey's method	(P.3~4)
(A-3) Omuralide-Salinosporamide A Hybrid	(P.4~6)
(A-4) Danishefsky's method	(P.7)
<b>(B) Ubiquitin-proteasome System.</b>	<b>(P.8~12)</b>
(B-1) Introduction	(P.8)
(B-2) Ubiquitin	(P.8)
(B-3) Ubiquitin system	(P.9)
(B-4) Proteasome	(P.10)
(B-5) Biological activity of Lactacystin and Omuralide	(P.11)
(B-6) Bortezomib (P-341)	(P.12)

## A: Total Synthesis of Salinosporamide A

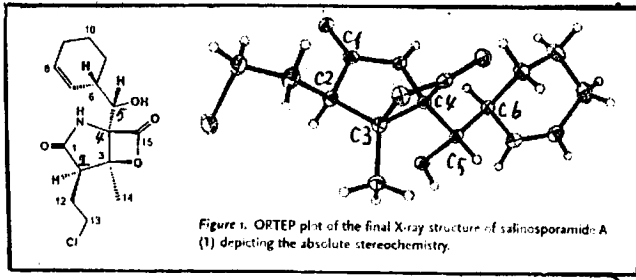


Fig. 2. Salinosporamide A

## \* Structural features:

- small molecule and it has five asymmetric carbons and functional groups.
- C2, C3, C4, C5, C6 ... asymmetric center.
- C3, C4 ... quaternary center.
- C5, C6 ... asymmetric center on side chain C4.

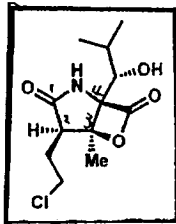


Fig 3. Omuralide-Salinosporamide A Hybrid.

\* The side chain on C4 was isopropyl group.

## \* Reported synthesis.

- E. J. Corey. JACS, 2004, 126, 6230
- Danishefsky. JACS, 2005, 127, 8298

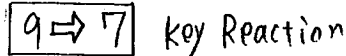
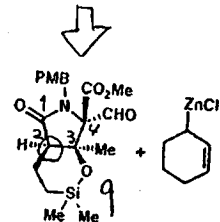
Fig. 3. \* Reported by E. J. Corey et al. JACS, 2005, 127, 8974

## (A-1) Retro Synthesis.

## Corey's method

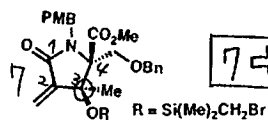


C5, C6 ... asymmetric center



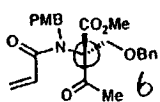
\* C2, C3: stereo relationship  
 $\rightarrow$  radical-chain cyclization

Key  $\downarrow$

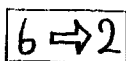


\* C(3): asymmetric center

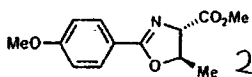
Key  $\downarrow$



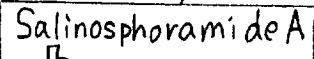
Baylis-Hilman  
 -aldol reaction



C(4): tertiary stereocenter

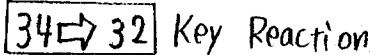
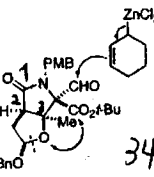


## Danishefsky's method



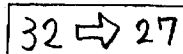
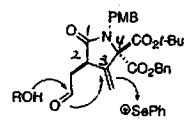
34 C5, C6 ... asymmetric center

same strategy as Corey's case.



C3: quaternary center  
 C2, C3: stereo relationship  
 $\rightarrow$  unusual cationic hemiacetal seleno cyclization

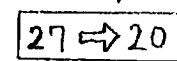
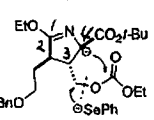
Key  $\downarrow$



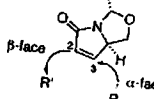
C4: quaternary center

$\rightarrow$  lactone formation and the use of nucleophilic selenium species.

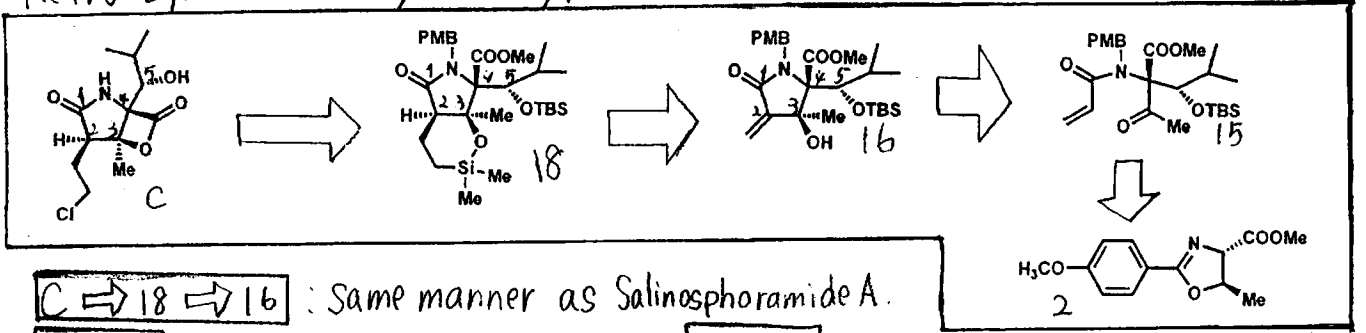
Key  $\downarrow$



C2, C3: asymmetric center.



# Retro Synthesis : Hybrid type.



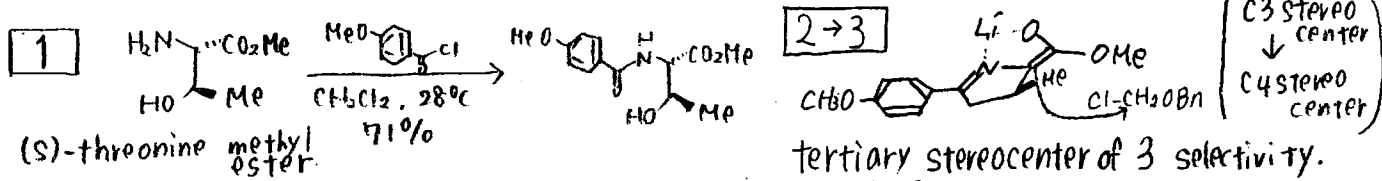
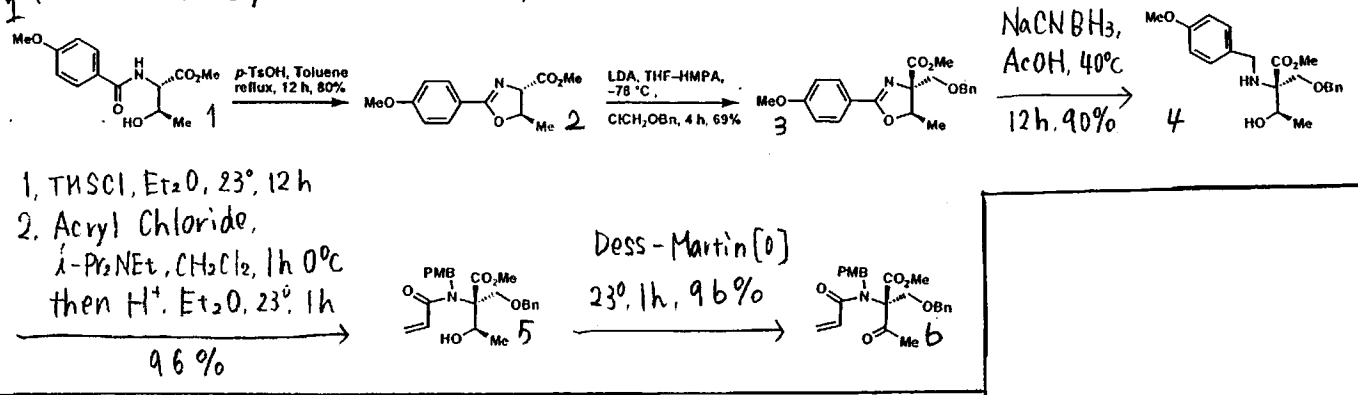
**C**  $\Rightarrow$  **18**  $\Rightarrow$  **16** : same manner as Salinosphoramide A.

**16**  $\Rightarrow$  **15** : more improved cyclization **15**  $\Rightarrow$  **2** diastereoselective aldol reaction.

## <A-2> Corey's Method

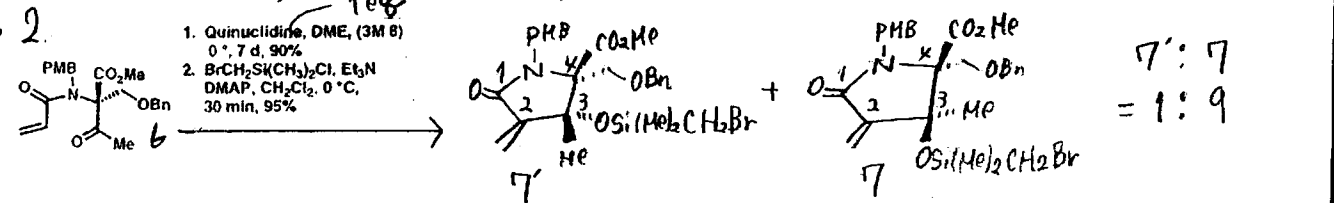
### <A-2-1> Synthesis of key intermediate b.

Scheme 1

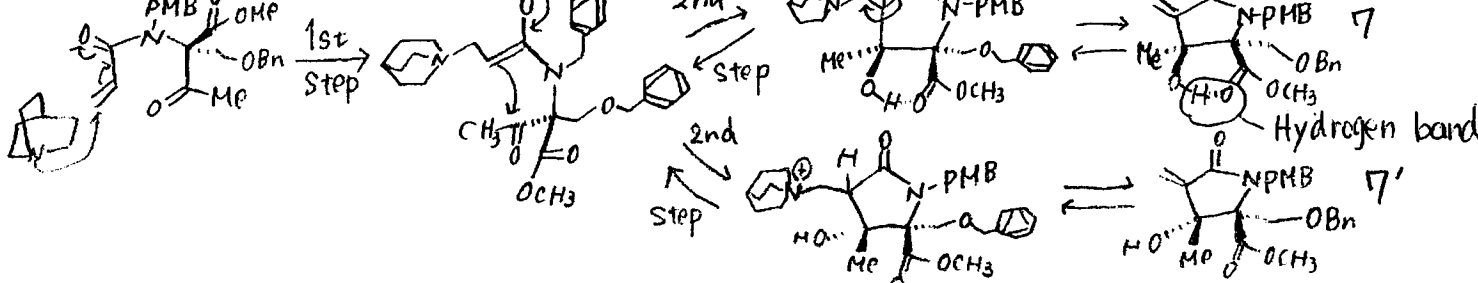


### <A-2-2> Key Reaction: Baylis Hilman-aldol Reaction

Scheme 2



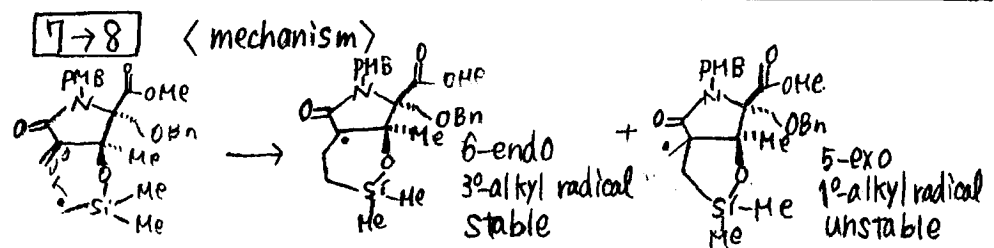
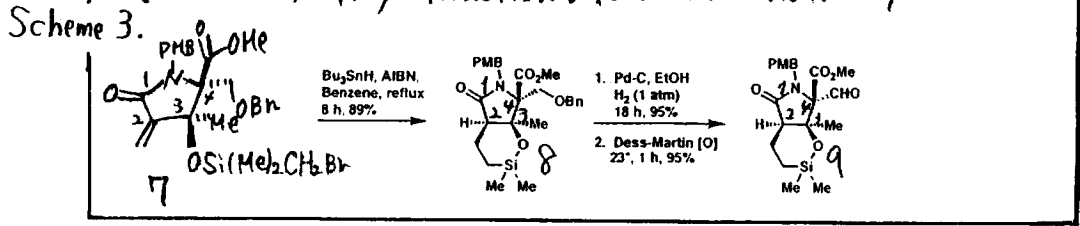
#### <mechanism>



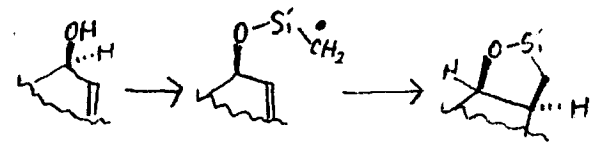
- \* C4 stereo center  $\rightarrow$  C3 stereo center
- \* 7, 7' : diastereomer were separated at this stage by column.
- \* Condition : 20°C for 9 h  $\Rightarrow$  7 : 7' = 1 : 4 90% yield
- 0°C for 7 day  $\Rightarrow$  7 : 7' = 1 : 9 90% yield

long time  $\Rightarrow$  Overcome this problem : P5, <A-3-2>

**(A-2-3) Key Reaction: Radical-Chain Cyclization**



\* cyclic allylic alcohols → cis bicyclic ether



(Ref: Stork, JACS: 1985, 107, 501  
Nishiyama, JOC: 1984, 49, 2299  
Chem Rev: 1995, 1253/1997, 2193)

\* Si-effect (Ref: Tetrahedron, 1985, 3979)

- Silicon-induced stereoelectronic factor
- Silyl radical were stabilized. And it would decrease their cyclization potency.

"Baldwin-Beckwith rules" (Radical Cyclization)

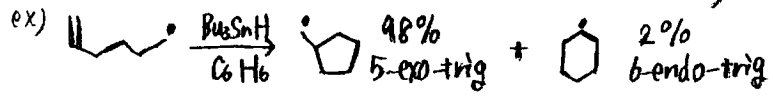
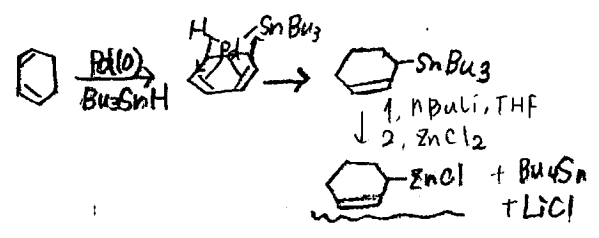
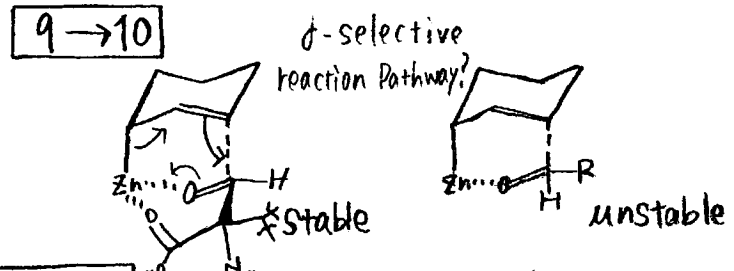
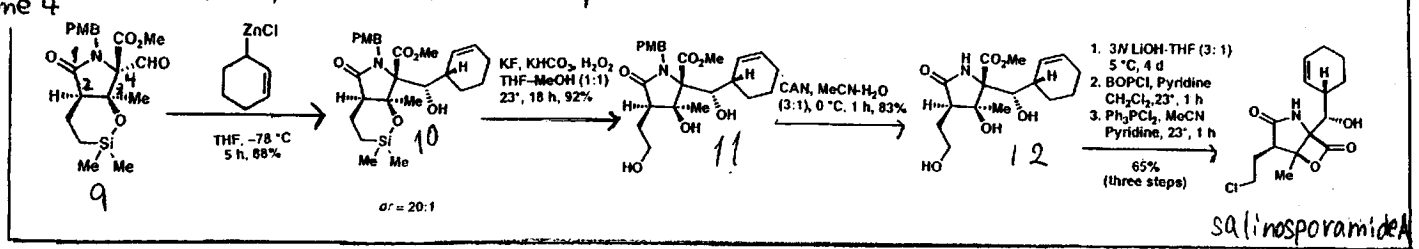


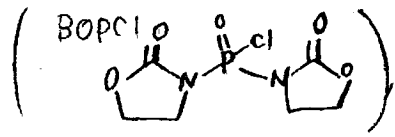
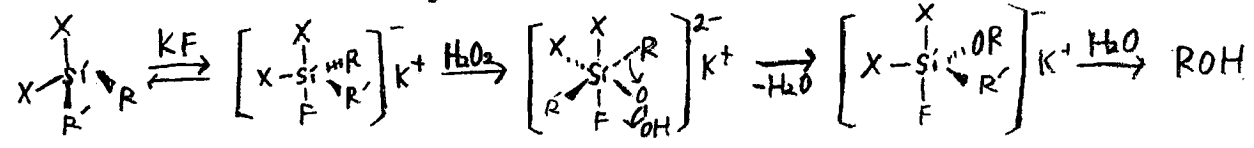
Table 2. Rate Constants for the Cyclization of 5-Hexenyl and Some Substituted 5-Hexenyl Radicals at 25 °C<sup>a</sup>

Radical	10 <sup>4</sup> k <sub>5-exo</sub>	10 <sup>4</sup> k <sub>6-endo</sub>
	~1	~0.1
	>200	~0.1
	0.7	1.4

**(A-2-4) Synthesis of Salinosporamide A**



**10 → 11** Tamao-Fleming oxidation

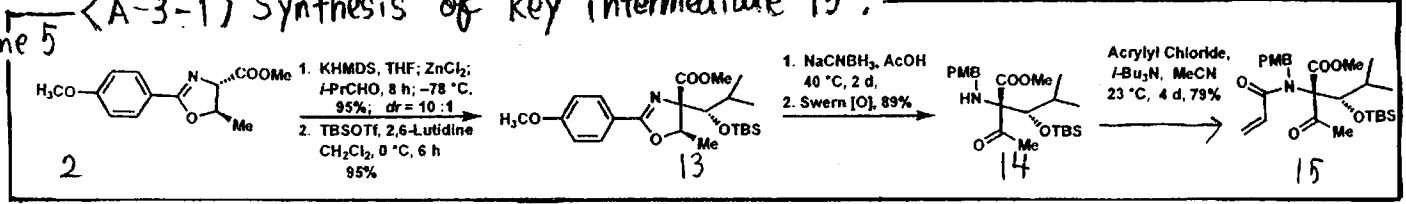


Total 17 steps, 13.7% yield.

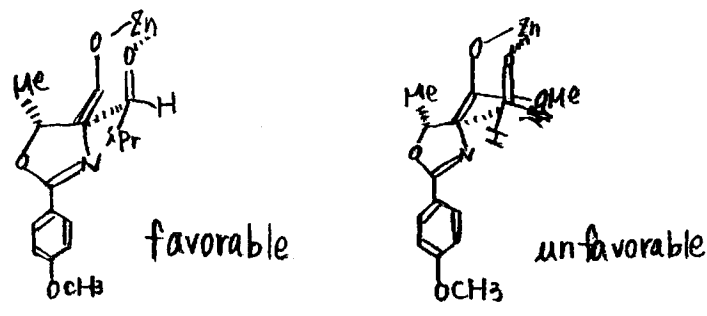
<A-3> Omuralide-Salinosporamide A Hybrid

<A-3-1> Synthesis of key intermediate 15.

Scheme 5

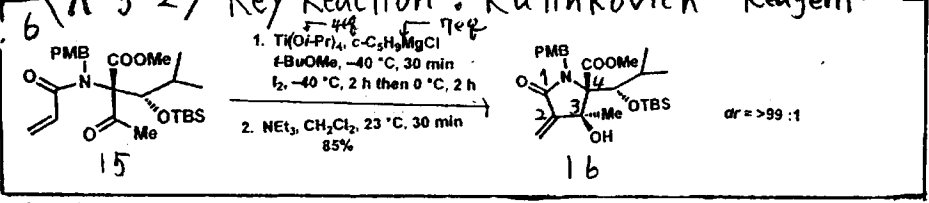


2 → 13 diastereo selective aldol reaction.

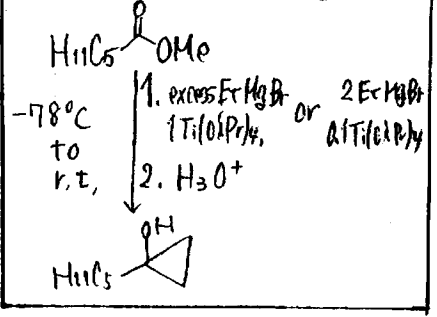


<A-3-2> Key Reaction: Kulinkovich Reagent

Scheme 6

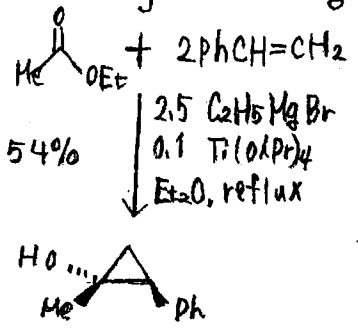


Kulinkovich Reagent

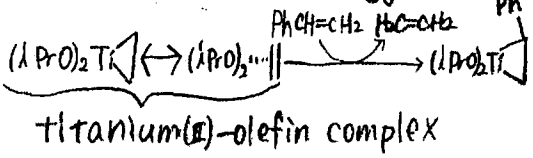


(Ref: Chem Rev, 2003, vol 103, 2597  
Adv. Synth. Catal, 2001, 343,  
JACS, 1996, 118, 3182)

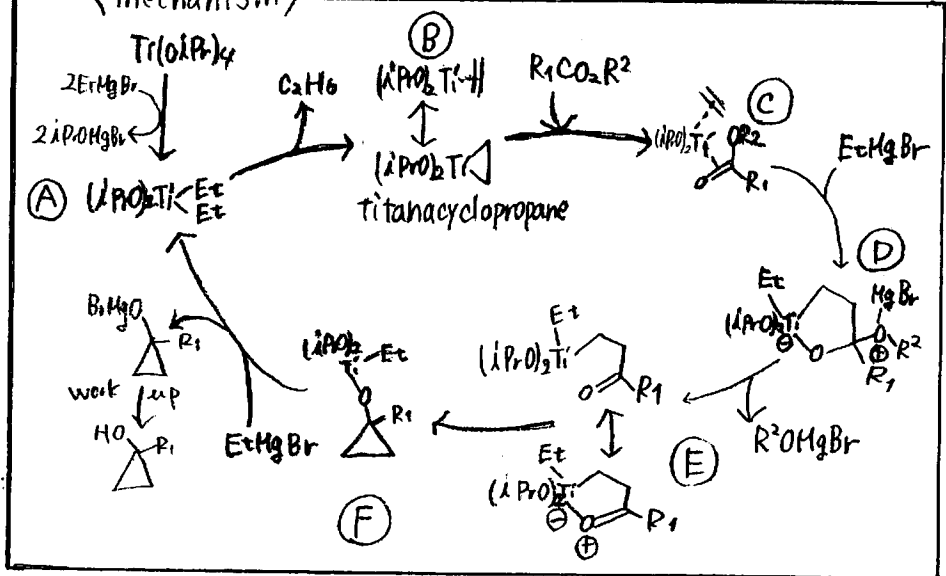
\* Olefin-ligand exchange



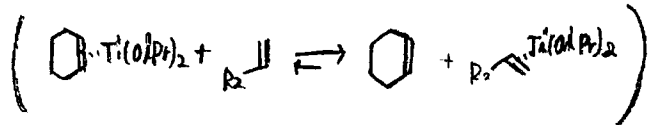
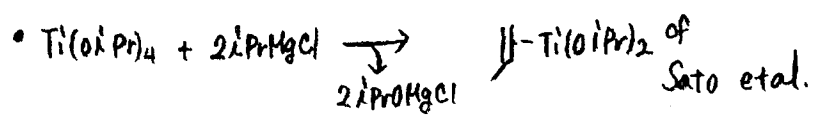
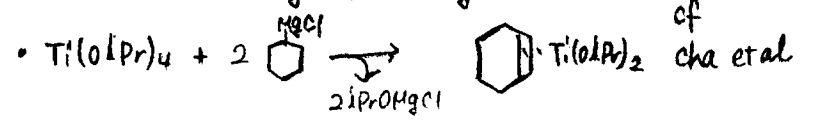
→ This result was suggested

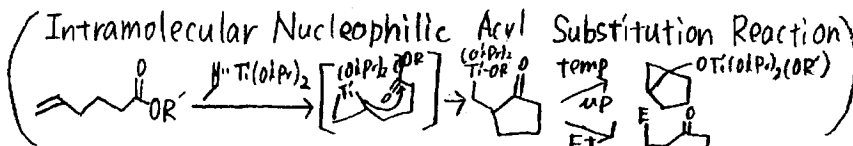
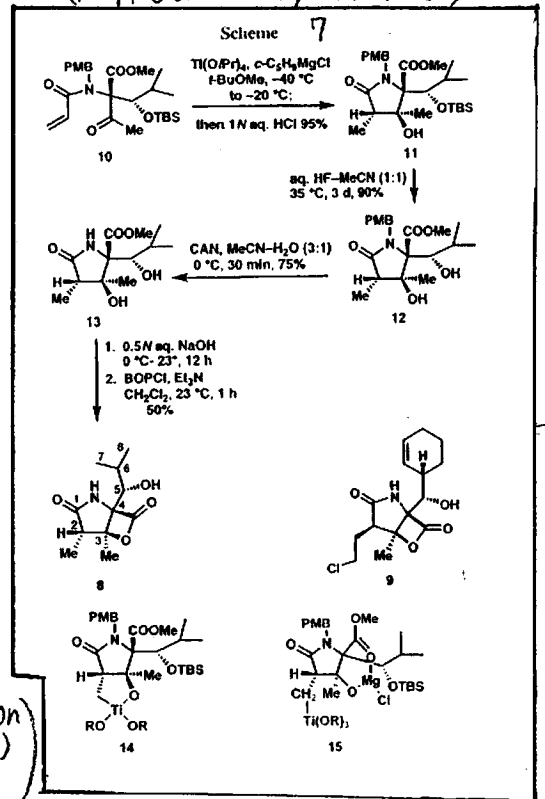
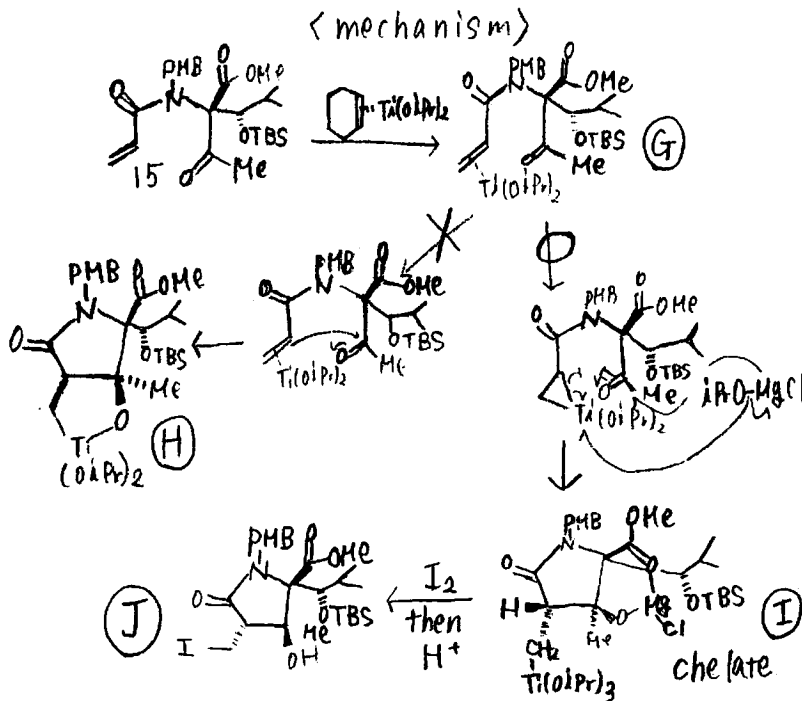


<mechanism>

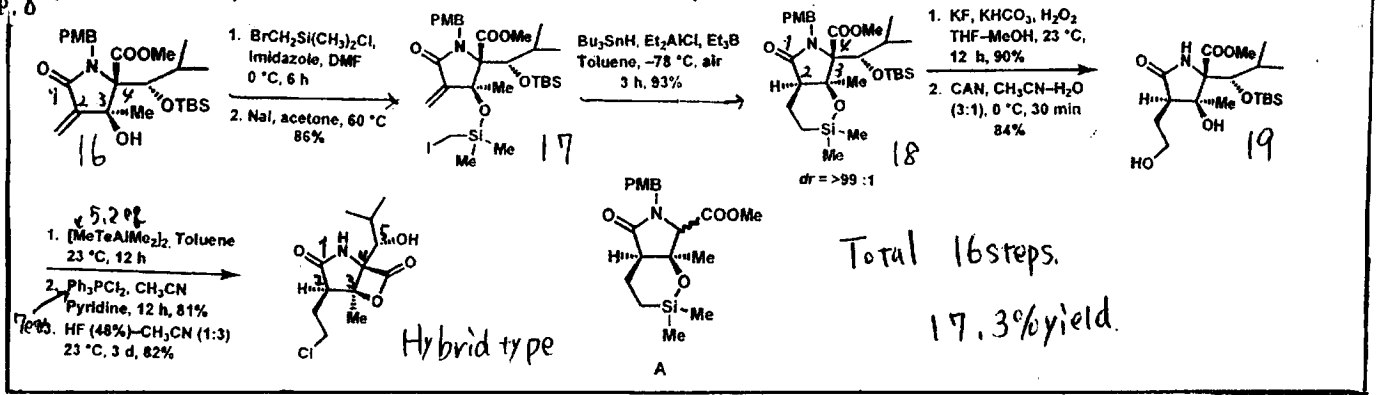


\* To facilitate ligand exchange.

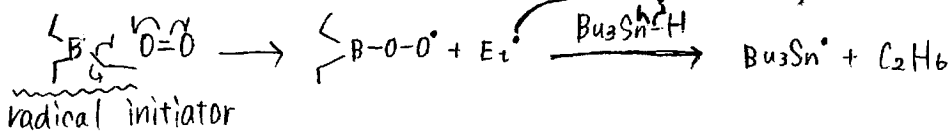




Scheme 8 (A-3-3) Synthesis of Omuralide-Salinosporamide A Hybrid.



17  $\rightarrow$  18 •  $\left\{ \begin{array}{l} \text{Bu}_3\text{SnH/AIBN condition: heat } 80\sim 100^\circ\text{C} \\ \text{Bu}_3\text{SnH/Et}_2\text{AlCl/air condition: } 0^\circ\text{C (mild condition (JAOS, 1987, 109, 2549))} \end{array} \right.$

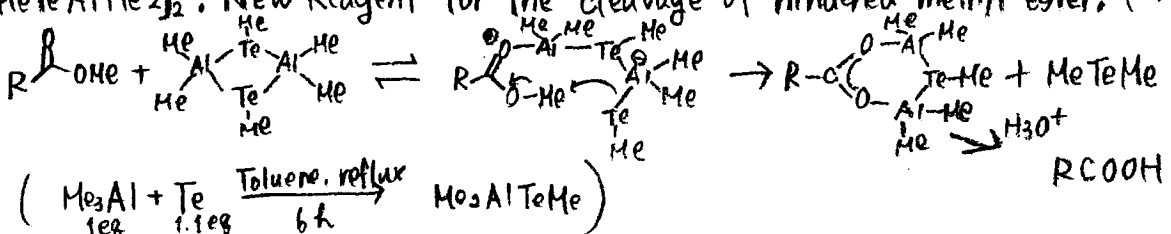


- $\text{Et}_2\text{AlCl}$  : Lewis Acid affects the reaction rates and stereoselectivity. (OL: 2001, 67)
- $\text{Et}_2\text{AlCl}$  increased the electron-withdrawing nature of the carbonyl group thus making the conjugate addition pathway via a low energy process.

19  $\rightarrow$  Hybrid type

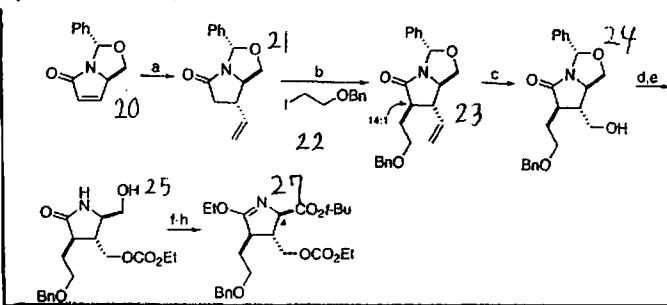
cleavage of methyl esters of 19 : complete failure under variety of conditions and compound A was obtained. (Ref, TL, 2005)

[MeTeAlMe<sub>2</sub>]<sub>2</sub> : New Reagent for the cleavage of hindered methyl ester. (46, 4589)

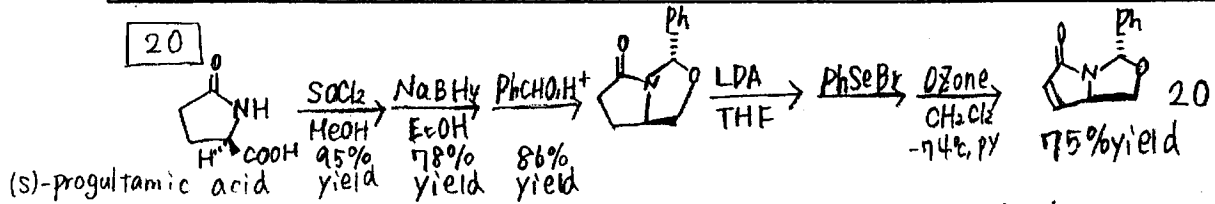


< A-4 > Danishefsky method

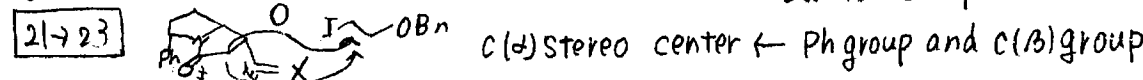
Scheme 9 (A-4-1) Synthesis of key intermediate 27



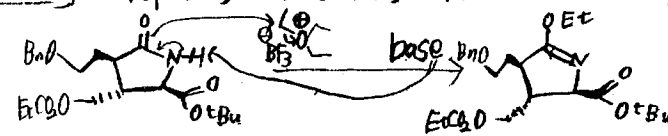
Key: (a) vinylmagnesium bromide, TMSCl, CuI, THF, -78 °C (75%); (b) **22**,<sup>10</sup> LDA, THF, room temperature (rt) (77%, dr = 14:1); (c) O<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>-MeOH (3:1), -78 °C then NaBH<sub>4</sub>, 0 °C (86%); (d) ClCO<sub>2</sub>Et, pyridine, rt (96%); (e) TfOH, THF-H<sub>2</sub>O (9:1), rt (quant); (f) Jones reagent, acetone, rt; (g) Me<sub>2</sub>NCH(O<sup>t</sup>-Bu)<sub>2</sub>, toluene, reflux (72% in two steps); (h) Et<sub>3</sub>OBf<sub>4</sub>, K<sub>2</sub>CO<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, rt (88%);



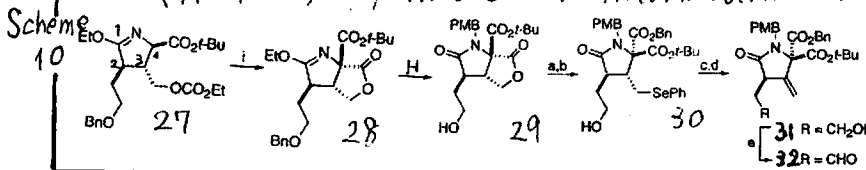
[20 → 21] \* TMSCl has effect on the rate and stereo chemical outcome of many cuprate C(β) stereo reactions. C(β) stereo center O-Si > O-Li. dππ\* complex → Cu<sup>II</sup> β-complex.



[25 → 27] Step (h): Et<sub>3</sub>OBf<sub>4</sub>; Meerwein Reagent



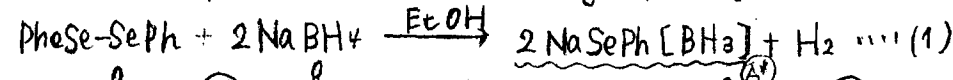
< A-4-2 > Synthesis of intermediate 32



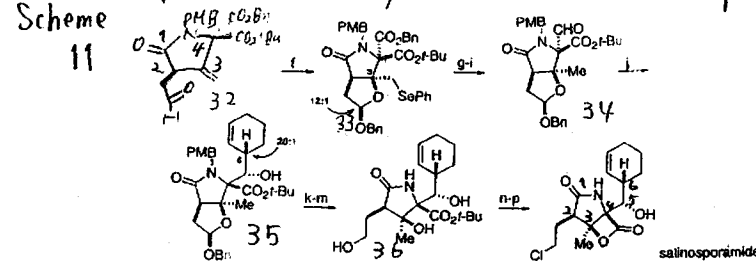
(i) LHMDS, THF, -20 °C (82%); (j) 1 M HCl aq, THF, 0 °C (90%); (k) PMBCl, NaH, DMF, rt (61%); (l) Pd(OH)<sub>2</sub>-C, H<sub>2</sub>, EtOH, rt (quant). (a) PhSePh, NaBH<sub>4</sub>, EtOH, 60 °C; (b) BnBr, K<sub>2</sub>CO<sub>3</sub>, DMF, rt (65% in 2 steps); (c) 30% H<sub>2</sub>O<sub>2</sub> aq, THF, rt; (d) toluene, 100 °C (94% in two steps, 72% **31** + 22% **32**); (e) Dess-Martin periodinane, CH<sub>2</sub>Cl<sub>2</sub>, rt (92%, 89% in three steps from 30);

[27 → 28] C(3) stereo → C(4) stereo : Using intramolecular acylation.

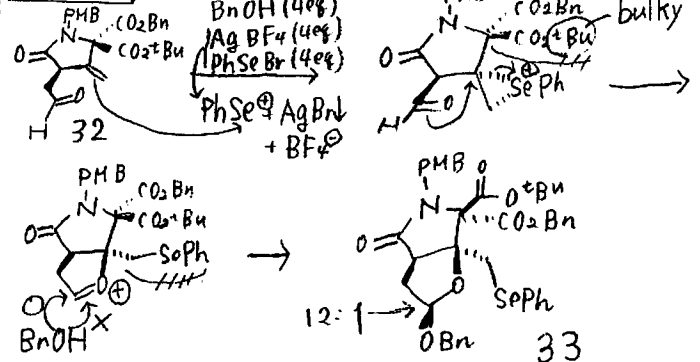
[29 → 30] Step (a): nucleophilic ring opening (Ref: JACS, 1980, 3904, JOC, 1981, 46, 2605)



< A-4-3 > Synthesis of Salinosporamide A



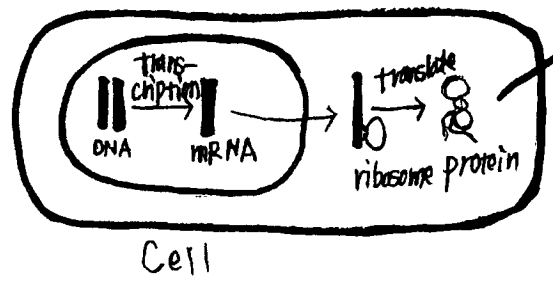
[32 → 33] acetal Mediate cationic cyclization



(f) PhSeBr, AgBF<sub>4</sub>, BnOH, CH<sub>2</sub>Cl<sub>2</sub>, -20 to 0 °C (74% as an anomeric mixture); (g) AIBN, n-Bu<sub>3</sub>SnH, toluene, 100 °C (98%); (h) NaBH<sub>4</sub>, THF-EtOH (3:1), rt (85%); (i) Dess-Martin periodinane, CH<sub>2</sub>Cl<sub>2</sub>, rt (95%); (j) 8% THF, -78 °C (88% for 35, dr = 20:1); (k) ceric ammonium nitrate (CAN), CH<sub>3</sub>CN-H<sub>2</sub>O, 0 °C (90%); (l) Na, liq NH<sub>3</sub>, -78 °C; (m) NaBH<sub>4</sub>, THF-H<sub>2</sub>O (2:1), rt (97% in two steps); (n) BCl<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, 0 °C; (o) BOPCl, TEA, CH<sub>2</sub>Cl<sub>2</sub>, rt; (p) Ph<sub>3</sub>PCL<sub>2</sub>, pyridine, CH<sub>3</sub>CN, rt (51% in three steps).

Total 28 steps, < 1.8% yield.

<B> Ubiquitin - Proteasome system  
 <B-1> Introduction.



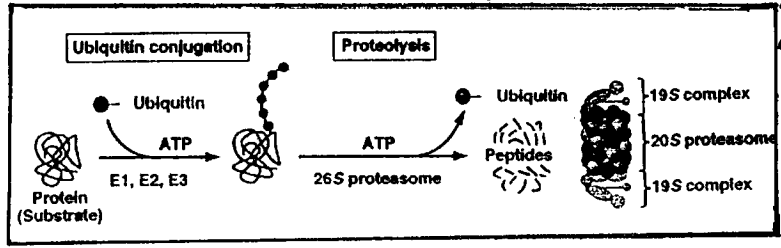
Protein control life phenomenon.  
 \* Abnormally folded protein cause human Diseases.  
 ex) Alzheimer's disease.  
 Parkinson's disease. etc....

- System of protein degradation.

Outside cell ... Protease  
 Inside cell } lysosome  
 Ubiquitin-Proteasome System ... selective protein degradation

- Ubiquitin-Proteasome system

Fig. 1

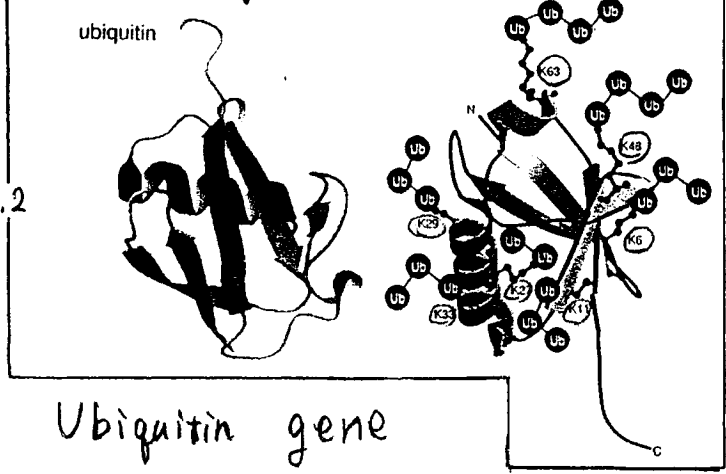


Ubiquitin ... Degradation-signal  
 Proteasome  
 a protein destruction machine  
 a complex protease  
 \* ATP 依存の 7 = 1107 質分解システム

- history - 1953 : Simpson 「The breakdown of the cell's own proteins does require energy」  
 1977 : Goldberg "A first step towards an explanation of this energy-dependent protein degradation"  
 late 1970 and early 1980s  
 Nobel prize winner succeeded in showing that protein was destroyed being labelled with the ubiquitin and this regulation requires energy.

<B-2> Ubiquitin

Fig. 2



- 76-amino-acid residue.
- Secondary structure ...  $\beta$   $\beta$   $\alpha$   $\beta$   $\beta$   $\beta$
- 熱安定性生体タンパク質
- Strongly conserved in evolution.
- Ubiquitin has seven Lys residue. (K6, K11, K27, K29, K33, K48, K63)

Ubiquitin gene  
 • code two type }  $(Ub)_n$  ...  $(Ub)_n$ : Polyubiquitin gene  
 •  $(Ub)_n$  LRP  
 •  $(Ub)_n$  SRP

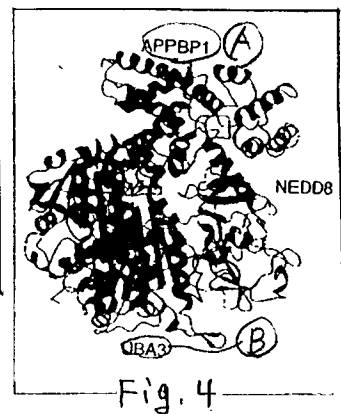
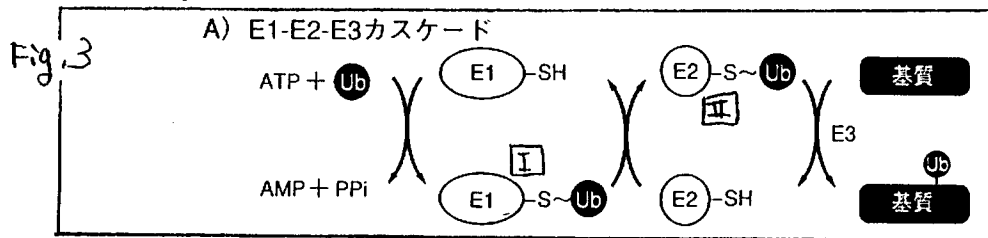
• エビキチン  
 • リボソームタンパク質  
 高虫合遺伝子

(LSP: 大リボソームタンパク質)  
 (SRP: 小リボソームタンパク質)

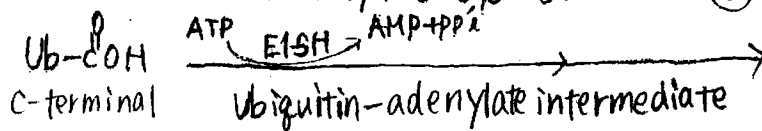
• 熱ショック応答遺伝子 • 脱ユビキチン化酵素により単体機能分子に成熟



# <B-3> Ubiquitin System

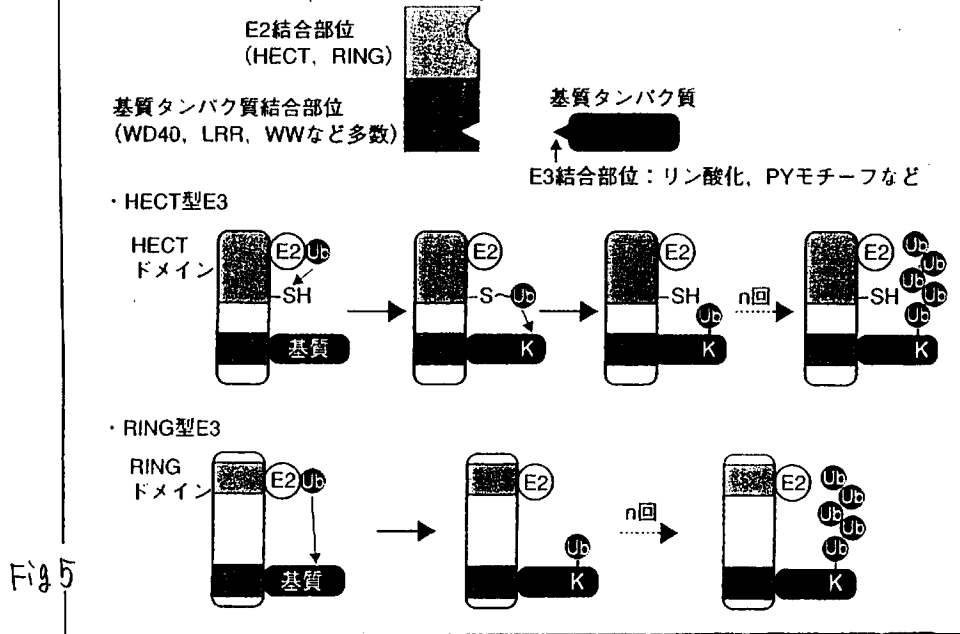


- E1 (Ubiquitin activating enzyme)
  - 110~120KDa
  - strongly conserved in evolution.
  - two domain
    - Adenylation domain → (A)
    - Catalytic Cys domain → (B)

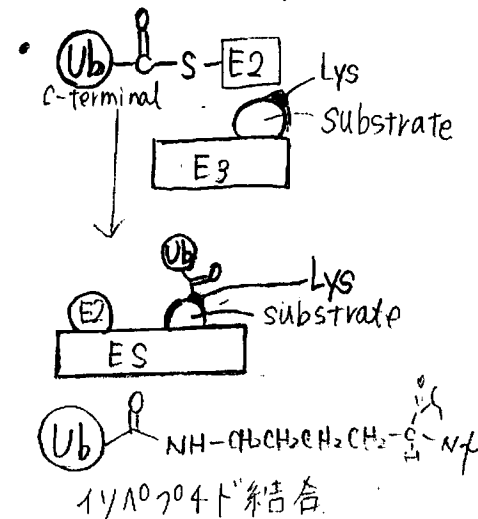


- E2 (Ubiquitin conjugating enzyme)
  - core domain ... 150 amino residue
  - E2 mediate the transfer of Ubiquitin from E1 to the protein substrate, with or without the participation of ubiquitin-protein ligase E3.

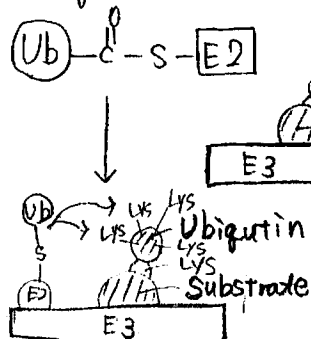
## E3 (Ub-protein ligase)



- hundreds of different E3 proteins existed.
- two type
  - HECT type
  - RING type



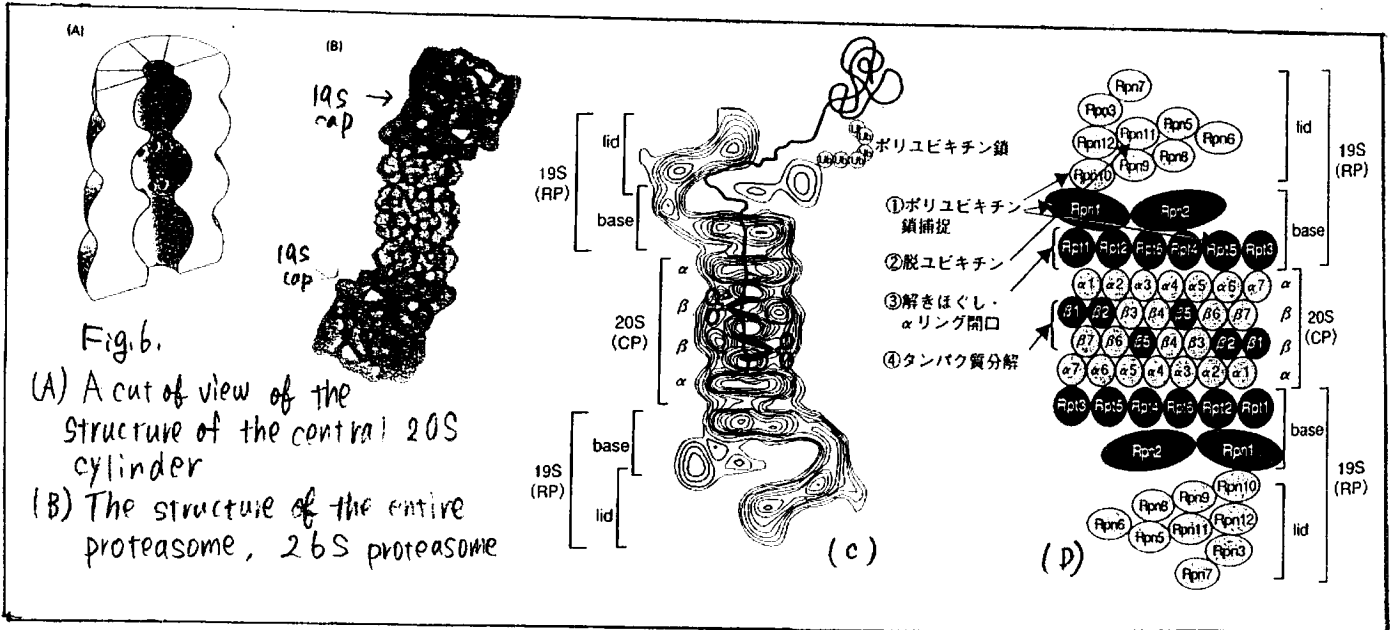
## Polyubiquitin



- \* Ubiquitin has seven Lys residue.
  - K48 → protein degradation.
  - K63 → DNAの修復 etc
  - not protein degradation.

→ エピキチンの修飾は多様性モリ

<B-4> Proteasome.



26S proteasome (Fig: 6 - (B), (C), (D))

- ~2500KDa complex
- made up of at least 32 different subunit
- highly conserved among all eukaryotes.
- two major subcomplexes

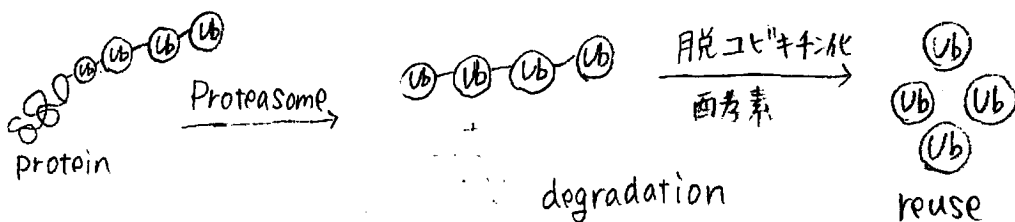
20S (CP): contains the protease subunit  
 19S (RP): regulatory particle (regulates the function of the former)

\* 20S (CP) (Fig: 6 - (A))

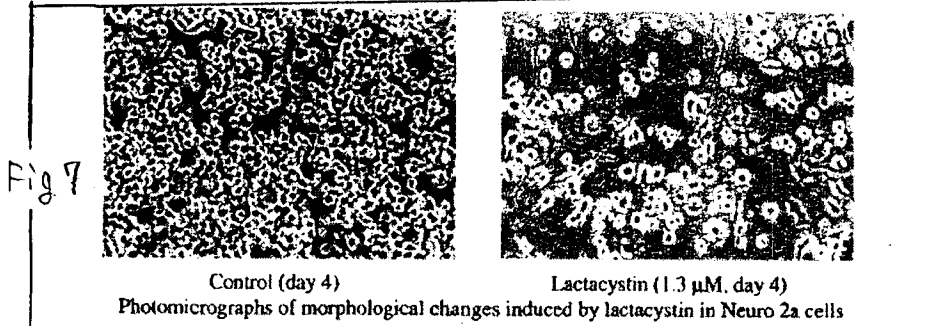
- barrel-shaped structure
- length: 15nm
- Diameter 11nm
- 700KDa
- made up of four rings of seven subunits each
- Two inner β-ring contain the proteolytic active sites facing inward into a sequestered proteolytic chamber.
- β1, β2, β5 subunit ... Active site  
 chymotrypsin-like, trypsin-like, caspase-like

\* 19S (RP) (Fig: 6 - (B), (C), (D))

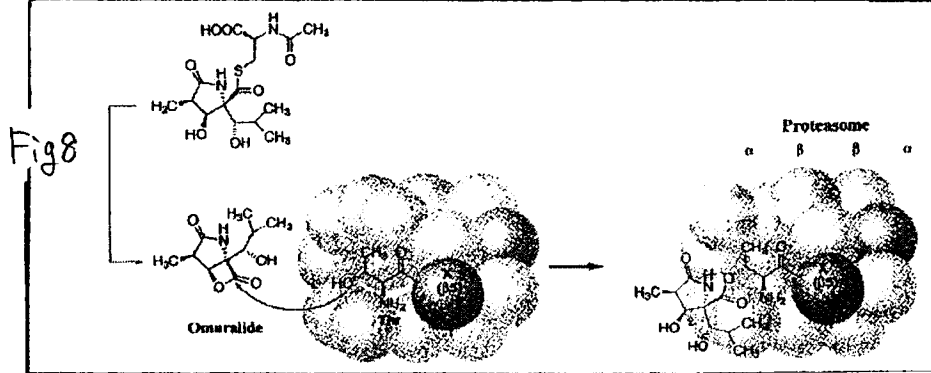
- two multisubunit ... lid (蓋部), base (基部)
- lid ... nine "Rpn" subunit  
 Rpn 11: de-ubiquitinating activity  
 Rpn 10: "Adhesive"
- base ... Rpt 1~6: six ATPase subunit  
 Rpn 1, 2: Non-ATPase subunit



<B-5> Biological activity of Lactacystin and Omuralide.



Bioactivity (PNAS, 1994, 91, 3358)  
 • induce Neurite outgrowth in Neuro 2A.  
 • arrest cell cycle at both G<sub>0</sub>/G<sub>1</sub> and G<sub>2</sub> phase

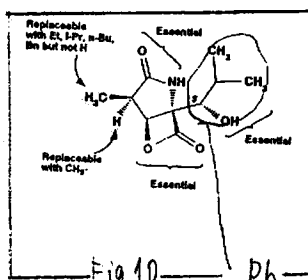
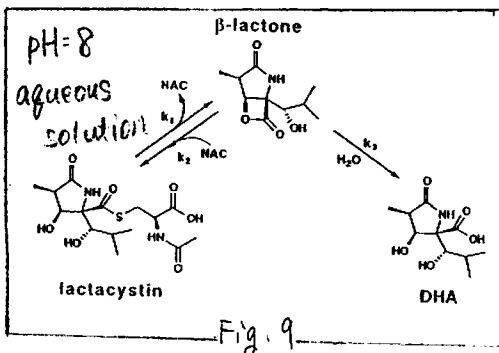


Inhibition of Proteasome Activities and Subunit-Specific Amino-Terminal Threonine Modification by lactacystin  
 (Science, 1995, vol 268, 726) (E.J. Corey)

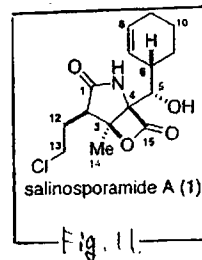
\* Lactacystin appears to modify covalently the highly conserved amino-terminal threonine of protease subunit β5.

⇒ Proteasome have NH<sub>2</sub> terminal threonine active site.

= Proteasomes are novel threonine protease.



Ph = no biological activity



\* Salinosporamide A may interact with proteasome in a manner that is different from Omuralide.

\* Application of Lactacystin to cell biology

lactacystin: Proteasome inhibitor has been utilized to dissect the proteasome function.

⇒ leading to unexpected finding about importance of Ubiquitin-proteasome pathway and proteasome function.

Ubiquitin-Proteasome } Cell cycle, apoptosis (細胞自滅)  
 Antigen presentation ... etc

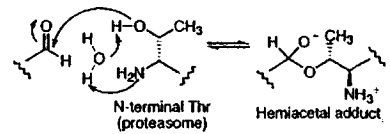
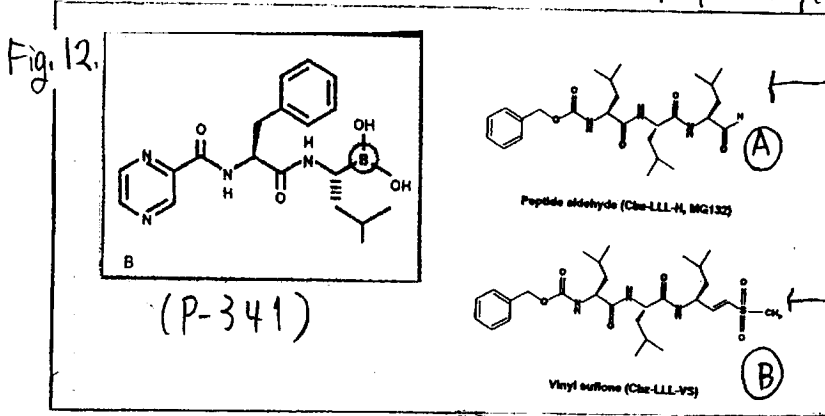
Ubiquitin-Proteasome system is involved in life-phenomenon.  
 → The Nobel Prize in 2004.

Tumorigenesis is deeply related to progression of Cell cycle and apoptosis.



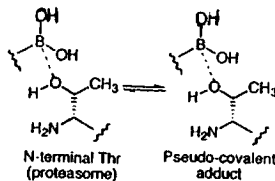
Proteasome inhibitors: Potential therapeutic agents.

<B-6> Bortezomib (P-341): First-in-class Proteasome inhibitor for the treatment of Multiple Myeloma (MM) (多发性骨髓瘤)



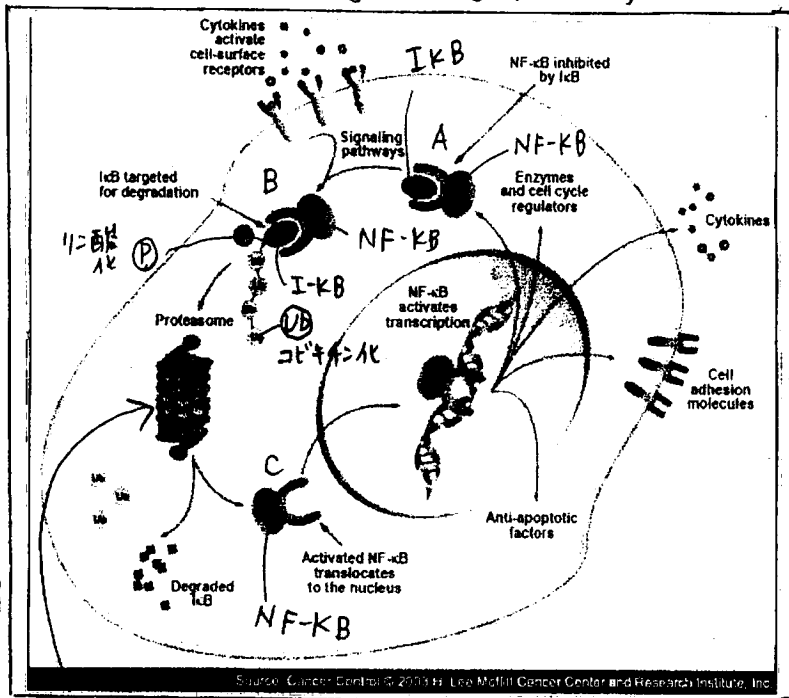
• Only short-lived proteasome inhibition

all bind to 20S core particle in irreversible manner.



dissociate more slowly from the proteasome stable inhibitor.

Intracellular signaling pathway



In Myeloma cell, Cell growth was related to transcription factor nuclear factor-κB (NF-κB). (転写因子)

NF-κB

• is normally sequestered in the cytoplasm and rendered inactive by its inhibitor protein (IκB) (阻害因子)

numerous stimuli, cell stress

↓ phosphorylation IκB

↓ Ubiquitin-IκB

↓ proteasome: degradation IκB and release activated NFκB

↓ NF-κB translated to the nuclear and initiates the transcription of wide range of genes, including involved in cell survival.

Proteasome inhibition

↓ stabilize IκB and prevent the activation NF-κB

↓ increased apoptosis in cancer cell. (細胞自滅)