

第4章 蛾類性フェロモンの生合成とその制御機構

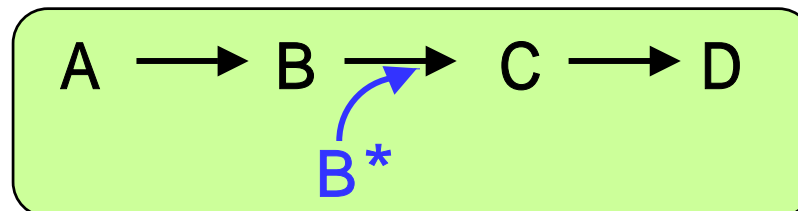
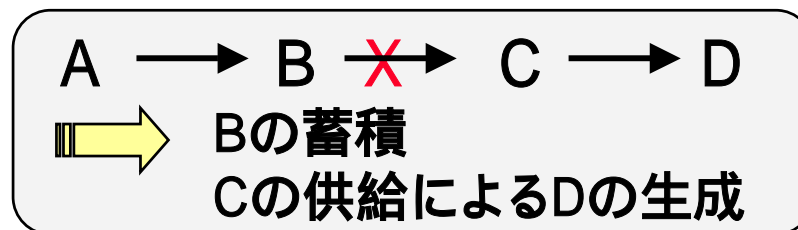
(A) 生合成の研究法

A-1) 実験法

1. 突然変異株(種)の利用

2. 標識前駆体の投与実験

- a) 安定同位元素 ^2H (D), ^{13}C
分析方法: GC-MS, NMR



フェロモン成分: 超微量物質
多量の前駆体の投与

問題点

本来の生合成系の攪乱の可能性

- b) 放射性同位元素 ^3H (T), ^{14}C , ^{11}C

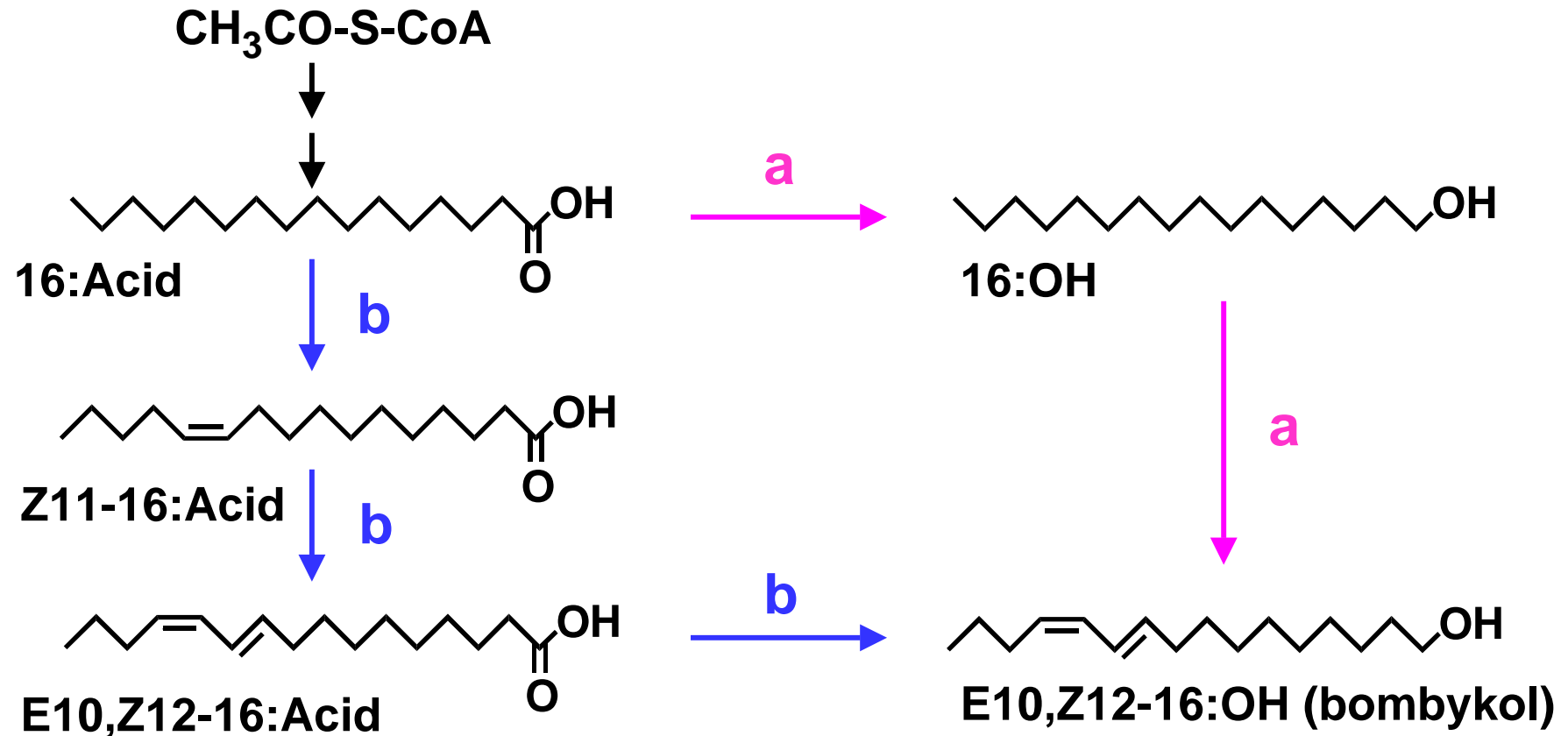
分析方法: オートラジオグラフィー 化学構造 OK?

問題点

個体: 再結晶

液体: クロマトグラフィー

A-2) Bombykol (カイコの性フェロモン) の生合成経路



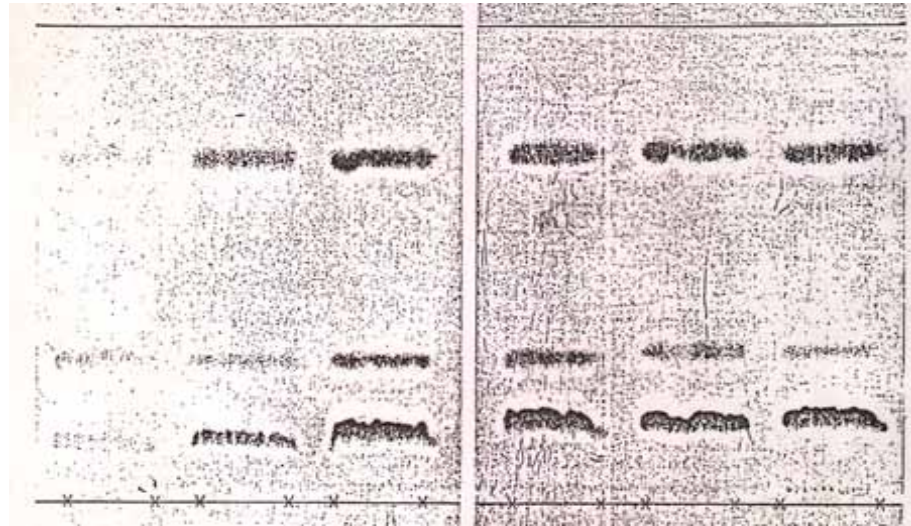
a Inoue & Hamamura, 1972. *Nippon Nogeikagaku Kaishi*, 46, 645

b Yamaoka *et al.*, 1984. *Experientia*, 48, 80

Bjostad & Roelofs, 1984. *Insect Biochem.*, 14: 275

A-3) パルミチン酸 (16:Acid) の変換

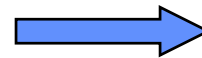
TLC (オートラジオグラフィー)



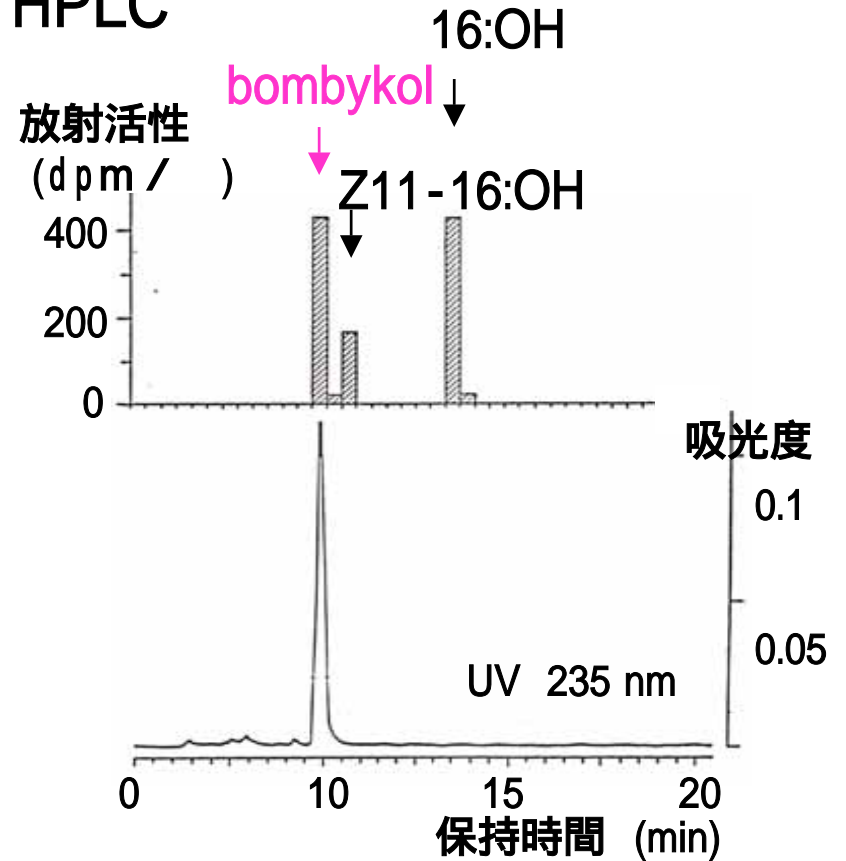
i ii iii iv v vi
 solvent: benzene + ethyl acetate (4:1)

- i [1-¹⁴C]12:Acid
- ii [1-¹⁴C]14:Acid
- iii [1-¹⁴C]16:Acid
- iv [16-¹⁴C]16:Acid
- v [1-¹⁴C]18:Acid
- vi [18-¹⁴C]18:Acid

A: triacylglycerols
 B: alcohols
 C: acids (recovery)

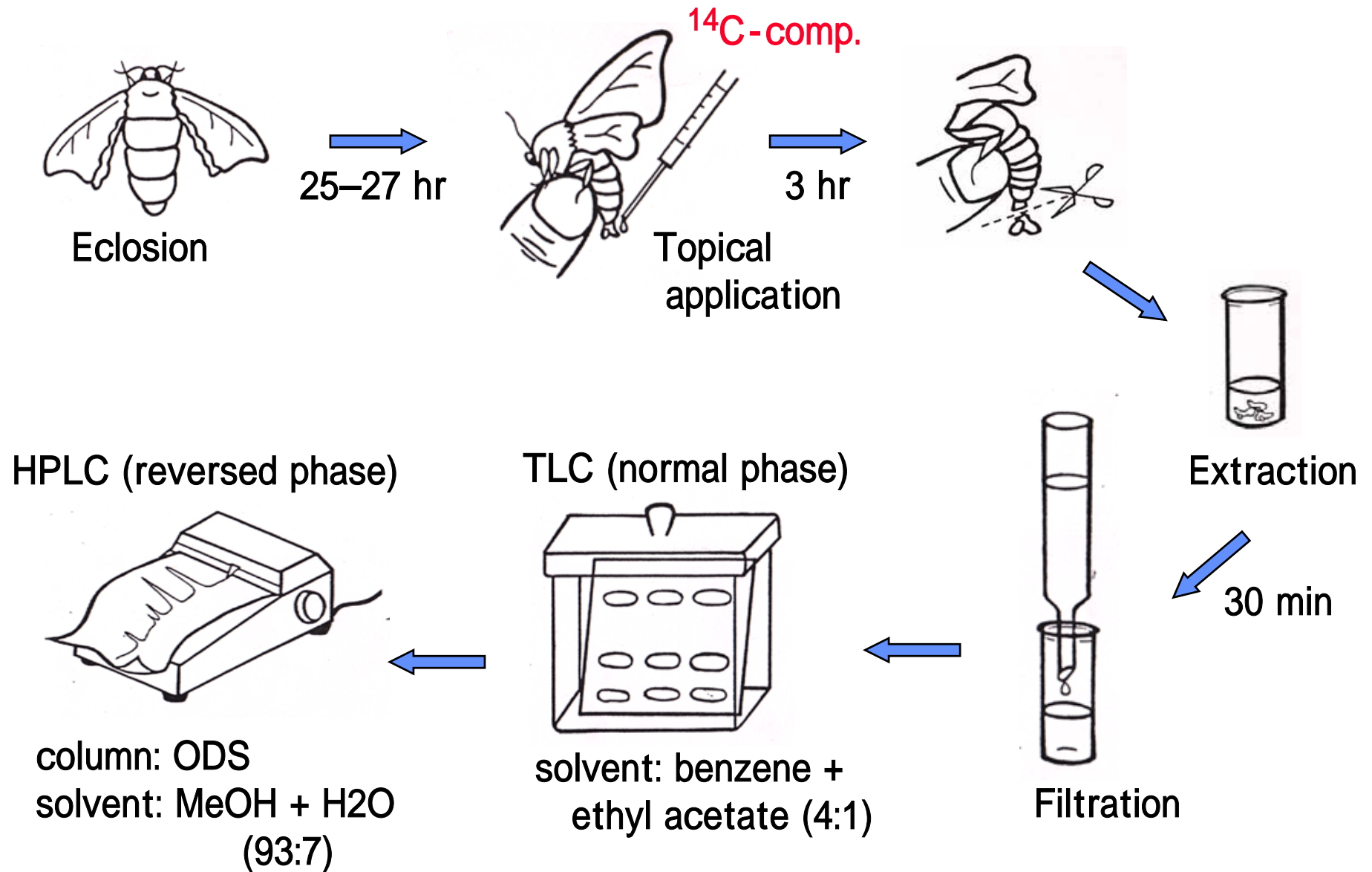


HPLC

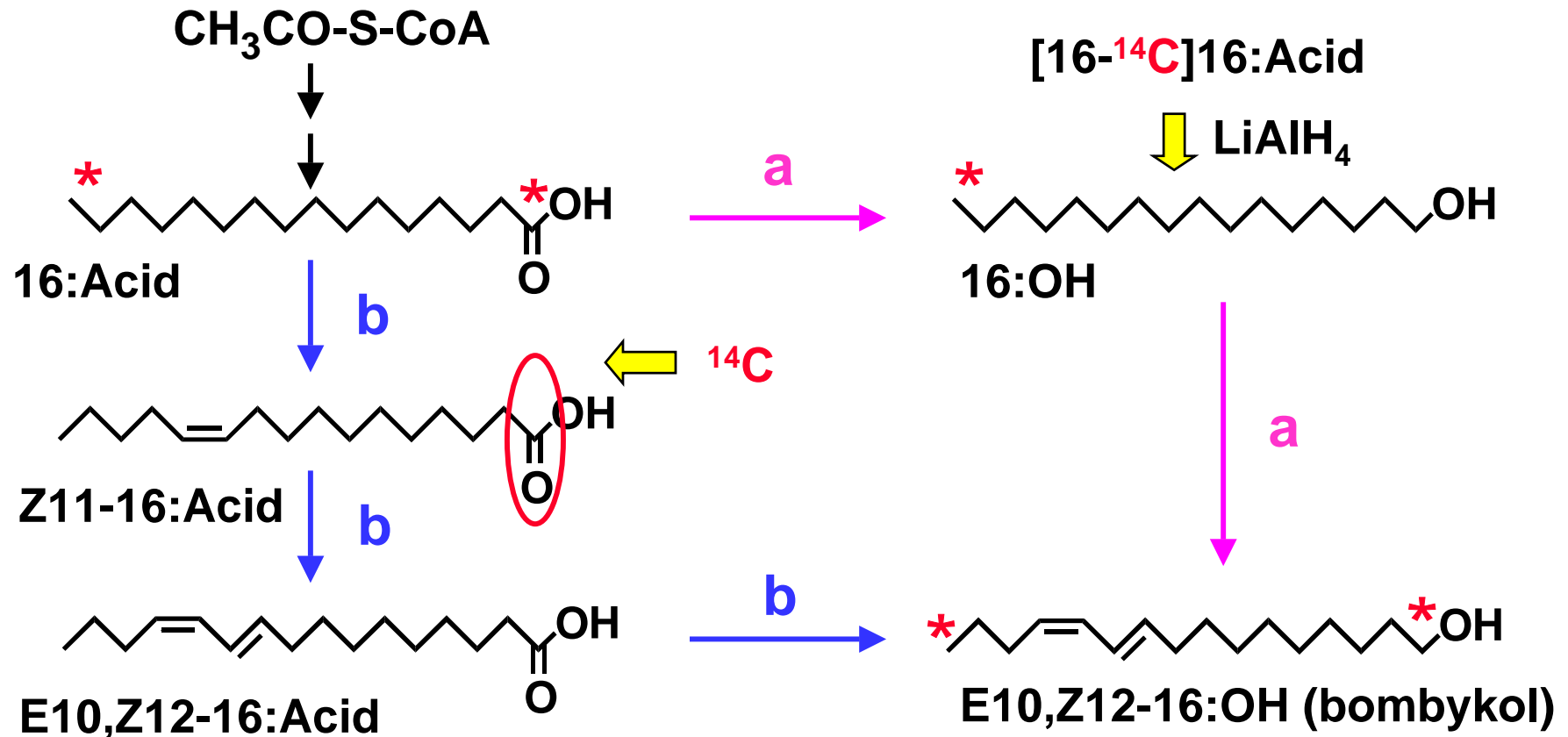


column: ODS (8 mm X 15 cm)
 solvent: MeOH + H₂O (93:7)

A-4) 取込み実験法



A-2) Bombykol (カイコの性フェロモン) の生合成経路



a Inoue & Hamamura, 1972. *Nippon Nogeikagaku Kaishi*, 46, 645

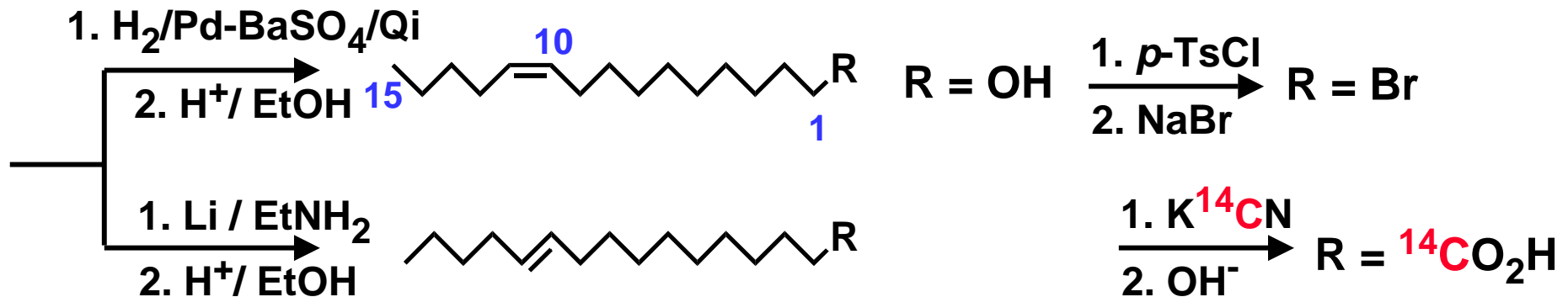
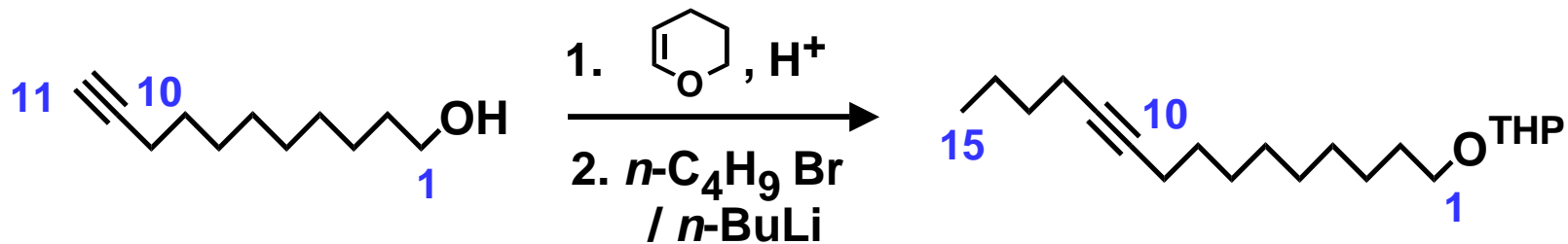
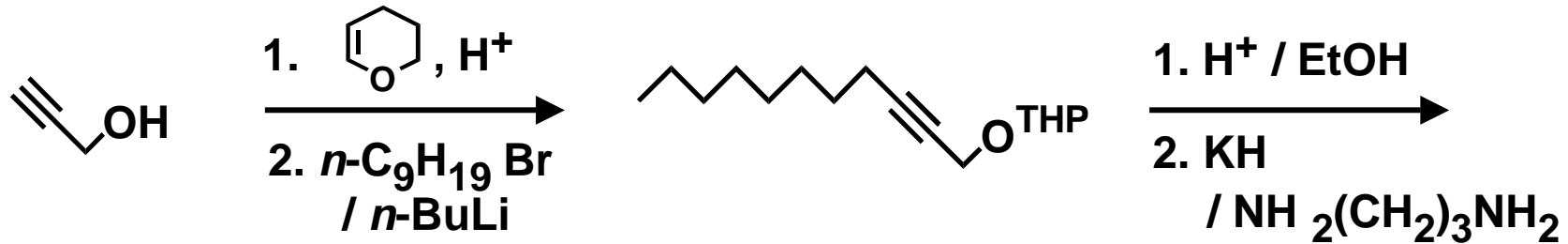
b Yamaoka *et al.*, 1984. *Experientia*, 48, 80

Bjostad & Roelofs, 1984. *Insect Biochem.*, 14: 275

A-5) 標識化合物の合成

鉄則: アイトープの導入は最終段階

Synthesis of [1-¹⁴C](Z)-11-hexadecenoic acid

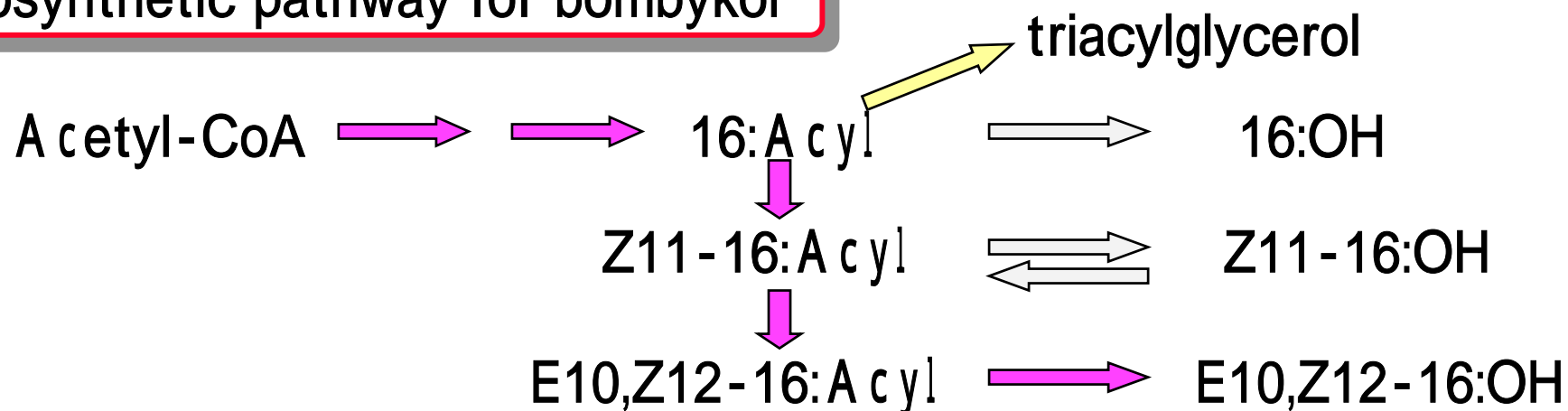


A-6) 取込み量の比較

¹⁴ C-Labeled compound	Incorporation ratio (%)
[a] 12:Acid	0 . 2
14:Acid	0 . 1
16:Acid	1 . 7
18:Acid	0 . 1
[b] 16:OH	0 . 2
Z11-16:OH	0 . 9
Z11-16:Acid	3 . 3

¹⁴ C-Labeled compound	Incorporation ratio (%)
[c] Z10-16:Acid	0 . 3
E10-16:Acid	0 . 4
E11-16:Acid	0 . 2
Z12-16:Acid	0 . 4
E12-16:Acid	0 . 5

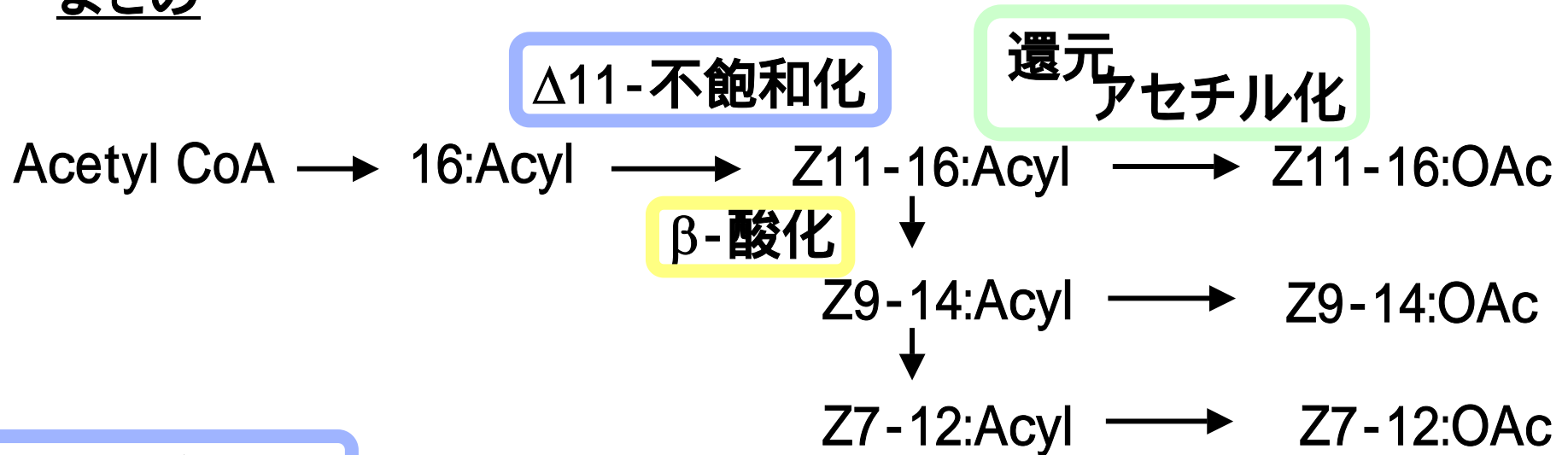
Biosynthetic pathway for bombykol



(B) 蛾類性フェロモン合成

B-1) Type I のフェロモン

まとめ



$\Delta 11$ -不飽和化

イラクサギンウワバにて遺伝子がクローニング

酵母内で発現 (Knipple et al., *ProNAS USA*, 95: 15287, 1998)

β -酸化酵素 ?

還元、アセチル化酵素 低い基質特異性

課題: 不飽和化の位置特異性のメカニズム、混合比の制御

不飽和化酵素

遺伝子の単離 (一覧、その1)

Z9	← 16	<i>Helicoverpa zea</i>	(Noct.)	Rosenfield <i>et al.</i> , 2001
		<i>Helicoverpa assulta</i>	(Noct.)	Jeong <i>et al.</i> , 2003
	← E11-14	<i>Spodoptera littoralis</i>	(Noct.)	Rosenfield <i>et al.</i> , 2004
Z10	← 14	<i>Planotortrix octo</i>	(Tort.)	Hao <i>et al.</i> , 2002(a)
Z11	← 16	<i>Trichoplusia ni</i>	(Noct.)	Knipple <i>et al.</i> , 2001
		<i>H. zea</i>	(Noct.)	Rosenfield <i>et al.</i> , 2001
		<i>H. assulta</i>	(Noct.)	Jeong <i>et al.</i> , 2003
		<i>Bombyx mori</i>	(Bomb.)	Moto <i>et al.</i> , 2004
		<i>Manduca sexta</i>	(Sphi.)	Matouskova <i>et al.</i> , 2007
E11	←14,16	<i>Epiphyas postvittana</i>	(Tort.)	Liu <i>et al.</i> , 2002
	← E9-14	<i>E. postvittana</i>	(Tort.)	Liu <i>et al.</i> , 2002
	← 14	<i>Choristoneura parallela</i>	(Tort.)	Liu <i>et al.</i> , 2004

不飽和化酵素

遺伝子の単離 (一覧、その2)

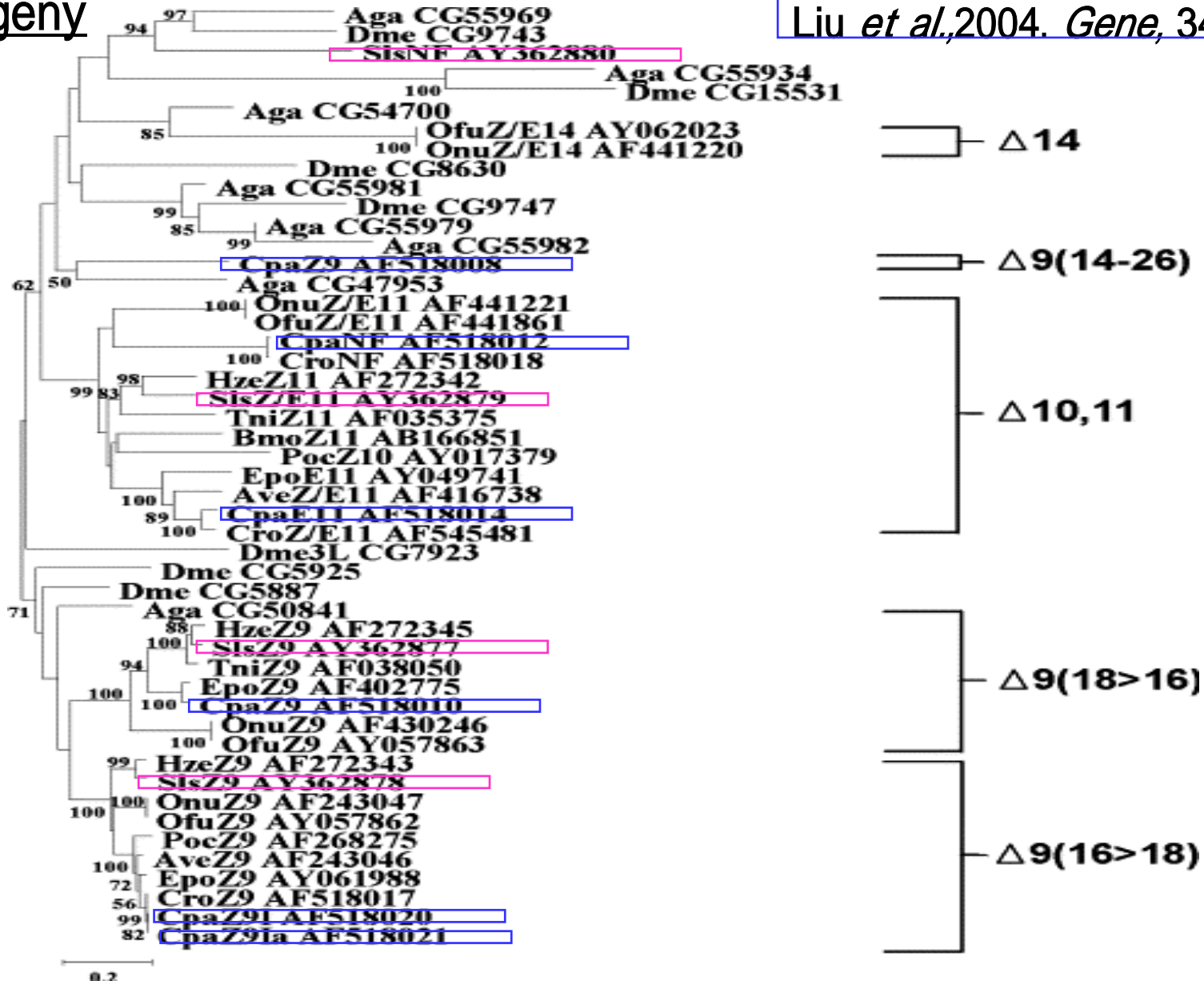
Z/E11	← 14	<i>Argyrotaenia velutinana</i>	(Tort.)	Liu <i>et al.</i> , 2002
		<i>Choristoneura rosaceana</i>	(Tort.)	Hao <i>et al.</i> , 2002(b)
		<i>Ostrinia furnacalis</i>	(Pyra.)	Roelofs <i>et al.</i> , 2002
		<i>Ostrinia nubilalis</i>	(Pyra.)	Roelofs <i>et al.</i> , 2002
		<i>S. littoralis</i>	(Noct.)	Rodriguez <i>et al.</i> , 2004
		<i>Ostrinia scapulalis</i>	(Pyra.)	Fukuzawa <i>et al.</i> , 2006
Z/E14	← 16	<i>O. furnacalis</i>	(Pyra.)	Roelofs <i>et al.</i> , 2002
		<i>O. nubilalis</i>	(Pyra.)	Roelofs <i>et al.</i> , 2002
E10,Z12	← 16	<i>B. mori</i>	(Bomb.)	Moto <i>et al.</i> , 2004
	← Z11-16	<i>M. sexta</i>	(Sphi.)	Matouskova <i>et al.</i> , 2007

不飽和化酵素

Rodriguez *et al.*, 2004. *IBMB* 34: 1315

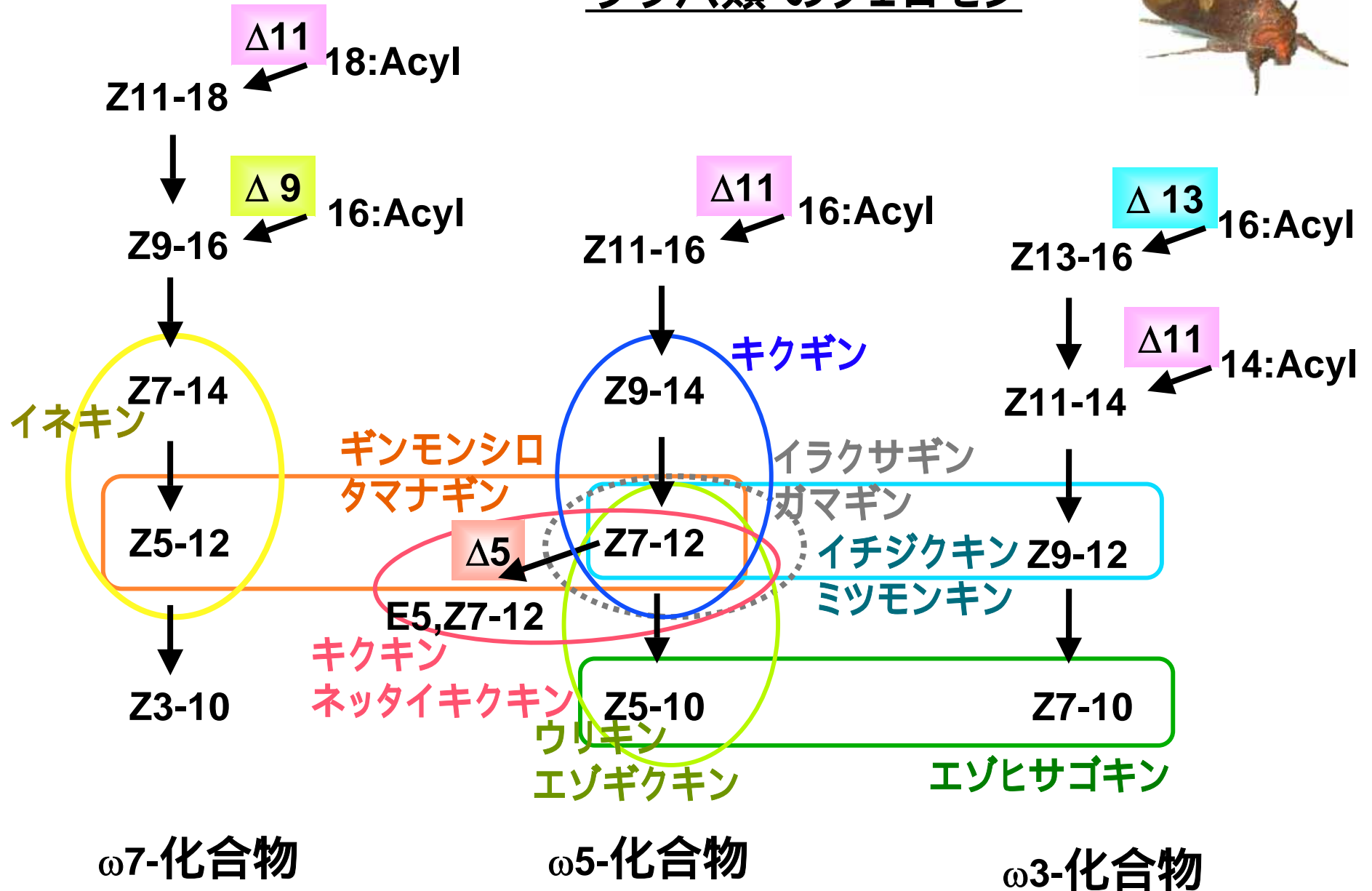
Liu *et al.*, 2004. *Gene*, 342: 303.

Phylogeny



B-1) Type I のフェロモン

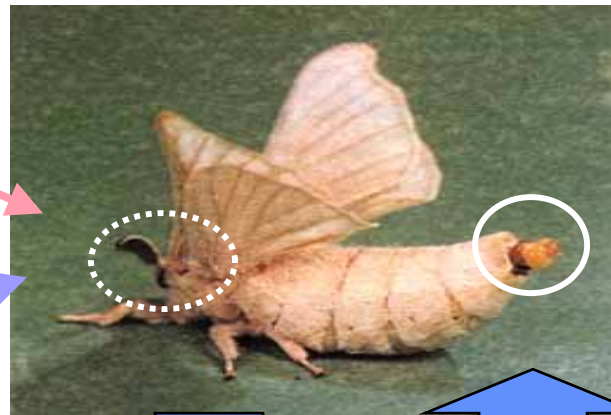
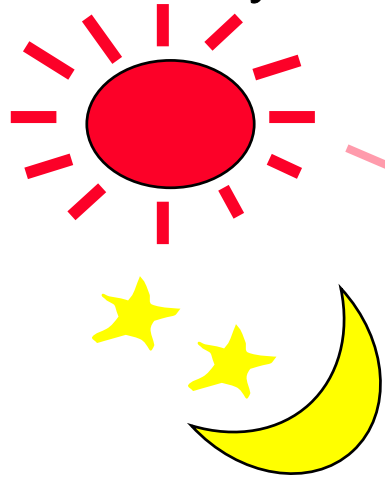
ウワバ類のフェロモン



(B) 生合成の制御機構

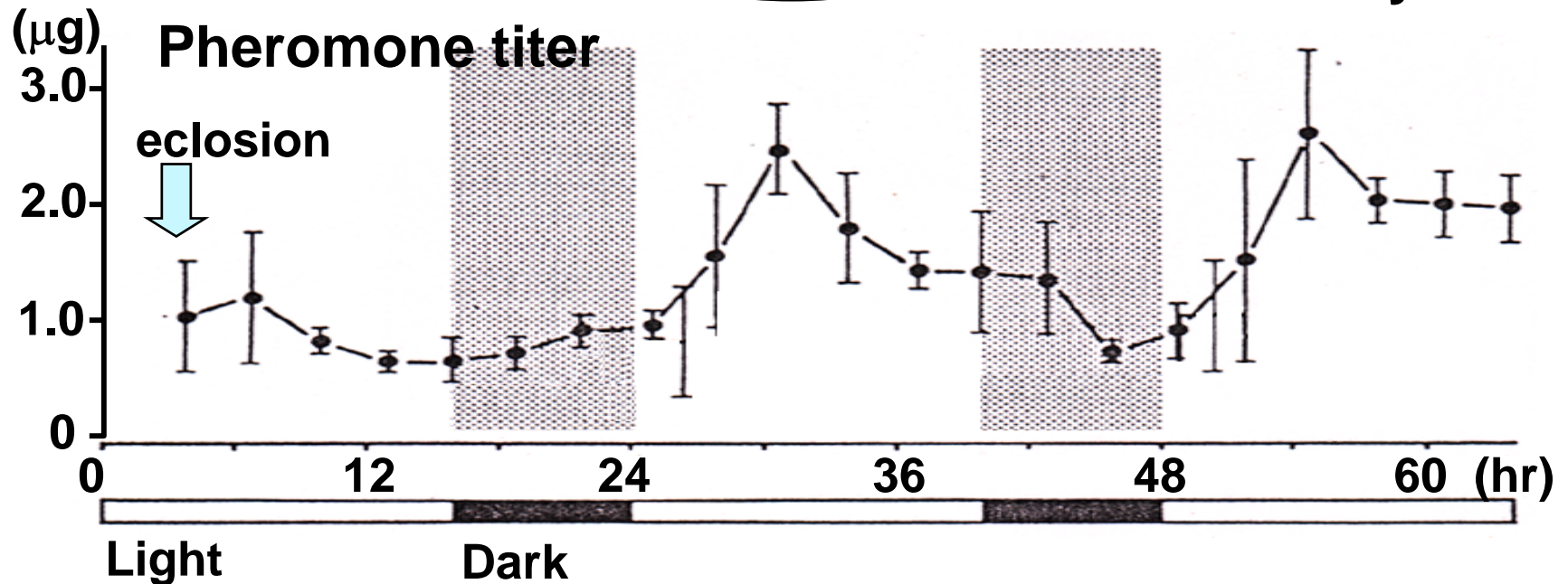
B-1) Introduction

Circadian rhythm



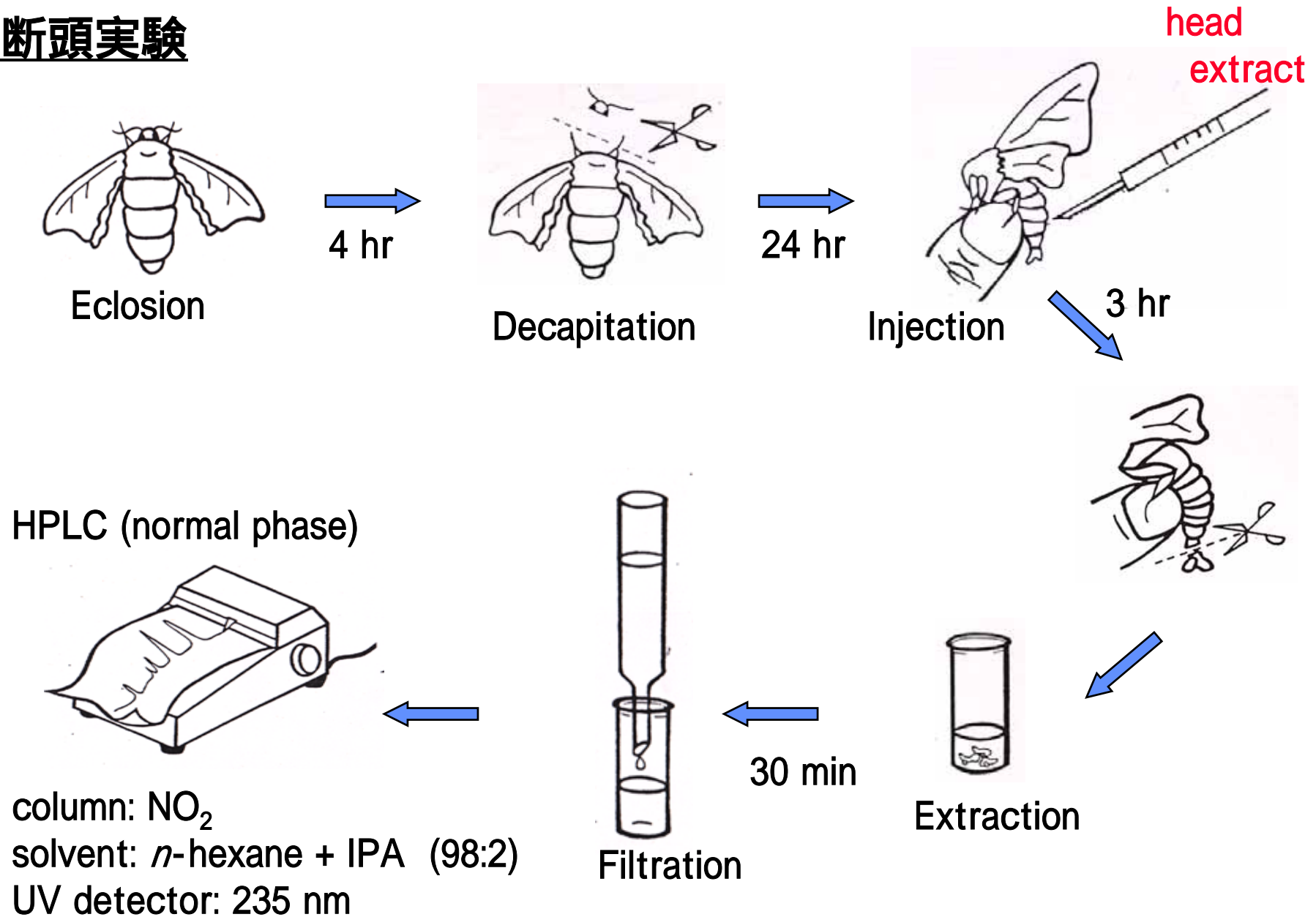
pheromone biosynthesis

nerve system ?
or
endocrine system ?



B-1) Introduction

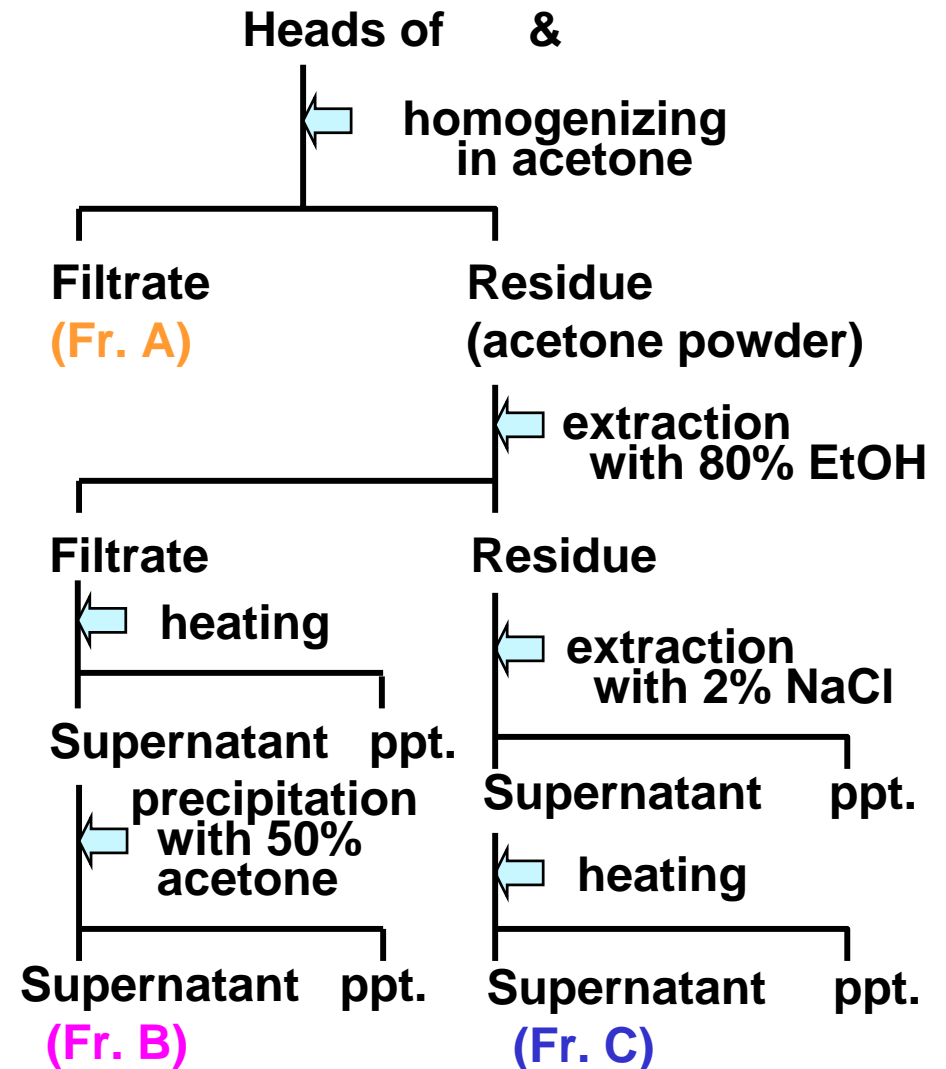
断頭実験



B-1) Introduction

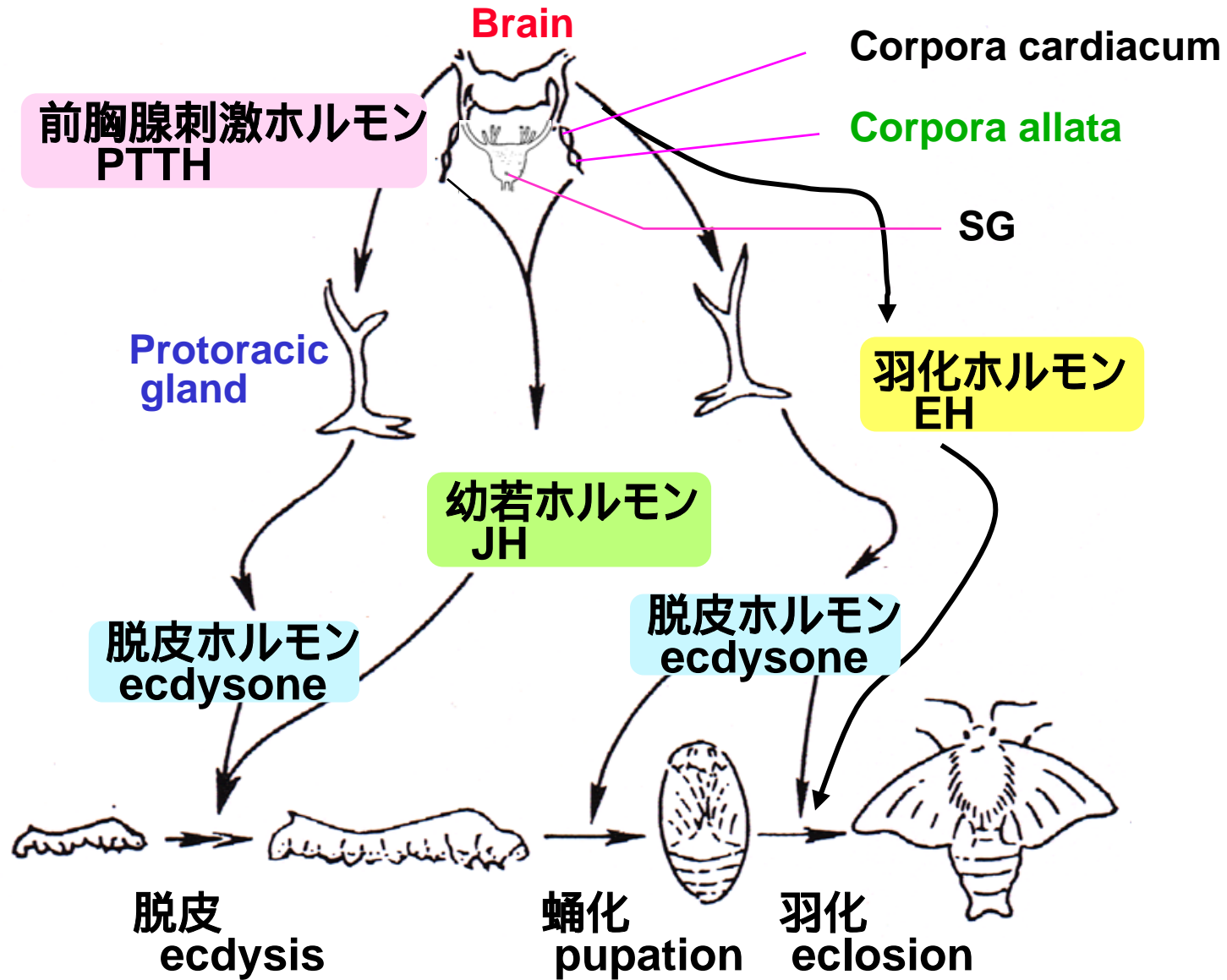
カイコでの予備実験

Treatment	Bombykol ($\mu\text{g/}$)
Exp. 1)	
None	1.31 \pm 0.05
Decapitated	
+ Buffer	0.00 \pm 0.00
+ heads	0.19 \pm 0.07
Exp. 2)	
None	0.87 \pm 0.20
Decapitated	
+ Buffer	0.00 \pm 0.00
+ heads	0.15 \pm 0.05



B-1) Introduction

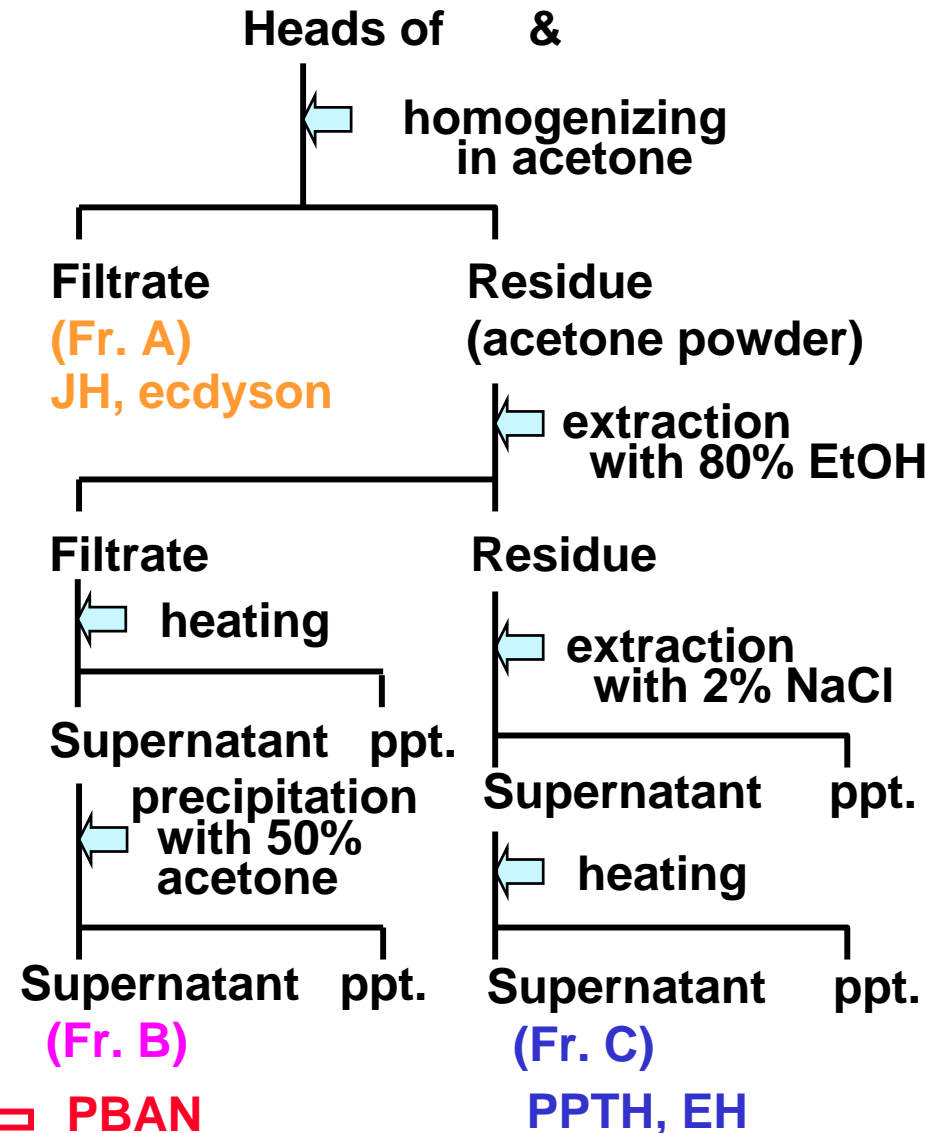
昆虫のホルモン



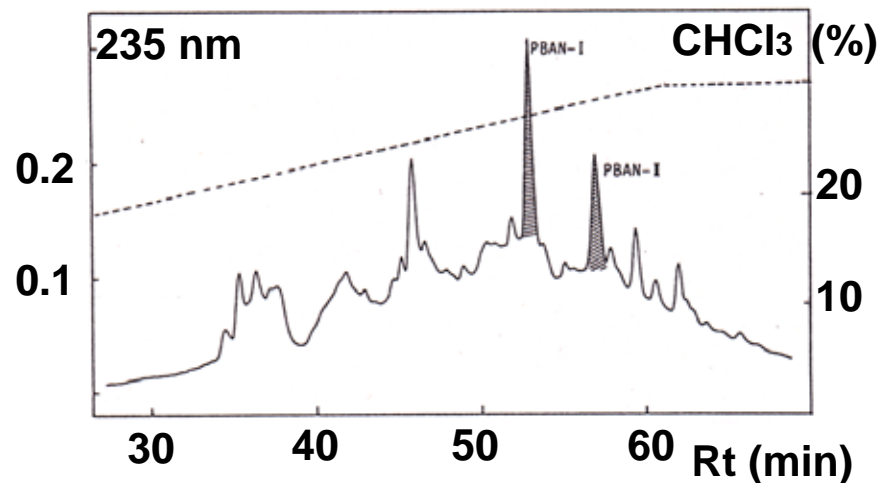
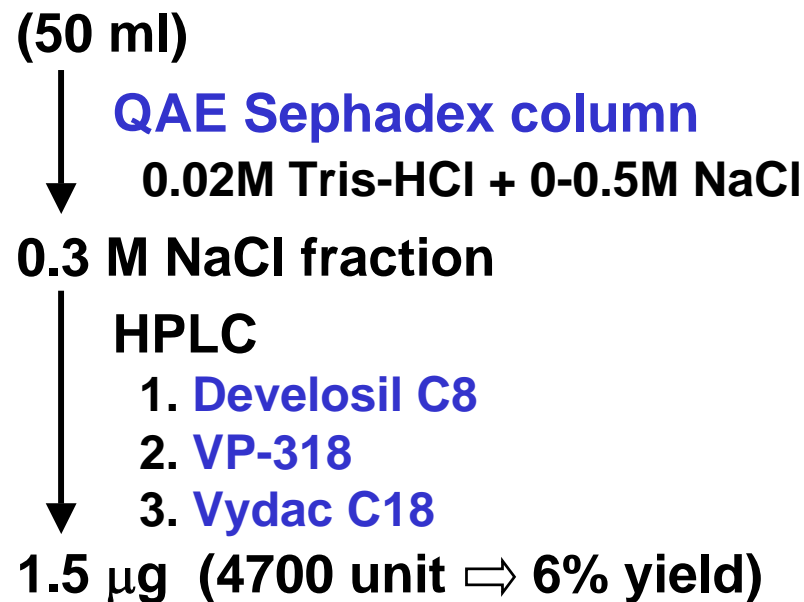
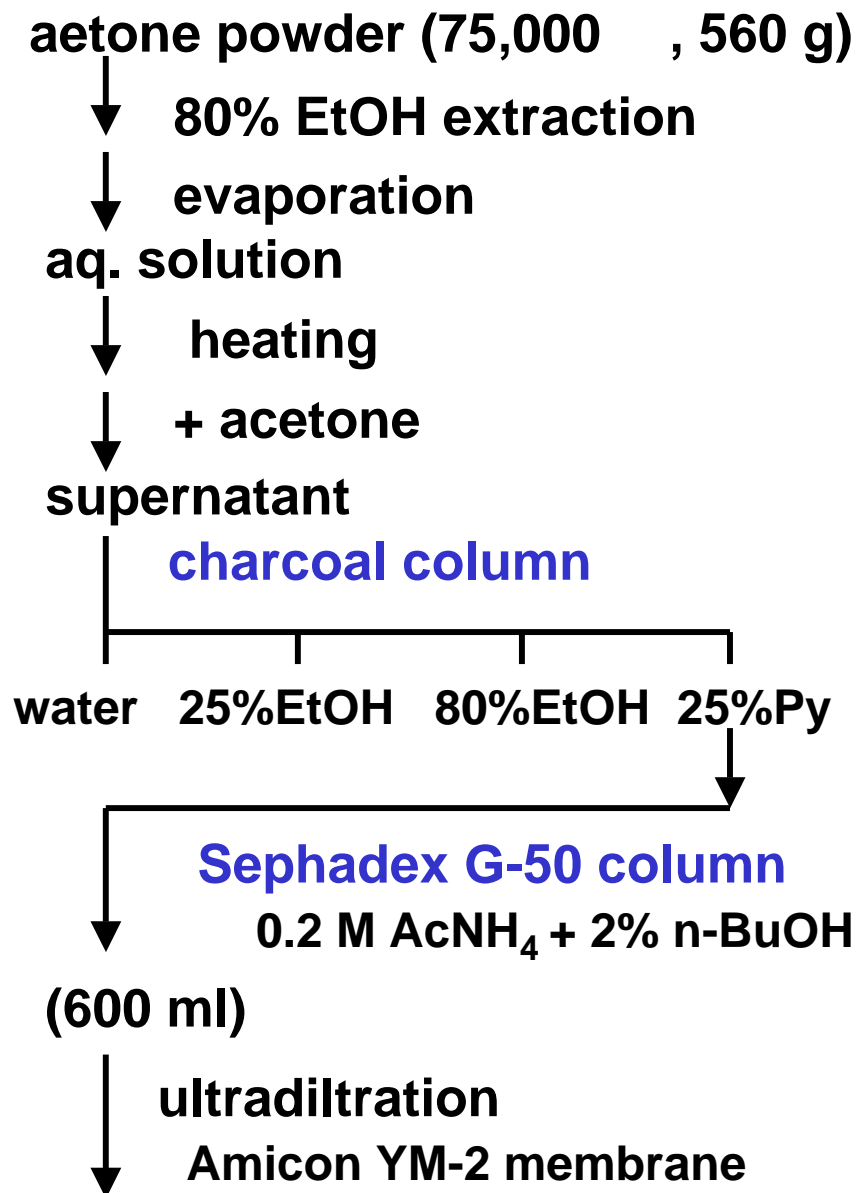
B-1) Introduction

カイコでの予備実験

Treatment	Bombykol ($\mu\text{g/}$)
Exp. 1)	
None	1.31 \pm 0.05
Decapitated	
+ Buffer	0.00 \pm 0.00
+ heads	0.19 \pm 0.07
Exp. 2)	
None	0.87 \pm 0.20
Decapitated	
+ Buffer	0.00 \pm 0.00
+ heads	0.15 \pm 0.05
Exp. 3)	
None	0.69 \pm 0.16
Decapitated	
+ Buffer	0.00 \pm 0.00
+ Fr. A	0.06 \pm 0.04
+ Fr. B	0.24 \pm 0.06
+ Fr. C	0.03 \pm 0.02



B-2) PBANの化学構造



B- 2) PBANの化学構造

FXPRL-amide

	1	5	10	15	
Hez-PBAN	H-Leu	Ser Asp Asp Met	Pro Ala Thr Pro	Ala Asp Gln Glu Met Tyr	Arg Gln-
Bom-PBAN I	H-Leu	Ser Glu Asp Met	Pro Ala Thr Pro	Ala Asp Gln Glu Met Tyr	Gln Pro-
Bom-PBAN II	H-Arg	Leu Ser Glu Asp Met	Pro Ala Thr Pro	Ala Asp Gln Glu Met Tyr	Gln Pro-
Lyd-PBAN	H-Leu	Ala Asp Asp Met	Pro Ala Thr Met	Ala Asp Gln Glu Val Tyr	Arg Pro-
Bom-DH				H -Thr Asp Met Lys Asp Glu Ser Asp-	
Pss-MRCH					H-Lsy Leu-

	20	25	30	33
Hez-PBAN	-Asp Pro Glu Gln Ile Asp	Ser Arg Thr Lys Tyr	Phe Ser Pro Arg	Leu-NH ₂
Bom-PBAN I	-Asp Pro Glu Glu Met Glu	Ser Arg Thr Arg Tyr	Phe Ser Pro Arg	Leu-NH ₂
Bom-PBAN II	-Asp Pro Glu Glu Met Glu	Ser Arg Thr Arg Tyr	Phe Ser Pro Arg	Leu-NH ₂
Lyd-PBAN	-Glu Pro Glu Gln Ile Asp	Ser Arg Asn Lys Tyr	Phe Ser Pro Arg	Leu-NH ₂
Bom-DH	-Arg Gly Ala His Ser Glu Arg	Gly Ala Leu Trp	Phe Gly Pro Arg	Leu-NH ₂
Pss-MRCH	-Ser Tyr Asp Asp Lys Val	Phe Glu Asn Val Glu	Phe Thr Pro Arg	Leu-NH ₂
Lom-MT I		H-Gly Ala Val Pro Ala Ala Gln	Phe Ser Pro Arg	Leu-NH ₂
Lem-PK			pGlu Thr Ser	Phe Thr Pro Arg Leu-NH ₂

Hez-PBAN (Raina *et al.*, 1989), Bom-PBAN I and II (Kitamura *et al.*, 1989, 1990),

Lyd-PBAN I (Masler *et al.*, 1994), Bom-DH (Imai *et al.*, 1991), Pss-MRCH (Matsumoto

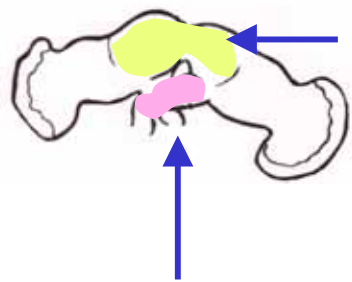
et al., 1992), Lom-MT I and Lem-PK (Nachman and Holman, 1991)

B-3) PBAN研究のまとめ

1. PBANの普遍性: 20種以上の蛾類昆虫で存在が示唆
2. 生産器官: SG (SOG, suboesophageal ganglion、食道下神経球)
3. 標的器官: フェロモン腺 (pheromone gland)
培養フェロモン腺を用いた実験により確認
例外) タバコヤガ *Helicoverpa zea*
VNC (ventral nerve cord)
TAG (terminal abdominal ganglion)
secondary messenger pheromone gland
Teal *et al.*, 1989. *ProNAS USA*, 86: 2488.
4. ペプチドの活性発現部位: アミド化C-末端5残基 (FXPRL)
5. レセプターたん白質: マウスneuromedin U のレセプター
*Drosophila*の遺伝子
タバコヤガ Choi *et al.*, 2004. *ProNAS USA*, 100: 9721.
カイコ Hall *et al.*, 2004. *J. Biol. Chem.*, 279: 51500.
6. 活性化する生合成ステップ: いろいろ?
7. 遺伝子: PBAN と DH の関係?

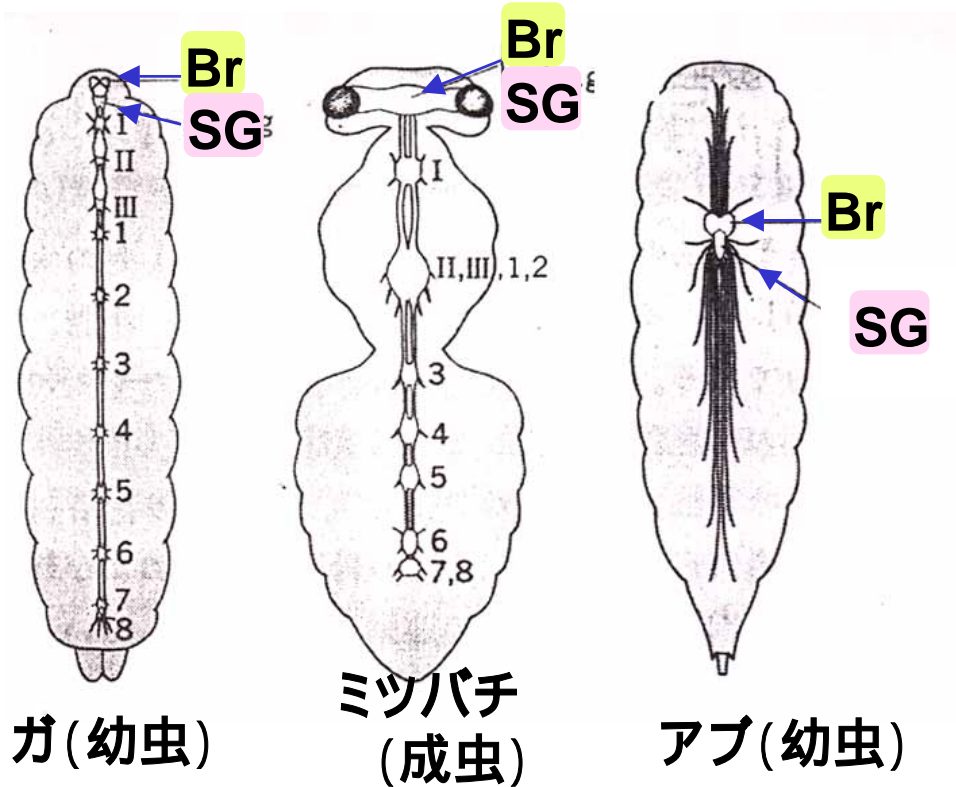
B-3) PBAN研究のまとめ

食道下神経球



脳 (Br)

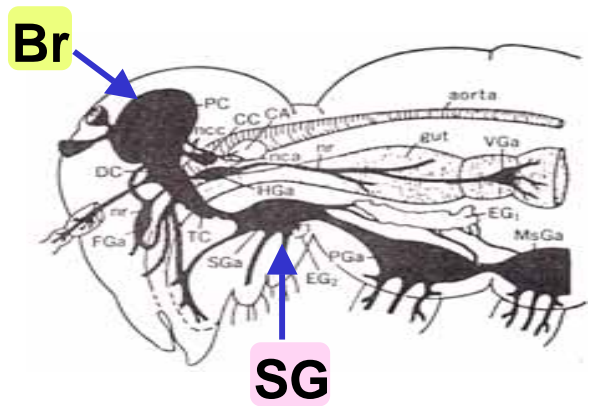
食道下神経球 (SG)



ガ(幼虫)

ミツバチ(成虫)

アブ(幼虫)

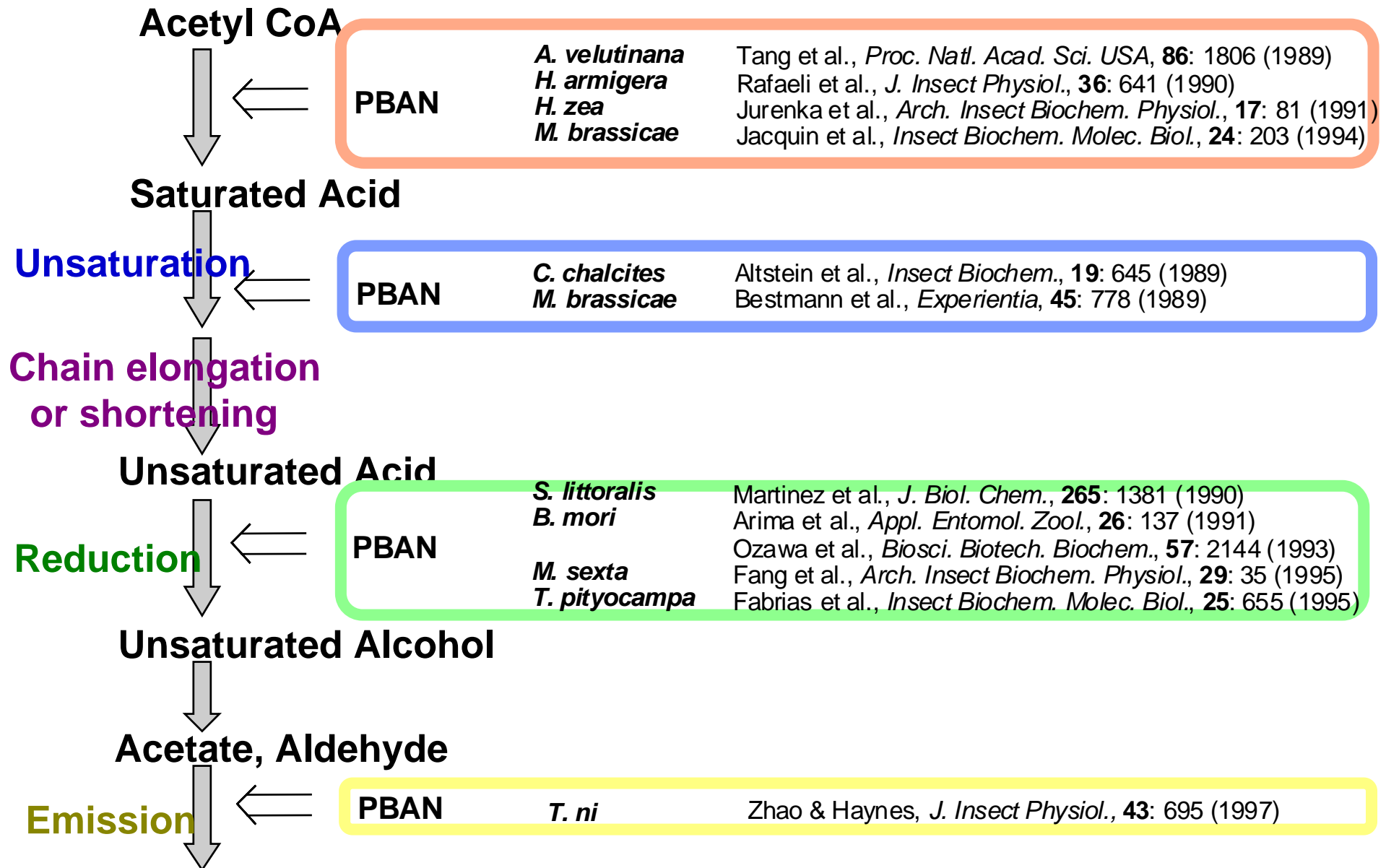


Br

SG

B-3) PBAN研究のまとめ

PBANの活性化ステップ



B-3) PBAN研究のまとめ

カイコPBANcDNAの構造

Kawano *et al.*, 1992. *B. B. R. C.*, 189: 221.

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-40 ATCACTTCGCCCTCCAACCACTGAAGGGACAACAACAAAA ATG TAT AAA ACC AAC ATT GTT TTC AAC GTT TTA GCT TTG
Met Tyr Lys Thr Asn Ile Val Phe Asn Val Leu Ala Leu
40 GCA TTG TTC AGT ATT TTC TTC GCG AGT TGC ACG GAT ATG AAG GAT GAA AGC GAC AGA GGA GCT CAC AGT
Ala Leu Phe Ser Ile Phe Phe Ala Ser Cys Thr Asp Met Lys Asp Glu Ser Asp Arg Gly Ala His Ser
109 GAG CGG GGC GCT CTC TGG TTC GGC CCC AGA CTC GGG AAG CGA TCA ATG AAG CCA TCC ACT GAA GAT AAC
Glu Arg Gly Ala Leu Trp Phe Gly Pro Arg Leu Gly Lys Arg Ser Met Lys Pro Ser Thr Glu Asp Asn
178 AGG CAA ACC TTC CTG AGG CTG CTC GAG GCG GCT GAT GCC CTC AAA TTT TAT TAC GAC CAG CTA CCT TAC
Arg Gln Thr Phe Leu Arg Leu Leu Glu Ala Ala Asp Ala Leu Lys Phe Tyr Tyr Asp Gln Leu Pro Tyr
247 GAG AGG CAA GCC GAT GAA CCG GAA ACC AAA GTA ACA AAG AAG ATC ATC TTC ACC CCC AAA CTC GGG AGG
Glu Arg Gln Ala Asp Glu Pro Glu Thr Lys Val Thr Lys Lys Ile Ile Phe Thr Pro Lys Leu Gly Arg
316 AGC GTC GCC AAA CCC CAG ACG CAT GAA AGC CTC GAA TTC ATC CCC CGG CTC GGA AGG CGG CTC TCT GAG
Ser Val Ala Lys Pro Gln Thr His Glu Ser Leu Glu Phe Ile Pro Arg Leu Gly Arg Arg Leu Ser Glu
385 GAC ATG CCT GCT ACG CCA GCT GAC CAG GAA ATG TAC CAA CCT GAC CCC GAA GAA ATG GAG TCA AGA ACA
Asp Met Pro Ala Thr Pro Ala Asp Gln Glu Met Tyr Gln Pro Asp Pro Glu Glu Met Glu Ser Arg Thr
454 AGA TAC TTC TCG CCC AGG CTG GGG CGC ACC ATG AGC TTT TCG CCC AGA CTG GGA AGG GAG CTT TCG TAC
Arg Tyr Phe Ser Pro Arg Leu Gly Arg Thr Met Ser Phe Ser Pro Arg Leu Gly Arg Glu Leu Ser Tyr
523 GAT TAC CCT ACA AAA TAT AGG GTT GCC AGA AGC GTT AAC AAG ACA ATG GAC AAC TAAACGAATTATGGTCCG
Asp Tyr Pro Thr Lys Tyr Arg Val Ala Arg Ser Val Asn Lys Thr Met Asp Asn ***
595 CTTGAGGTACCTCATTGAGGTCTCGATCGACTCCGACGAACGGTTACGGGTAAACGGCGACAATGTTAATGTTTTGGACGAAACAATTG
685 TTAATTATAAAATTCATGTGATTTTGTAAATTGTAATTTATAAGTATAAAATAAACTATTTAAAAT(polyA)
```

DH

α

β

PBAN

γ

タバコガ Davis *et al.*, 1992. *ProNAS USA*, 89: 142.

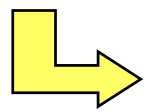
B-4) シャクガでのPBAN研究

末端官能基がない (約 15%)

ヨモギエダシャク : Japanese giant looper (*Ascotis selenaria cretacea*)

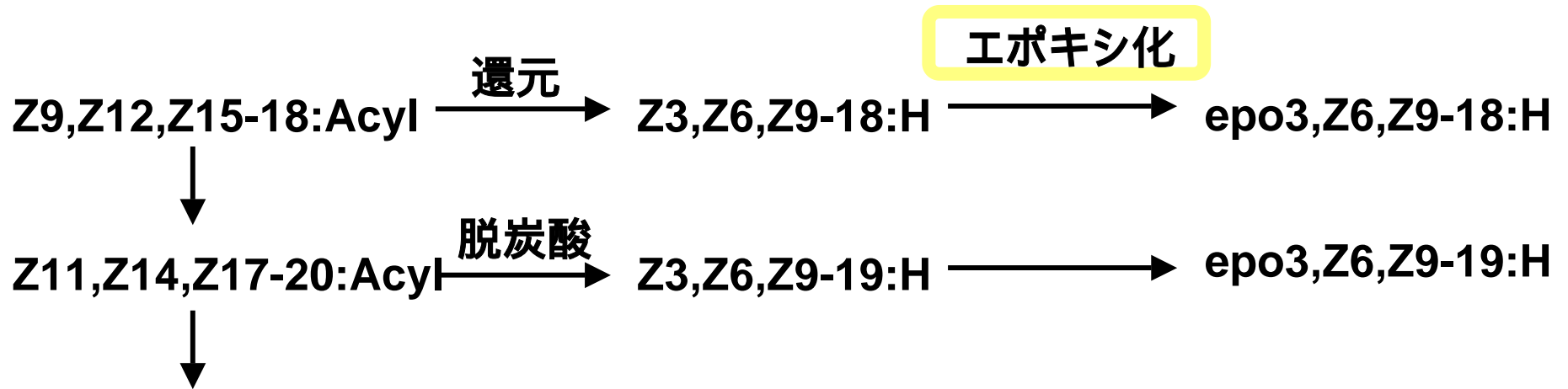


スキバドクガ : clear-winged tussock moth (*Perina nuda*)



- i 不飽和炭化水素とそのエポキシ化物
- ii 直鎖の炭素数: 17 ~ 23
- iii 二重結合数: 1 ~ 4
- iv リノール酸、リノレン酸から生合成される

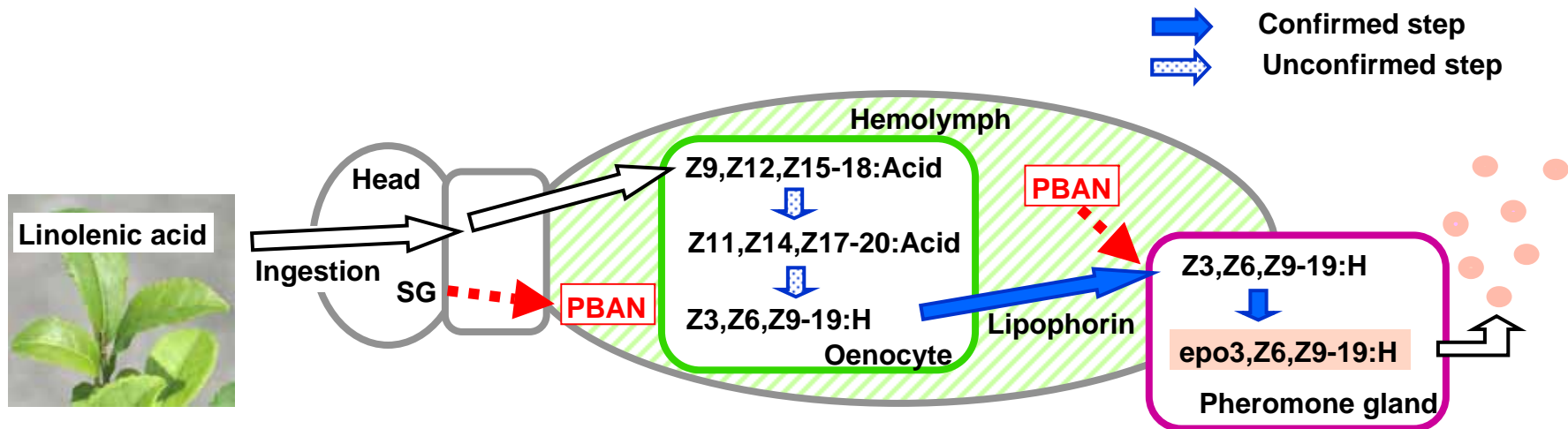
B-4) シャクガでのPBAN研究



還元酵素 ? 脱炭酸酵素 ? 長鎖脂肪酸: 未同定

エポキシ化酵素 (P₄₅₀-dependent mono-oxygenase)

反応の位置選択性: 高い 基質特異性: 低い



B-4) シャクガでのPBAN研究

Degenerateプライマーを用いたPCR

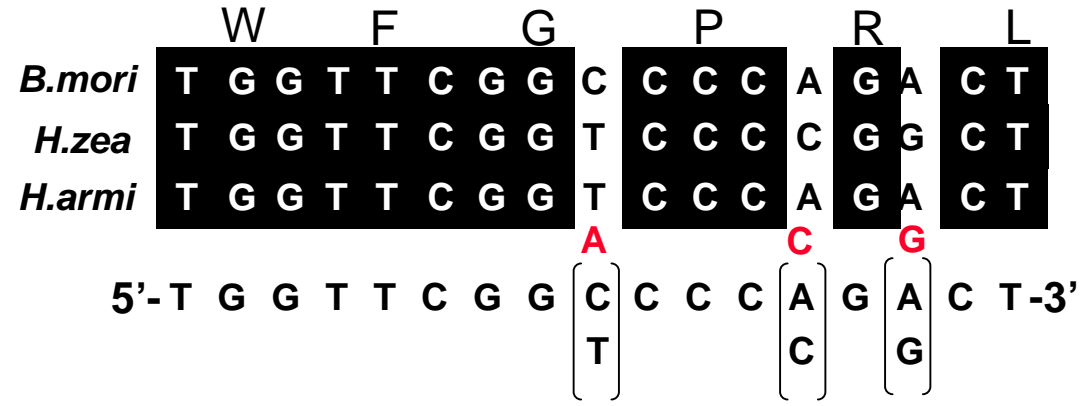
DH

MYKTNIVFNVLALALFSIFFASCTDMKDESDRGAHSERGALW**FGPRL**GKR
 SMKPSTEDNRQTFLRLLEAADALKFYDQLPYERQADEPETK VTKK **II FT**
PKLGRSVAKPQTHESLE**FIPRL**GRRLSEDMPATPADQEMYQPDPEEMESR

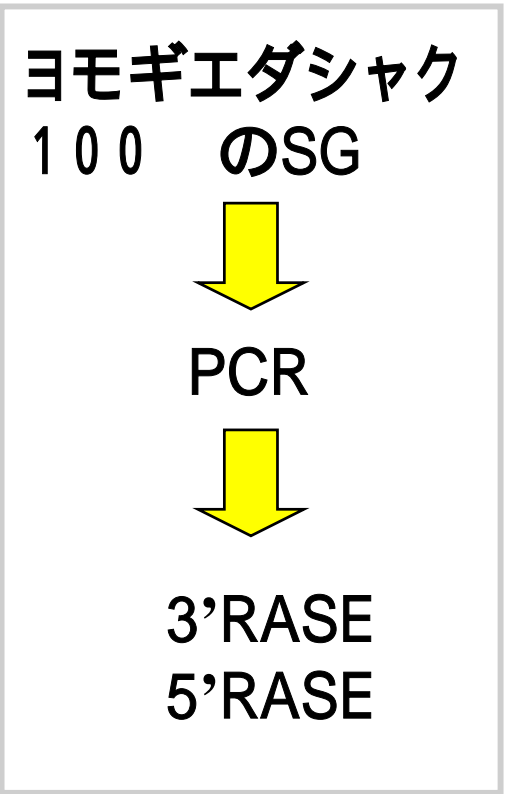
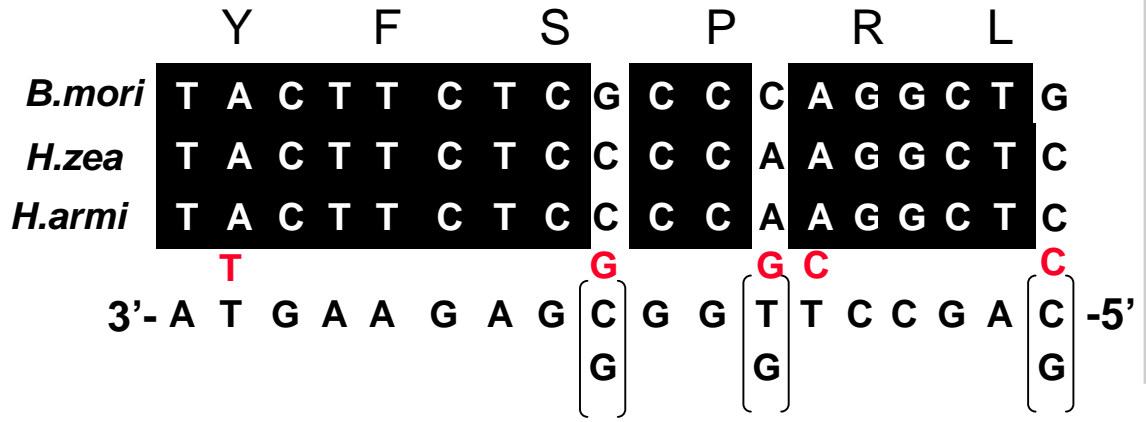
PBAN

TRY**FSPRL**GRTMS**FSPRL**GRELSYDYPTKYRVARSVNKTMDN

DH



PBAN



B-4) シャクガでのPBAN研究

ヨモギエダシャクのPBANの構造

DH homologue

Assc	NDLKEDG-EREANSDRQLWFGPRL	
Bom	TDMKDES-DRGAHSERGLWFGPRL	59%
Hez	NDVKDGA-ASGAHSDRLLGLWFGPRL	68% (62%)
Hev	NDDKDGA-ASGAHSDRLLGLWFGPRL	68% (62%)
Mas	NDIKDEG-DRGAHSDRGALWFGPRL	69% (83%)
Ads	N-FKEENFDRNIRSGRANVVEKPIIL	37% (40%)

α-SGNP

Assc	VIFTPKL	
Bom	IIFTPKL	85%
Hez	VIFTPKL	100% (85%)
Hev	VIFTPKL	100% (85%)
Mas	VIFTPKL	85% (71%)
Ads	VIFTPKL	100% (85%)

β-SGNP

Assc	S-----VDFTPRL	
Bom	SVA-----KPQTHESLEFIIPRL	35%
Hez	SLA--YDDK-SF-ENVEFTPRL	38% (45%)
Hev	SLS--YDDK-SF-ENVEFTPRL	38% (40%)
Mas	SLDDSTQEKRVFYENEFTPRL	31% (36%)
Ads	SMEDPYEEKRSY-D-VDFTPRL	40% (38%)

PBAN homologue

Assc	QLVDDVPQRQIEEDRL-----GSRTRFSPRL	
Bom	-LSEDMPATPADQE--IYQPDPEEMESR-TRYFSPRL	44%
Hez	-LSDMPATPADQE--MYRQDPEQIDSR-TKYFSPRL	46% (76%)
Hev	-LADDMPATPADQE--MYRQDPEQIDSRRTKYFSPRL	45% (71%)
Mas	-ISEDMPATPSDQEYPMYHPDPEQIDTR-TRYFSPRL	32% (71%)
Ads	Q-----SEAVTSSDEQVYRQDMSPVDGR-LKYFSPRL	40% (45%)

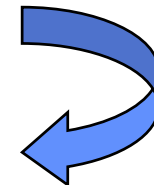
γ-SGNP

Assc	TTMNFSPRL	
Bom	T-MSFSPRL	77%
Hez	T-MNFSPRL	88% (87%)
Hev	T-MNFSPRL	88% (87%)
Mas	T--HFSPRL	66% (75%)
Ads	ANVVEKPIIL	36% (37%)

**b-SGNP、PBAN homologue とともに異常に短い
両者の開裂部はGR (他種はすべてGRR)**

PBAN

Assc	SVDFTPRLGRQLVDDVPQRQIEEDRLGSRTRFSPRL	
Bom (PBAN I)	LSEDMPATPADQE--IYQPDPEEMESR-TRYFSPRL	42%
Hez	LSDMPATPADQE--MYRQDPEQIDSR-TKYFSPRL	44% (76%)
Lyd	LADDMPATPADQE--VYRPEPEQIDSRN-KYFSPRL	42% (66%)



B-4) シャクガでのPBAN研究

PBAN遺伝子に関する系統樹

