Directed Biosynthesis of Peptaibol Antibiotics in Two Trichoderma Strains

I. Fermentation and Isolation

GREGORY LECLERC, SYLVIE REBUFFAT*, CHRISTOPHE GOULARD and BERNARD BODO

Laboratoire de Chimie des Substances Naturelles, URA 401 CNRS, GDR 790 CNRS, Muséum National d'Histoire Naturelle, 63 rue Buffon, 75231 Paris Cedex 05, France

(Received for publication August 15, 1997)

Peptaibols are linear α -aminoisobutyric acid-containing peptide antibiotics originating from soil fungi mainly of the genus Trichoderma and biosynthesized in complex mixtures of closely related analogues by a polyenzymatic pathway. Addition of amino acids such as α -aminoisobutyric acid (Aib), glutamic acid or arginine, to the fermentation medium of two Trichoderma strains, T. harzianum and T. longibrachiatum, has been shown to result in the simplification of the natural peptaibol mixtures, leading in each case to the almost exclusive biosynthesis of a single peptide. Surprisingly, the obtained peptides are Aibenriched, whether the added amino acid is Aib, Glu or Arg. By adding Aib to the fermentation medium of T. harzianum, two new Aib-rich peptaibols were isolated. Moreover, adding glutamic acid to the culture medium of T. longibrachiatum, which produces both neutral and acidic 20-residue peptaibols with either glutamine or glutamic acid at position 18, increases the production of the acidic peptides. However, arginine which is a positively charged amino acid generally absent from peptaibol sequences, is not incorporated in trichorzins when added to the fermentation medium of T. harzianum.

Trichoderma are widespread soil fungi characterized by the production of peptides which display antifungal and antibacterial activities. Belonging to the peptaibol class, these peptides interact with phospholipid bilayers, increasing their permeability properties^{1,2)} and forming voltage-dependent ion channels.2~4) With a high proportion of α,α -dialkylated amino acids, such as α aminoisobutyric acid (Aib, U) and isovaline (Iva, J), peptaibols are linear peptides which exhibit an Nterminal acylated residue and a C-terminal amino alcohol. According to their chain length and chemical characteristics, they are classified into three subclasses: the long-sequence peptaibols including the well-known alamethicin⁵⁾ with $18 \sim 20$ residues, $6^{\circ 9}$ the shortsequence peptaibols with $11 \sim 16$ residues^{9 ~ 12}) and the lipopeptaibols with 7 or 11 residues and an N-terminal lipid chain. 13,14)

As frequently observed, peptide antibiotics originating from microbial organisms, bacteria and filamentous fungi, arise from a nonribosomal biosynthetic pathway implying large protein templates termed peptide synthetases. 15~17) The alamethicin synthetase extracted

from *T. viride* has a molecular weight around 480 kDa; the *N*-acetyl and the *C*-terminal amino alcohol linkage steps have been studied in detail. Such biosyntheses proceed by formation of a linear peptidyl intermediate, the termini of which may be either enzymatically modified or transformed into cyclic structures, as observed for the potent immunosuppressant, cyclosporin A. Seneral observation for such biosynthesis mechanism is that the peptides are produced in mixtures of closely related sequence analogues, termed microheterogeneous mixtures and particularly exemplified by peptaibols. 11,20)

Previous studies on cyclosporins biosynthesis have shown that the course of their formation is strongly influenced by an exogenous supply of amino acids to the fermentation medium, resulting either in enhanced yields of some of the natural components at the expense of others, or in the incorporation of foreign amino acids, depending on the added amino acid. ^{21,22} Similarly, the biosynthesis of the Aib-containing efrapeptins by *Tolypocladium geodes* is affected by the amino acid composition of the culture medium. ²³ However, such experiments have never been tried with peptaibols. We

thus analyzed the influence of amino acid supplies on the biosynthesis course of *Trichoderma* strains, in an attempt to incorporate exogenous amino acid and to synthesize new analogues, which could exhibit modified membrane properties.

For the present study, we selected two Trichoderma strains which had been previously studied for their peptide content and considered the influence of an Aib-supply on the biosynthesis of the 20-residue longibrachins by Trichoderma longibrachiatum²⁴ (Fig. 1-A) and of the 18-residue trichorzins PA (Fig. 1-B) and the 14-residue harzianins PC by Trichoderma harzianum. 25) We here describe the modifications of the microheterogeneous peptide mixtures composition arising from Aib-directed biosynthesis in these two Trichoderma strains. We also report on the effect of supplementing the standard growth medium of T. longibrachiatum and T. harzianum with charged amino acids such as glutamic acid and arginine, glutamic acid being a natural component of peptaibols, while the positively charged arginine is usually absent in these peptides.

Results and Discussion

Aib-directed Biosynthesis of Peptaibols

As Aib is the peculiar amino acid responsible for the special characteristics and properties of peptaibols, the Aib-directed biosynthesis of the selected peptaibols was

first examined. Fermentation of T. longibrachiatum (M-853431) on a synthetic medium led to a crude peptaibol mixture, isolated from the culture broth extract by exclusion chromatography over Sephadex LH20. When submitted to silica gel chromatography, the crude peptide fraction provided two peptide groups, designated as longibrachins LGA (neutral) and LGB (acidic) (Table 1). The LGA and LGB HPLC profiles obtained on a C_{18} reversed-phase column showed four (LGA I \sim IV) and five (LGB I \sim V) components, respectively (Fig. 2-A, 2-C). The sequences of the major longibrachins (LGA I \sim IV and LGB II, III) have been determined previously (Fig. 1-A). 24)

When *T. longibrachiatum* was grown on a 0.8% w/v Aib-supplied medium and the peptide mixture separated according to the usual procedure, two peptide groups were also obtained (Table 1) and termed longibrachins LGA_U and LGB_U. Regarding their HPLC profile, they appeared much simpler, each characterized by one main peptide, LGA_U 1 (Fig. 2-B) and LGB_U 2 (Fig. 2-D). Both longibrachins were isolated by repetitive semi-preparative HPLC and submitted to a co-chromatography HPLC study in order to assign LGA_U 1 and LGB_U 2, as compared to the LGA and LGB natural mixtures obtained on the normal culture medium (Table 2). The HPLC results indicated LGA_U 1 to be LGA I and LGB_U 2 to be LGB II, in agreement with the NMR and MS data.

Fig. 1. Amino acid sequences.

A: Longibrachins LGA and LGB from T. longibrachiatum.

```
LGA I Ac Aib Ala Aib Ala Aib Ala Gln Aib Val Aib Gly Leu Aib Pro Val Aib Aib Gln Gln Pheol LGA II Ac Aib Ala Aib Ala Aib Ala Gln Aib Val Aib Gly Leu Aib Pro Val Aib Iva Gln Gln Pheol LGA IV Ac Aib Ala Aib Ala Aib Ala Aib Ala Gln Aib Val Aib Gly Leu Aib Pro Val Aib Iva Gln Gln Pheol LGA IV Ac Aib Ala Aib Ala Aib Aib Gln Aib Val Aib Gly Leu Aib Pro Val Aib Iva Gln Gln Pheol LGB II Ac Aib Ala Aib Ala Aib Ala Gln Aib Val Aib Gly Leu Aib Pro Val Aib Iva Gln Gln Pheol LGB III Ac Aib Ala Aib Ala Aib Ala Gln Aib Val Aib Gly Leu Aib Pro Val Aib Aib Glu Gln Pheol LGB III Ac Aib Ala Aib Ala Aib Ala Gln Aib Val Aib Gly Leu Aib Pro Val Aib Iva Glu Gln Pheol LGB III Ac Aib Ala Aib Ala Aib Ala Gln Aib Val Aib Gly Leu Aib Pro Val Aib Iva Glu Gln Pheol LGB III Ac Aib Ala Aib Ala Aib Ala Gln Aib Val Aib Gly Leu Aib Pro Val Aib Iva Glu Gln Pheol
```

B: Trichorzins PA and PA_U 4 and harzianin PC_U 4 from T. harzianum.

```
PA II
PA IV
PA V
PA V
PA VI
PA VI
PA VI
PA VI
PA VII
PA VII
PA VIII
PA
```

PAU 4 Ac Aib Ser Ala Aib Aib Gln Aib Val Aib Gly Leu Aib Pro Leu Aib Aib Gln Trpol

PCU 4 Ac Aib Asn Leu Aib Pro Ser Ile Aib Pro Aib Leu Aib Pro Valol

Bold-faced letters indicate those amino acids which differ in the sequences; Aib, U: α-aminoisobutyric acid, Iva, J: isovaline, Pheol, Fol: phenylalaninol, Trpol, Wol: tryptophanol, Valol, Vol: valinol.

Table 1. Weights of peptaibols (mg) for typical T. longibrachiatum (M-853431) 20 liter cultures performed on normal culture medium (NM) and on the same culture medium supplemented either with Aib $(NM+U)^a$ or with Glu $(NM+E)^a$.

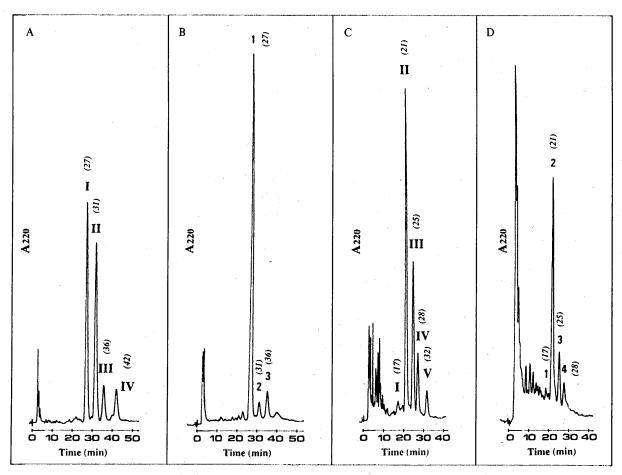
	NM (mg)	NM+U (mg)	NM + E (mg)
Crude extract	2120	1500	1610
Crude peptide mixture ^b	140	95	51
LGA	40	13	9
LGB	58	50	22

^a The culture medium was supplemented with 8 g/liter of the concerned amino acid.

Using the same multistep chromatography procedure as described above, fermentation of *T. harzianum* (M-902608) led to three main peptide groups of different polarity designated as PC, PD and trichorzins PA (Table 3). HPLC analysis of the PA mixture was performed on a C₁₈ reversed-phase column, exhibiting at least nine components (Fig. 3-A, Table 4), while PC and PD mixtures, which are not described here, were more complex. The sequence of the isolated trichorzins PA (Fig. 1-B) have been determined in a previous work.²⁵⁾

When the medium was Aib-enriched, the T. harzianum strain led only to two peptide groups, a minor group termed harzianins PC_U and a major group termed trichorzins PA_U (Table 3), revealing the simplification of the biosynthesized peptide mixture and the complete inhibition of the PD group production. HPLC analysis of the PA_U mixture showed mainly one peptide PA_U 4 and few minor components (Fig. 3-B). After purification by repetitive semi-preparative HPLC, the different tri-

Fig. 2. HPLC chromatograms of the longibrachins LGA (A), LGA_U (B), LGB (C), LGB_U (D), from *T. longibrachiatum*.



Kromasil C₁₈ (5 μ), 4.6 × 250 mm, flow rate 1 ml/minute, absorption monitored at 220 nm. Solvent systems for A and B: MeOH - H₂O, 85:15; for C and D: MeOH - H₂O, 86:14. Retention times of the peptides are given in brackets.

b The crude peptide mixtures contained both peptides and others metabolites.

Table 2. HPLC retention times (minute) of the longibrachins LGA, LGA_U, LGA_E and LGB, LGB_U, LGB_E from *T. longibrachiatum* (M-853431).

Peptides		LGA ^a			LGA _U /LGA _E ^a				LGB ^b					LGB _U /LGB _E ^b			
replices	I	II	III	IV	1	2	3	Ι.	II	III	IV	V	1	2	3	4	
Rt (minute)		31	36	42	27		36			25	28	32	17	21	25	28	

^a Solvent system: MeOH-H₂O (85:15).

b Solvent system: MeOH-H₂O, TFA 0.05% (86:14).

Table 3. Weights of peptaibols (mg) obtained for typical T. harzianum (M-902608) 20 liter cultures performed on normal culture medium (NM) and on the same culture medium supplemented either with Aib $(NM+U)^a$ or with Arg $(NM+R)^a$.

	NM (mg)	NM+U (mg)	NM + R (mg)
Crude extract	1670	1460	1370
Crude peptide mixture ^b	920	233	255
PC	170	43	
PA	347	115	76
PD	168	-	-

^a The culture medium was supplemented with 8 g/liter of the concerned amino acid.

chorzins PA_U were co-injected with the natural mixture of known trichorzins PA obtained on the standard culture medium. That indicated PA_U 4 and PA_U 6 to exhibit the same retention time as PA I and PA III respectively (Table 4), as shown by a significant enhancement in their peak intensity. As both PA I and PA III peptides have been previously shown to be heterogeneous and each composed of at least two homologous peptides, from MS and NMR data, 25) the present data suggested PAU 4 and PA_U 6 to be one of the two components making up the PA I and PA III peaks, respectively. In addition, new peptides were produced; three of them, PA_U 1, PA_U 2 and PA_U 3, exhibited shorter retention time than those of the trichorzins PA obtained on the normal medium (Fig. 3-B, Table 4); less hydrophobic than trichorzins PA, they appeared original but were not isolated in sufficient amount to examine their structures. Moreover, HPLC analysis of the PC_U mixture showed more than

12 peaks (Fig. 4); among the 12 peptide fractions isolated by further semi-preparative HPLC, only one corresponding to PC_U 4 was homogeneous, as shown by further analyses (see the following paper).

As exemplified by the two studied *Trichoderma* strains, these results give evidence that an exogenous Aibsupplied medium results in peptaibol biosynthesis modifications, leading to peptide mixture simplification. Indeed, the addition of Aib to the *T. harzianum* strain cultures led to the almost exclusive biosynthesis of a single peptide, in which all Iva residues have been replaced by Aib.

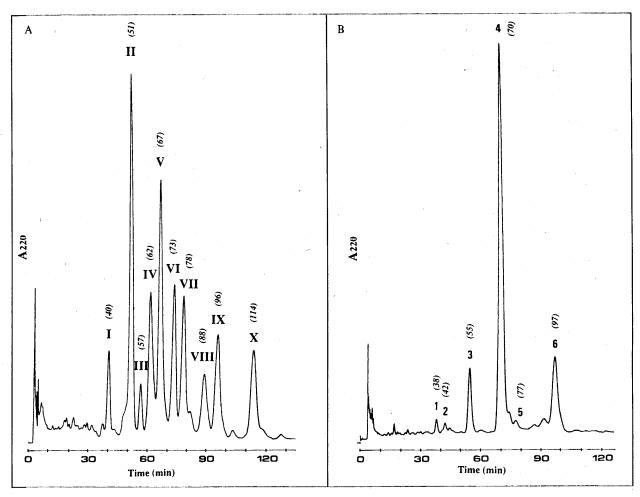
Charged Amino Acid-directed Biosynthesis of Peptaibols

In a second step, the feasibility of incorporating charged amino acids into peptaibol sequences has been investigated by examining the effects of adding glutamic acid and arginine to T. longibrachiatum and T. harzianum fermentations, respectively. As the negatively charged glutamic acid naturally occurs in the LGB group of longibrachins biosynthesized by T. longibrachiatum, the influence of a Glu-supplied medium on the course of the longibrachin biosynthesis was examined, expecting an increased or exclusive production of acidic longibrachins LGB. Indeed, a larger production of longibrachins LGB_E to the detriment of the neutral LGA_E (Table 1) was observed. The composition of each peptide group was also affected by the Glu-supply, as a single peptide was synthesized inside each longibrachin group. These two peptides, designated as LGA_E 1 and LGB_E 1 were assigned to LGA I and LGB II, respectively (Table 2) by HPLC co-chromatography confirmed by further NMR and MS data.

The positively charged arginine is normally absent from any peptaibol sequence and the possibility of incorporation of this amino acid in a peptaibol could affect considerably the membrane and antibiotic prop-

The crude peptide mixtures contained both peptides and others metabolites.

Fig. 3. HPLC chromatograms of the trichorzins PA (A) and PA_U (B), from T. harzianum.



Kromasil C_{18} (5 μ), 4.6 × 250 mm, flow rate 1 ml/minute, absorption monitored at 220 nm. Solvent system for A: MeOH - H_2O , 84:16; for B: MeOH - H_2O , 83:17.

Retention times of the peptides are given in brackets.

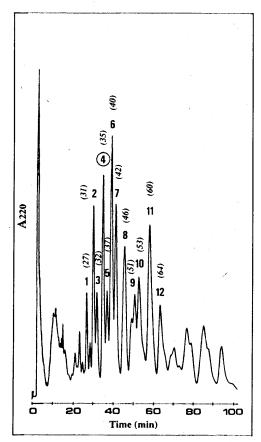
Table 4. HPLC retention times (minute) of the trichorzins PA, PA_U and PA_R from T. harzianum (M-902608).

Peptides	PA ^a										
	I	II	III	IV		v	VI	VII	VIII	IX	X
Rt (minute)	40	51	57	62		67	73	78	88	96	114
Peptides		AND THE PROPERTY OF THE PARTY O	PA	יט				-	PA _R		
	1	2	3	4	5	6	1	2	3	4	. 5
Rt (minute)	24 ^a 38 ^b	26 ^a 42 ^b	30 ^a 55 ^b	40 ^a 70 ^b	41 ^a 77 ^b	57ª 97 ^b	32	2 4	51	57	7

Solvent system: MeOH-H₂O (84:16).

b Solvent system: MeOH-H₂O (83:17).

Fig. 4. HPLC chromatogram of the harzianins PC_U from *T. harzianum*.



Kromasil C_{18} (5 μ), 4.6×250 mm, flow rate 1 ml/minute, absorption monitored at 220 nm. Solvent system: MeOH - H_2O , 80:20.

Retention times of the peptides are given in brackets.

erties of these peptides. The effect of adding arginine was studied on the *T. harzianum* strain. Upon the Argenrichment of the medium, *T. harzianum* only produced one peptide group, designated as trichorzins PA_R, showing that biosynthesis of the PC and PD groups was more strongly inhibited than with an Aib-supply (Table 3). The HPLC profile of trichorzins PA_R exhibited mainly two peptides, PA_R 2 and PA_R 3, and three minor compounds, PA_R 1, PA_R 4 and PA_R 5 (Table 4). A HPLC co-chromatography indicated PA_R 2 and PA_R 3 to be PA_U 4 and PA II respectively, which was confirmed by further NMR and MS data. Similarly, the minor compounds, PA_R 1, PA_R 4 and PA_R 5, were shown to be PA_U 3, PA III and PA VI, respectively.

As observed with the Aib-supply, the adjunction of charged amino acids to the fermentation medium led to the simplification of the peptaibol mixtures. The Glu-supply favored the synthesis of acidic longibrachins,

as expected from the presence of glutamic acid in their sequences. However and more surprisingly, it also affected the microheterogeneous mixture composition, leading to the biosynthesis of a single peptide inside each longibrachin class, this peptide being the same as that obtained with the Aib-supply. At last, the arginine incorporation by *T. harzianum* failed, as it only resulted in the synthesis of trichorzins PA already described and in the absence of Arg-containing peptides.

Conclusion

The microheterogeneous mixtures of peptaibols produced by two Trichoderma strains, T. harzianum and T. longibrachiatum are composed of closely related sequence analogues differing from each other by amino acid substitutions, such as Aib/Iva, Glu/Gln or Trpol/ Pheol at fixed positions. In the present study, we have given evidence that an exogenous amino acid supply results in simplification of the microheterogeneous peptide mixtures, as illustrated by both Trichoderma strains when Aib, Glu, or Arg is added to the fermentation medium. Furthermore, the amino acid supplies lead to enhanced yields of Aib-rich peptides, as unexpectedly observed for the Glu- and Arg-supplies. Our attempt to incorporate into the trichorzin sequences an amino acid normally absent from the sequences of peptaibols, such as the positively charged arginine, failed. However, a number of amino acids unknown in the sequences of cyclosporins have been previously incorporated by similar feeding experiments.²²⁾ Further experiments are now in progress in our laboratory towards understanding the mechanisms by which the multienzymatic complexes responsible for the trichorzin and longibrachin production generate the corresponding peptide products.

Materials and Methods

Fermentation

The *T. longibrachiatum* (M-853431) and *T. harzianum* (M-902608) strains were obtained from the "Collection de souches fongiques du Muséum National d'Histoire Naturelle", Paris (France). Stocked as lyophilized samples, the preparations were transfered to agar slants that contained 2% w/v malt-agar at 27°C for 5 days. Fermentations were performed in Roux flasks (1 liter), each containing 170 ml of sterilized synthetic medium (glucose 0.5%; KH₂PO₄ 0.08%; KNO₃ 0.07%; Ca(H₂-PO₄)₂ 0.02%; MgSO₄·7H₂O 0.05%; MnSO₄·5H₂O

0.001%; ZnSO₄·7H₂O 0.001%; CuSO₄·5H₂O 0.0005%; FeSO₄·7H₂O 0.0001%). The exogenous supply of α -aminoisobutyric acid, glutamic acid or arginine, was 0.8%. The pH was controlled and set to 4.5. The stationary 20 liter cultures were incubated at 27°C, for 13 days in the case of *T. harzianum* and for 20 days in the case of *T. longibrachiatum*.

Isolation

Separated from the mycelia by filtration, the filtrated broths were extracted three times with n-butanol. The crude extracts were evaporated to dryness. The residues were then submitted to gel chromatography over Sephadex LH20 with MeOH as eluent. Crude peptide mixtures, which contained peptides and others metabolites such as carbohydrates, were then chromatographed on silica gel (Kieselgel 60H, Merck, Darmstadt) with CH_2Cl_2 -MeOH (9:1 \rightarrow 3:7) as eluent. In the case of T. longibrachiatum cultivated on normal medium, the LGA group eluted first with CH₂Cl₂-MeOH (8:2), followed by LGB with CH₂Cl₂-MeOH (3:7); for the amino acid-supplemented media, two groups eluted in similar conditions, LGA_U and LGB_U for the Aib supply, LGA_E and LGB_E for the Glu supply (E and U designate the Glu- and Aib-supplies by the amino acid one-letter code). In the case of T. harzianum cultivated on normal medium, the PC group eluted first, followed by PA and PD, with CH₂Cl₂-MeOH (8:2); for the Aib-supplemented medium PC_U and PA_U groups eluted with CH_2Cl_2 - MeOH (8:2), whereas the PA_R group was only present with the Arg-supply, eluting with CH₂Cl₂-MeOH (8:2) (Tables 1 and 2).

HPLC Separations

High-Performance Liquid Chromatography was carried out with a Waters liquid chromatograph (717 plus Autosampler, 600E Multisolvent Delivery System and 486 Tunable Absorbance Detector) on an analytical C_{18} column (Kromasil C_{18} , $5\,\mu$, $4.6\,\mathrm{mm}\times250\,\mathrm{mm}$; AIT France).

Retention times of longibrachins (LGA, LGA_U, LGA_E and LGB, LGB_U, LGB_E) and of trichorzins (PA, PA_U, PA_R) are given in Table 2 and Table 4, respectively.

Harzianins PC_U : eluent, MeOH - H₂O (80:20); flow rate, 1 ml/minute. Rt (minute): PC_U 1=27, PC_U 2=31, PC_U 3=32, PC_U 4=35, PC_U 5=37, PC_U 6=40, PC_U 7=42, PC_U 8=46, PC_U 9=51, PC_U 10=53, PC_U 11=60, PC_U 12=64.

The different trichorzins PA_U and PA_R and harzianin PC_U were further purified by repetitive HPLC on a

semi-preparative C_{18} column (Kromasil C_{18} , $5\,\mu$, 7.8 mm × 300 mm; AIT France). Regarding the *T. harzianum* strain, three cultures were performed independently with an Aib-supply and the results were shown to be reproducible.

Acknowledgments

We thank Pr. M. F. ROQUEBERT (Laboratoire de Cryptogamie, Muséum National d'Histoire Naturelle, Paris, France) for providing the *Trichoderma* strains.

References

- 1) EL HAJJI, M.; S. REBUFFAT, T. LE DOAN, G. KLEIN, M. SATRE & B. BODO: Interaction of trichorzianins A and B with model membranes and with the amoeba *Dictyostelium*. Biochim. Biophys. Acta 978: 97 ~ 104, 1989
- LUCACIU, M.; S. REBUFFAT, C. GOULARD, H. DUCLOHIER, G. MOLLE & B. BODO: Interaction of the 14-residue peptaibols, harzianins HC, with lipid bilayers: permeability modifications and conductance properties. Biochim. Biophys. Acta 1323: 85~96, 1997
- Rebuffat, S.; H. Duclohier, C. Auvin-Guette, G. Molle, G. Spach & B. Bodo: Membrane-modifying properties of the pore-forming peptaibols saturnisporin SA IV and harzianin HA V. FEMS Microbiol. Immunol. 105: 151~160, 1992
- Sansom, M. S. P.: The biophysics of peptide models of ion channels. Prog. Biophys. Mol. Biol. 55: 139~235, 1991
- 5) Pandey, R. C.; J. C. Cook Jr. & K. L. Rinehart Jr.: High resolution and field desorption mass spectrometry studies and revised structures of alamethicins I and II. J. Am. Chem. Soc. 99: 8469~8483, 1977
- 6) REBUFFAT, S.; L. CONRAUX, M. MASSIAS, C. AUVIN-GUETTE & B. BODO: Sequence and solution conformation of the 20-residue peptaibols, saturnisporins SA II and SA IV. Int. J. Peptide Protein Res. 41: 74~84, 1993
- DORNBERGER, K.; W. IHN, M. RITZAU, U. GRÄFE, B. SCHLEGEL, W. F. FLECK & J. W. METZGER: Chrysospermins, new peptaibol antibiotics from *Apiocrea chrysosperma* Ap101. J. Antibiotics 48: 977~989, 1995
- 8) Jung, G.; W. A. König, D. Leibfritz, T. Ooka, K. Janko & G. Вонеім: Structural and membrane modifying properties of suzukacillin, a peptide antibiotic related to alamethicin; Part A: sequence and conformation. Biochim. Biophys. Acta 45: 164~181, 1976
- 9) BECKER, D.; M. KIESS & H. BRÜCKNER: Structures of peptaibol antibiotics hypomurocin A and B from the ascomycetous fungus *Hypocrea muroiana* Hino et Katsumoto. Liebigs Annal. 4: 767~772, 1997
- 10) KARLE, I. L.; J. L. FLIPPEN-ANDERSON, M. SUKUMAR & P. BALARAM: Conformation of a 16-residue zervamicin IIA analog peptide containing three different structural features: 3₁₀-helix, α-helix and β-bend ribbon. Proc. Natl. Acad. Sci. USA 84: 5087~5091, 1987
- 11) Rebuffat, S.; C. Goulard & B. Bodo: Antibiotic peptides from *Trichoderma harzianum*: harzianins HC, proline-rich 14-residue peptaibols. J. Chem. Soc. Perkin Trans. 1: 1849 ~ 1855, 1995

- 12) Rebuffat, S.; S. Hlimi, Y. Prigent, C. Goulard & B. Bodo: Isolation and structural elucidation of the 11-residue peptaibol antibiotic, harzianin HK VI. J. Chem. Soc. Perkin Trans. 1: 2021 ~ 2027, 1996
- 13) AUVIN-GUETTE, C.; S. REBUFFAT, Y. PRIGENT & B. BODO: Trichogin A IV, an 11-residue lipopeptaibol from Trichoderma longibrachiatum. J. Am. Chem. Soc. 114: 2170~2174, 1992
- 14) FUJITA, T.; S. WADA, A. IIDA, T. NISHIMURA, M. KANAI & N. TOYAMA: Isolation and structural elucidation of new peptaibols, trichodecenins-I and -II, from *Trichoderma viride*. Chem. Pharm. Bull. 42: 489~494, 1994
- 15) KLEINKAUF, H. & H. VON DÖHREN: A non-ribosomal system of peptide biosynthesis. Eur. J. Biochem. 236: 335~351, 1996
- 16) STEIN, T.; J. VATER, V. KRUFT, A. OTTO, B. WITTMANN-LIEBOLD, P. FRANKE, M. PANICO, R. MCDOWELL & H. R. MORRIS: The multiple carrier model of nonribosomal peptide biosynthesis at modular multienzymatic templates. J. Biol. Chem. 271: 15428~15435, 1996
- 17) STACHELHAUS, T.; A. SCHNEIDER & M. A. MARAHIEL: Engineered biosynthesis of peptide antibiotics. Biochem. Pharmacol. 52: 177~186, 1996
- 18) MOHR, H. & H. KLEINKAUF: Alamethicin biosynthesis, acetylation of the amino terminus and attachment of phenylalaninol. Biochim. Biophys. Acta 526: 375~386, 1978

- 19) ZOCHER, R.; N. MADRY, H. PEETERS & H. KLEINKAUF. Biosynthesis of Cyclosporin A. Phytochemistry 23: 549 ~ 551, 1984
- 20) BRÜCKNER, H. & M. PRZYBYLSKI: Isolation and structural characterization of polypeptide antibiotics of the peptaibol class by high-performance chromatography with high field desorption and fast atom bombardment mass spectrometry. J. Chromatogr. 296: 263~275, 1984
- KOBEL, H. & R. TRABER: Directed biosynthesis of Cyclosporins. Eur. J. Appl. Microbiol. Biotechnol. 14: 237~240, 1982
- 22) TRABER, R.; H. HOFFMANN & H. KOBEL: Cyclosporins— New analogues by precursor directed biosynthesis. J. Antibiotics 42: 591 ~ 597, 1988
- 23) Krasnoff, S. B. & S. Gupta: Identification and directed biosynthesis of efrapeptins in the fungus *Tolypocladium geodes* GAMS (Deuteromycotina: Hyphomycetes). J. Chem. Ecol. 17: 1953~1962, 1991
- 24) HLIMI, S.: Peptides antibiotiques d'origine fongique, isolement de nouveaux peptaïbols, études structurales, relations structure/activité membranaire et antibiotiques. Ph.D. Thesis, Paris VI University, 1997
- 25) DUVAL, D.; S. REBUFFAT, C. GOULARD, Y. PRIGENT, M. BECCHI & B. BODO: Isolation and sequence analysis of the peptide antibiotics trichorzins PA from *Trichoderma harzianum*. J. Chem. Soc. Perkin Trans. 1: 2147~2153, 1997