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## Poster Presentation

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### Chemoselective One-Step Purification Method for Peptides Synthesized by the Solid Phase Technique

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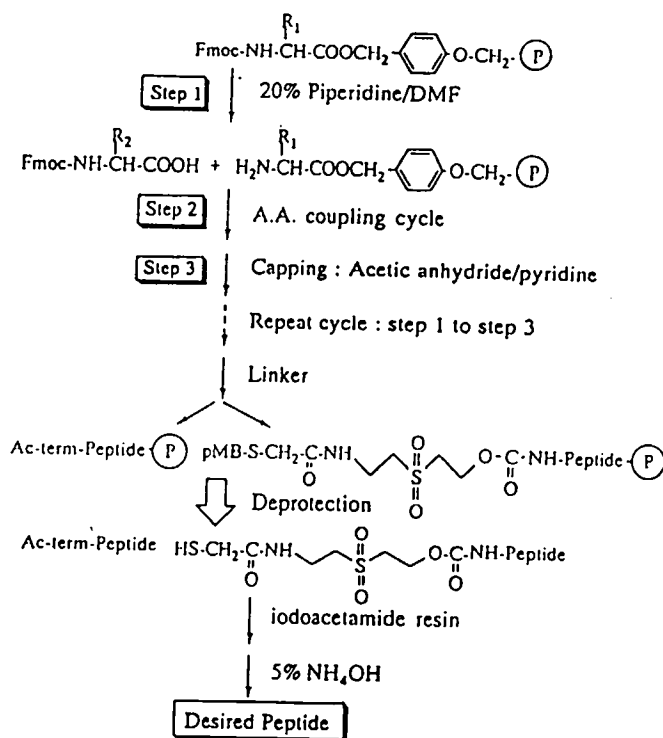
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*W. E. H.*

## Abstract

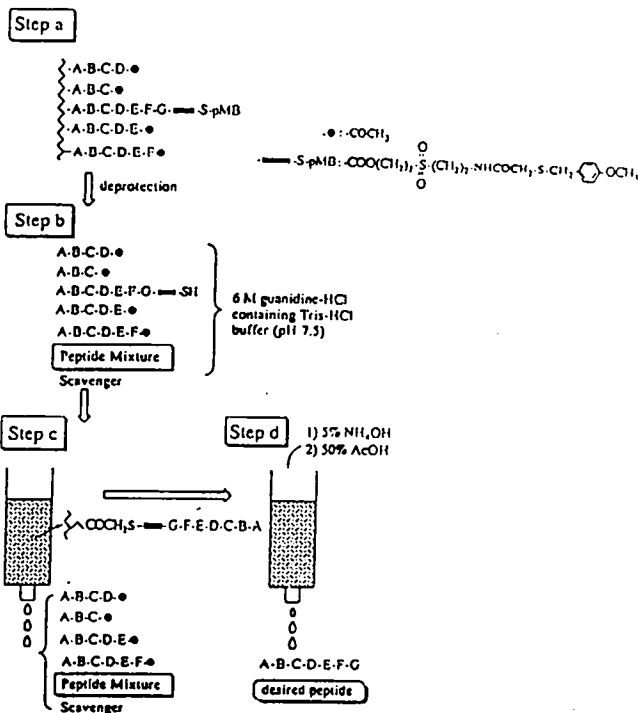
Based upon the specific reaction between SH and iodoacetamide groups, we have explored a new affinity-type purification procedure for peptides synthesized by the solid-phase technique. For this affinity-type purification procedure, we synthesized a new SH precursor reagent bearing an acid-labile S-protecting group, pMB-S-CH<sub>2</sub>CO-NHCH<sub>2</sub>CH<sub>2</sub>SO<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>OCOOpNP [1]. Using this reagent, the procedure involves the following sequence of 4 reactions. 1. Attachment of the SH function of [1] to the α-amino group of a peptide-resin through a base-labile sulfonylthio-carbonyl linkage in the final step of solid-phase peptide synthesis. 2. Acid treatment to remove the S-pMB and side chain protecting groups employed and cleave the modified peptide from the resin. 3. Immobilization of the derived SH-peptide on an iodoacetamide resin column. 4. Base (5% NH<sub>4</sub>OH) treatment to release the desired peptide from the resin in nearly pure form. To facilitate this purification procedure, unreacted amino groups were acetylated in each step during solid-phase synthesis. The usefulness of this method was demonstrated by the purification of several peptides (18–44 amino acids in length) synthesized by the Fmoc-based solid-phase technique. The principle of this affinity-type purification procedure may also be applied to Boc-based solid-phase technique.

## General scheme for Fmoc-based solid-phase peptide synthesis.



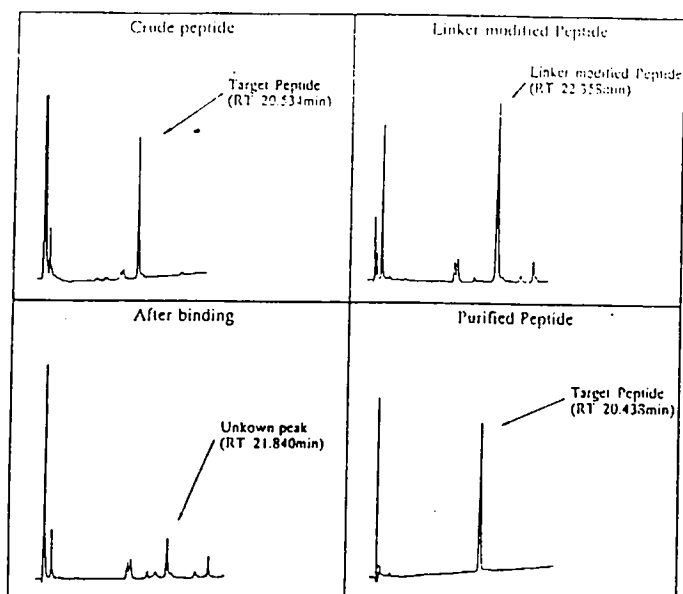
Synthesis was carried out automatically according to the principles of Sheppard *et al.*, using the following side chain protected Fmoc-amino acids: Arg(Mtr), Lys(tBoc), His(tBoc), Glu(O<sup>t</sup>Bu), Asp(O<sup>t</sup>Bu), Ser(tBu), and Tyr(tBu). Resin: polystyrene resin cross-linked with 1% divinyl benzene was used. Deprotection of the Fmoc group: resin was treated with 20% piperidine in DMF for 10 min. Coupling: coupling of protected amino acids was carried out in DMF for 30 min using several acylating reagents. Capping: capping of truncated peptides was performed with acetic anhydride (0.5 M)-pyridine (0.5 M) in DMF for 10 min. The S-protected base-labile SH linker was introduced manually by the *p*-nitrophenyl active ester-HOBt method.

## Deprotection, and purification of peptides synthesized by the solid-phase technique.



The base-labile SH introducing reagent and an iodoacetamide column were used for the isolation of peptides synthesized by the solid-phase technique. After each step in which an Fmoc-amino acid group was added (A,B,C,D,...), the residual unreacted amino groups were blocked with an acetyl group. In the last step of synthesis, the S-protected base-labile SH introducing reagent was added to the N-terminal residue (step a). After removal of the synthesized peptide from the solid support (step b), the mixture was passed through an iodoacetamide column to remove all of the truncated peptides and scavengers (step c). After washing the column, the resin was treated with 5% NH<sub>4</sub>OH to release the desired peptide (step d).

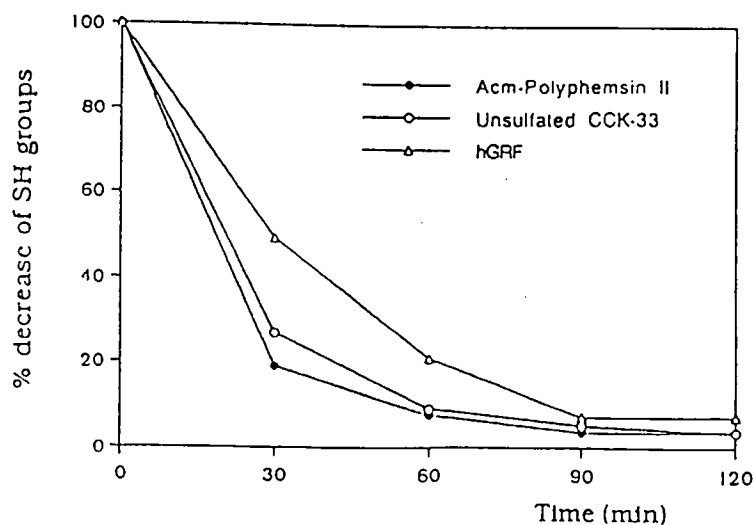
### Identification of the target peptide during purification



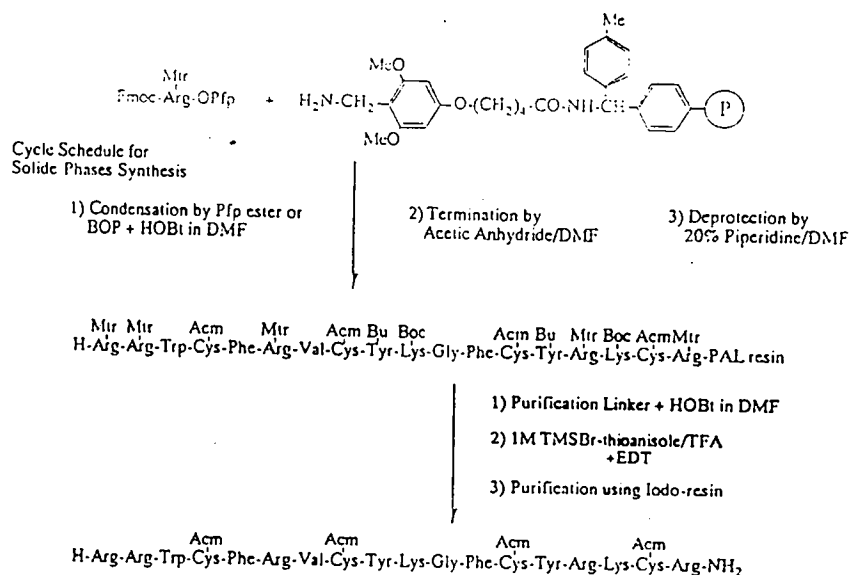
This new purification method can easily identify the target peptide by comparison of HPLC assays of the purified peptide, the linker modified peptide and the crude peptide. The linker modified peptide was characterized by its longer retention time due to the hydrophobicity of the linker. Therefore the peak with its retention time extended by introduction of the linker can be identified as that of the target peptide.

HPLC was performed on a  $\mu$  Bondasphere 5C18 (100Å) column (3.9 x 150mm), using a 0.1% TFA/acetonitril solvent system at a flow rate of 1.0ml/min. Absorbance was monitored at 220nm and the column was eluted with a linear gradient of 20-30% solvent B over 30min.

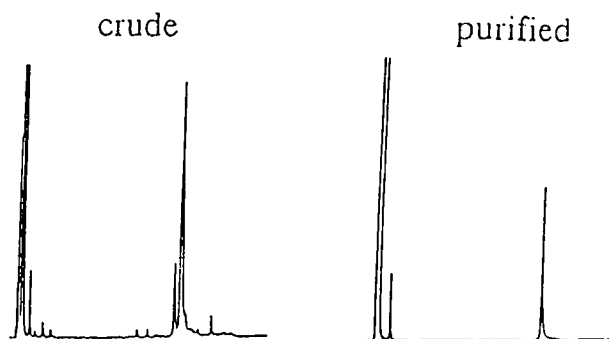
### Binding of the linker-modified peptide to an iodoacetamide resin column monitored by the Ellman method.



## Fmoc-Based Solid-Phase Synthesis of Acm-Polyphemusin II



## HPLC elution profile of synthetic Acm-polyphemusin II.



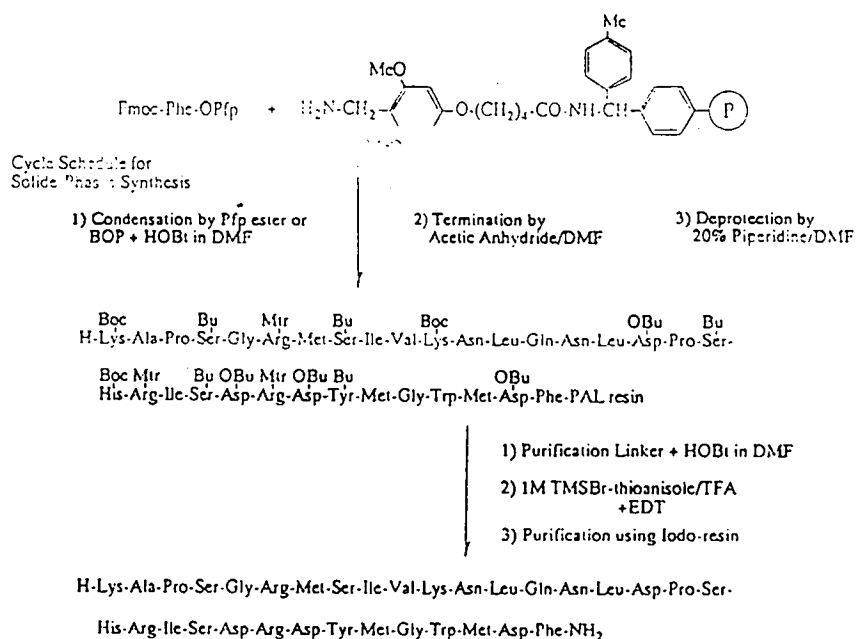
## Structure of Acm-polyphemusin II

II-Arg-Arg-Trp-Cys(Acm)-Phe-Arg-Val-Cys(Acm)-Tyr-Lys-Gly-Phe-  
 Cys(Acm)-Tyr-Arg-Lys-Cys(Acm)-Arg-NH<sub>2</sub>

FAB-MS: cal. 2712.3 (C<sub>120</sub>H<sub>185</sub>N<sub>41</sub>O<sub>24</sub>S<sub>4</sub>), ob. 2714.1 (M+H<sup>+</sup>)

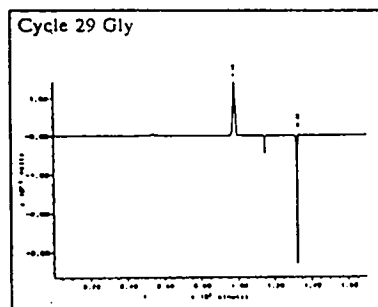
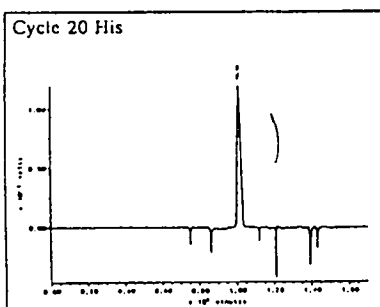
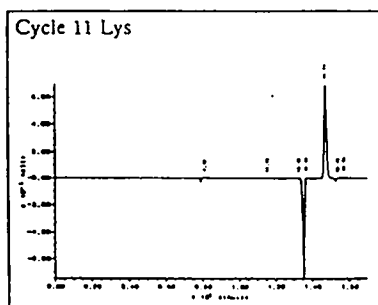
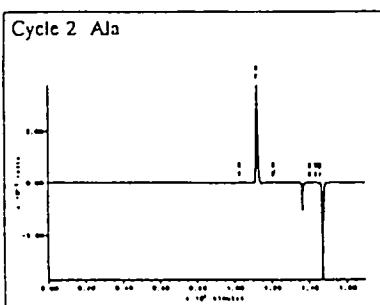
HPLC was performed on a  $\mu$ Bondasphere 5C18 (100 Å) column (3.9 x 150 mm), using a 0.1% TFA/acetonitrile solvent system at a flow rate of 1.0 ml/min. Absorbance was monitored at 220 nm and the column was eluted with a linear gradient of 10-40% solvent B over 30 min.

# Fmoc-Based Solid-Phase Synthesis of unsulfated Human CCK-33

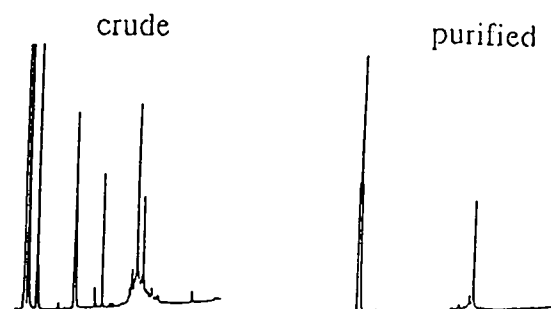


## Amino Acid Sequence Analysis of unsulfated Human CCK-33

Sequence ; H-KAPSGRMSI<sup>10</sup>VKNLQNLDP<sup>20</sup>SHRISDRDYM<sup>30</sup>GWMD<sup>30</sup>F-NH<sub>2</sub>



## HPLC elution profile of synthetic unsulfated CCK-33.



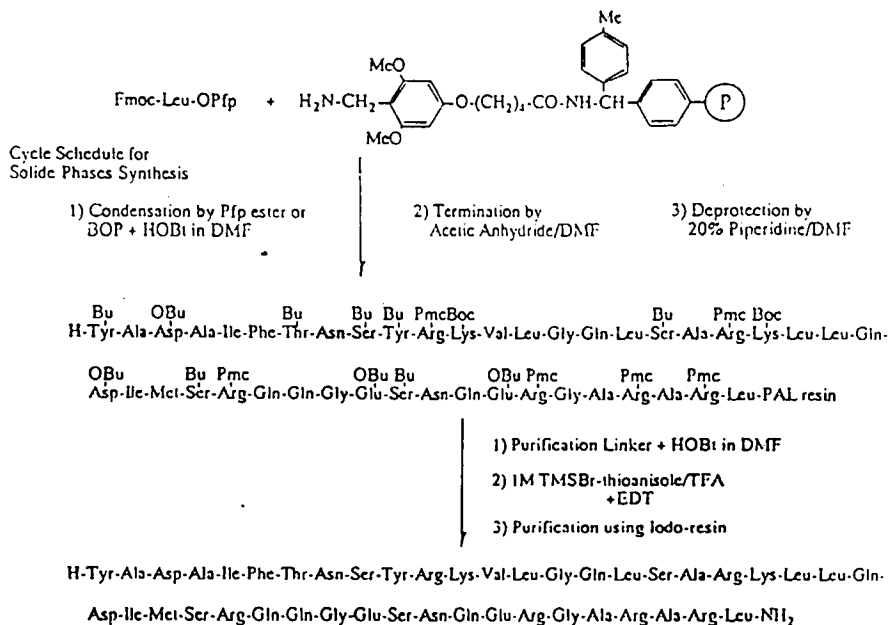
## Structure of unsulfated human CCK-33

H-Lys-Ala-Pro-Ser-Gly-Arg-Met-Ser-Ile-Val-Lys-Asn-Leu-Gln-Asn-Leu-Asp-Pro-Ser-His-Arg-Ile-Ser-Asp-Arg-Asp-Tyr-Met-Gly-Trp-Met-Asp-Phe-NH<sub>2</sub>

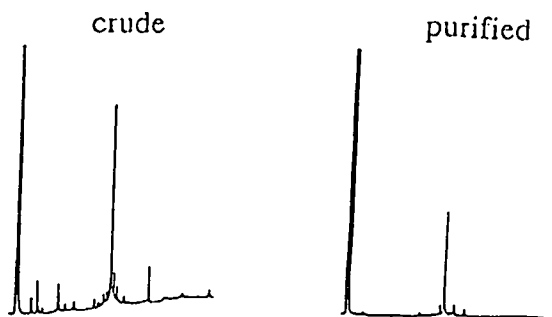
FAB-MS: cal. 3862.9 (C<sub>167</sub>H<sub>263</sub>N<sub>51</sub>O<sub>49</sub>S<sub>3</sub>), ob. 3865.2 (M+H<sup>+</sup>)

HPLC was performed on a  $\mu$ Bondasphere 5C18 (100Å) column (3.9 x 150 mm), using a 0.1% TFA/acetonitrile solvent system at a flow rate of 1.0 ml/min. Absorbance was monitored at 220 nm and the column was eluted with a linear gradient of 10-60% solvent B over 30 min.

# Fmoc-Based Solid-Phase Synthesis of Human Growth hormone releasing Factor



HPLC elution profile of synthetic human growth hormone releasing factor.



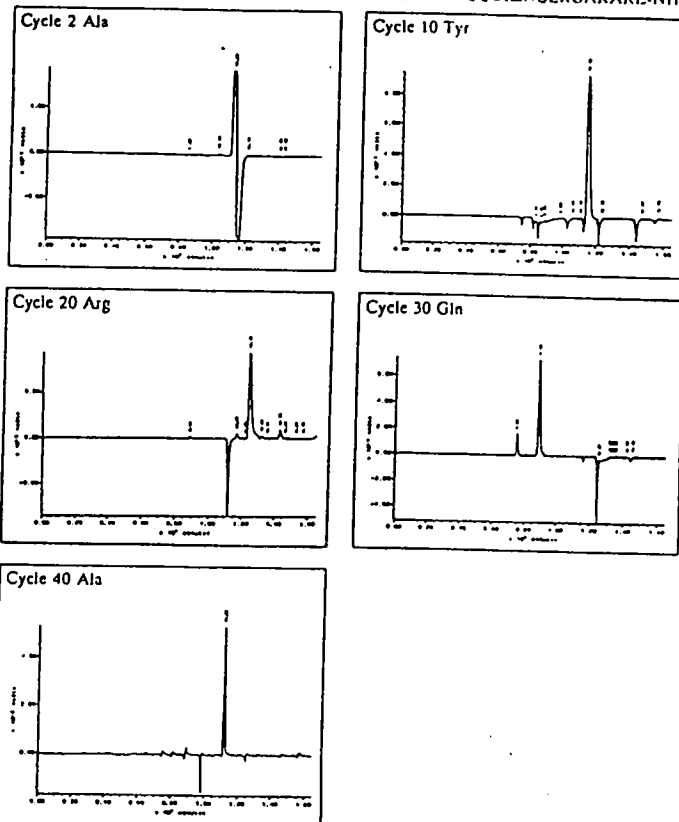
## Structure of human growth hormone releasing factor

$\text{H-Tyr-Ala-Asp-Ala-Ile-Phe-Thr-Asn-Ser-Tyr-Arg-Lys-Val-Leu-Gly-}$   
 $\text{Gln-Leu-Ser-Ala-Arg-Lys-Leu-Leu-Gln-Asp-Ile-Met-Ser-Arg-Gln-}$   
 $\text{Gln-Gly-Glu-Ser-Asn-Gln-Glu-Arg-Gly-Ala-Arg-Ala-Arg-Leu-NH}_2$   
 FAB-MS: cal. 5036.7 ( $\text{C}_{215}\text{H}_{358}\text{N}_{72}\text{O}_{66}\text{S}$ ), ob. 5037.8 ( $\text{M}+\text{H}^+$ )

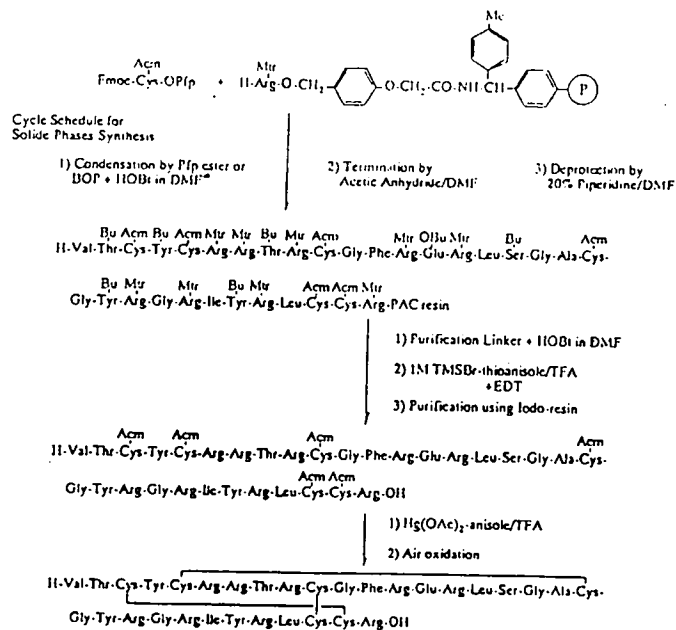
HPLC was performed on a  $\mu\text{Bondasphere 5C18}$  (100Å) column (3.9 x 150 mm), using a 0.1% TFA/acetonitrile solvent system at a flow rate of 1.0 ml/min. Absorbance was monitored at 220 nm and the column was eluted with a linear gradient of (a) 23-40% solvent B over 30 min or (b) 25-40% solvent B over 30 min.

## Amino Acid Sequence Analysis of Human Growth hormone releasing Factor

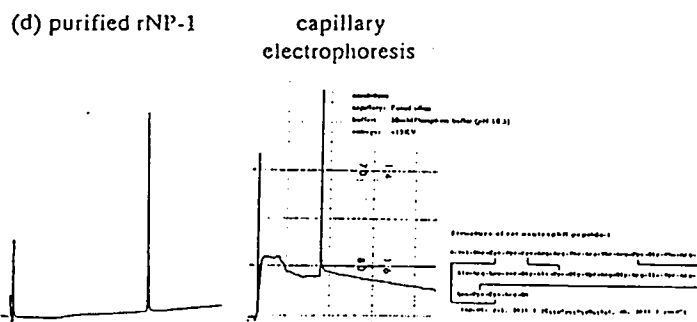
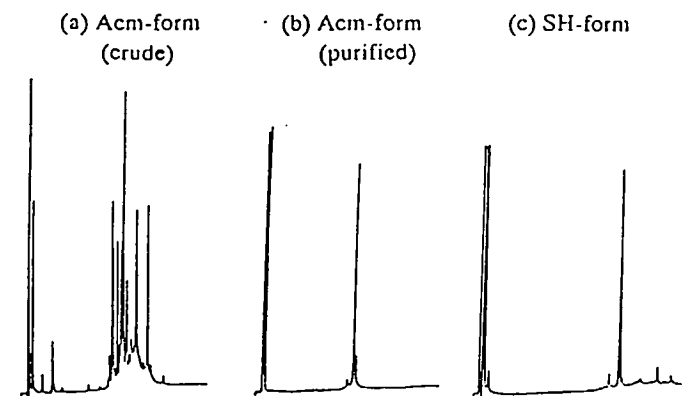
Sequence ; H-YADAIFTNSYRKVLGQLSARKLLQDIMSROOGESNOERGARL-NH<sub>2</sub>



# Fmoc-Based Solid-Phase Synthesis of Rat Neutrophil Peptide-1



## HPLC elution profile and capillary electrophoresis of synthetic rat neutrophil peptide-1 (rNP-1)



HPLC was performed on a  $\mu$ Bondasphere SC18 (100Å) column (3.9 x 150 mm), using a 0.1% TFA/acetonitrile solvent system at a flow rate of 1.0 ml/min. Absorbance was monitored at 220 nm and the column was eluted with a linear gradient of (a,b,d) 10-40% solvent B over 30 min (c) 15-25% solvent B over 40 min.

## Conclusion

*The advantages of our method consists of the purity of the product and its easiness. Even peptides which have shoulder peaks can be purified with this method. Compared with conventional HPLC purification, this method can make it possible for any researcher without special technique to obtain pure peptides easily at one step in short time by solid-phase technique. The easily prepared cross-linking reagent introduced here will be of wide applicability for the one-step purification of peptides synthesized by the solid-phase technique. Additionally, a more readily obtainable supply of high-purity or constant-purity peptides should facilitate research in the fields of biochemistry, physiology, and medicine with respect to the physiological activities and mechanisms of action of peptides and proteins.*

## Acknowledgement

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