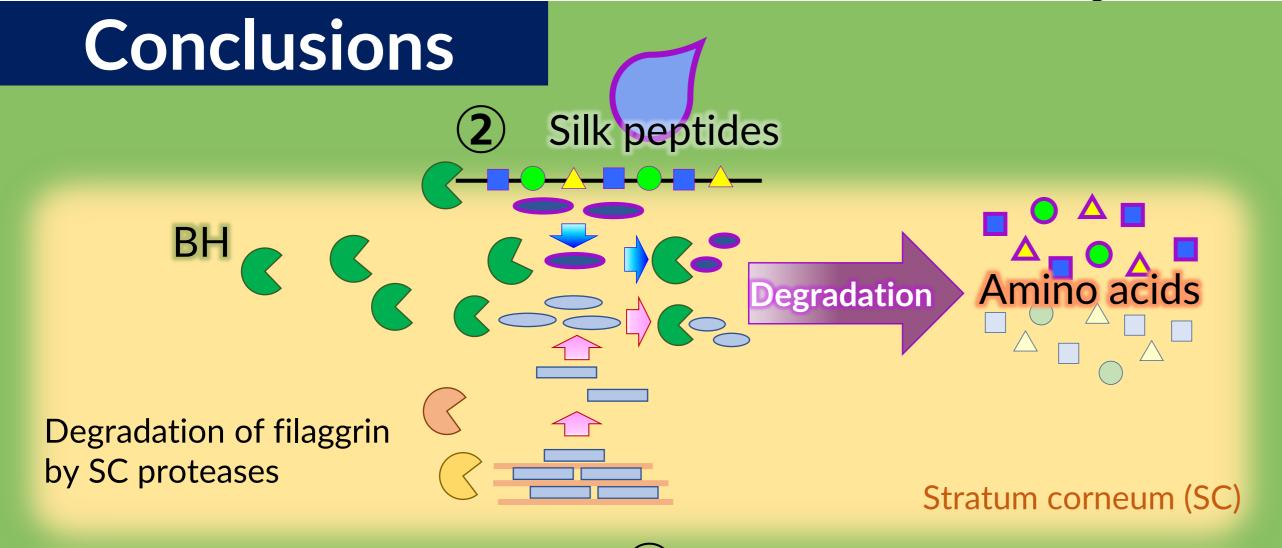
Poster ID 35 **Biodegradation of exogenously applied silk peptides**

iKeda **R&D** Center

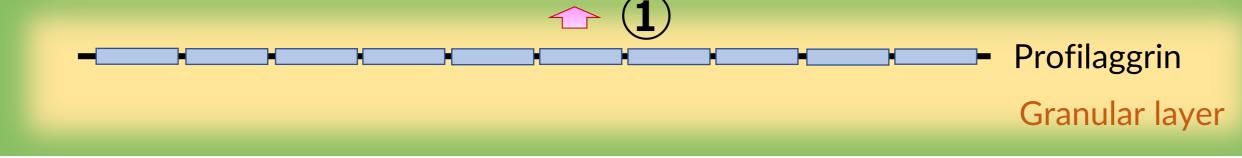
32nd **IFSCC** Congress

by stratum corneum cysteine proteases

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We found that exogenously applied silk peptides are degraded by cysteine proteases such as **bleomycin hydrolase** (BH), which is known to degrade filaggrin peptides into amino acids. This suggests that amino acids of NMFs can be supplemented not only endogenously within the skin but also exogenously.



This concept of peptide degradation upon application to the SC provides a novel approach to skin moisturization!!

Introduction

"Conventional degradation route"

Previous studies revealed that filaggrin is degraded into amino acids by proteases such as BH and these amino acids may serve as NMFs.

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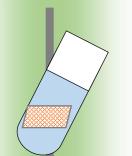
We hypothesized "New biodegradation route"

We aimed to know "whether exogenous peptides would be hydrolyzed to amino acids by SC proteases" and "what kind of proteases are involved?"

Materials & Methods



Collected by tape-stripping



Silk peptide solution

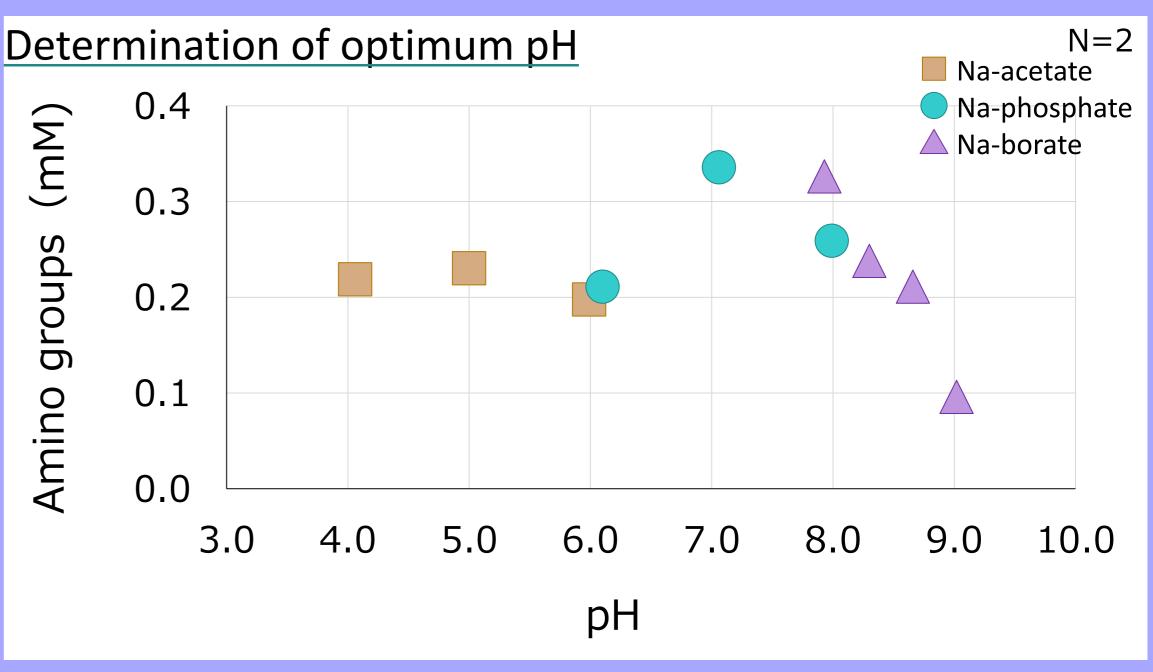
37°C for 48h

Biodegradation by SC proteases Materials - Hydrolyzed silk : Mixture of peptides

(Average molecular weight 500 Da, Consists of high levels of Gly and Ala residues, Concentration 10%)

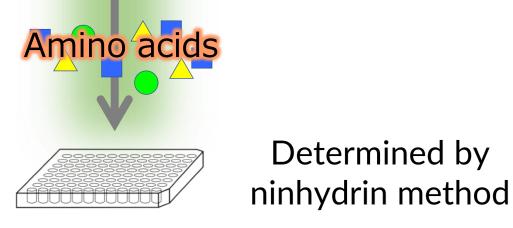
SILKPRO® F, IKEDA CORPORATION SC collection

Results & Discussion



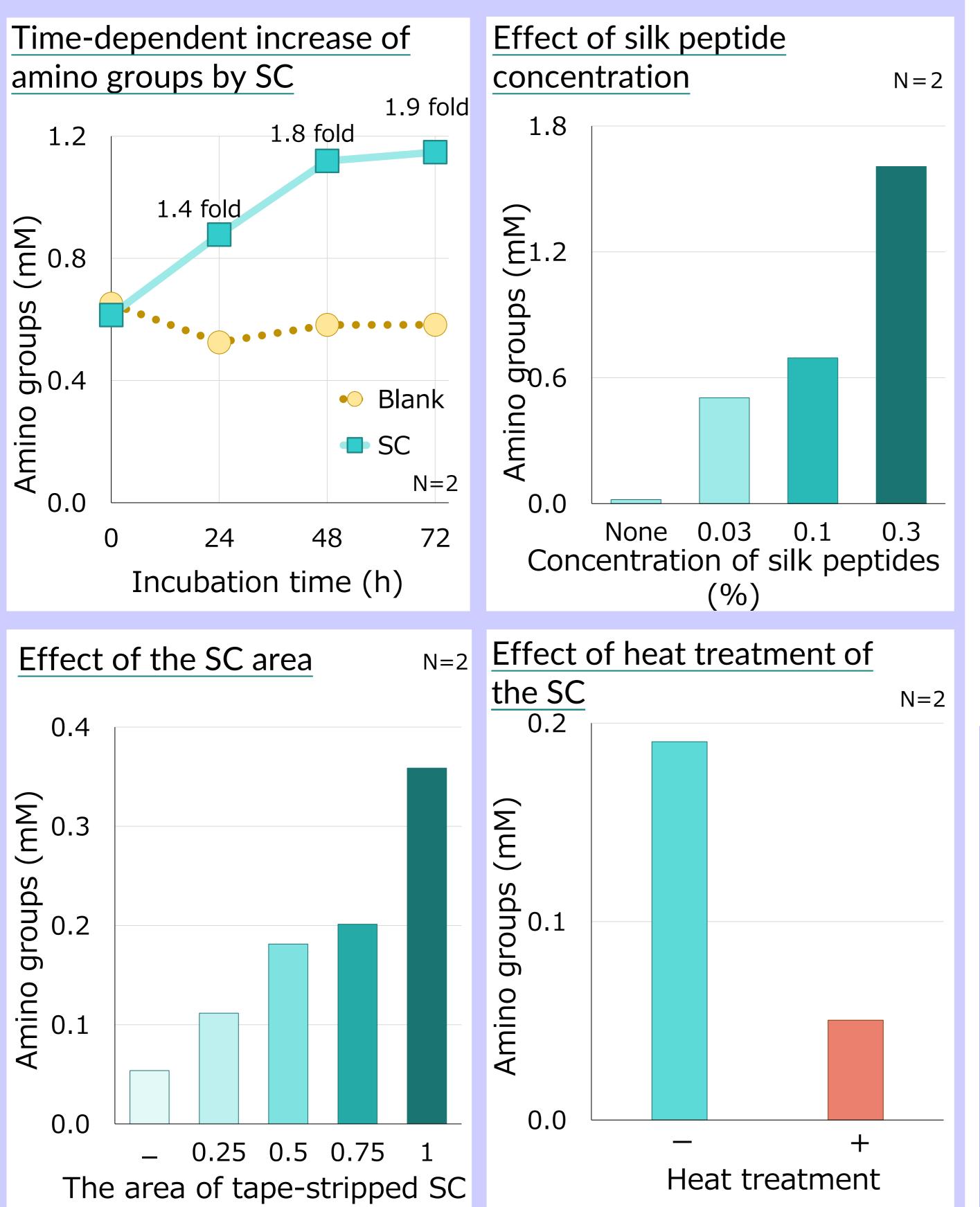
Effect of protease inhibitors and reducing agent

		Concentration	Relative activity (%)			
	Substances		Silk peptide		Arg-MCA	
			degradation		degradation	
			pH5.5	pH7.0	pH5.5	pH7.0
Inhibitor	None		100	100	100	100
	E-64	160 μΜ	40	23	30	14
	Leupeptin	120 μM	55	20	77	70
	Pepstatin	120 μM	97	81	43	112
	AEBSF	1 mM	112	132	81	92
	NEM	1 mM	75	72	45	15
	Iodoacetic acid	1 mM	16	50	24	19
	EDTA	1 mM	105	120	114	94
Reducing agent	DTT	10 mM	255	159		

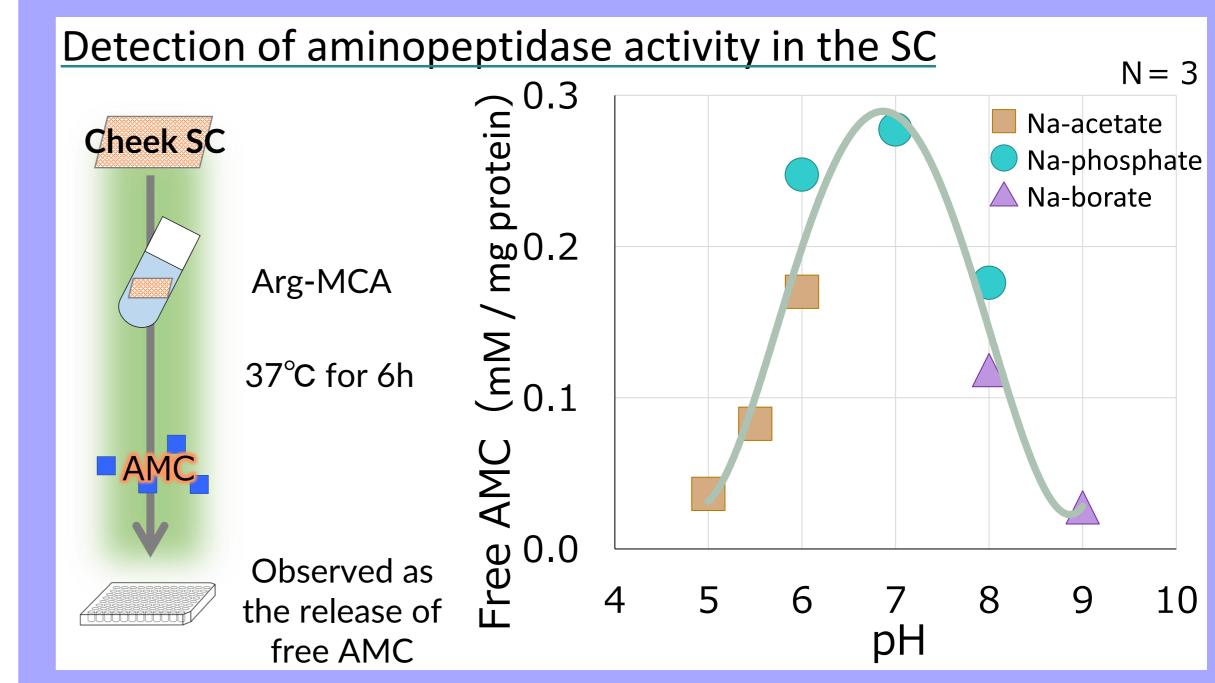


- Collected from the cheek by tape-stripping
- Monitoring of biodegradation of silk
 - The ninhydrin method,
 - MTP-900 Lab microplate reader (Corona Electric)

The protocol was approved by the ethics committee of Mukogawa Women's Univ.

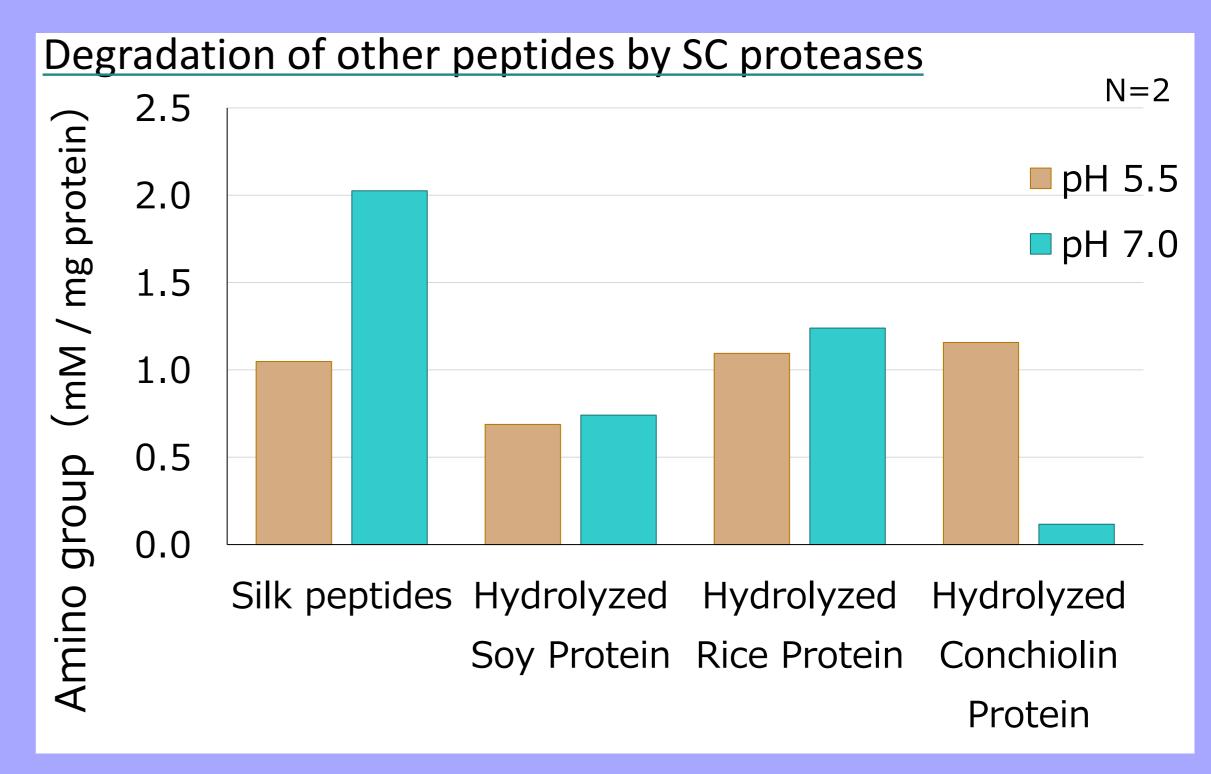


These results of silk peptide degradation showed the involvement of cysteine proteases, and is in agreement with the previous report [1].



Biochemical characterization of silk peptide degradation by SC, including a time-course experiment, dependence on substrate dose, dependence on SC dose, and heat inactivation, clearly showed the involvement of certain proteases in the degradation.

The pH dependence of the aminopeptidase activity suggested that BH is involved in the degradation of peptide.



Each cosmetic ingredients were also degraded by SC proteases.

Reference

[1] Takeda A, Higuchi D, Yamamoto T, Nakamura Y, Masuda Y, Hirabayashi T, Nakaya K (1996) Purification and characterization of bleomycin hydrolase, which represents a new family of cysteine proteases, from rat skin. J Biochem 119:29–36.

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