

<sup>4</sup>Department of Pulmonary Medicine, Institute of Clinical Medicine, Graduate School of Comprehensive Human Sciences, University of Tsukuba, Tsukuba, Ibaraki, Japan and <sup>5</sup>University Hospital, University of Tsukuba, Tsukuba, Ibaraki, Japan  
E-mail: akiyama@med.hokudai.ac.jp

## REFERENCES

- Fallon PG, Sasaki T, Sandilands A et al. (2009) A homozygous frameshift mutation in the mouse *Flg* gene facilitates enhanced percutaneous allergen priming. *Nat Genet* 41: 602–8
- Hanifin JM, Rajka G (1980) Diagnostic features of atopic dermatitis. *Acta Derm Venereol* 92:44–7
- Henderson J, Northstone K, Lee SP et al. (2008) The burden of disease associated with filaggrin mutations: a population-based, longitudinal birth cohort study. *J Allergy Clin Immunol* 121:872–7
- Isada A, Konno S, Hizawa N et al. (2010) A functional polymorphism (-603A → G) in the tissue factor gene promoter is associated with adult-onset asthma. *J Hum Genet* 55: 167–74
- Nemoto-Hasebe I, Akiyama M, Nomura T et al. (2010) *FLG* mutation p.Lys4021X in the C-terminal imperfect filaggrin repeat in Japanese atopic eczema patients. *Br J Dermatol* 161:1387–90
- Nomura T, Akiyama M, Sandilands A et al. (2008) Specific filaggrin mutations cause ichthyosis vulgaris and are significantly associated with atopic dermatitis in Japan. *J Invest Dermatol* 128:1436–41
- Nomura T, Sandilands A, Akiyama M et al. (2007) Unique mutations in the filaggrin gene in Japanese patients with ichthyosis vulgaris and atopic dermatitis. *J Allergy Clin Immunol* 119:434–40
- Rodríguez E, Baurecht H, Herberich E et al. (2009) Meta-analysis of filaggrin polymorphisms in eczema and asthma: robust risk factors in atopic disease. *J Allergy Clin Immunol* 123:1361–70
- Sandilands A, Terron-Kwiatkowski A, Hull PR et al. (2007) Comprehensive analysis of the gene encoding filaggrin uncovers prevalent and rare mutations in ichthyosis vulgaris and atopic eczema. *Nat Genet* 39:650–4
- van den Oord RA, Sheikh A (2010) Filaggrin gene defects and risk of developing allergic sensitisation and allergic disorders: systematic review and meta-analysis. *BMJ* 339:b2433

See related commentary on pg 2703

# RNase 7 Protects Healthy Skin from *Staphylococcus aureus* Colonization

*Journal of Investigative Dermatology* (2010) 130, 2836–2838; doi:10.1038/jid.2010.217; published online 29 July 2010

## TO THE EDITOR

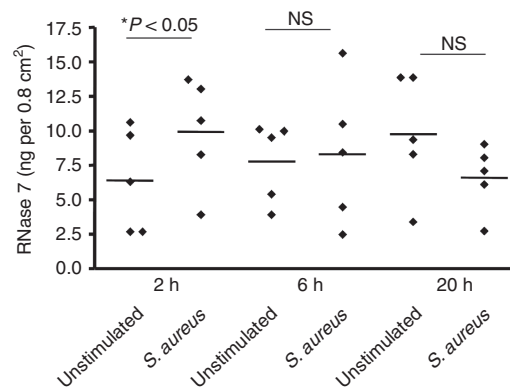
The Gram-positive bacterium *Staphylococcus aureus* is an important pathogen that causes various skin infections (Miller and Kaplan, 2009). However, healthy skin is usually not infected by *S. aureus*, despite the high carrier rates in the normal population (Noble, 1998). This suggests that the cutaneous defense system has the capacity to effectively control the growth of *S. aureus*. There is increasing evidence that antimicrobial proteins are important effectors of the cutaneous defense system (Harder et al., 2007). A recent study reported that keratinocytes contribute to cutaneous innate defense against *S. aureus* through the production of human  $\beta$ -defensin-3 (Kisich et al., 2007). In addition to human  $\beta$ -defensin-3, other antimicrobial proteins may also participate in cutaneous defense against *S. aureus*. One candidate is RNase 7, a potent antimicrobial ribonuclease that is highly expressed in healthy skin (Harder and Schröder, 2002; Köten et al., 2009).

To investigate the hypothesis that RNase 7 may contribute to protect

healthy skin from *S. aureus* colonization, we first incubated natural RNase 7 isolated from stratum corneum skin extracts (Harder and Schröder, 2002) with *S. aureus* (ATCC 6538). In concordance with our initial report about RNase 7 (Harder and Schröder, 2002), we verified that RNase 7 exhibited

a high killing activity against *S. aureus* (lethal dose of 90% = 3–6  $\mu\text{g ml}^{-1}$ ).

Recently, we reported a moderate induction of RNase 7 mRNA expression in primary keratinocytes treated with heat-killed *S. aureus* (Harder and Schröder, 2002). To assess the induction of RNase 7 by *S. aureus* in the



**Figure 1. Induced secretion of RNase 7 on the skin surface on treatment with living *S. aureus*.**

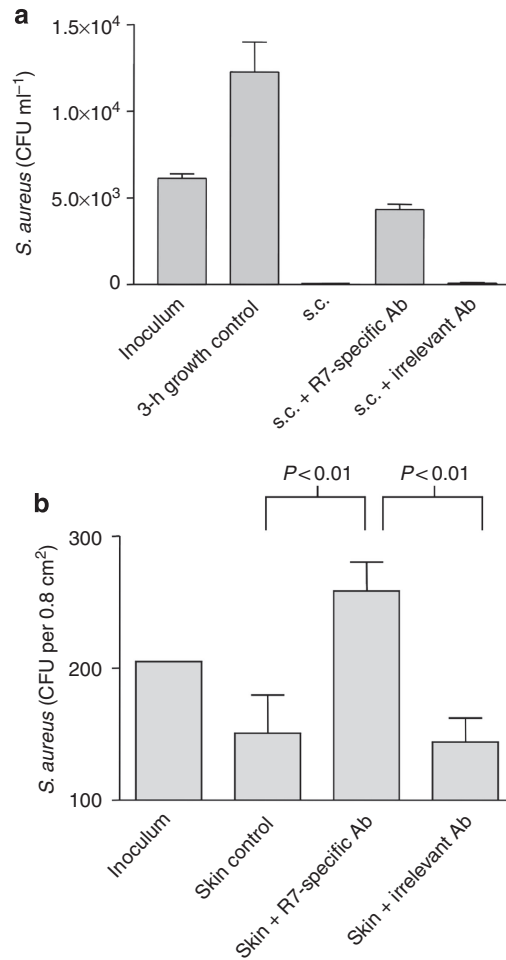
Defined areas (0.8 cm<sup>2</sup>) of skin explants derived from plastic surgery were incubated with or without approximately 1,000 colony-forming units of *S. aureus* (ATCC 6538) in 100  $\mu\text{l}$  of sodium phosphate buffer. After 2, 6, and 20 hours, the concentration of secreted RNase 7 was determined by ELISA. Stimulation with *S. aureus* for 2 hours revealed a significant induction as compared with the unstimulated control after 2 hours (\*P < 0.05, Student's *t*-test; n.s. = not significant). Data shown are means of triplicates of five skin explants derived from five donors.

context of more physiological conditions, we used human skin explants and incubated them for 2, 6, and 20 hours with living *S. aureus* (the use of skin material for this study was approved by the ethical committee of the University Kiel (A 104/06) in accordance with the Declaration of Helsinki Principles). As shown in Figure 1, only 2 hours of incubation with living *S. aureus* significantly induced the secretion of RNase 7 on the skin surface as measured by an RNase 7-specific ELISA (Köten *et al.*, 2009). This indicates a fast release of preformed material. The reduced secretion after 20 hours could be a result of cytotoxic effects from *S. aureus*, as well as of a decreased viability of skin explants. Secretion of RNase 7 after 2 hours was enhanced by increasing concentrations of *S. aureus* (Supplementary Figure S1 online).

To further assess the physiological relevance of RNase 7 in cutaneous defense against *S. aureus*, we investigated whether the killing activity of skin extracts derived from stratum corneum was inhibited through blocking of the antimicrobial activity of RNase 7 by RNase 7-specific antibodies as previously described (Köten *et al.*, 2009). First, we analyzed whether RNase 7 antibodies neutralized the antibacterial activity of RNase 7 against *S. aureus*. For this purpose, we tested the activity of RNase 7 against *S. aureus* in an antibacterial microdilution assay in the presence of RNase 7 antibodies. This approach revealed that the application of RNase 7 antibodies completely blocked the *S. aureus*-killing activity of RNase 7. In contrast, the antimicrobial activity was not inhibited when equivalent concentrations of irrelevant antibodies (derived from goat pre-immune serum; Köten *et al.*, 2009) were used (data not shown). Having established that RNase 7-specific antibodies neutralized the antimicrobial effect of RNase 7 against *S. aureus*, we used this approach to test the functional role of RNase 7 in cutaneous defense against *S. aureus*. To this end, we first incubated a stratum corneum skin extract with *S. aureus*. The skin extract was diluted to a ratio of 1:200 in 25  $\mu$ l of sodium phosphate buffer (pH 7.4) without or with RNase 7-blocking

antibody (4 mg ml<sup>-1</sup>) or an equivalent concentration of an irrelevant antibody. This dilution leads to a final concentration of 6  $\mu$ g ml<sup>-1</sup> RNase 7 in the assay system, which corresponds to the determined range of the lethal dose<sub>90</sub> of RNase 7 against *S. aureus* (see above). After 45 minutes, 25  $\mu$ l of *S. aureus* (ATCC 6538) in sodium phosphate buffer (pH 7.4) containing 2% tryptone soya broth was added and incubated for 3 hours. Killing activity was analyzed by plating serial dilutions of the incubation mixture and determining colony-forming units the following day. The skin extract exhibited a potent

*S. aureus*-killing activity, as incubation of *S. aureus* with the skin extract resulted in a complete killing of *S. aureus* (Figure 2a). The application of RNase 7-blocking antibodies to the skin extract before inoculation with *S. aureus* resulted in an outgrowth of *S. aureus* of approximately 35% as compared with the growth control (Figure 2a). In contrast, an irrelevant antibody did not inhibit the killing activity of the skin extract (Figure 2a). These data indicate an important contribution of RNase 7 to the *S. aureus*-killing activity of the skin extracts. Other skin-derived antimicrobial proteins may be responsible for the



**Figure 2. RNase 7 contributes to the killing activity of human skin against *S. aureus*.** (a) Stratum corneum extracts were incubated without (subcutaneous (s.c.)) or with RNase 7-blocking antibody (s.c. + R7-specific Ab) or an irrelevant antibody (s.c. + irrelevant Ab). After 45 minutes, *S. aureus* was added and colony-forming units (CFUs) were determined after 3 hours of incubation. Data shown are means  $\pm$  SD of a representative result of three independent experiments. (b) Defined areas (0.8 cm<sup>2</sup>) of skin explants were incubated without (skin control) or with RNase 7 antibody (skin + R7-specific Ab) or with an equivalent concentration of an irrelevant antibody (skin + irrelevant Ab). After 30 minutes, *S. aureus* was added and CFU were determined after 2 hours of incubation. Data show means  $\pm$  SD of triplicate experiments of the same donor. A representative result of three independent experiments from three donors is shown. Student's *t*-test was used for statistics.

*S. aureus*-killing activity, which remained after blocking the antimicrobial activity of RNase 7.

To obtain additional insight into the functional role of RNase 7 in cutaneous defense against *S. aureus*, we studied the role of RNase 7 using human skin explants. In this experimental *ex vivo* model, *S. aureus* was applied to defined areas (0.8 cm<sup>2</sup>) of the surface of skin explants. Preincubation of skin samples with 1.5 mg ml<sup>-1</sup> RNase 7-blocking antibody in 100 μl of sodium phosphate buffer (pH 7.4) led to a significant outgrowth of *S. aureus* as compared with pretreatment of skin with an equivalent concentration of irrelevant antibody (Figure 2b). These data further indicate a physiological relevance of RNase 7 for the control of *S. aureus* growth on the skin surface.

In summary, data present herein show that skin infected with living *S. aureus* responds with an increased release of RNase 7, which contributes to limit the growth of *S. aureus*. These

data are in concordance with a recently published study reporting that lower RNase 7 expression in healthy skin is associated with a higher risk of *S. aureus* skin infection (Zanger *et al.*, 2009). Future studies have to evaluate whether skin infections caused by *S. aureus* may be associated with an impaired expression or function of RNase 7.

#### CONFLICT OF INTEREST

The authors state no conflict of interests.

#### ACKNOWLEDGMENTS

We thank H Hinrichs, E Jeske, C Martensen-Kerl, and C Wilgus for excellent technical assistance. This study was supported by a grant from the Federal Ministry of Education and Research (BMBF, Skin Staph) to R Gläser and J Harder and by a Heisenberg program of the Deutsche Forschungsgemeinschaft (DFG) to J Harder.

**Maren Simanski<sup>1</sup>, Stefanie Dressel<sup>1</sup>,  
Regine Gläser<sup>1</sup> and Jürgen Harder<sup>1</sup>**

<sup>1</sup>Department of Dermatology, University Hospital Schleswig-Holstein, Campus Kiel, Kiel, Germany  
E-mail: jharder@dermatology.uni-kiel.de

#### SUPPLEMENTARY MATERIAL

Supplementary material is linked to the online version of the paper at <http://www.nature.com/jid>

#### REFERENCES

- Harder J, Gläser R, Schröder JM (2007) Human antimicrobial proteins effectors of innate immunity. *J Endotoxin Res* 13:317–38
- Harder J, Schröder JM (2002) RNase 7, a novel innate immune defense antimicrobial protein of healthy human skin. *J Biol Chem* 277: 46779–84
- Kisich KO, Howell MD, Boguniewicz M *et al.* (2007) The constitutive capacity of human keratinocytes to kill *Staphylococcus aureus* is dependent on beta-defensin 3. *J Invest Dermatol* 127:2368–80
- Köten B, Simanski M, Gläser R *et al.* (2009) RNase 7 contributes to the cutaneous defense against *Enterococcus faecium*. *PLoS One* 4:e6424
- Miller LG, Kaplan SL (2009) *Staphylococcus aureus*: a community pathogen. *Infect Dis Clin North Am* 23:35–52
- Noble WC (1998) Skin bacteriology and the role of *Staphylococcus aureus* in infection. *Br J Dermatol* 139(Suppl 53):9–12
- Zanger P, Holzer J, Schleucher R *et al.* (2009) Constitutive expression of the antimicrobial peptide RNase 7 is associated with *Staphylococcus aureus* infection of the skin. *J Infect Dis* 200:1907–15

## Generation of Increased Numbers of HLA-DR<sup>high</sup> IgG<sup>+</sup> Plasma Cells in the Peripheral Blood of Patients with Bullous Pemphigoid: NC16a-Specific Cells Belong to the Short-Lived Plasma Blast Population

*Journal of Investigative Dermatology* (2010) 130, 2838–2841; doi:10.1038/jid.2010.213; published online 29 July 2010

#### TO THE EDITOR

Bullous pemphigoid (BP) is a rare, usually severe and potentially life-threatening disease. Immunologically, this disorder is characterized by the presence of circulating IgG autoantibodies targeting distinct adhesion molecules of the dermoepidermal basement membrane zone (Hertl, 2000). Binding of autoantibodies to adhesion structures

of the skin leads to a massive loss of function, resulting in blister formation. The majority of BP patients possesses high-affinity IgG autoantibodies directed against the 180 kDa BP autoantigen (BP180; Diaz *et al.*, 1990). The immunodominant site of BP180 is located within the short NC16a domain, which is recognized by more than 90% of BP patients. In the previous studies we have

identified the presence of memory B cells specific for the NC16a domain, which can be induced *in vitro* to synthesize autoantibodies (Leyendeckers *et al.*, 2003). Also, it was shown that the serum level of autoantibodies to NC16a correlates with the severity of BP (Schmidt *et al.*, 2000). The bulk of antibodies measurable in the serum are synthesized by terminally differentiated B lymphocytes, that is, plasma cells (PCs), mostly residing in the bone

Abbreviations: BP, bullous pemphigoid; BM, bone marrow; PB, plasma blast; PC, plasma cell