of the exome was affected by CNVs in KNSTRN-mutated tumors versus 1.4% (SD 3.0%) in KNSTRN-wild-type tumors (P=0.74). Although this was not statistically significant, our ability to detect differences may have been limited by a small sample size.

In SCC, KNSTRN p.S24F is present in 19% of tumor precursors, suggesting that it arises early in disease progression (Lee et al., 2014). To determine whether KNSTRN mutagenesis is an early event in BCC development as well, we screened 30 early stage BCCs for KNSTRN mutations. We identified a nonsynonymous KNSTRN mutation in only 1/30 (3%) early stage BCCs, suggesting that, unlike in SCC, mutant KNSTRN in BCC appears to be acquired later in disease and is possibly a marker of aggressive behavior (Figure 1c and f). The mutation, pH284D, is absent from dbSNP137 and ESP6500 and is predicted to be deleterious by SIFT; however, it has not previously been reported in the COSMIC database and is not predicted to be damaging by Polyphen.

These findings are the first to implicate KNSTRN in BCC tumorigenesis. Alongside recent data offering a role for KNSTRN in SCC and melanoma, our work supports the classification of KNSTRN as an oncogene and an important contributor to the pathogenesis of malignancies related to UV-exposure. In both SCC and BCC, mutant KNSTRN disrupts sister chromatid cohesion and promotes genomic instability in functional assays. However, unlike in SCC, KNSTRN mutations in BCC appear to occur late in disease progression and are preferentially found in advanced tumors. Further exploration of the role of KNSTRN in skin cancer and genomic stability is warranted.

# CONFLICT OF INTEREST

The authors state no conflict of interest.

#### ACKNOWLEDGMENTS

This work was funded by the V Foundation Translational Award, NIAMS (5ARO54780, 2ARO46786), NIH Pathway to Independence Award 1K99CA176847 (SXA), the Damon Runyon Clinical Investigator Award (JYT), Stanford Medical Scholars Program (PDJ) and the Dermatology Foundation Career Development Award (KYS).

# Prajakta D. Jaju<sup>1</sup>, Christine B. Nguyen<sup>1</sup>, Angela M. Mah<sup>1</sup>, Scott X. Atwood<sup>1</sup>, Jiang Li<sup>1</sup>, Amin Zia<sup>2</sup>, Anne Lynn S. Chang<sup>1</sup>, Anthony E. Oro<sup>1</sup>, Jean Y. Tang<sup>1</sup>, Carolyn S. Lee<sup>1</sup> and Kavita Y. Sarin<sup>1</sup>

<sup>1</sup>Department of Dermatology, Stanford University School of Medicine, Stanford, California, USA and <sup>2</sup>Stanford Center for Genomics and Personalized Medicine, Stanford University School of Medicine, Stanford, California, USA. E-mail: ksarin@gmail.com

#### SUPPLEMENTARY MATERIAL

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

#### REFERENCES

- Atwood SX, Sarin KY, Whitson RJ *et al.* (2015) Smoothened variants explain the majority of drug resistance in basal cell carcinoma. *Cancer Cell* 27:342–53
- Epstein EH (2008) Basal cell carcinomas: attack of the hedgehog. *Nat Rev Cancer* 8:743–54
- Fang L, Seki A, Fang G (2009) SKAP associates with kinetochores and promotes the metaphase-to-anaphase transition. *Cell Cycle* 8:2819–27
- Jayaraman SS, Rayhan DJ, Hazany S et al. (2014) Mutational landscape of basal cell carcinomas by whole-exome sequencing. J Investig Dermatol Symp Proc 134:213–20
- Lee CS, Bhaduri A, Mah A *et al.* (2014) Recurrent point mutations in the kinetochore gene KNSTRN in cutaneous squamous cell carcinoma. *Nat Genet* 46:1060–2
- Lomas A, Leonardi-Bee J, Bath-Hextall F (2012) A systematic review of worldwide incidence of nonmelanoma skin cancer. Br J Dermatol 166: 1069–80
- Ng PC, Henikoff S (2001) Predicting deleterious amino acid substitutions. *Genome Res* 11: 863–74
- Sunyaev S, Ramensky V, Bork P (2000) Towards a structural basis of human non-synonymous single nucleotide polymorphisms. *Trends Genet* 16:198–200
- Xi R, Hadjipanayis AG, Luquette LJ et al. (2011) Copy number variation detection in wholegenome sequencing data using the Bayesian information criterion. Proc Natl Acad Sci USA 108:E1128–36

# Estrogen Upregulates Slug to Enhance the Migration of Keratinocytes

Journal of Investigative Dermatology (2015) 135, 3200-3203; doi:10.1038/jid.2015.315; published online 24 September 2015

## **TO THE EDITOR**

Declining estrogen levels in postmenopausal women are related to delayed wound healing. It is known that estrogen influences skin re-epithelialization (Ashcroft *et al.*, 1997) and may promote skin wound healing via estrogen receptor- $\beta$  (ER- $\beta$  Campbell *et al.*, 2010). However, the cellular and subcellular mechanisms of estrogen's action on keratinocyte migration are still poorly understood. Epithelial–

mesenchymal transition (EMT) is an important process in which epithelial cells lose their epithelial characteristics and acquire mesenchymal traits. This significant phenotypic conversion leads to the acquisition of motile behavior. In the skin, EMT has been shown to have a role in skin morphogenesis (Kong *et al.*, 2006), skin cancers (Shimokawa *et al.*, 2013; Wong *et al.*, 2013; Magnoni

Abbreviations: E2, 17β-estradiol; EMT, epithelial-mesenchymal transition; HaCaT, immortalized human skin keratinocyte cell line; mRNA, messenger RNA; PHK, primary human keratinocyte Accepted article preview online 20 August 2015; published online 24 September 2015



**Figure 1. Upregulated Slug, Twist, and vimentin and downregulated E-cadherin in HaCaT cells after treatment with E2.** (a) Quantitative real-time PCR (qRT-PCR) analysis of E-cadherin, vimentin, Slug, and Twist in HaCaT cells after treatment with 0.1, 1, 10, 100, and 1,000 nm E2 for 48 h. (b) Western blot analysis of E-cadherin, vimentin, Slug, and Twist in HaCaT cells. (c) Estrogen receptor (ER) mediates E2-induced E-cadherin, vimentin, Slug, and Twist expression in HaCaT cells. Inhibition of ER, but not GPR30, blocked E2-induced Slug, Twist, and vimentin messenger RNA (mRNA) expression. (d) Slug knockdown impairs the migration of HaCaT cells. HaCaT cells were transfected with siRNA oligo specific for Slug (Slug siRNA) or a scrambled siRNA oligo (scrambled siRNA) before stimulations with E2, and the migration of cells were assessed using scratch assay and are represented as percentage (%) of wound remaining open (Scale bar = 100 µm). All results are representative or means  $\pm$  SD of three independent experiments. Differences were determined by one-way analysis of variance (ANOVA) or two-way ANOVA. \**P*<0.05, \*\**P*<0.001 versus vehicle-treated or E2 100 nm-treated HaCaT cells.

et al., 2014; Qu et al., 2014), wound healing (Savagner et al., 2005; Arnoux et al., 2008; Hudson et al., 2009), and skin fibrosis (Postlethwaite et al., 2004; Takahashi et al., 2013). In this study, we tested the hypothesis that estrogeninduced acceleration of wound healing is mediated by EMT during keratinocyte migration. This study was approved by the institutional review board and written informed consent was obtained from all subjects before they participated in the study. All experimental details are given in Supplementary Materials and Methods online.

As the expression of ER in human skin remains contentious, the expression of ERs in HaCaT (immortalized human skin keratinocyte cell line) cells and primary human keratinocytes (PHK) was analyzed by immunofluorescence. T47D, a human ductal breast epithelial tumor cell line, was used as a positive control of ERs. Whereas expression of both ER- $\alpha$  and ER- $\beta$  was observed in T47D cells, HaCaT cells, and PHK expressed only ER- $\beta$  (Supplementary Figures S1a and c online). The effect of 17 $\beta$ -estradiol (E2) treatment on viability of HaCaT cells was measured using an MTT assay. Although E2 had little effect on cell viability at lower concentrations, there was a statistically significant reduction in cell viability after treatment with 1  $\mu$ M E2 (P<0.0001, Supplementary Figure S1d online).

To investigate the role of EMT in E2-induced keratinocyte migration, we first analyzed whether E2 could modulate the expression of EMT-biomarkers. HaCaT cells were cultured to 80% of confluency and then incubated with 0.1, 1, 10, 100 nm, or 1,000 nm E2 for 48 h. We found that concentrations of E2 ranging from 1 to 100 nm significantly induced the expression of Slug, Twist, and vimentin and significantly repressed the expression of E-cadherin, whereas treatment with 1,000 nm E2 had little or no effect (Figure 1a and b). We next utilized ICI, a potent ER antagonist, and G15, an antagonist of another estrogen-related receptor, GPR30, to evaluate the E2 mechanism of action on HaCaT cells. E2 altered expression of the EMT-biomarkers, Slug, Twist, vimentin, and E-cadherin, was reversed following treatment with ICI; however, G15 could not abrogate the changes in expression of EMT-biomarkers (Figure 1c). These data indicate that estrogen regulates expression of EMT-related molecules through ERs but not GPR30. Next, to determine the role of Slug in the E2-dependent modulation of EMT-related molecules, cells were transfected with Slug siRNA before stimulation with E2. We observed that expression of E-cadherin was increased, whereas that



**Figure 2.** Upregulated Slug and Twist, and downregulated E-cadherin in an *ex vivo* skin explant model after E2 treatment. Immunofluorescence confocal images were obtained for Slug (green), Twist (red), and E-cadherin (red) in vehicle-treated (upper), 10 nm E2-treated (middle), and 100 nm E2-treated (lower) skin. The image is representative data of three independent experiments using nine skin explants obtained from two donors. Nuclei were stained with DAPI (blue). Scale bar = 50 µm.

of vimentin was reduced (Supplementary Figure S2 online), as compared with scrambled siRNA control.

To investigate whether estrogen can alter the rate of cell migration, a scratch assay was performed. Confluent monolayers of HaCaT cells were scratched with a sterile P200 pipette tip, and cellular debris was completely removed by washing. Cells were then treated with 0.1, 1, 10, or 100 nm E2, and images of the same field were captured at 0, 24, 48, and 72 h after scratching. As shown in Supplementary Figure S3a online, HaCaT cells incubated with all concentrations of E2 migrated toward the center of the wound to a greater extent than those treated with vehicle. In order to confirm results obtained from the scratch assay, transwell assays were performed. When various concentrations of E2 (0.1, 1, 10, and 100 nm) were added to the lower chambers, HaCaT cells in the upper chambers significantly migrated towards E2, as compared with those treated with vehicle (Supplementary Figure S3b online). To further determine whether Slug is involved in the E2-induced migration of HaCaT cells, scratch assay after Slug siRNA transfection was performed. The migration of HaCaT cells was markedly inhibited by transfection with Slug siRNA, compared with cells as

transfected with scrambled siRNA control (Figure 1d). Collectively, these results suggest a critical role for Slugmediated EMT in the E2-mediated enhancement of keratinocyte migration.

Because HaCaT cells are immortalized keratinocytes that may behave differently from normal human skin, we investigated the effects of E2 on expression of Slug in an ex vivo skin explant model. As the endogenous estrogen levels and response to exogenous estrogen are different depending on subjects' age and gender, skin explants were obtained from postmenopausal donors (Supplementary Table 1 online). At 2 days post-treatment with 10 or 100 nm E2, epidermal thickness was significantly increased in E2-treated epidermis compared with that in vehicletreated epidermis. (Figure 2). Immunofluorescence labeling together with confocal laser microscopy using specific mABs revealed higher cellular expression of Slug and lower expression of E-cadherin in the 10 nm E2-teated (Figure 2, middle image) and 100 nm E2-treated (Figure 2, lower image) epidermis compared with vehicle-treated skin control (Figure 2, upper image). In addition, Slug was expressed mostly in the lower epidermis of vehicle-treated skin, however, it was expressed in the entire epidermis of 10 nm E2- and 100 nm E2-treated skin. Also, at 2 days post-treatment with 10 or 100 nm E2, epidermal thickness was significantly increased in E2-treated epidermis compared with that in vehicle-treated epidermis (Supplementary Figure S4 online). This suggests that E2 can modulate EMT in normal epidermis.

The Snail family of transcription factors is best known for its ability to trigger EMT through repression of E-cadherin (Batlle et al., 2000; Cano et al., 2000; Bolos et al., 2003), which is important for the strength of cellular adhesion in a tissue and cellular motility. Savagner et al. (2005) analyzed the expression of Slug during wound healing using in situ hybridization and reported a crucial role for Slug in keratinocyte migration in vitro. Hudson et al. (2009) also showed that re-epithelialization of excisional wound healing was significantly impaired in Slug-deficient mice. In addition, EGF, the most potent effector of keratinocyte migration, induced activation of ERK5 and upregulated Slug (Arnoux et al., 2008).

The estrogen-induced molecular changes demonstrated in this study are consistent with changes seen during EMT. Estrogen also accelerated keratinocyte migration, and this effect was lost in Slug knockdown cells. Our data indicate that estrogen might promote re-epithelialization in wound healing via an EMT process. This highlights the possibility that further exploring the role of estrogen-induced EMT in the skin may yield novel insights into critical processes such as aging, fibrosis, and cancer metastasis.

### **CONFLICT OF INTEREST**

The authors state no conflict of interest.

#### ACKNOWLEDGMENTS

This research was supported by Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Ministry of Science, ICT, and future Planning (NRF-2010-0010475). This study was also supported by an Amore-Pacific Research Grant in 2011 (Korea). We thank Dong-Su Jang, (Medical Illustrator, Medical Research Support Section, Yonsei University College of Medicine, Seoul, Korea) for his help with the illustrations.

# Jung U. Shin<sup>1,8</sup>, Ji Y. Noh<sup>1,8</sup>, Shan Jin<sup>1,2,3</sup>, Seo H. Kim<sup>1,2</sup>, Dong K. Rah<sup>4</sup>, Dong W. Lee<sup>4</sup>, Jong S. Yoo<sup>5,6</sup>, Kunhong Kim<sup>7</sup>, Yun S. Lee<sup>1</sup>, Inhee Jung<sup>1</sup>, Ju H. Lee<sup>1</sup> and Kwang H. Lee<sup>1,2</sup>

<sup>1</sup>Department of Dermatology, Severance Hospital, Cutaneous Biology Research Institute, Yonsei University College of Medicine, Seoul, Korea; <sup>2</sup>Brain Korea 21 PLUS Project for Medical Science, Yonsei University College of Medicine, Seoul, Korea; <sup>3</sup>Department of Dermatology, Yanbian University Hospital, Yanji, Jilin, China; <sup>4</sup>Department of Plastic and Reconstructive Surgery, Severance Hospital, Institute for Human Tissue Restoration, Yonsei University College of Medicine, Seoul, Korea; Biochemistry and Molecular Biology, Yonsei University College of Medicine, Seoul, Korea E-mail: juhee@yuhs.ac or kwanglee@yuhs.ac <sup>8</sup>The first two authors contributed equally to this work.

#### SUPPLEMENTARY MATERIAL

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

## REFERENCES

- Arnoux V, Nassour M, L'Helgoualc'h A et al. (2008) Erk5 controls Slug expression and keratinocyte activation during wound healing. *Mol Biol Cell* 19:4738–49
- Ashcroft GS, Dodsworth J, van Boxtel E et al. (1997) Estrogen accelerates cutaneous wound healing associated with an increase in TGFbeta1 levels. Nat Med 3:1209–15
- Batlle E, Sancho E, Franci C et al. (2000) The transcription factor snail is a repressor of E-cadherin gene expression in epithelial tumour cells. Nat Cell Biol 2:84–9
- Bolos V, Peinado H, Perez-Moreno MA *et al.* (2003) The transcription factor Slug represses E-cadherin expression and induces epithelial to mesenchymal transitions: a comparison with Snail and E47 repressors. *J Cell Sci* 116: 499–511
- Campbell L, Emmerson E, Davies F et al. (2010) Estrogen promotes cutaneous wound healing via estrogen receptor beta independent of its antiinflammatory activities. J Exp Med 207: 1825–33
- Cano A, Perez-Moreno MA, Rodrigo I et al. (2000) The transcription factor snail controls epithelial-mesenchymal transitions by

repressing E-cadherin expression. *Nat Cell Biol* 2:76–83

- Hudson LG, Newkirk KM, Chandler HL et al. (2009) Cutaneous wound reepithelialization is compromised in mice lacking functional Slug (Snai2). J Dermatol Sci 56:19–26
- Kong W, Li S, Liu C *et al.* (2006) Epithelialmesenchymal transition occurs after epidermal development in mouse skin. *Exp Cell Res* 312: 3959–68
- Magnoni C, Giudice S, Pellacani G et al. (2014) Stem cell properties in cell cultures from different stage of melanoma progression. *Appl Immunohistochem Mol Morphol* 22: 171–81
- Postlethwaite AE, Shigemitsu H, Kanangat S (2004) Cellular origins of fibroblasts: possible implications for organ fibrosis in systemic sclerosis. *Curr Opin Rheumatol* 16:733–8
- Qu X, Shen L, Zheng Y et al. (2014) A signal transduction pathway from TGF-beta1 to SKP2 via Akt1 and c-Myc and its correlation with progression in human melanoma. J Investig Dermatol Symp Proc 134:159–67
- Savagner P, Kusewitt DF, Carver EA et al. (2005) Developmental transcription factor slug is required for effective re-epithelialization by adult keratinocytes. J Cell Physiol 202:858–66
- Shimokawa M, Haraguchi M, Kobayashi W et al. (2013) The transcription factor Snail expressed in cutaneous squamous cell carcinoma induces epithelial-mesenchymal transition and down-regulates COX-2. Biochem Biophys Res Commun 430:1078–82
- Takahashi M, Akamatsu H, Yagami A et al. (2013) Epithelial-mesenchymal transition of the eccrine glands is involved in skin fibrosis in morphea. J Dermatol 40:720–5
- Wong CE, Yu JS, Quigley DA et al. (2013) Inflammation and Hras signaling control epithelial-mesenchymal transition during skin tumor progression. Genes Dev 27:670–82