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Mrgprs activation is required for chronic itch conditions in mice

Yuyan Zhu, PhD^a, Claire E. Hanson, BSc^a, Qin Liu, PhD^b, Liang Han, PhD^{a,*}

Introduction: Chronic itch has been drawing much attention due to its clinical significance and the complexity of its mechanisms. To facilitate the development of anti-itch strategies, it is necessary to investigate the key players in itch sensation under chronic itch conditions. Several members of the Mrgpr family were identified as itch receptors that detect cutaneous pruritogens in primary sensory neurons. However, the role of Mrgprs in chronic itch conditions has not been well described.

Methods: Scratching behaviors of Wildtype and *Mrgpr-clusterΔ^{-/-}* mice were examined in dry skin model and contact dermatitis model to examine the role of *Mrgpr* genes in mediating chronic itch sensation. Scratching behaviors of the mice were also examined in allergic itch model. Real-time polymerase chain reaction were performed to examine the expression level of MrgprA3 and MrgprC11 under naive and dry skin conditions. The MrgprA3⁺ itch-sensing fibers were labeled by tdTomato fluorescence in *Mrgpra3^{GFP-Cre}*; *ROSA26^{tdTomato}* mice, and the morphology and density of those fibers in the epidermis were analyzed under dry skin condition.

Results: We showed that deleting a cluster of *Mrgpr* genes in mice reduced scratching behavior severely under 2 chronic itch conditions, namely dry skin and contact dermatitis, and the allergic itch condition. Moreover, the gene expressions of itch receptors MrgprA3 and MrgprC11 in dorsal root ganglia were upregulated significantly under dry skin condition. Consistently, the percentage of MrgprA3⁺ itch-sensing neurons was increased as well. We also observed hyperinnervation of MrgprA3⁺ itch-sensing fibers in the epidermis of the skin under dry skin condition.

Discussion: We demonstrate that Mrgprs play important roles in mediating chronic itch and allergic itch. These findings enrich our knowledge of itch mechanism and may lead to the development of novel therapeutic approach to combat itch.

Keywords: Chronic itch, MrgprA3, MrgprC11, Skin

Introduction

Chronic itch accompanying cutaneous disease or systemic disorders is a serious condition^[1–3]. In older adults, xerosis (aka dry skin), a common skin problem, usually causes pruritus, and the scratching response results in a high risk of secondary infection in skin^[4]. Patients with advanced renal disease and liver disease frequently report pruritus as a common symptom^[2,3]. The persistent itching significantly affects patients' quality of life^[5–7]. Despite its clinical importance, the molecular and cellular mechanisms of chronic itch are largely unknown.

Recent studies have identified several members of the Mrgpr (mas-related G-protein coupled receptor) family as new itch receptors^[8–12]. The Mrgpr family consists of >20 functional members that are grouped into several subfamilies: MrgprA1–22, MrgprB1–13, MrgprC1–14, and MrgprD–G. Among them, MrgprA3, MrgprC11, and MrgprD have been identified as itch receptors that are specifically expressed in small-diameter sensory neurons and directly detect pruritogens in the skin^[9,11,12]. MrgprA3 is the receptor for chloroquine, a drug used to treat malaria and generates itch as a side effect. MrgprC11 is the receptor for pruritogen bovine adrenal medulla 8–22 peptide (BAM8–22) and SLIGRL-β-alanine, a muscle building supplement that frequently elicits itch in human, acts through MrgprD. Moreover, the expression of MrgprA3 defines a subset of nociceptors that specifically mediate itch, but not pain^[13]. These neurons express multiple itch receptors including MrgprA3, MrgprC11, and histamine receptor 1, and correspondingly respond to multiple pruritogens. Deletion of MrgprA3⁺ sensory neurons significantly inhibits itch behavior not only in acute itch conditions^[14], but also in chronic itch conditions^[13,15], demonstrating that this neuronal population is crucial for mediating itch sensation. However, whether the chronic itch response requires the activation of Mrgprs in MrgprA3⁺ neurons is unknown.

Here we show that the scratching behaviors evoked under 2 chronic itch conditions, namely dry skin and contact dermatitis, were significantly suppressed in Mrgprs-deficient mice. We also found that allergen-induced scratching behaviors were reduced after the deletion of Mrgprs. Moreover, dry skin chronic itch condition increased the expression of itch receptors MrgprA3 and

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MrgprC11. We observed for the first time that dry skin condition induced the hyperinnervation of MrgprA3⁺ itch-sensing fibers in the epidermis. Taken together, we demonstrate that Mrgprs play a critical role in mediating chronic itch and allergic itch.

Experimental procedures

Animals

All experiments were performed with approval from the Georgia Institute of Technology Animal Use and Care Committee. The generations of *Mrgpr-clusterΔ^{-/-}* mice and *Mrgpra3^{GFP-Cre}* mice were described previously^[12,13]. *ROSA26^{tdTomato}* mouse line was purchased from Jackson Laboratory. All the mice used had been backcrossed to C57BL/6 mice for at least 10 generations. 2- to 3-month-old males (20–30 g) were used for all experiments. Mice were housed in the vivarium with 12-hour-light/dark cycle, and all the behavioral tests were performed from 9 AM to 1 PM in the light cycle. The housing group was 5 at maximum with food and water ad libitum.

Scratch assay

All procedures were performed using protocols approved by the Animal Care and Use Committee of Georgia Institute of Technology. The experiments were performed as previously described^[13]. Briefly, the day before the scratch assay, all the mice were acclimated for at least 30 minutes to their testing environment. During scratch assay, mice were kept in plastic cylinders individually and the behavioral responses were video recorded for 30–60 minutes. The video recording was subsequently played back in a slow motion and the number of bouts of scratching with the hindpaw were counted. All behavioral tests were performed by an experimenter blind to the genotypes of the mice.

Three itch conditions

Dry skin pruritus^[16]

The shaved back skin of a mouse was treated with an acetone/ether (1:1) mixture for 15 seconds followed by distilled water (Acetone, Ether and Water) twice daily for 6 days to generate skin dehydration. On day 7, mice were recorded for 60 minutes to quantify the spontaneous scratching.

Contact dermatitis^[17]

A mouse model of contact dermatitis is produced by treating the skin with an allergen squaric acid dibutylester (SADBE, 25 μL, 0.5% in acetone). On days 1–3, SADBE was applied to the shaved abdomen once a day to initiate T-lymphocyte sensitization. On days 8–10, SADBE was applied to the shaved back skin to induce dermal inflammation and spontaneous scratching. On day 12, the spontaneous scratching behavior was recorded for 60 minutes.

Allergic itch^[18]

Ovalbumin (OVA) (50 μg) dissolved in phosphate-buffered saline (PBS) was administered intraperitoneally together with 2 mg of inject Alum twice at a 10-day interval. One week after the second sensitization, 50 μg of OVA was injected subcutaneously into the nape of a mouse's neck to induce acute allergic itch. The scratching behavior was recorded for 30 minutes.

Real-time polymerase chain reaction (RT-PCR)

Dorsal root ganglia (DRG) innervating the dry skin-treated area (C3–T10) were collected from the trunk region of mice and total RNA was extracted from the ganglia using the RNeasy Micro Kits (Qiagen) according to the manufacturer protocol. RNA was reverse-transcribed into cDNA using SuperScript III First-Strand Synthesis System (Invitrogen). Quantitative RT-PCR was performed using a StepOnePlus Real-Time PCR System (Thermo Fisher Scientific) with the SYBR Green detection method. The cycle threshold (C_T) values were analyzed by the 2^{-ΔC_T} method to determine the normalized expression ratio of target genes. β-actin expression level was used as the internal control. The intron-spanning primers used are listed here: β-actin-F, AGCTGTGC TATGTTGCTCTAG, β-actin-R, AGGTCCTTTACGGATGTCAA CG, MrgprA3-F, ACACAAGCCAGCAAGCTACA, MrgprA3-R, ACTTCCAGGGATGGTTTCGT, MrgprC11-F, AGCATCCACA ACCCCAGAAG, MrgprC11-R, TGGAGTGCAGTTGGGATGAC. TRPV1-F, ATCATCAACGAGGACCCAGG. TRPV1-R, TGCTAT GCCTATCTCGAGTGC.

Quantification of itch-sensing neurons and fibers in the skin

After 6 days of dry skin treatment, *MrgprA3^{GFP-Cre}*; *ROSA26^{tdTomato}* mice were anesthetized with pentobarbital and perfused with PBS (pH 7.4, 4°C) followed with 4% paraformaldehyde (4°C). DRG innervating the treated skin (C3–T10) were dissected from the perfused mice. DRG were postfixed in 4% paraformaldehyde at 4°C for 30 minutes. Treated skin were dissected and fixed in 2% paraformaldehyde at 4°C overnight. All tissues were cryoprotected in 20% sucrose in PBS at 4°C for 24 hours, and frozen in Optimal cutting temperature at –80°C. Tissues were sectioned at 20 μm with a Cryostat NX50 Cryostat (Thermo Fisher scientific). The sections on slides were dried at 37°C for 30 minutes, and fixed with 4% paraformaldehyde at room temperature for 10 minutes. The slides were washed with PBS, and the fluorescence labeled neurons or skin fibers were visualized and quantified under fluorescent microscope. Totally 3000–8000 sensory neurons from each mouse were counted to quantify the percentage of tdTomato labeled MrgprA3⁺ neurons. Cell nuclei in the skin sections were stained with 4',6-diamidino-2-phenylindole to visualize the epidermis. Skin sections (10–22) were analyzed to quantify the intraepidermal density of tdTomato labeled MrgprA3⁺ skin fibers. n = 3 for both naive group and dry skin-treated group.

Histologic analysis

The treated skin were collected and fixed in 10% neutralized formalin for >24 hours. The skin samples were embedded in paraffin, sectioned at 5 μm, and stained with hematoxylin and eosin. The hematoxylin and eosin-stained images were taken using Nikon Ni-U microscopy and DS-Fi2 color camera. At least 10 sections per sample were analyzed. The area of each section to measure epidermal thickness was randomly picked. The skin thickness and epidermal thickness were analyzed using Image J.

Statistical analysis

Data are presented as means ± SEM. Statistical analysis was performed using the Student 2-tailed *t* test to compare between groups. *P*-value <0.05 was considered statistically significant.

Figure 1. *Mrgpr* genes are required for dry skin itch. A, Hematoxylin and eosin staining of the skin sections collected from WT and *Mrgpr-clusterΔ^{-/-}* mice after dry skin treatment. B, Dry skin treatment induced comparable increase of the epidermal thickness in WT and *Mrgpr-clusterΔ^{-/-}* mice ($n \geq 10$ sections from 3 to 5 mice per group). C, *Mrgpr-clusterΔ^{-/-}* mice exhibited reduced scratching behavior under dry skin condition. $n \geq 5$ for each group. *** $P < 0.005$. Two-tailed unpaired Student *t* test. Error bars represent SEM. Scale bar = 100 μm. WT indicates Wildtype.

Results

*Mrgpr*s mediate chronic itch and allergic itch

A straightforward way to determine the role of *Mrgpr*s in chronic itch is to delete *Mrgpr* genes and see whether scratching behavior is affected. As many *Mrgpr* genes in mice are clustered together, to avoid compensatory effects, *Mrgpr-clusterΔ^{-/-}* mouse has been generated previously in which 12 *Mrgpr* genes including 2 identified itch receptors, *MrgprA3* and *MrgprC11*, have been deleted in the mouse genome^[12]. We compared the scratching behavior of wild-type and *Mrgpr-clusterΔ^{-/-}* mice under 2 different chronic itch conditions: dry skin^[16] and contact dermatitis^[17]. Skin dehydration is always accompanied by persistent itchiness of the skin, especially in the elderly^[4]. In the mouse model, the dry skin condition was generated by repetitive exposure of the skin to the mixture of acetone and ether^[16]. The treatment induced pronounced epidermal hyperplasia and parakeratosis, but did not induce infiltration of the inflammatory cells^[16]. Histologic analysis demonstrated that the treated skin

collected from Wildtype (WT) and *Mrgpr-clusterΔ^{-/-}* mice exhibited similar morphologic changes (Fig. 1A) and comparable increase of the epidermal thickness (Fig. 1B). These results suggest that the deletion of *Mrgpr* genes did not affect the development of the dry skin model. Excitingly, *Mrgpr-clusterΔ^{-/-}* mice exhibited significant decrease in scratching behavior compared with WT mice (Fig. 1C), suggesting that the activation of *Mrgpr* genes are required for dry skin itch.

Contact dermatitis is a condition of skin inflammation produced by unwanted contact with irritant or allergic chemicals. The mouse model of contact dermatitis is generated by treating the skin with the allergen SADBE. The treatment generated acanthosis (thickening of the skin), parakeratosis, and dense inflammatory cells infiltration in both WT and *Mrgpr-clusterΔ^{-/-}* mice (Fig. 2A). The skin thickness is comparable between the 2 groups after the treatment (Fig. 2B). However, *Mrgpr-clusterΔ^{-/-}* mice showed a severe reduction in scratching behavior compared with WT mice suggesting that *Mrgpr* genes mediate itch sensation in the contact dermatitis model (Fig. 2C).

Figure 2. *Mrgpr* genes are required for itch sensation in contact dermatitis model. A, Hematoxylin and eosin-stained skin sections collected from WT and *Mrgpr-clusterΔ^{-/-}* mice after contact dermatitis treatment. B, WT and *Mrgpr-clusterΔ^{-/-}* mice showed similar increase of skin thickness after contact dermatitis treatment ($n \geq 10$ sections from 3 to 5 mice per group). C, *Mrgpr-clusterΔ^{-/-}* mice exhibited reduced scratching behavior in contact dermatitis model. $n \geq 5$ for each group. *** $P < 0.005$. Two-tailed unpaired Student *t* test. Error bars represent SEM. Scale bar = 100 μm. WT indicates Wildtype.

Figure 3. *Mrgprs clusterΔ^{-/-}* mice exhibited reduced allergic itch behavior. $n \geq 5$ for each group. * $P < 0.05$. Two-tailed unpaired Student *t* test. Error bars represent SEM.

We next examined whether *Mrgprs* are involved in acute allergen-induced itch sensation. In the allergic itch mouse model, the allergen OVA was administered intraperitoneally twice, 10 days apart, to sensitize the mice. Injection of OVA subcutaneously 1 week after the second sensitization induced acute allergic responses and scratching behavior directed to the injected site. We found that *Mrgpr-clusterΔ^{-/-}* mice exhibited significantly less scratching behavior compared with WT littermates (Fig. 3) under the allergic itch condition. In summary, our data showed that *Mrgprs* play a critical role in mediating the itch sensation in 3 tested itch conditions.

Dry skin condition upregulates the expression of *Mrgprs*

MrgprA3 and *MrgprC11*, 2 previously identified itch receptors, were among the 12 genes that were deleted in *Mrgpr-clusterΔ^{-/-}* mice. Both of them are specifically expressed in the small-diameter sensory neurons^[8,12,13]. *MrgprA3* gene expression has been shown to be upregulated in sensory neurons in the dry skin model^[19]. The percentage of sensory neurons responding to

chloroquine and SLIGRL were increased in the dry skin model^[20,21]. Consistently, we found that the mRNA levels of *MrgprA3* and *MrgprC11* were significantly increased after the dry skin treatment (Figs. 4A, B). In the contrary, the expression of *TRPV1*, a sensor for noxious heat and pain sensation^[22,23], was not changed in the dry skin model.

Next, we examined whether the percentage of *MrgprA3⁺* neurons, the previously identified specific itch-sensing neurons^[13], increased after the dry skin treatment. Our previous study generated a BAC transgenic mouse line *MrgprA3^{GFP-Cre}* to express GFP-Cre fusion protein in *MrgprA3⁺* neurons^[13]. *MrgprA3⁺* neurons were labeled with tdTomato fluorescence by crossing the *MrgprA3^{GFP-Cre}* line with a Cre-dependent *ROSA26^{tdTomato}* line (Fig. 5A). We collected the DRG sections from the *MrgprA3^{GFP-Cre}*; *ROSA26^{tdTomato}* mice with or without dry skin treatment, and found that the percentage of tdTomato labeled *MrgprA3⁺* neurons from dry skin-treated mice was significantly higher than that from control animals (Fig. 5B). Collectively, our data demonstrated that the dry skin condition enhanced the expression of *MrgprA3* and *MrgprC11* and expanded the expression of *MrgprA3* into a broader sensory neuron population.

Dry skin condition induces hyperinnervation of *MrgprA3⁺* itch-sensing fibers in the epidermis

Pruritic skin is often accompanied by alteration of the sensory innervations in the skin. Skin biopsies from atopic dermatitis patients showed enhanced intraepidermal fiber density labeled by PGP9.5^[24,25]. Dry skin pruritus has been shown to be associated with an increase of Ret-positive intraepidermal fiber density^[20]. *MrgprA3⁺* sensory neurons, which constitute about 8% of sensory neurons and exclusively innervate the epidermis of the skin, are required for mediating itch sensation induced by multiple pruritogens^[13]. Whether and how the *MrgprA3⁺* itch-sensing fibers are changed in the skin by the chronic itch conditions have not yet been investigated. With the tdTomato fluorescence labeling in the neuron, we were able to visualize the periphery projection of *MrgprA3⁺* itch-sensing fibers in the skin (Figs. 6A–D). Epidermis in the skin sections was identified by the morphology of the keratinocytes labeled by 4',6-diamidino-2-phenylindole. Consistent

Figure 4. Dry skin condition upregulates the expression of *Mrgpr* genes. Real-time polymerase chain reaction results showing that the expression of *MrgprA3* (A) and *MrgprC11* (B) were upregulated in the dorsal root ganglia under the dry skin condition. The expression of *TRPV1* was not changed (C). $n = 3$ for *MrgprA3* and *TRPV1* and $n = 6$ for *MrgprC11*. * $P < 0.05$. Two-tailed unpaired Student *t* test. Error bars represent SEM. n.s. indicates not significant.

Figure 5. Dry skin condition increases the percentage of MrgprA3⁺ itch-sensing neurons in the dorsal root ganglia (DRG). A, DRG sections of *Mrgpra3*^{GFP-Cre}; *ROSA26*^{tdTomato} mice showing the MrgprA3⁺ neurons labeled by tdTomato fluorescence under control or dry skin condition. tdTomato fluorescent images were merged with phase contrast images to show the morphology of the DRG sections. B, The percentage of MrgprA3⁺ neurons in the DRG under control or dry skin condition. n = 3 for each experiment. ***P < 0.005. Two-tailed unpaired Student *t* test. Error bars represent SEM. Scale bar = 100 μ m.

with previous results, MrgprA3⁺ fibers were free nerve endings in the epidermis and terminated in the stratum granulosum, the superficial layer of the epidermis^[13] (Figs. 6A–D). Epidermal hyperplasia induced by the dry skin treatment can be observed in dry skin sections but not in control sections (Figs. 6B, D). We found that the morphology of MrgprA3⁺ fibers was different under the chronic itch condition compared with the naive condition. The fibers from the naive condition normally penetrated the epidermis to reach the stratum granulosum and did not show fiber branches in the epidermis. However, about 13% of the MrgprA3⁺ fibers under chronic itch condition generated branches in the epidermis (Figs. 6A–E). Moreover, MrgprA3⁺ intraepidermal fiber density increased significantly under dry skin conditions (Fig. 6F). These data collectively demonstrate that the dry skin condition induced

hyperinnervation of the MrgprA3⁺ itch-sensing fibers in the epidermis.

Discussion

Itch sensation is initiated by the activation of sensory nerves in the skin. Itch receptors expressed in the nerve endings are responsible for first sensing the pruritogens. Among all the itch receptors, Mrgprs are relatively new but play a critical role in itch sensation. In this study, we showed that deleting 12 *Mrgpr* genes in mice suppressed scratching behavior in 2 chronic itch conditions and allergic itch condition. We further demonstrate that dry skin chronic itch condition triggered the active expression of itch receptors MrgprA3 and MrgprC11 in the DRG sensory neurons

Figure 6. Dry skin condition induces hyperinnervation of itch-sensing fibers in the epidermis. A–D, Skin sections of *Mrgpra3*^{GFP-Cre}; *ROSA26*^{tdTomato} mice with (A, B) and without (C, D) DAPI staining showing the MrgprA3⁺ fibers labeled by tdTomato fluorescence in the epidermis. The dashed line labels the boundary between dermis and epidermis. Arrow point to a branched fiber in the epidermis of a dry skin section. E, The percentage of MrgprA3⁺ fibers with branches in the epidermis under control and dry skin conditions. F, The MrgprA3⁺ intraepidermal nerve fiber (IENF) density under control and dry skin conditions. n = 3 for each experiment. ***P < 0.005. Two-tailed unpaired Student *t* test. Error bars represent SEM. Scale bar = 20 μ m. DAPI indicates 4',6-diamidino-2-phenylindole.

and the morphologic expansion of itch-sensing nerves in the skin. Our data highlight the critical role of Mrgpr3 in mediating chronic itch and allergic itch.

Clinically, itch is categorized into 4 groups: dermatologic, systemic, neurological, and psychiatric^[26]. Currently, only a handful of mouse models for itch conditions have been established^[14]. We have used 3 representative dermatologic mouse itch models in this study, including a nonallergic chronic itch model (Acetone, Ether and Water-dry skin), an allergic chronic itch model (SADBE-contact dermatitis), and an allergic acute itch model (OVA-allergic itch). These 3 models involve distinct signaling pathways. Dry skin treatment decreases stratum corneum hydration, induced epidermal hyperplasia, and parakeratosis, but does not induce skin inflammation^[16] (Figs. 1A, B). SADBE-contact dermatitis model represents the allergic skin inflammation condition. The challenged skin is infiltrated with inflammatory cells and soaked in an inflammatory soup of cytokines and chemokines^[27]. OVA-allergic itch model evokes IgE-dependent mast cell-mediated acute allergic response. OVA challenge induces the secretion of a variety of mast cell mediators in the skin to evoke cutaneous anaphylactic symptoms including itch sensation^[28,29]. It is intriguing that our data suggest that Mrgpr3 can be activated in 3 different models. As 12 Mrgpr genes were deleted in the *Mrgpr-clusterΔ^{-/-}* mice, it is possible that different Mrgpr genes mediate itch sensation in different conditions. Alternatively, a certain Mrgpr gene may be activated by different pruritogens produced in different conditions. For example, MrgprC11 has multiple identified agonists such as neuropeptide FF (NPFF), Cathepsin S, and Der p1^[12,30–32], all of which can be good candidates for evoking itch. NPFF is a neuropeptide that can be released from mouse bone marrow-derived mast cells^[30]. It is possible that NPFF is released by the skin mast cells in the OVA-allergic itch condition and activate MrgprC11⁺ nerves. Cathepsin S is associated with inflammatory processes and chronic itch conditions. The expression of cathepsin S is upregulated in psoriatic keratinocytes^[33]. Mice carrying a transgene expressing cathepsin S spontaneously develop atopic dermatitis^[34]. Der p1 is a prominent allergen from house dust mite and has been involved in the allergic inflammatory processes in both skin^[35,36] and the respiratory system^[37]. It will be interesting to examine whether cathepsin S and Der p1 are endogenous agonists for MrgprC11 under the itch conditions we tested.

Patients with chronic itch often display heightened sensitivity to itchy stimuli, which can be partly attributed to peripheral sensitization of sensory neurons^[38–41]. MrgprA3⁺ sensory neurons are a major sensory neuron population mediating itch sensation^[13]. Consistently, MrgprA3⁺ neurons exhibit enhanced excitability in a mouse model of inflammatory itch^[15]. Here, we used the dry skin chronic itch model to examine how chronic itch condition change the MrgprA3⁺ itch-sensing neurons. We showed that the dry skin treatment increased the gene expression of *MrgprA3* and *MrgprC11*, and increased the percentage of MrgprA3⁺ neurons in DRG (Fig. 2). The dry skin treatment mainly generates skin barrier disruption and stratum corneum dehydration. It is very likely that the mediators secreted by keratinocytes during the treatment promote upregulation of itch receptors. In addition, neuronal activity-dependent transcription regulation is a common mechanism used by many neuron types including DRG sensory neurons for proper development and adaptation during adulthood^[42,43]. It is also possible that the continuous activation of MrgprA3⁺ itch-sensing neurons under dry skin condition induced the

activity-dependent transcription of itch receptors. These hypotheses need to be tested with further studies.

We have analyzed the morphology and density of itch-sensing nerve fibers in the epidermis under dry skin condition. MrgprA3⁺ itch-sensing neurons only constitute a small percentage of DRG sensory neurons. Their intraepidermal nerve density is high enough to sense the potential stimuli in every spot in the skin, but not too high to prevent the clear visualization and quantification of the nerve fibers and the fiber branches. Using *Mrgpra3^{GFP-Cre}*; *ROSA26^{tdTomato}* mouse, we were able to observe the hyperinnervation of the MrgprA3⁺ itch-sensing fibers under dry skin conditions (Fig. 6). Notably, this is the first time that the morphologic changes of MrgprA3⁺ itch-specific fibers under chronic itch condition have been observed. Both upregulation of itch receptors and hyperinnervation of itch-sensing fibers might contribute to the hypersensitivity of itchy skin to pruritogens.

Taken together, our data suggest that Mrgpr3 play an important role in mediating itch sensation in multiple conditions. Human MrgprX1 is the functional receptor for both CQ and BAM8–22 and is the functional orthologue for MrgprA3 and MrgprC11^[12,44]. Further studies are required to investigate the involvement of hMrgprX1 in chronic itch conditions in human patients and to test the possibility to antagonize hMrgprX1 for anti-itch treatment.

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Conflict of interest disclosure

The authors declare that they have no financial conflict of interest with regard to the content of this report.

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