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Evoked itch perception is associated with changes in functional brain connectivity

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Abstract

Chronic itch, a highly debilitating condition, has received relatively little attention in the neuroimaging literature. Recent studies suggest that brain regions supporting itch in chronic itch patients encompass sensorimotor and salience networks, and corticostriatal circuits involved in motor preparation for scratching. However, how these different brain areas interact with one another in the context of itch is still unknown. We acquired BOLD fMRI scans in 14 atopic dermatitis patients to investigate resting-state functional connectivity before and after allergen-induced itch exacerbated the clinical itch perception in these patients. A seed-based analysis revealed decreased functional connectivity from baseline resting state to the evoked-itch state between several itch-related brain regions, particularly the insular and cingulate cortices and basal ganglia, where decreased connectivity was significantly correlated with increased levels of perceived itch. In contrast, evoked itch increased connectivity between key nodes of the frontoparietal control network (superior parietal lobule and dorsolateral prefrontal cortex), where higher increase in connectivity was correlated with a lesser increase in perceived itch, suggesting that greater interaction between nodes of this executive attention network serves to limit itch sensation via enhanced top-down regulation. Overall, our results provide the first evidence of itch-dependent changes in functional connectivity across multiple brain regions.

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1. Introduction

The brain circuitry supporting itch sensation is an active topic of research. While numerous studies have investigated and described pain-related brain mechanisms (Apkarian et al., 2005; Henry et al., 2011), much less is known about the brain circuitry involved with itch in spite of the clinical relevance of this highly debilitating symptom, which is present across multiple chronic itch conditions (Yosipovitch et al., 2003). Moreover, recent studies suggest that pain perception is associated with altered resting-state functional brain connectivity, which refers to stereotypical patterns of co-activation across multiple brain regions that show both trait and pain-state specificity (e.g., Napadow et al., 2010; Napadow et al., 2012; Loggia et al., 2013; reviewed in Fomberstein et al., 2013). However, itch-related functional brain connectivity, especially in chronic itch patients, has never been reported.

Atopic dermatitis (AD), also known as atopic eczema, is a pruritic chronic inflammatory disease of the skin, affecting approximately 17.8 million persons in the United States, in which patients experience highly debilitating itch (Finlay, 2001; Lupidus, 2001; Yosipovitch et al., 2003; Berke et al., 2012). Previous studies indicate that AD involves not only peripheral sensitization for itch (Lee and Yu, 2011; Rahman et al., 2011) but also central sensitization, in an analogous fashion as central sensitization for pain in chronic pain patients (Koltzenburg, 2000), such that stimuli that are perceived as painful in healthy subjects are
experienced as itching in AD patients (Ikoma et al., 2004; Ikoma et al., 2006; Schmelz, 2010). Not only specific allergens, but also stress and psychosocial factors, can exacerbate AD symptoms (Gil and Sampson, 1989; Koblenez, 1999; Rahman et al., 2011). This complex symptomatology, and its bidirectional relationship to stress and psychosocial factors, which can be both a cause and a consequence of itching and scratching (Gil and Sampson, 1989; Lapidus, 2001; Ark and Paus, 2006; Chida et al., 2008; Mizawa et al., 2013; Yaghaime et al., 2013; Schut et al., 2014), point to an essential role of the central nervous system in AD (Ikoma et al., 2004; Misery, 2011; Pfab et al., 2012a; Darlenski et al., 2014).

In healthy subjects, a number of studies indicate that experimentally-induced itch activates a brain network which includes pre-motor and supplementary motor area, thalamus, and cingulate, insular, inferior parietal, and prefrontal cortices (reviewed in Pfab et al., 2012a). However, few studies have investigated the brain circuitry supporting itch in AD. Studies from our group and others suggest that brain response to experimentally induced itch in AD differs from healthy subjects, consistent with abnormal itch sensitivity in AD (Ikoma et al., 2004; Schneider et al., 2008; Ishiiju et al., 2009; Pfab et al., 2010; Pfab et al., 2011; Pfab et al., 2012b; Napadow et al., 2014). In a recent functional MRI (fMRI) study, we evoked clinically-relevant itch in AD patients by using individually-matched allergens and our validated temperature-modulated itch induction procedure (Napadow et al., 2014). In this procedure, a thermode is used to modulate skin temperature between 32 °C (warm) and 25 °C (cool), in 20-s blocks (Pfab et al., 2006). This procedure has reliably been used to experimentally modulate itch, such that cooling the skin gradually increases itch sensation over the first 10 s, followed by a phase in which itch sensation reaches a peak plateau (Pfab et al., 2006; Pfab et al., 2010). We found that different brain regions were activated during these different phases of itch induction (i.e., increasing itch and peak itch plateau). During the increasing itch phase, the right anterior insula, right anterior middle cingulate cortex, bilateral striatum (putamen and caudate), right globus pallidus, and right ventrolateral prefrontal cortex were activated. Conversely, during the peak itch plateau phase the right dorsolateral prefrontal cortex, bilateral premotor cortex, and left superior parietal lobule were activated. Notably, reduction of itch after non-pharmacological (acupuncture) therapy was associated with reduced activation in the putamen and right anterior insula (Napadow et al., 2014), further highlighting the important role of these regions in itch processing.

In the present study, we investigated functional brain connectivity before and after experimentally inducing (or exacerbating) itch, using allergen skin prick in nonlesional skin while AD patients were resting in the MRI scanner. We used functional connectivity magnetic resonance imaging (fcMRI) with seed-based correlation methods to evaluate changes in functional brain connectivity associated with itch. To our knowledge, this is the first study to investigate how functional brain connectivity is modulated by itch, particularly the clinically-relevant itch state—a common cause of suffering in AD patients.

2. Materials and methods

2.1. Study participants

Patients aged 18–60 with a clinical diagnosis of AD and a score greater than 18 on the SCORing Atopic Dermatitis (SCORAD) scale (European Task Force on Atopic Dermatitis 1993; Schmitt et al., 2013) were enrolled. Recruitment was by print and e-mail advertisement as well as referrals from physician colleagues in the Department of Dermatology at the Massachusetts General Hospital. Volunteers who met eligibility criteria via phone screening were invited to a preliminary session in the laboratory during which a licensed dermatologist (F.P.) assessed their eligibility to the study via medical examination and focused history. In addition, eligible participants had to show type-I sensitivity to grass pollen and/or dust mites, as demonstrated by wheal and flare formation upon skin prick testing. For each participant, the most effective itch-inducing allergen was determined among Timothy grass pollen (100,000 bioequivalent allergen units per milliliter) and the two most common types of house dust mites in North America and Europe (Dermatophagoides pteronyssinus,10,000 allergen units per milliliter, and Dermatophagoides farinae,10,000 allergen units per milliliter, Allergy Laboratories, Oklahoma City, OK). Participants also completed the Edinburgh Inventory for handedness (Oldfield, 1971). The study protocol was approved by the Human Research Committee of Massachusetts General Hospital and all study participants gave informed consent. Participants were required to stop all immunosuppressive medications at least 10 days prior to the study to avoid potential suppression of itch perception.

2.2. Data acquisition

As part of a previously published study of the anti-pruritic effects of acupuncture (Napadow et al., 2014), each participant took part in an initial clinical screening session followed by two separate fMRI sessions (at least 1 week apart) during which resting-state scans were performed before and after allergen itch induction (but before any verum or placebo treatment was applied), with corresponding self-report ratings of perceived itch. The baseline resting-state scan (REST) occurred at the beginning of the session. The allergen-evoked itch resting-state scan (ITCH) was performed 15–17 min after allergen skin prick, within the period of time when allergen induced itch sensation is known to persist after skin prick (Pfab et al., 2010). A single drop of the most pruritogenic allergen (as determined during the clinical screening session) was applied on a non-lesion site on the left volar forearm. The skin was then punctured with a plastic MR-compatible lancet (Duotip Test II, Lincoln Diagnostics, Decatur, IL) such that the allergen solution was deposited at the dermal–epidermal junction, where the terminals of itch-related C-fibers are located (Shelley and Arthur, 1957). Using this procedure, an itch sensation develops with a median latency of 35 s and persists more than 15 min after application (Pfab et al., 2010). An absorbent gauze pad was then used to carefully remove the drop of allergen solution 120 s after application. With this method, the skin puncture is sufficiently shallow to avoid sustained pain, as previously reported (Pfab et al., 2006), and can be considered an induction of pure itch rather than a combination of itch and pain.

Brain imaging was performed on a Siemens Trio 3 Tesla MRI scanner (Siemens AG, Erlangen, Germany) with vendor-supplied 32-channel multi-array head coil. A high-resolution T1-weighted anatomical scan was collected using an isotropic multi-echo MPRAGE pulse sequence (TR/TE1/TE2 = 2530/1.64/1200 ms, 256 × 256 matrix, 256 mm field-of-view (FOV), 7° flip angle) (van der Kouwe et al., 2008). Functional imaging (BOLD fMRI) was performed using a gradient echo T2*-weighted pulse sequence (TR/TE = 2 s/30 ms, 32 anterior commissure–posterior commissure (AC–PC) aligned slices, slice thickness 3.6 mm, 64 × 64 matrix, 200 mm FOV, 90° flip angle). Each resting-state fMRI scan had a duration of 6 min (180 time points). Electrocardiography (ECG) and respiratory activity were simultaneously recorded throughout the scans using a Powerlab system (ML880, ADInstruments, Colorado Springs, CO) at a 400 Hz sampling rate. ECG data were acquired and filtered using an MR-compatible physiological monitor (Magnitude 3150 MRI Patient Monitor, In vivo, Gainesville, Florida) designed to minimize radio frequency and gradient switching artifacts generated during the MRI scan. Respiratory data were collected using a custom-built system based on that devised by Binks et al. (2007) which consists of two MR-compatible pneumobelt placed around the chest and abdomen and connected to air pressure transducers (PX138-0.3D5V, Omegaadyne, Inc., Sunbury, Ohio).

Immediately before the baseline scan and immediately after the induced-itch scan, subjects were asked to rate the intensity of experienced itch on a visual analog scale (VAS) from 0 (no itch) to 100 (most intense itch imaginable), with 33 corresponding to an “urge to
temporal fieldmap correction (dewarping); (2) physiological noise correction with RETROICOR; (3) slice timing correction (‘slicetimer’, FSL); (4) head motion correction (MCFLIRT, FSL (Smith et al., 2002)); (5) skull stripping (BET, FSL (Smith, 2002)); (6) nonlinear registration to the Montreal Neurological Institute (MNI) template (FLIRT/FNIRT, FSL); and (7) 0.008 Hz band-pass temporal filtering (‘1dBandpass’, AFNI). The translation parameters resulting from the motion correction step were then used to compute the relative mean motion during each scan. Relative mean motion was defined as the average over time of the absolute displacement of each brain volume as compared to the previous volume in time, where the absolute displacement at each time point was computed as the root-mean-square of the absolute displacement at each time point was computed as the root-

### 2.3. Data analysis

#### 2.3.1. Preprocessing

BOLD data preprocessing and analysis were performed using tools from the FMRIB Software Library (FSL) [http://www.fmrib.ox.ac.uk/fsl](http://www.fmrib.ox.ac.uk/fsl) (Smith et al., 2004; Woolrich et al., 2009), the FreeSurfer (V. 5.2) suite [http://surfer.nmr.mgh.harvard.edu/](http://surfer.nmr.mgh.harvard.edu/) (Dale et al., 1999; Fischl et al., 1999; Fischl et al., 2002; Fischl et al., 2004), AFNI (Cox, 1996; Cox and Hyde, 1997), and RETROICOR (Glover et al., 2000) for retrospective correction of physiological motion artifacts using our peripheral measures of cardiac and respiratory activities as independent assessments of physiological noise in the BOLD signal.

Preprocessing consisted of: (1) $B_0$ fieldmap correction (dewarping); (2) physiological noise correction with RETROICOR; (3) slice timing correction (‘slicetimer’, FSL); (4) head motion correction (MCFLIRT, FSL (jenkinson et al., 2002)); (5) skull stripping (BET, FSL (Smith, 2002)); (6) nonlinear registration to the Montreal Neurological Institute (MNI) template (FLIRT/FNIRT, FSL); and (7) 0.008–0.1 Hz band-pass temporal filtering (‘1dBandpass’, AFNI). The translation parameters resulting from the motion correction step were then used to compute the relative mean motion during each scan. Relative mean motion was defined as the average over time of the absolute displacement of each brain volume as compared to the previous volume in time, where the absolute displacement at each time point was computed as the root-mean-square of the $x$, $y$, and $z$ translation parameters (Van Dijk et al., 2012).

#### 2.3.2. Seed-based functional connectivity analyses

Functional connectivity was computed using a conventional seed-based correlation analysis method. This method consists in (1) choosing a specific voxel or cluster of voxels in the brain, referred to as the seed, (2) extracting the time series of the BOLD signal in this seed region, and (3) using this time series as a regressor in a general linear model (GLM) analysis to calculate whole-brain, voxel-wise functional connectivity maps of covariance with this seed region (Biswal et al., 1995; Cole et al., 2010). Our seed regions of interest were chosen as the clusters significantly activated during the “increasing itch” or “peak itch plateau” phase from an evoked, allergen-induced itch block fMRI scan in the same AD subjects, as previously reported (Napadow et al., 2014). Importantly, activation clusters came from different fMRI runs than those investigated with our resting fcMRI approach, assuring independence between the seed fMRI time series during rest and that during temperature-modulated evoked itch scanning. The brain regions from which ROI seeds were chosen were the right anterior insula (MNI coordinates in the Talairach–Tournoux Atlas: $X = 30$ mm, $Y = 18$ mm, $Z = 2$ mm), right anterior mid-cingulate cortex (aMCC) ($X = 8, Y = 34, Z = 32$), right putamen ($X = 24, Y = 18, Z = 4$), left putamen ($X = −22, Y = 10, Z = 10$), right caudate ($X = 20, Y = 18, Z = 8$), left caudate ($X = −20, Y = 10, Z = 16$), right globus pallidus (X = 18, Y = 4, Z = 4), right ventrolateral prefrontal cortex (vPFC) ($X = 34, Y = 40, Z = 6$), right dorsolateral prefrontal cortex (dlPFC) ($X = 36, Y = 20, Z = 52$), right premotor cortex (PMC) ($X = 22, Y = −4, Z = 62$), left PMC ($X = −20, Y = −20, Z = 62$), and left superior parietal lobule (SPL) ($X = −28, Y = −44, Z = 64$) (see Table 1). For each seed region, the BOLD signal was averaged across a 4-mm radius sphere centered on each cluster’s peak voxel. The resulting time series was then used as a regressor in a standard GLM analysis using FSL, along with 10 regressors of no interest: 6 motion correction parameters, the average time series from cerebrospinal fluid voxels in the lateral ventricles, the average time series from white matter voxels in deep parietal white matter, and cardiac and respiratory time series obtained by convolving the cardiac and respiratory signals with hemodynamic response functions from the ‘RVHR’ model (Chang et al., 2009). The output of this GLM analysis for each seed region of interest was a 3D volume (“brain map”) in which each voxel value represented the strength of functional connectivity between this seed and every other voxel in the brain.

#### 2.3.3. ITCH–REST difference map for each subject

For each subject, a difference map was computed by calculating the difference in parameter estimates for connectivity taken from REST

### Table 1

Seed-based functional connectivity results. For each seed, statistically significant clusters (with a cluster-forming threshold of $Z > 2.3$ and a cluster significance level of $p < 0.05$) are listed. For each cluster, the Z-value, MNI coordinates, and atlas label of the peak voxel are listed, along with Pearson’s correlation coefficient $r$ and corresponding $p$-value of the correlation with level of perceived itch.

<table>
<thead>
<tr>
<th>Seed region</th>
<th>$Z$-value</th>
<th>X</th>
<th>Y</th>
<th>Z</th>
<th>Atlas label</th>
<th>Correlation with perceived itch</th>
</tr>
</thead>
<tbody>
<tr>
<td>R premotor cortex</td>
<td>3.56</td>
<td>−9</td>
<td>−64.5</td>
<td>66</td>
<td>L precuneus</td>
<td>n.s.</td>
</tr>
<tr>
<td>R insula</td>
<td>3.16</td>
<td>8</td>
<td>−48</td>
<td>61</td>
<td>R superior parietal lobule (5L)</td>
<td>n.s.</td>
</tr>
<tr>
<td>R putamen</td>
<td>3.92</td>
<td>−30</td>
<td>−44</td>
<td>−14</td>
<td>L fusiform gyrus</td>
<td>n.s.</td>
</tr>
<tr>
<td>R putamen</td>
<td>3.81</td>
<td>−18</td>
<td>−36</td>
<td>−2</td>
<td>L parahippocampal gyrus</td>
<td>$r = −0.39, p = 0.045$</td>
</tr>
<tr>
<td>R putamen</td>
<td>3.61</td>
<td>−12</td>
<td>−60</td>
<td>14</td>
<td>L posterior cingulate cortex</td>
<td>$r = −0.42, p = 0.025$</td>
</tr>
<tr>
<td>R putamen</td>
<td>3.62</td>
<td>−52</td>
<td>−66</td>
<td>6</td>
<td>L middle temporal gyrus</td>
<td>n.s.</td>
</tr>
<tr>
<td>R putamen</td>
<td>3.38</td>
<td>−4</td>
<td>−18</td>
<td>40</td>
<td>L mid-cingulate cortex</td>
<td>n.s.</td>
</tr>
<tr>
<td>R putamen</td>
<td>3.78</td>
<td>−60</td>
<td>−20.5</td>
<td>22.5</td>
<td>L postcentral gyrus</td>
<td>n.s.</td>
</tr>
<tr>
<td>R putamen</td>
<td>3.95</td>
<td>−61</td>
<td>−17.5</td>
<td>11</td>
<td>L transverse (or superior) gyrus, BA42</td>
<td>n.s.</td>
</tr>
<tr>
<td>R putamen</td>
<td>3.68</td>
<td>−2</td>
<td>−22</td>
<td>48</td>
<td>L medial frontal gyrus, BA6 or L paracentral lobule</td>
<td>n.s.</td>
</tr>
<tr>
<td>R superior parietal lobule</td>
<td>3.92</td>
<td>−53</td>
<td>−24.5</td>
<td>32</td>
<td>L inferior parietal lobule</td>
<td>n.s.</td>
</tr>
<tr>
<td>R superior parietal lobule</td>
<td>3.69</td>
<td>−41.5</td>
<td>29</td>
<td>27.5</td>
<td>L dIPFC (Middle frontal gyrus)</td>
<td>$r = −0.40, p = 0.036$</td>
</tr>
<tr>
<td>R superior parietal lobule</td>
<td>3.61</td>
<td>43.5</td>
<td>−10.5</td>
<td>15</td>
<td>R posterior insula</td>
<td>n.s.</td>
</tr>
<tr>
<td>R superior parietal lobule</td>
<td>4.24</td>
<td>29.5</td>
<td>−14</td>
<td>11.5</td>
<td>R putamen (not the same locus as the R putamen seed)</td>
<td>n.s.</td>
</tr>
<tr>
<td>R anterior mid-cingulate cortex</td>
<td>4.32</td>
<td>34</td>
<td>−21.5</td>
<td>10</td>
<td>R posterior insula</td>
<td>n.s.</td>
</tr>
<tr>
<td>R anterior mid-cingulate cortex</td>
<td>4.04</td>
<td>28.5</td>
<td>−12.5</td>
<td>9.5</td>
<td>R putamen</td>
<td>$r = −0.40, p = 0.036$</td>
</tr>
<tr>
<td>R caudate</td>
<td>4.03</td>
<td>58</td>
<td>−26.5</td>
<td>22</td>
<td>R inferior parietal lobule</td>
<td>n.s.</td>
</tr>
<tr>
<td>R caudate</td>
<td>4.39</td>
<td>−10.5</td>
<td>−73</td>
<td>23.5</td>
<td>L cuneus, BA18</td>
<td>n.s.</td>
</tr>
<tr>
<td>R globus pallidus</td>
<td>3.86</td>
<td>−51.5</td>
<td>19.5</td>
<td>−4.5</td>
<td>L inferior frontal gyrus (pars orbitalis)</td>
<td>r = −0.37, p = 0.050</td>
</tr>
<tr>
<td>R globus pallidus</td>
<td>3.98</td>
<td>−5</td>
<td>−22.5</td>
<td>44</td>
<td>L mid-cingulate cortex or paracentral lobule</td>
<td>n.s.</td>
</tr>
<tr>
<td>R globus pallidus</td>
<td>4.01</td>
<td>35.5</td>
<td>27</td>
<td>49</td>
<td>R superior/middle frontal gyrus (dIPFC)</td>
<td>n.s.</td>
</tr>
</tbody>
</table>
versus ITCH scan runs using ‘fslmaths’ (FSL). The variance map for this difference map was estimated by adding the parameter estimate variances computed from the REST and ITCH scan runs. These maps, calculated from all subjects, were then passed up to a group level analysis using FSL, as described below.

2.3.4. Group-level analysis of main effect of itch on functional connectivity

A group-level analysis yielded a paired difference map between REST and ITCH across subjects and therefore indicated the main effect of itch induction on functional connectivity. Multiple comparisons correction was performed with cluster correction, with a cluster-forming threshold of Z > 2.3 and a cluster significance level of p < 0.05. Finally, for visualization purposes, the resulting brain maps were spatially smoothed using the cubic resampling function in AFNI and overlaid on the 0.5 mm MNI152 template.

2.3.5. Correlation between perceived itch and level of functional connectivity

Using only the brain regions in which a main effect was found, we then performed a linear regression analysis between the average main effect within the region and the itch perception score difference between the REST and ITCH conditions. This analysis specifically indicated which functional connectivity changes between REST and ITCH were correlated with perceived change in itch, thus linking changes in functional connectivity with itch more directly than the main effect analysis alone, and taking advantage of the intersubject variability in itch responses. Therefore, in this final analysis, changes in functional connectivity were deemed related to itch only in brain regions where there was both (i) a main effect of itch and (ii) a significant correlation across subjects between change in functional connectivity and change in perceived itch (from REST to ITCH).

3. Results

Fourteen AD patients (8 females, 13 right-handed, age: 25.4 ± 9.1 years old, SCORAD: 38.7 ± 14.9, mean ± standard deviation) participated in the study, with itch induction performed using the most pruritogenic allergen for each patient among Timothy grass pollen (N = 8), D. pteronyssinus house dust mite (N = 3), or D. farinae house dust mite (N = 3).

3.1. Itch rating levels

Perceived itch increased significantly after allergen induction (REST: 6.4 ± 11.7; ITCH: 32.0 ± 16.6, mean ± standard deviation on a 0–100 visual analog scale; paired t-test, p < 10^{-7}; Fig. 1), as expected. There was no significant correlation between perceived itch and age across subjects, neither during REST (r = 0.14, p = 0.63), nor during ITCH (r = 0.18, p = 0.54).

3.2. Head motion and respiration during the scan

Since subject motion during the scan can significantly affect measures of functional connectivity, it is important to measure and report head motion in any study of functional connectivity (Power et al., 2012; Van Dijk et al., 2012). Here we verified that mean motion ranged within previously published values in a healthy population (Van Dijk et al., 2012) and did not significantly differ between REST (0.100 ± 0.043 mm, mean ± standard deviation) and ITCH (0.097 ± 0.037 mm; paired t-test, p = 0.9). We further verified that gross translational motion never exceeded 3 mm on any axis. We also verified that the participants’ breathing rates did not change between the REST condition (0.35 Hz ± 0.11, mean ± standard deviation) and the ITCH condition (0.34 Hz ± 0.10, paired t-test, p = 0.19).

3.3. Correlation between perceived itch and level of functional connectivity

While functional connectivity changed significantly from REST to ITCH for several of the seed regions of interest (see Table 1), the change in functional connectivity was significantly correlated with change in perceived itch for a smaller subset of regions. Among these regions, functional connectivity decreased during ITCH, compared to REST, in most cases. Specifically, connectivity was decreased and negatively correlated with change in itch for connectivity between the right anterior insula and left hippocampal formation (MNI coordinates of the peak voxel: X = −18, Y = −36, Z = −2, r = −0.39, p = 0.045) and left posterior cingulate cortex (PCC, X = −12, Y = −60, Z = 14, r = −0.42, p = 0.025), between the right aMCC and right putamen (X = 28.5, Y = −12.5, Z = 9.5, r = −0.40, p = 0.036), and between the right globus pallidus and left posterior mid-cingulate cortex (pMCC, X = −5, Y = −25.5, Z = 44, r = −0.37, p = 0.050; Fig. 2 and Table 1). Thus, for these connectivity changes, a higher increase in perceived itch was associated with a greater decrease in connectivity. The exception to this pattern was for connectivity between the left SPL and left dlPFC connectivity also reported lower change in ITCH, and a negative correlation between change in connectivity and change in itch (r = −0.40, p = 0.036; Fig. 3 and Table 1). In other words, a higher increase in perceived itch was associated with a lower increase in SPL–dlPFC connectivity.

4. Discussion

In this fcMRI study of allergen-induced itch in AD patients, we found that an increase in perceived itch was significantly correlated with a decrease in functional connectivity between brain regions previously demonstrated to activate in response to phasic itch provocation (Napadow et al., 2014). These brain regions included the insula, cingulate cortex, and basal ganglia. Specifically, greater itch was associated with reduced connectivity between the right anterior insula and both hippocampal formation and posterior cingulate cortex. Similarly, itch was associated with reduced connectivity between the anterior mid-cingulate cortex and putamen, and between the globus pallidus and posterior mid-cingulate cortex. While itch was mostly associated with reduced functional connectivity, increased connectivity between the superior parietal lobule and dorsolateral prefrontal cortex was associated with an increase in perceived itch. Moreover, a negative correlation was found between SPL–dlPFC connectivity and itch rating; that is, subjects showing greater increase in SPL–dlPFC connectivity also reported lower increase in itch sensation. This result suggests that greater interaction between the SPL and dlPFC serves to limit itch sensation via enhanced top-down regulation. Our study clearly demonstrates for the first time that “resting” functional connectivity is altered when AD patients...
Fig. 2. Changes in functional connectivity from the baseline resting state (‘REST’) to the allergen-evoked itch state (‘ITCH’). A. Seed-based functional connectivity from the right anterior insula (MNI coordinates: [30, 18, 1]). Top: Significant cluster in the left hippocampal formation (peak voxel coordinates: [−30, −44, −14] in the fusiform gyrus). Bottom: Significant cluster in the left posterior cingulate cortex (PCC) (peak voxel coordinates: [−12, −60, 14]). In each of the two panels, the graph on the left shows $z$-score levels of functional connectivity (main effect), in the resting state (‘REST’; white bar) and 15–17 min after allergen-based itch induction (‘ITCH’; black bar). The graph on the right shows for each scan session ($N = 28$) the difference in connectivity from ‘REST’ to ‘ITCH’ as a function of the difference in perceived itch intensity from ‘REST’ to ‘ITCH’, as reported by participants; linear regression across subjects is shown (red line) along with corresponding $r$ and $p$ values. B. Seed-based functional connectivity from the right anterior mid-cingulate cortex (coordinates: [8, 34, 28]), showing a significant cluster in the right putamen (peak voxel coordinates: [28, −12, 9]). C. Seed-based functional connectivity from the right globus pallidus (coordinates: [18, 4, 3]), showing a significant cluster in the left posterior mid-cingulate cortex (peak voxel coordinates: [−5, −25, 44]).
experience itch, and these state-specific alterations can be directly linked with the intensity of perceived itch.

Overall, we found that a sustained itch state altered functional connectivity for nodes known to be activated by evoked itch. For instance, we found that functional connectivity between the basal ganglia (caudate and globus pallidus) and cingulate cortex, normally present at rest (Postuma and Dagher, 2006), decreased with increased itch (see Fig. 2B, 2C). This finding is reminiscent of our recent pain neuroimaging study in which we found that experimentally-induced sustained myofascial pain in healthy adults reduced functional connectivity between the somatotopic cortical representation within the primary sensorimotor (SI/M1) area contralateral to the pain stimulus site and other regions of the primary sensorimotor network (Kim et al., 2013). Hence, in both studies, brain areas activated by an evoked stimulus (whether itch or pain) showed decreased connectivity within their intrinsic connectivity network.

Evoked itch in our AD patients was also associated with a decrease in functional connectivity between the right anterior insula and both left hippocampal formation and left PCC (see Fig. 2A), both of which are nodes of the default mode network (DMN) (Buckner et al., 2008). Interestingly, while the anterior insula is not usually connected with DMN regions, chronic pain patients show greater DMN-insula connectivity at rest than healthy controls (Napadow et al., 2010; Loggia et al., 2013), and even in healthy subjects, DMN-insula connectivity increases with state anxiety (Dennis et al., 2011). In our AD subjects, functional connectivity between the right anterior insula and these two DMN nodes (left hippocampal formation and left PCC) was present during a resting state and decreased after itch induction, and this decrease was significantly correlated with an increase in perceived itch. The allergen-induced itch may therefore have disrupted the resting connectivity patterns that support chronic (versus evoked) itch perception in these patients, perhaps by shifting attention to an evoked stimulus. Together, these results reinforce the notion that unpleasant interoceptive experiences such as itch or pain may decrease functional connectivity between brain regions that are functionally connected at rest.

In contrast, while connectivity between itch-processing brain areas was mostly decreased during the sustained itch state, increased connectivity was noted between SPL and dIPFC. In fact, we found a negative correlation between itch intensity and the change in connectivity between the SPL and dIPFC. In other words, a higher increase in functional connectivity between SPL and dIPFC was associated with a lower increase in perceived itch after allergen itch induction (see Fig. 3). This suggests that increased SPL–dIPFC connectivity may be a protective mechanism, limiting perceived itch severity after allergen itch induction in AD. Both the SPL and dIPFC sub-regions in our results are key nodes of the frontoparietal control network, which supports executive processing of cognitive control (Vincent et al., 2008). Hence, our finding suggests that subjects able to regulate increased information exchange within this network were better able to limit the severity of sustained itch sensation after allergen itch induction. Interestingly, we previously found that SPL and dIPFC were activated by evoked allergen itch during the “peak itch” plateau phase (Napadow et al., 2014), an experimental condition likely to be more closely related to sustained itch than the preceding “increasing itch” phase, when itch sensation was steadily increasing. Of note, the brain regions that were activated during the “increasing itch” phase (anterior insula, cingulate, putamen) instead showed decreased connectivity in response to sustained evoked itch. This suggests that different brain networks become engaged when itch sensation first develops versus when itch is a steadily ongoing perception.

The continuous, sustained itch state induced in our study was associated with altered functional connectivity between several brain regions which previous studies have reported as being activated by evoked itch stimuli. In healthy subjects, brain activation during itch has been investigated by performing functional brain imaging – with either positron emission tomography (PET), BOLD fMRI, or arterial spin labeling (ASL) fMRI – while inducing itch experimentally using various means such as histamine prick, allergen prick, cowhage spicule application, or electrical stimulation (Walter et al., 2005; Herde et al., 2007; Leknes et al., 2007; Mochizuki et al., 2007; Pfab et al., 2008; Valet et al., 2008; Yosipovitch et al., 2008; Mochizuki et al., 2009; Vierow et al., 2009; Jeffry et al., 2011; Kleyn et al., 2012; Papoiu et al., 2012; Pfab et al., 2012a). These studies, most of them conducted with healthy subjects rather than chronic itch patients, highlight the involvement of a network of brain regions which includes ipsilateral pre-motor and supplementary motor areas and contralateral cingulate cortex, insular cortex, thalamus, inferior parietal cortex, and dorsolateral prefrontal cortex (reviewed in Pfab et al., 2012a). In addition, the above studies found that itch stimuli activate habit-encoding motor planning areas, including the putamen, probably in relation to the motor act of scratching which is strongly associated with itch (Yosipovitch et al., 2008; Vierow et al., 2009). In the present study, itch perception was associated with reduced connectivity between the insula, cingulate cortex, and basal ganglia (i.e., putamen, globus pallidus) – regions whose activation
was strongly associated with itch in previous evoked-itch neuroimaging studies.

Notably, functional connectivity for basal ganglia areas such as the putamen and globus pallidus was significantly altered during sustained itch. These areas are critical in regulating both voluntary and involuntary movements, and may play an important role in the scratch response to itch. Putamen activation has been reported not only during scratching, with greater activation in the presence of itch than in its absence, but also during itching without scratching, suggesting that putamen activation may represent the urge and/or intention to scratch (Vierow et al., 2009). We previously reported that putamen activation was maximal during an increasing itch phase in AD patients subjected to allergen itch (Napadow et al., 2014). The putamen is also a key structure within the striato-thalamo-cortical circuitry implicated in action motivation and initiation and in habitual, repetitive behavior (Graybiel, 2008). This circuit is dysfunctional in various psychopathologies, including addiction (Koob and Volkow, 2010), which is consistent with its implication in the “urge” to scratch (Leknes et al., 2007; Vierow et al., 2009).

Here we found that functional connectivity between the basal ganglia (caudate and globus pallidus) and cingulate cortex decreased with increased itch (see Fig. 2B, 2C). The cingulate cortex is part of the salience network (caudate and globus pallidus) and cingulate cortex decreased with inactivation in the motor cortex of resting human brain using echo-planar MRI. Magn. Reson. Med. 34 (4), 537–541. http://dx.doi.org/10.1002/mrm.1910340409.8524021.


