



Abstract

In the present study, we used functional MRI in awake rats to follow the pain response that accompanies intradermal injection of capsaicin into the hindpaw. To this end, we used BOLD imaging together with a 3D segmented, annotated rat atlas and computational analysis to identify the integrated neural circuits involved in capsaicin-induced pain. The specificity of the pain response to capsaicin was tested in a transgenic knock-out of the TRPV1 receptor. This model contains a biallelic deletion of the TRPV1 gene, encoding for the transient receptor potential cation channel subfamily V member 1. Capsaicin is an exogenous ligand for TRPV1. As expected, capsaicin did not elicit the activation pattern in the TRPV1 knockout rats observed in wild-type controls. However, the intradermal injection of formalin elicited a significant activation of the putative pain pathway as represented by such areas as the anterior cingulate, somatosensory cortex, parabrachial nucleus and periaqueductal gray. Capsaicin in wild-type controls also activated the putative pain pathway in addition to brain areas involved in emotion and cognition.

Materials & Methods

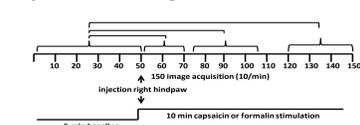
Study and Imaging Protocols

Awake, male Sprague-Dawley (n=36; Charles River Laboratories) and TRPV1-KO rats (n=12; SAGE Laboratories) were challenged with capsaicin (1 mg/ml) (Material # M2028 Sigma-Aldrich, St. Louis, MO) subdermal in the right hindpaw in a volume of 50 µl via a 27-gauge needle connected to PE 20 tubing extending from the bore of the magnet to a calibrated syringe. The same volume in the same hindpaw was given for formalin (3% (source) or vehicle of 20% Captisol® (a modified β-cyclodextrin) in buffered saline (Ligand Pharmaceuticals, Inc. La Jolla, CA). Injections were separated by two weeks. All injection were performed in the magnet, under awake conditions during image acquisition. Experiments were conducted using a Bruker Biospec 7.0T/20-cm USR horizontal magnet (Bruker, Billerica, Massachusetts) and a 20-G/cm magnetic field gradient insert (ID = 12 cm) capable of a 120-µs rise time (Bruker). At the beginning of each imaging session, a high-resolution anatomical data set was collected using the RARE pulse sequence (22 slice; 1.0 mm; field of vision [FOV] 3.0 cm; 256 × 256; repetition time [TR] 2.5 sec; echo time [TE] 12.4 msec; NEX 6; 6.5 minute acquisition time. Functional images were acquired using a multi-slice HASTE pulse sequence (Half Fourier Acquisition Single Shot Turbo Spin Echo). A single scanning session acquired 22 slices, 1.0 mm thick, every 6.0 seconds (FOV 3.0 cm, matrix size 96 × 96, ETL 36, NEX 1) repeated 150 times for a total time of 15 minutes. Scanning session was continuous, starting with 50 baseline image acquisitions, then hindpaw injection followed by another 100 image acquisitions.

Imaging Analysis

Images were aligned and registered to a 3D rat brain atlas, which is segmented and labeled with 152 discrete anatomical regions. The alignment process was facilitated by an interactive graphic user interface. The registration process involved translation, rotation and scaling independently and in all three dimensions. Matrices that transformed each subject's anatomy were used to embed each slice within the atlas. All pixel locations of anatomy that were transformed were tagged with major and minor regions in the atlas. This combination created a fully segmented representation of each subject within the atlas. The inverse transformation matrix [T]_i-1 for each subject (i) was also calculated.

Experimental Design

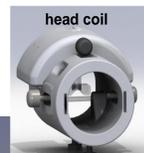


Shown are the statistical comparisons of different image acquisitions compared to baseline. A non-parametric Kruskal-Wallis test statistic was used to compare the average signal intensity in each of ca 15,000 voxel for their first 5 minutes baseline (acquisitions 1-50) to minutes 1-2 (acquisitions 51-71), minutes 3-5 (acquisitions 75-105), and min 8-10 (acquisitions 121-150) post capsaicin and formalin.

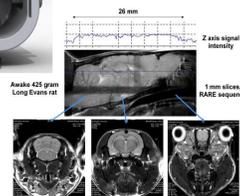
Bruker BioSpec 7T / 20cm USR



Quad transmit/receive volume coil built into the head holder



Optimal signal-to-noise and field homogeneity

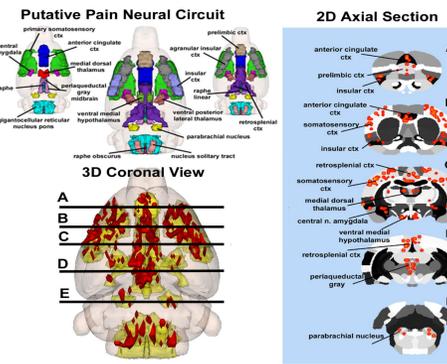


Studies were performed with a quad transmit/receive head coil and rat restrainer developed by Animal Imaging Research, Holden, Massachusetts

Main Results

| Region of Interest | Volume of Activation - Positive BOLD 3-5 Min | | | | | |
|-----------------------------------|--|-----|-----------|-----|---------|-------|
| | Vehicle | | Capsaicin | | P value | |
| | med | max | min | med | max | |
| central amygdala | 0 | 1 | 0 | 10 | 19 | 0.001 |
| tegmental nucleus | 0 | 1 | 0 | 3 | 9 | 0.001 |
| temporal ctx | 3 | 9 | 0 | 25 | 46 | 0.001 |
| parietal ctx | 1 | 2 | 0 | 4 | 8 | 0.001 |
| entorhinal ctx | 15 | 23 | 5 | 99 | 168 | 0.001 |
| basal amygdala | 0 | 2 | 0 | 6 | 23 | 0.001 |
| medial amygdala | 0 | 3 | 0 | 9 | 13 | 0.001 |
| insular ctx | 2 | 6 | 0 | 15 | 38 | 0.001 |
| ventral medial hypothalamus | 3 | 8 | 0 | 16 | 21 | 0.001 |
| auditory ctx | 1 | 6 | 0 | 22 | 36 | 0.001 |
| lateral amygdala | 1 | 2 | 0 | 6 | 10 | 0.001 |
| visual ctx | 10 | 30 | 2 | 49 | 96 | 0.001 |
| somatosensory ctx primary | 8 | 41 | 0 | 97 | 195 | 0.001 |
| somatosensory ctx secondary | 2 | 13 | 0 | 26 | 66 | 0.002 |
| motor secondary ctx | 30 | 57 | 8 | 78 | 130 | 0.002 |
| agranular insular ctx | 3 | 13 | 0 | 21 | 36 | 0.002 |
| gustatory ctx | 1 | 6 | 0 | 17 | 41 | 0.003 |
| periaqueductal gray midbrain | 10 | 23 | 1 | 28 | 46 | 0.003 |
| medial dorsal thalamus | 3 | 11 | 0 | 12 | 15 | 0.003 |
| bed nucleus stria terminalis | 0 | 11 | 0 | 5 | 25 | 0.004 |
| habenula thalamus | 1 | 3 | 0 | 5 | 9 | 0.004 |
| lateral hypothalamus | 7 | 14 | 0 | 34 | 54 | 0.005 |
| infralimbic ctx | 1 | 6 | 0 | 9 | 19 | 0.005 |
| ventral lateral striatum | 1 | 18 | 0 | 17 | 34 | 0.009 |
| parabrachial nucleus | 2 | 6 | 0 | 11 | 15 | 0.005 |
| dentate gyrus hippocampus | 10 | 16 | 2 | 38 | 59 | 0.005 |
| paraventricular thalamic nuclei | 3 | 9 | 0 | 9 | 11 | 0.006 |
| motor ctx primary | 17 | 44 | 2 | 37 | 82 | 0.006 |
| midbrain reticular nucleus | 8 | 16 | 1 | 24 | 47 | 0.007 |
| medial preoptic area | 3 | 10 | 0 | 11 | 27 | 0.008 |
| retrosplenial ctx | 13 | 42 | 0 | 40 | 73 | 0.009 |
| lateral preoptic area | 0 | 3 | 0 | 3 | 8 | 0.009 |
| superior vestibular nucleus | 3 | 18 | 0 | 20 | 33 | 0.01 |
| CA3 hippocampus ventral | 6 | 16 | 0 | 19 | 35 | 0.012 |
| pontine reticular nucleus | 5 | 14 | 0 | 19 | 28 | 0.012 |
| lateral septal nucleus | 11 | 23 | 2 | 25 | 36 | 0.013 |
| subiculum hippocampus | 13 | 29 | 4 | 50 | 87 | 0.015 |
| substantia nigra reticularis | 6 | 11 | 2 | 15 | 20 | 0.016 |
| dorsal medial striatum | 4 | 9 | 0 | 13 | 19 | 0.017 |
| dorsal lateral striatum | 2 | 16 | 0 | 11 | 24 | 0.021 |
| ventral medial striatum | 1 | 14 | 0 | 7 | 15 | 0.023 |
| ventral cingulate ctx | 5 | 14 | 0 | 14 | 31 | 0.023 |
| ventral tegmental area | 1 | 7 | 0 | 7 | 12 | 0.055 |
| gigantocellular reticular n. pons | 14 | 57 | 4 | 33 | 76 | 0.064 |

Areas comprising the putative pain circuit (yellow/gold) and areas comprising the habenular system (blue).

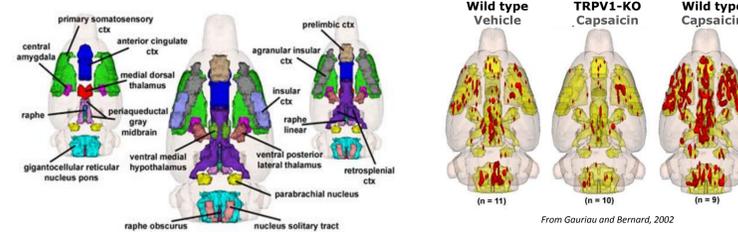


The putative pain neural circuit of the rat

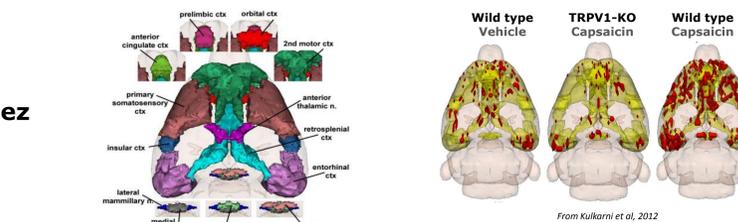
Above The illustrations above show confluence of the segmented brain areas that make up the putative pain circuit (yellow/gold) as well as the location of the average significant increase (red) in BOLD signal for nine rats, 3-5 minutes after capsaicin injection into the right hind-foot. The panel of 2D axial images on the far right depict the location of significant increase in BOLD signal (red) in brain slices approximating the positions A-E shown in the 3D illustration.

Left A truncated list of 152 brain areas activated at 3-5 min post subdermal injection of vehicle and capsaicin in wild-type rats. The brain areas are rank ordered by significance. The voxel numbers were analyzed using a Newman-Keuls multiple comparisons test statistic.

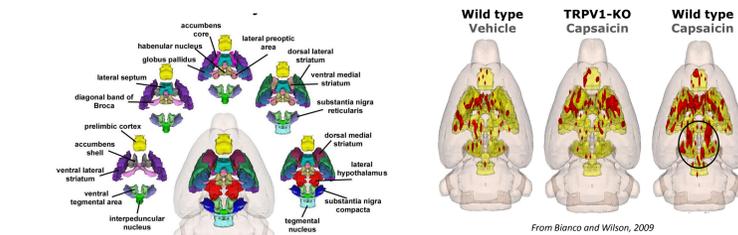
Pain Neural Circuit



Cortical Loop of Papez



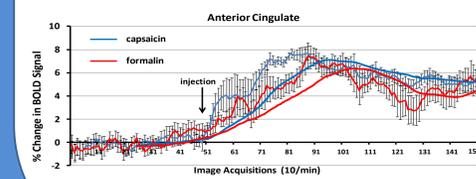
Habenular System



Results

| Region of Interest (ROI) | Vehicle | | Capsaicin | | Formalin | | P value | |
|--|---------|-----|-----------|-----|----------|-----|---------|-------|
| | med | max | min | med | max | min | | |
| culmen cerebellum | 21 | 68 | 3 | 1 | 4 | 0 | 96 | 0.002 |
| lateral dorsal thalamus | 1 | 9 | 0 | 0 | 0 | 0 | 3 | 0.002 |
| medial dorsal thalamus | 2 | 12 | 0 | 0 | 1 | 0 | 9 | 0.003 |
| CA1 hippocampus ventral | 4 | 34 | 0 | 0 | 1 | 0 | 13 | 0.003 |
| paraventricular thalamic nuclei | 5 | 11 | 3 | 0 | 2 | 0 | 5 | 0.004 |
| superior vestibular nucleus | 3 | 27 | 0 | 0 | 0 | 0 | 7 | 0.004 |
| simple lobule cerebellum | 8 | 49 | 0 | 0 | 6 | 0 | 25 | 0.005 |
| decive cerebellum | 8 | 47 | 0 | 0 | 0 | 0 | 21 | 0.005 |
| anterior olfactory nucleus | 0 | 14 | 0 | 0 | 2 | 0 | 5 | 0.006 |
| paraflocculus cerebellum | 11 | 44 | 0 | 0 | 0 | 0 | 1 | 0.006 |
| periaqueductal gray midbrain | 6 | 37 | 1 | 0 | 4 | 0 | 10 | 0.009 |
| paraflocculus cerebellum | 8 | 23 | 2 | 0 | 5 | 0 | 14 | 0.009 |
| auditory ctx | 2 | 9 | 0 | 0 | 0 | 0 | 6 | 0.009 |
| motor ctx primary | 10 | 68 | 2 | 8 | 15 | 0 | 31 | 0.011 |
| nucleus lateral lemniscus | 3 | 11 | 0 | 0 | 2 | 0 | 7 | 0.016 |
| motor secondary ctx | 18 | 57 | 1 | 6 | 24 | 0 | 59 | 0.017 |
| anterior thalamic nuclei | 5 | 11 | 0 | 0 | 3 | 0 | 8 | 0.017 |
| CA3 hippocampus dorsal | 11 | 16 | 0 | 1 | 2 | 0 | 8 | 0.018 |
| parietal ctx | 1 | 5 | 0 | 0 | 0 | 0 | 4 | 0.018 |
| aniform cerebellum | 10 | 76 | 0 | 0 | 4 | 0 | 22 | 0.021 |
| habenula thalamus | 1 | 5 | 0 | 2 | 0 | 3 | 9 | 0.021 |
| visual ctx | 7 | 86 | 2 | 0 | 23 | 0 | 40 | 0.021 |
| nucleus colliculus | 23 | 77 | 1 | 2 | 9 | 0 | 38 | 0.024 |
| diagonal band of Broca | 0 | 2 | 0 | 1 | 0 | 0 | 2 | 0.024 |
| temporal ctx | 2 | 12 | 0 | 0 | 4 | 0 | 10 | 0.027 |
| somatosensory ctx primary | 14 | 119 | 1 | 5 | 7 | 1 | 62 | 0.031 |
| somatosensory ctx secondary | 1 | 11 | 0 | 0 | 1 | 0 | 14 | 0.034 |
| retrosplenial ctx | 16 | 72 | 0 | 2 | 6 | 0 | 22 | 0.035 |
| CA3 hippocampus ventral | 9 | 22 | 0 | 1 | 2 | 0 | 9 | 0.036 |
| accumbens shell | 1 | 4 | 0 | 0 | 2 | 0 | 3 | 0.037 |
| prelimbic ctx | 0 | 19 | 0 | 3 | 6 | 0 | 6 | 0.041 |
| subiculum hippocampus | 3 | 41 | 0 | 1 | 9 | 0 | 20 | 0.043 |
| tenia tecta ctx | 1 | 5 | 0 | 1 | 7 | 0 | 6 | 0.045 |
| dentate gyrus hippocampus | 12 | 29 | 0 | 0 | 9 | 0 | 11 | 0.045 |
| lateral septal nucleus | 2 | 23 | 0 | 0 | 5 | 0 | 10 | 0.048 |
| olfactory tubercles | 9 | 21 | 0 | 3 | 8 | 1 | 22 | 0.049 |
| central lobule cerebellum | 13 | 81 | 0 | 3 | 22 | 0 | 43 | 0.055 |
| gigantocellular reticular nucleus pons | 12 | 93 | 0 | 0 | 9 | 0 | 8 | 0.061 |
| piriform ctx | 26 | 86 | 2 | 5 | 32 | 2 | 18 | 0.066 |
| orbital ctx | 0 | 8 | 0 | 1 | 2 | 0 | 5 | 0.065 |
| interpeduncular nucleus | 0 | 2 | 0 | 0 | 0 | 1 | 6 | 0.066 |
| anterior cingulate ctx | 3 | 33 | 0 | 4 | 5 | 0 | 8 | 0.071 |

Areas comprising the putative pain circuit (yellow/gold) and areas comprising the habenular system (blue).



TRPV1-KO rats' response to injections

A truncated list of 152 brain areas at 3-5 min post intradermal injection of vehicle, capsaicin and formalin into the hindpaw of TRPV1 knockout rats. The brain areas are rank ordered by significance. The voxel numbers for all time points were analyzed using a Newman-Keuls multiple comparisons test statistic.

BOLD timecourse in anterior cingulate cortex

Percent change in BOLD signal over a 10 min period (acquisitions 51-150) following capsaicin in wild-type controls (blue) and formalin injection in TRPV1 KO rats (red). The anterior cingulate cortex, a key area in the pain neural circuit, shows a maximal change in BOLD signal within 3-5 min post injection of capsaicin or formalin. Trend lines (solid blue and red). Vertical lines denote SEM.

Conclusions

Functional neuroimaging of BOLD responses to painful stimuli in un-anesthetized, awake animals effectively identified brain regions that have previously been demonstrated to mediate neural responses to pain, corroborating past work with a novel experimental approach.

Our data suggest that neuroimaging of pain in awake, conscious animals has the potential to inform the neurobiological basis of full and integrated perceptions of pain which, as we show here, recruit both affective and learning systems in addition to areas subserving sensorimotor aspects of pain perception.

Areas comprising the cortical loop described by Papez, and thought to play an important role in communicating affective information with subcortical limbic areas, were activated by capsaicin in wild-type rats, but displayed low activation levels equivalent with vehicle-treated wild-type rats in capsaicin-treated TRPV1-KO rats.

Areas comprising the habenular system, and thought to play an integrative role in aversive learning, were activated by capsaicin in wild-type rats, but displayed low activation levels equivalent with vehicle-treated wild-type rats in capsaicin-treated TRPV1-KO rats.

Formalin, which activates TRPA1 pain receptors, was capable of activating BOLD responses in areas that regulate conscious pain perception in TRPV1-KO rats, demonstrating that the specificity of a genetic manipulation of pain, i.e. eliminating only TRPV1-mediated pain, can be corroborated by differential BOLD activation patterns in the brain.