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#### SUPPLEMENTARY MATERIALS

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Materials and Methods

Figs. S1 to S11  
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## NEUROSCIENCE

# The TRPM2 channel is a hypothalamic heat sensor that limits fever and can drive hypothermia

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Body temperature homeostasis is critical for survival and requires precise regulation by the nervous system. The hypothalamus serves as the principal thermostat that detects and regulates internal temperature. We demonstrate that the ion channel TRPM2 [of the transient receptor potential (TRP) channel family] is a temperature sensor in a subpopulation of hypothalamic neurons. TRPM2 limits the fever response and may detect increased temperatures to prevent overheating. Furthermore, chemogenetic activation and inhibition of hypothalamic TRPM2-expressing neurons *in vivo* decreased and increased body temperature, respectively. Such manipulation may allow analysis of the beneficial effects of altered body temperature on diverse disease states. Identification of a functional role for TRP channels in monitoring internal body temperature should promote further analysis of molecular mechanisms governing thermoregulation and foster the genetic dissection of hypothalamic circuits involved with temperature homeostasis.

Core body temperature ( $T_{\text{core}}$ ) is normally maintained very accurately in mammals within a narrow range around 37°C and serves as an important vital sign that is routinely monitored in hospitalized patients. Pathological conditions such as infections and systemic inflammation cause fever (1, 2), an increase in the body's temperature that is thought to be mediated by prostaglandin  $E_2$  (PGE<sub>2</sub>)-induced inhibition of warm-sensitive neurons (WSNs) in the hypothalamic thermoregulatory center (3). Adverse drug-induced fluctuations in  $T_{\text{core}}$  can have severe and even fatal consequences (4). Conversely, clinically controlled modulation of  $T_{\text{core}}$  has beneficial effects and can prevent tissue damage, promote trauma recovery (5, 6), or help decrease obesity (7).

The preoptic area (POA) of the hypothalamus serves as a thermostat, integrating temperature information to orchestrate the autonomous and behavioral adaptations needed to achieve thermal

homeostasis (3, 8). POA neurons receive and integrate information from peripheral temperature sensors located in the skin, spinal cord, and viscera. Local POA warming and cooling experiments (9–13) and genetic manipulations (14) have shown that this region of the central nervous system also detects local brain temperature changes, which have a strong impact on  $T_{\text{core}}$ . In fact, temperature changes in the POA can exert a dominant effect on  $T_{\text{core}}$  and can override lower-priority temperature inputs from peripheral temperature sensors, such as those located in the skin (15, 16). Warming of the POA induces heat dissipation mechanisms such as cutaneous vasodilation, evaporative heat loss mechanisms, and behavioral adaptations, whereas cooling of the POA triggers heat conservation and thermogenesis.

POA neurons exhibiting pronounced temperature sensitivity have been described in *in vivo* models and *in vitro* preparations (17–19). WSNs enhance their firing rate upon warming, a process that is thought to relay temperature information to peripheral organs to promote heat loss (3). Conversely, a drop in deep brain temperature inhibits WSNs, a process that correlates with induction of thermogenesis and heat gain.

However, the genetic identity of WSNs and the molecular basis of their temperature sensitivity have remained elusive. Our work defined a population of WSNs that controls temperature homeostasis and identified a transient receptor

potential (TRP) ion channel that subserves thermo-detection in hypothalamic neurons.

## TRPM2 is a heat sensor in a subset of POA neurons

To identify thermosensory molecules in the thermoregulatory center of the hypothalamus, we established calcium imaging as a means to monitor warming-induced responses of primary POA neuronal cultures (fig. S1, A to F). We applied a stringent criterion to identify WSNs by fura-2-mediated calcium imaging. We only counted neurons as WSNs if their relative increase in signal on receiving a thermal stimulus (up to 45°C) was larger than the mean fura-2 signal plus 5 times the standard deviation (5SD) of thermally unresponsive human embryonic kidney-293 cells subjected to the same temperature stimulus (fig. S1, E and F). This procedure allowed us to account and correct for the small unspecific temperature effect on the fluorescent fura-2 signal (20) (see fig. S1F and table S1 for details). We found that  $16.3 \pm 4.7\%$  (mean  $\pm$  SEM;  $n = 6$  mice) of cultured POA neurons responded to warming, a fraction that is in the same range (10 to 16%) observed in electrophysiological studies of rodent POA cultures (21, 22) and slightly lower than WSN percentages (20 to 40%) observed in POA slice preparations (17, 23). Extracellular calcium appeared to be the source of the temperature-triggered increase in intracellular concentrations of free calcium, because chelation of external calcium abolished the response, whereas depletion of intracellular calcium stores did not have any effect on warm sensitivity (fig. S1, D and G).

Several ion channels are thermosensitive, exemplified by a steep temperature dependence ( $Q_{10}$  coefficient) of their opening probabilities. In particular, cationic TRP ion channels have been identified as thermosensors in the peripheral nervous system and function in the detection of ambient temperature changes (24–27).

To identify candidate channels mediating the observed warm sensitivity in neuronal POA cultures, we tested various inhibitors and agonists of TRPs and other thermosensitive ion channels (fig. S1, H to L). The calcium response appeared to be mediated by direct  $\text{Ca}^{2+}$  flux through a thermosensitive ion channel and not an indirect readout of voltage-gated channels, because nifedipine and tetrodotoxin, which are blockers of voltage-gated calcium and sodium channels, respectively, had only a small effect on the responses to increased temperature (fig. S1, H, J, K, and L). 2-aminoethoxydiphenyl borate (2-APB), a potent inhibitor of several ion channels and activator of TRPV-type channels (28–30), abolished the temperature response of POA neurons in a reversible manner (Fig. 1A).

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Among known temperature-responsive ion channels, TRPC5, TRPM2, TRPM3, TRPM8, TRPV1, TRPV2, TRPV3, and STIM1-ORAI1 are sensitive to 2-APB (fig. S1M) (24–26). Because TRPC5 and TRPM8 are cold-sensitive channels (31–33), we eliminated them from our list of candidates. The

STIM1-ORAI1 channel complex responds to cooling subsequent to a heat stimulus (34), a characteristic that differs from the POA responses described here.

The TRPV1 to -3 channels are activated rather than inhibited by 2-APB (28, 35), and capsaicin, the cognate agonist of heat-sensitive TRPV1, did

not have any effect on POA neurons (fig. S1N), suggesting that TRPV1 to -3 do not serve as temperature sensors in POA neurons. Moreover, these channels are sensitive to the pore blocker ruthenium red (36), a substance that had only a subtle effect on the temperature sensitivity of POA neurons (fig. S1, I and L).

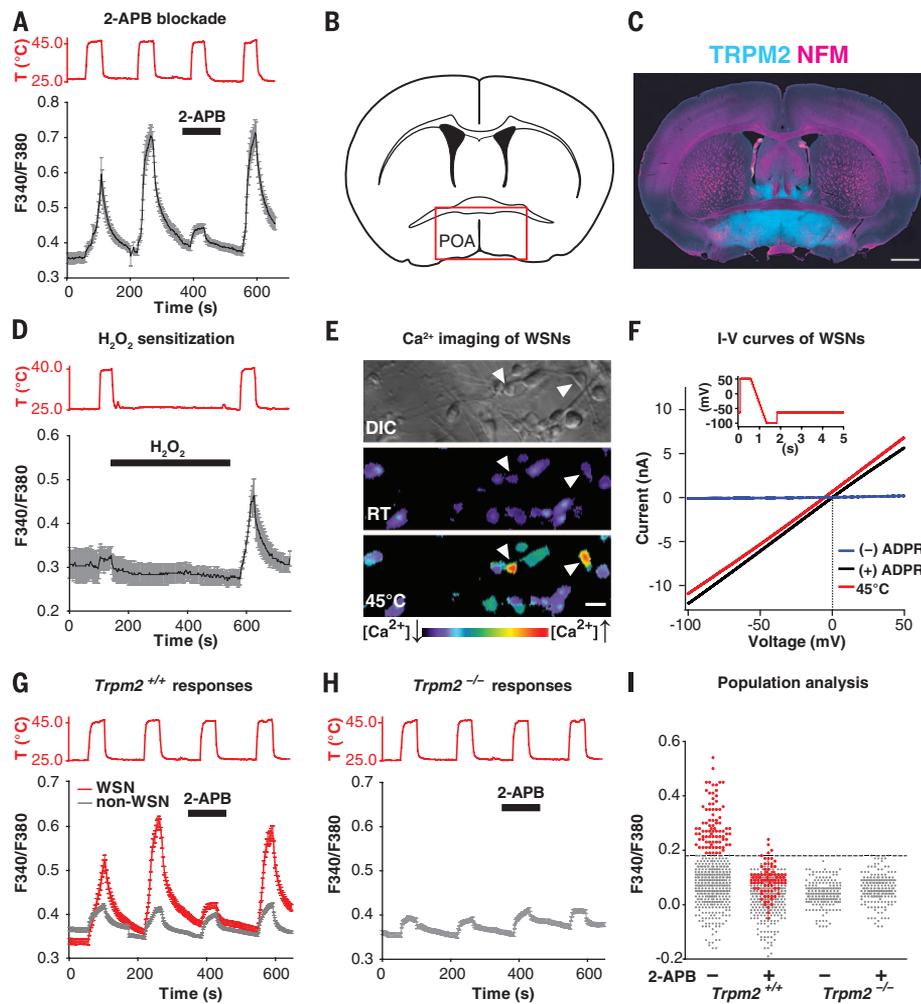
TRPM3, one of the two remaining candidates, appeared to be expressed in the POA, albeit in low amounts, as assessed by in situ hybridization (fig. S1O). However, the TRPM3 agonist pregnenolone sulfate did not produce calcium responses in neuronal POA cultures (fig. S1N). It did so in dorsal root ganglia sensory neurons that express the receptor (37), which we used as a positive control.

Immunohistochemistry and in situ hybridization for TRPM2 showed prominent labeling within the mouse POA (Fig. 1, B and C, and fig. S3, B and C).  $H_2O_2$ , a sensitizer of TRPM2 channel activity (38), shifted the warm sensitivity of POA cultures to a lower activation temperature (Fig. 1D). Moreover, adenosine diphosphate ribose (ADPR), an intracellular agonist of TRPM2 (39, 40), induced large currents in POA neurons that were identified by calcium imaging to be warm-sensitive (Fig. 1, E and F, and fig. S1, P to S). The ADPR-induced current showed a TRPM2-characteristic linear current-voltage relationship (39, 40). Additionally, a similar current was induced in these neurons by increased temperatures (Fig. 1F).

Thus, TRPM2 may mediate responses to temperature increases in the POA. To test this, we used cultures obtained from *Trpm2*<sup>-/-</sup> mice (41) and compared their thermal responses with those from wild-type (*Trpm2*<sup>+/+</sup>) littermate controls. The magnitude of the response of POA neurons to increased temperature was decreased in the absence of TRPM2, and these neurons were indistinguishable from warm-insensitive (non-WSN) *Trpm2*<sup>+/+</sup> POA neurons (Fig. 1, G and H). Using our stringent definition of warm sensitivity (fig. S1F), no responses to increased temperatures were detected in cultures obtained from *Trpm2*<sup>-/-</sup> mice (Fig. 1I).

The activation temperature of cultured WSNs—in the absence of a sensitizer such as  $H_2O_2$  or any other reactive oxygen species—was in the range of 45°C (fig. S2A). Although this temperature is similar to that previously reported for TRPM2 activation (38), it exceeds any physiologically relevant deep brain temperature, even when compared with extreme forms of fever and hyperthermia (42). However, similar to other thermosensitive TRP channels, the activation temperature for TRPM2 appears not to be a fixed threshold temperature. Rather, it ranges from 33° to 47°C, depending on the cellular context (38, 43). We therefore tested whether the activation temperature of POA neurons shifted to lower, more physiological temperatures when calcium imaging was performed on acutely obtained brain slices from 7- to 8-week-old mice. In this experimental setting, the neurons largely retained their native cellular environment and had attained maturity.

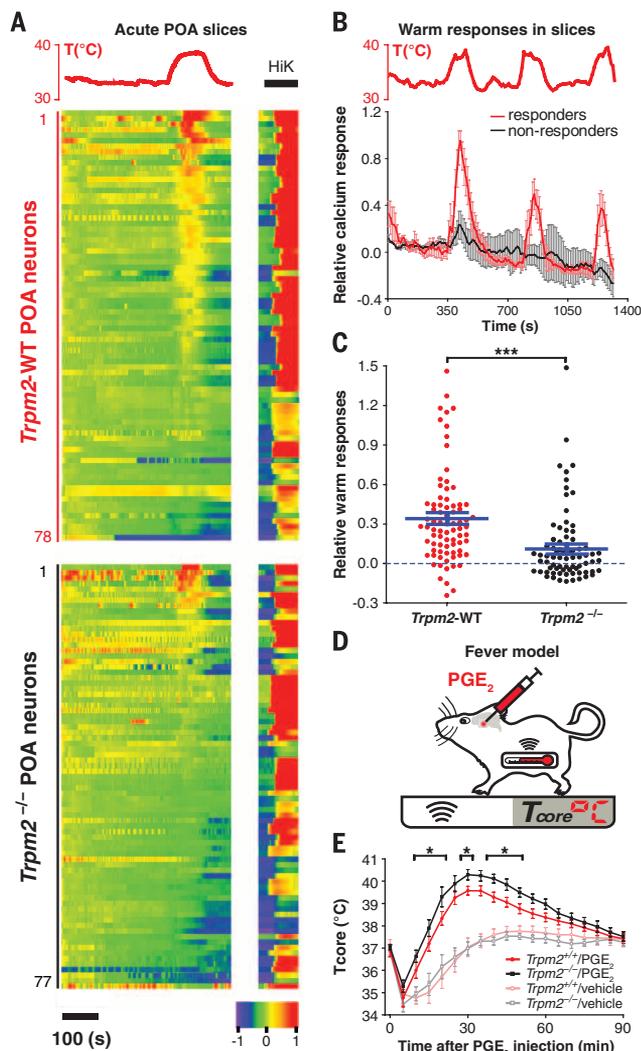
To this end, we injected virus particles into the mouse POA for Cre-dependent expression of the virally encoded intracellular calcium indicator GCaMP6, which shows increased fluorescence



**Fig. 1. TRPM2 mediates heat responses in a subset of POA neurons.** (A) Cultured mouse POA neurons were examined for calcium responses [black trace; plotted are non-normalized fura-2 fluorescence intensity ratios after excitation at 340 and 380 nm (F340/F380)] to repetitive increases in temperature (red trace). 2-APB (30  $\mu$ M, black bar) reversibly blocked thermal responses in POA neurons. (B) Cartoon depicting a coronal section of a mouse brain (Bregma level anterior-posterior, 0.14 mm). (C) Immunostaining of a representative mouse brain section corresponding to the cartoon in (B), using antibodies against TRPM2 (cyan) and neurofilament medium (NFM, magenta). Scale bar, 1 mm. (D)  $H_2O_2$  (1 mM, black bar) sensitized POA neurons to respond to lower activation temperatures (40°C), as assessed by fura-2–based calcium imaging (black trace). (E) Differential interference contrast (DIC) image (top) and fura-2 ratio images before (middle; RT, room temperature) and during (bottom) a temperature stimulus of cultured POA neurons. Arrowheads indicate WSNs. (F) Current-voltage (I-V) relationships of voltage-clamped WSNs shown in (E) in response to a voltage ramp shown at the top, in the absence (blue) or presence (black) of ADPR (100  $\mu$ M) or at a temperature stimulus of 45°C (red). (G and H) POA cultures from *Trpm2*<sup>+/+</sup> (G) and *Trpm2*<sup>-/-</sup> (H) mice were examined by calcium imaging for responses to repetitive thermal stimuli (upper red traces) in the presence or absence of 2-APB (30  $\mu$ M, black bar). In *Trpm2*<sup>+/+</sup> cultures, both WSNs (red) and non-WSNs (light gray) could be detected, whereas WSNs were absent in cultures of *Trpm2*<sup>-/-</sup> mice. (I) Population analysis of thermal responses of POA neurons from *Trpm2*<sup>+/+</sup> and *Trpm2*<sup>-/-</sup> mice as shown in (G) and (H). Plotted are the peak responses of individual neurons at the second (without 2-APB) and third (with 2-APB) temperature stimulus. POA neurons of both genotypes had similar responses to a high-potassium solution (100 mM KCl, not shown). WSNs are colored in red and non-WSNs in gray. The dashed line demarcates the cutoff for warm sensitivity as defined in fig. S1F and table S1. Throughout, error bars indicate SEM.

## Fig. 2. TRPM2 mediates heat responses in POA brain slices and modulates fever temperature in vivo.

(A) Acute POA slice preparations from *Nestin-Cre* mice with (*Trpm2*-WT; upper panel) or without (*Trpm2*<sup>-/-</sup>; lower panel) functional TRPM2 were consecutively subjected to a temperature stimulus of 38 ± 1°C (red) and a high-potassium stimulus (HiK, black). Neuronal activity was recorded by means of a Cre-induced, virally encoded GCaMP6 calcium sensor that was stereotactically introduced into the POA 2 weeks before slice preparation. A nonlinear pseudocolor scale (bottom) indicates relative calcium responses in individual neurons. Numbering on the left refers to individual neurons. The black bar indicates the time scale. (B) Average relative calcium responses of exemplary GCaMP6<sup>+</sup> WSNs (red; n = 5) and non-WSNs (black; n = 3) from *Nestin-Cre* mice to repetitive temperature stimuli of up to 40°C (top, red trace). (C) Population analysis of cellular responses shown in (A), demonstrating that relative heat-induced calcium signals are reduced in POA slices from mice lacking functional TRPM2 (\*\*\*P < 0.0001, two-tailed Mann-Whitney test). Blue bars show the mean and SEM. (D) Cartoon depicting the experimental paradigm used to examine the fever response by thermotelemetrical measurements of *T*<sub>core</sub> upon preoptic injection of PGE<sub>2</sub> into the POA. (E) Enhanced PGE<sub>2</sub>-induced fever in *Trpm2*<sup>-/-</sup> mice (black trace; n = 9) compared with littermate *Trpm2*<sup>+/+</sup> controls (red trace; n = 10). Mice were briefly anesthetized to allow stereotactic injection of PGE<sub>2</sub> (or vehicle control), resulting in a transient drop of *T*<sub>core</sub> for all conditions and genotypes. \*P < 0.05, two-way analysis of variance (ANOVA) (*F*<sub>1,318</sub> = 40.78) followed by post hoc Fisher's least significant difference (LSD) test.



at 500 to 530 nm upon an increase in cytosolic calcium levels (44). To test temperature sensitivity in POA neurons, we used the neuron-specific *Nestin-Cre* mice (45) that express Cre recombinase, and thus GCaMP6, in POA neurons. Subsequent temperature stimulation of brain slices prepared from these animals revealed calcium responses in POA neurons at 38 ± 1°C (Fig. 2, A and B). Moreover, POA neurons in slices derived from mice lacking TRPM2 (*Nestin-Cre Trpm2*<sup>-/-</sup>) had attenuated responses to the same temperature stimulus [mean relative heat responses ± SEM: *Nestin-Cre Trpm2*<sup>-/-</sup>, 0.111 ± 0.033 (n = 78); *Nestin-Cre Trpm2*-WT, 0.341 ± 0.040 (n = 79); P < 0.0001, two-tailed Mann-Whitney test; *Trpm2*-WT signifies *Trpm2*<sup>+/+</sup> and *Trpm2*<sup>+/-</sup> mice] (Fig. 2, A and C, and fig. S2B). General excitability, as assessed by high potassium stimulation, was indistinguishable in the presence or absence of

TRPM2 (fig. S2C). These results demonstrate that preoptic TRPM2 mediates responses to heat stress at temperatures of (or exceeding) 38°C.

### Preoptic TRPM2 receptors modulate the magnitude of fever temperature

Fever is a cardinal response to infection and systemic inflammation. The POA, as part of its thermoregulatory function, is the key brain structure in controlling the increased body temperature in fever (2, 3). The TRPM2-mediated activation of POA neurons at stimulation temperatures slightly above the physiological set point of 37°C (sometimes referred to as the balance point), might allow hypothalamic TRPM2 to detect and modulate inflammatory fever temperature.

As a consequence of infection and downstream of the resulting immune response, PGE<sub>2</sub> is the

final mediator that triggers fever by acting directly on POA neurons (2, 46). To address a putative fever-modulatory role for TRPM2 directly in the POA, we microinjected PGE<sub>2</sub> into this brain area in mice (Fig. 2D and fig. S2D), an established fever model (46–48). Telemetrically measured *T*<sub>core</sub> revealed similar diurnal temperatures in the presence and absence of TRPM2 at normothermic (nonfever) conditions (fig. S2E). We found that injection of a low dose of PGE<sub>2</sub> (0.4 nmol per mouse) produced a mild fever response that was similar in both genotypes (fig. S2, F and I). However, a high dose of PGE<sub>2</sub> (4 nmol per mouse) caused an increased fever temperature in *Trpm2*<sup>-/-</sup> mice compared with that of *Trpm2*<sup>+/+</sup> littermate controls (Fig. 2E and fig. S2I). In the absence of functional TRPM2 protein, the maximal fever temperature reached 40.4 ± 0.2°C, compared with 39.6 ± 0.2°C in the presence of the receptor (n = 9 *Trpm2*<sup>-/-</sup> and 10 *Trpm2*<sup>+/+</sup> mice; P < 0.001, two-tailed unpaired t test).

We also tested the effects of the interleukins IL-1β and IL-6, which have been shown to synergistically induce fever at central sites (49). Again, significantly higher fever was detected in *Trpm2*<sup>-/-</sup> mice only for the stronger of the two stimuli, when both interleukins were co-injected into the POA, but not when IL-1β was injected alone (fig. S2, G to I). Together, these results indicate that preoptic TRPM2 limits the magnitude of the temperature increase in strong fever responses, presumably to prevent excessive temperature increases and tissue damage.

### *Trpm2*-expressing POA neurons are part of a circuit controlling body temperature homeostasis

Our results indicated that *Trpm2*-expressing (*Trpm2*<sup>+</sup>) neurons might be part of a thermoregulatory circuit and that TRPM2-driven activity might counteract body heating and perhaps promote peripheral heat loss at normothermic conditions as well. In the absence of specific agents to modulate TRPM2 function in vivo, we generated *Trpm2*-2A-Cre knock-in mice (referred to hereafter as *Trpm2*-Cre mice) that express Cre recombinase exclusively in *Trpm2*<sup>+</sup> cells (fig. S3A). We thereby gained genetic access to *Trpm2*<sup>+</sup> POA neurons and ascertained the potential thermoregulatory role of this specific cell population in unrestrained, conscious mice. We first assessed proper Cre expression in *Trpm2*<sup>+</sup> neurons, using in situ hybridization and Cre reporter mice. Cre was expressed in the POA of *Trpm2*-Cre mice, and its expression pattern recapitulated that of native *Trpm2* transcripts (fig. S3, B to F). Two-color in situ hybridization analysis of the POA revealed essentially complete cellular overlap of *Trpm2* and Cre (fig. S3F), with 97% of *Trpm2*<sup>+</sup> neurons (335 out of 344 cells) expressing the Cre-induced reporter.

We then tested whether modulating the activity of *Trpm2*<sup>+</sup> POA neurons influenced body temperature in freely moving mice. We chose a chemogenetic, DREADD (designer receptors exclusively activated by designer drugs) receptor-assisted approach that enabled us to remotely turn the activity of *Trpm2*<sup>+</sup> neurons on and off.

DREADDs are engineered heteromeric guanine nucleotide-binding protein (G protein)-coupled receptors that are activated by the otherwise inert drug clozapine-*N*-oxide (CNO) and couple to different G protein signaling cascades that either activate (Gq-DREADD) or inhibit (Gi-DREADD) neuronal activity (50). We injected viral constructs encoding Cre-dependent (“flexed”) DREADD receptor expression cassettes (51) into the POA of *Trpm2-Cre* mice (Fig. 3A). This allowed the specific activation or inhibition of *Trpm2-Cre*<sup>+</sup> POA neurons through systemic application of CNO. No expression of DREADDs occurred in the absence of Cre recombinase.

To verify the capacity of CNO to exert control over *Trpm2*<sup>+</sup> neurons in vitro, we co-infected POA neurons by stereotaxic injection of Cre-inducible DREADD receptors and GCaMP6 viral constructs into *Trpm2-Cre* mice. In brain slices derived from these mice, the respective DREADD receptor and GCaMP6 are coexpressed in *Trpm2*<sup>+</sup> neurons. Calcium imaging of POA neurons expressing Gq-DREADD or Gi-DREADD together with GCaMP6 confirmed that CNO application activated or inhibited *Trpm2*<sup>+</sup> neurons, respectively (fig. S3G).

We implanted telemetric temperature probes in the peritoneal cavity of DREADD-virus-injected *Trpm2-Cre* mice to permit recording of *T*<sub>core</sub> in conscious, unrestrained animals (Fig. 3B). Diurnal body temperature regulation of DREADD-infected *Trpm2-Cre* mice in the absence of CNO was similar to that of control animals (fig. S3H). In contrast, CNO-mediated activation of *Trpm2*<sup>+</sup> POA neurons resulted in a sustained decrease in body temperature (Fig. 3, C to E, and fig. S3I). *T*<sub>core</sub> dropped to 27.4 ± 0.6°C when *Trpm2*<sup>+</sup> neurons were activated in Gq-DREADD-infected mice but not in control-infected or saline-injected mice.

The temperature decrease correlated with a reduction in activity of the animals and was also reflected by a temperature drop of the body shell, as visualized by infrared (IR) imaging (Fig. 3F and movie S1). The hypothermic state lasted several hours (0.3 mg/kg of CNO induced hypothermia for 12.5 ± 2.4 hours) and could be induced repetitively without causing any apparent long-term adverse effects (Fig. 3E).

The animal’s tail, an important thermal effector organ controlling heat dissipation (3, 52), showed an increase in surface temperature of 3.7 ± 0.5°C that coincided with the overall heat loss (Fig. 3, F and G; fig. S3J; and movie S1), demonstrating that *Trpm2*<sup>+</sup> POA neurons drive hypothermia by triggering cutaneous vasodilation. Additionally, stimulation of *Trpm2*<sup>+</sup> neurons prevented activation of interscapular brown adipose tissue (BAT; fig. S3K), which normally mediates thermogenesis when the body cools.

Conversely, Gi-DREADD receptor-mediated inhibition of *Trpm2*<sup>+</sup> POA neurons resulted in a significant temperature increase, reaching a *T*<sub>core</sub> of up to 39.1 ± 0.1°C (Fig. 3H and fig. S3L), suggesting that under normothermic conditions, *Trpm2*<sup>+</sup> neurons in the POA are tonically active. This is in agreement with previous

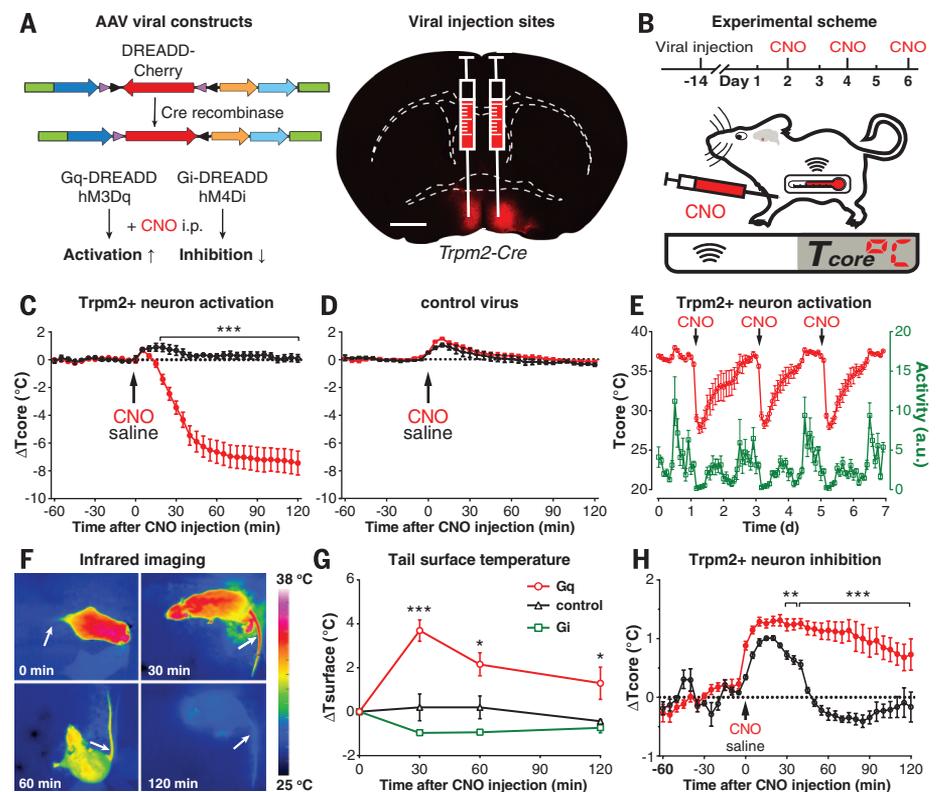
electrophysiological recordings that show tonic ongoing activity of WSNs (17–19). Gi-DREADD-mediated neuronal inhibition had no significant effect on tail vasomotor tone at ambient temperature (~20°C), likely because the prominent vasoconstriction that prevails under these conditions (53) precludes detection of any further reduction of the IR signal (Fig. 3G). These results demonstrate that *Trpm2*<sup>+</sup> neurons in the POA are part of a circuit that orchestrates thermal effector processes to modulate *T*<sub>core</sub> homeostasis.

### Excitatory *Trpm2*<sup>+</sup> neurons drive hypothermia

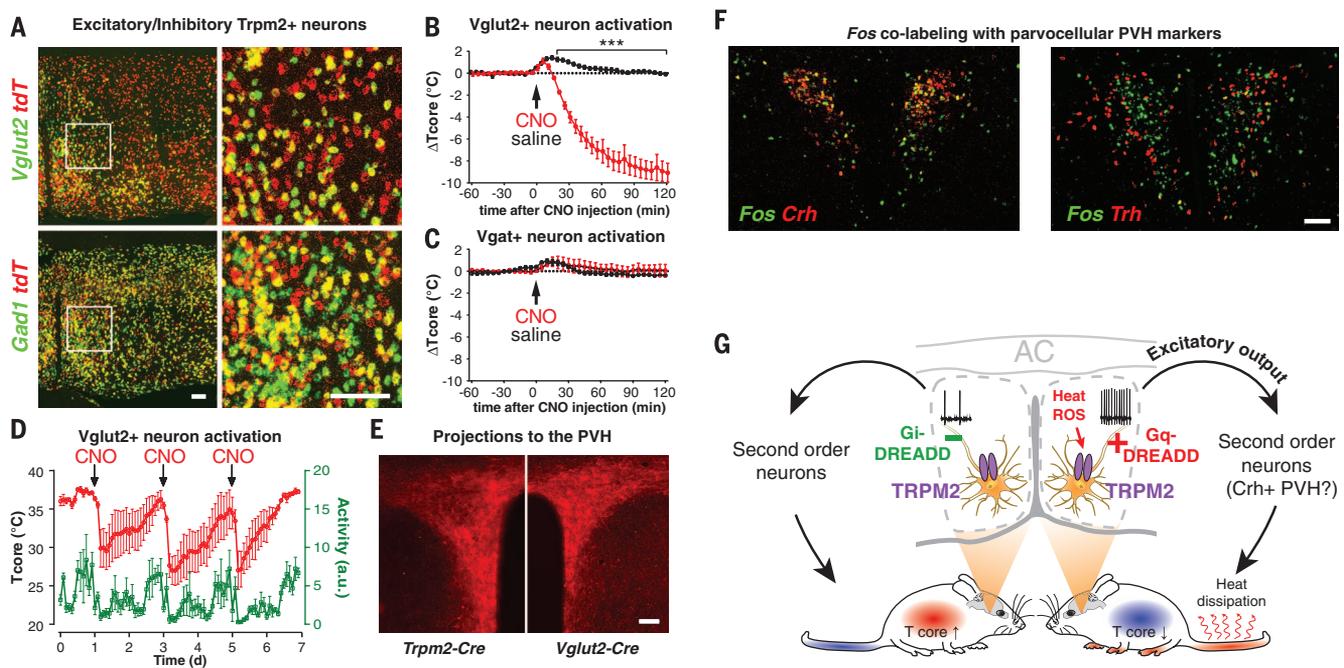
WSNs in the POA have been proposed to be  $\gamma$ -aminobutyric acid (GABA)-releasing neurons that exert their effect largely by inhibiting down-

stream components of the thermoregulatory circuit (3). We therefore used two-color in situ hybridization on *Trpm2-Cre* reporter mice to ascertain whether *Trpm2*<sup>+</sup> neurons are of inhibitory or excitatory origin. We used in situ probes for transcripts encoding the vesicular glutamate transporter 2 (VGLUT2) and glutamate decarboxylase 1 (GAD1), which are excitatory and inhibitory markers, respectively (54) (Fig. 4A).

By this analysis, 30 ± 1% of *Trpm2*<sup>+</sup> POA neurons were excitatory (*Vglut2*<sup>+</sup> neurons, whereas 57 ± 7% coexpressed the inhibitory marker *Gad1* (mean ± SEM; *n* = 3 mice) (Fig. 4A and fig. S4A). We therefore tested whether activation of the excitatory or inhibitory neuronal population within the POA recapitulated the strong thermoregulatory phenotype observed on activation of *Trpm2*<sup>+</sup> neurons (Fig. 3, C to G). *Vgat-Cre* (54) and



**Fig. 3. Control of core body temperature by *Trpm2*<sup>+</sup> POA neuron activity.** (A) Virally encoded, Cre-inducible DREADD-receptor–cherry fusion proteins (left panel) were bilaterally injected into the POA of *Trpm2-Cre* mice and visualized by fluorescent microscopy (right panel). The red arrows indicate the orientation of the DREADD–cherry cDNA reading frame; i.p., (injected) intraperitoneally. (B) Illustration detailing the experimental paradigm for examining the impact of *Trpm2*<sup>+</sup> neuronal activity on *T*<sub>core</sub>. (C to E) CNO (0.3 mg/kg; red) or vehicle (saline; black) was intraperitoneally injected into mice infected with Gq-DREADD [(C) and (E)] or control virus (D), and *T*<sub>core</sub> and activity [green trace in (E)] were measured telemetrically. Arrows indicate the time of injection. a.u., arbitrary units. (F) IR thermal imaging of *Trpm2-Cre* mice preoptically expressing Gq-DREADD in *Trpm2*<sup>+</sup> neurons, before (0 min) or at indicated time points after CNO injection, revealed pronounced cutaneous heat dissipation through tail vasodilation (arrows). (G) Change in tail temperature over time [(arrows in (F))] in *Trpm2-Cre* mice infected with Gq-DREADD (red), saline control (black), or Gi-DREADD (green) upon CNO treatment. (H) CNO-mediated inhibition of *Trpm2*<sup>+</sup> neurons in Gi-DREADD-infected *Trpm2-Cre* mice showed an oppositely directed effect compared with that in Gq-DREADD-infected animals (C) and resulted in a significant increase in *T*<sub>core</sub>. The arrow indicates the time of injection. Telemetry data [(C), (D), and (H)]: Gq-DREADD *F*<sub>16</sub> = 75.6, \*\*\**P* < 0.0001; control virus *F*<sub>16</sub> = 1.55; Gi-DREADD *F*<sub>16</sub> = 38.5, \*\**P* < 0.01, \*\*\**P* < 0.0001. IR imaging (G): Gq-DREADD *F*<sub>15</sub> = 9.3, \**P* < 0.05, \*\*\**P* < 0.001; Gi-DREADD *F*<sub>14</sub> = 5.2; two-way ANOVA followed by post hoc Fisher’s LSD test (*n* = 4 mice per group). *P* values indicate significance of the post hoc test.



**Fig. 4. Stimulation of preoptic *Vglut2*<sup>+</sup> neurons recapitulates *Trpm2*<sup>+</sup> neuron-driven hypothermia and activates *Crh*<sup>+</sup> neurons in the PVH.** (A) Representative two-color in situ hybridizations of coronal brain sections from *Trpm2-Cre* mice crossed with a *Cre* reporter expressing *tdTomato* (*tdT*) with probes detecting *tdT* (red) and *Gad1* or *Vglut2* (green). The right panels show close-up views indicated by the boxes in the left panels. (B to D) *Vglut2-Cre* mice and *Vgat-Cre* mice were bilaterally injected with Gq-DREADD virus and subjected to the same experimental paradigm as shown in Fig. 3B. Arrows indicate the time of injection. CNO-mediated activation of *Vglut2-Cre* mice [(B) and (D)], but not *Vgat-Cre* mice (C), recapitulated *Trpm2*<sup>+</sup> neuron-driven hypothermia both in magnitude (B) and kinetics (D). *Vglut2-Cre*  $F_{1,6} = 86.08$ ,  $***P < 0.0001$ ; *Vgat-Cre*  $F_{1,4} = 0.44$ ; two-way ANOVA followed by post hoc Fisher's LSD test ( $n = 4$  *Vglut2-Cre* and 3 *Vgat-Cre* mice). (E) Representative fluorescent images of brain sections

from *Vglut2-Cre* and *Trpm2-Cre* mice preoptically expressing Gq-DREADD–cherry fusion protein (Fig. 3A), showing cherry<sup>+</sup> fibers projecting to the PVH. (F) Two-color fluorescent in situ hybridizations of brain sections from *Vglut2-Cre* mice preoptically expressing Gq-DREADD, using probes to detect *Fos* (green) and *Crh* or *Trh* (red). The animals were injected with CNO 30 minutes before preparation of the tissue sections. Neuronal activation of *Crh*<sup>+</sup> neurons but not *Trh*<sup>+</sup> neurons in the PVH was detected. (G) Model depicting *Trpm2*<sup>+</sup> POA neurons and their bidirectional effect on  $T_{core}$ . Heat (and reactive oxygen species, ROS) activate TRPM2 to depolarize WSNs, a process that can be mimicked by exogenous Gq-DREADD activation. Increased WSN activity (black traces) triggers an excitatory pathway that promotes heat loss and hypothermia, possibly through a circuit involving *Crh*<sup>+</sup> neurons in the PVH. Conversely, inhibiting *Trpm2*<sup>+</sup> neurons results in an elevation of  $T_{core}$ . AC, anterior commissure. *P* values indicate significance of the post hoc test. All scale bars, 100  $\mu$ m.

*Vglut2-Cre* (55) mice have been used previously to label inhibitory and excitatory hypothalamic neurons, respectively. Thus,  $T_{core}$  recordings of Gq-DREADD-injected *Vgat-Cre* and *Vglut2-Cre* mice were acquired, and viral expression in the respective POA neurons was later verified (fig. S4, B and C). CNO activation of *Vglut2*<sup>+</sup> neurons but not *Vgat*<sup>+</sup> neurons reproduced the prolonged drop in  $T_{core}$  (Fig. 4, B to D), indicating that excitatory, *Trpm2*<sup>+</sup> neurons mediate the strong thermoregulatory effect. *Vglut2*<sup>+</sup> neuron-driven hypothermia could be induced repetitively and was well tolerated by the animals (Fig. 4D), similar to observations for in vivo chemogenetic activation of preoptic *Trpm2*<sup>+</sup> neurons (Fig. 3E). In situ hybridization revealed that *Cre* recombinase in the *Vglut2-Cre* line was expressed only in ~20% of all *Vglut2*<sup>+</sup> POA neurons (fig. S4D). This indicates that activation of a subpopulation of *Vglut2*<sup>+</sup> POA neurons, making up only ~6% of all preoptic *Trpm2*<sup>+</sup> neurons (fig. S4A), is sufficient to mediate the strong hypothermic effect (Fig. 4, B and D).

Thermoregulatory signals arising from the POA are channeled to different peripheral effector organs through discrete hypothalamic and

extrahypothalamic output nuclei (3). Given the overlapping expression pattern of *Vglut2* and *Trpm2* in the POA and a similar capacity of both neuron groups to trigger hypothermia on activation, we reasoned that axons of both neuron cohorts might converge on the same downstream output nucleus. Projections of *Cre*-expressing POA neurons were labeled with fluorescent reporters that were introduced together with the DREADD expression cassette by viral infection of the POA area (Fig. 3A). POA infections of *Trpm2-Cre* and *Vglut2-Cre* mice resulted in prominent labeling of fibers terminating in the paraventricular nucleus of the hypothalamus (PVH) (Fig. 4E), a region not previously identified as a major thermoregulatory outlet (3). Moreover, CNO administration in *Trpm2-Cre* and *Vglut2-Cre* mice harboring Gq-DREADD-infected POA neurons resulted in DREADD-specific expression of *Fos* in the PVH, demonstrating that *Trpm2*<sup>+</sup> *Vglut2*<sup>+</sup> POA neurons can excite PVH neurons (fig. S4E). Corticotropin-releasing hormone-positive (*Crh*<sup>+</sup>) neurons in the PVH are part of the hypothalamic-pituitary-adrenal (HPA) stress response system, which is also activated in response to fever (56). We found that DREADD-mediated activation of

*Vglut2*<sup>+</sup> POA neurons specifically excites parvocellular *Crh*<sup>+</sup> neurons but not neighboring thyrotropin-releasing hormone-positive (*Trh*<sup>+</sup>) neurons or magnocellular vasopressin<sup>+</sup> and oxytocin<sup>+</sup> PVH neurons (Fig. 4F and fig. S4F). Of all *Fos*<sup>+</sup> neurons, we found that 88.4 ± 1.5% (mean ± SEM;  $n = 3$  mice) are *Crh*<sup>+</sup>.

Collectively, these results indicate that *Trpm2*<sup>+</sup> *Vglut2*<sup>+</sup> POA neurons detect increased body temperature and initiate thermoregulatory defense mechanisms, putatively through activation of stress-responsive *Crh*<sup>+</sup> neurons in the PVH (Fig. 4G).

## Discussion

Explanations of the molecular mechanisms mediating thermal detection in preoptic WSNs have remained speculative (57, 58). We identified TRPM2 as a thermal sensor in a subset of POA neurons that are part of a circuit controlling  $T_{core}$ .

Preoptic TRPM2 was activated at temperatures above the physiological set point of 37°C, arguing for a function of this receptor at conditions of elevated core temperatures and heat stress. Our results demonstrate a role for preoptic TRPM2 in regulating the magnitude of the fever response, suggesting that heightened temperatures, potentially

together with reactive oxygen species that are produced under these conditions (59), activate the channel to promote heat loss and prevent overheating.

Fever temperatures activate the immune system to modulate inflammatory responses (60). TRPM2 is expressed in the immune system (61), suggesting that fever-activated TRPM2 might modulate immune cell function. TRPM2 activity has been shown to inhibit and counteract inflammatory (pyrogenic) signaling in phagocytes (62), paralleling our results and emphasizing a convergent TRPM2-based mechanism used by both the immune and nervous systems to limit the extent of an inflammatory response and the mediation of antipyresis.

We observed a TRPM2-mediated reduction in  $T_{\text{core}}$  only when strong fever responses were induced, suggesting that TRPM2 limits the upper fever range and acts as an “emergency break” to prevent overheating and tissue damage. Alternatively, it is possible that TRPM2 plays a more prominent role in fever regulation, but that this function is masked by compensatory mechanisms triggered in *Trpm2*<sup>-/-</sup> mice that ubiquitously lack this ion channel throughout development.

The current model of thermoregulation predicts that the action potential firing rate of WSNs in the POA underlies  $T_{\text{core}}$  homeostasis and controls peripheral autonomous thermal responses (3). We found that chemogenetic activation and inhibition of *Trpm2*<sup>+</sup> WSNs in mice results in hypo- and hyperthermia, respectively, providing direct in vivo evidence for the validity of this hypothesis.

Twenty to 40% of POA neurons are sensitive to temperature and respond to stimuli around 37°C in electrophysiological experiments (17). Using calcium imaging, we found that only around 16% of neurons responded to a temperature increase. It is possible that our approach only allowed the detection of neurons that robustly responded to a thermal stimulus. Buffering of intracellular free calcium by the calcium indicators used in our experiments is one possible explanation for the reduced sensitivity and might explain the lower percentage of responsive neurons. Our data indicate that the TRPM2 receptor is unlikely to account for temperature sensitivity at normothermia but rather detects heat stress above 37°C. Accordingly, in POA brain slices, some thermal responses were still detectable in the absence of TRPM2, implying that additional mechanisms mediate preoptic responses to warming. Given the relatively broad expression of TRPM2 in the POA, it is plausible that the receptor covers detection of the upper, pathological temperature range in many or even all WSNs.

Previous single-cell transcriptome analysis has shown that preoptic WSNs are unexpectedly heterogeneous (63). Nevertheless, WSNs have been found to largely be inhibitory (GABAergic) neurons, which is in agreement with our histochemical analysis (summarized in fig. S4A). However, because the hypothermic phenotype that we observed in *Trpm2-Cre* mice was recapitulated by activating the smaller subpopulation of excitatory *Vglut2*<sup>+</sup> neurons but not by activating inhibitory

(*Vgat*<sup>+</sup>) neurons, we focused on the excitatory branch of the *Trpm2*<sup>+</sup> population. We found that preoptic *Vglut2*<sup>+</sup> neurons project to and excite *Crh*<sup>+</sup> neurons in the PVH, a hypothalamic cell population that is not primarily associated with thermoregulation (3), but rather controls stress reactions including fever responses by activation of the HPA axis. Excitation of PVH neurons inhibits PGE<sub>2</sub>-induced fever (47). Moreover, corticosterone release by HPA axis activation has been found to counteract and reduce fever responses (64, 65). These findings are in agreement with our data. The TRPM2 receptor may curtail PGE<sub>2</sub>-triggered fever and induce hypothermia by mediating monosynaptic excitation of *Crh*<sup>+</sup> neurons in the PVH through an excitatory thermoregulatory pathway that connects the POA with the HPA axis.

Our study delineates a genetic framework for dissecting the central pathways controlling temperature homeostasis and provides a way to remotely control core body temperature in conscious, unrestrained mice by chemogenetic manipulation of the hypothalamic thermostat. These hypo- and hyperthermic model systems may help in exploring the effects of altered core body temperature on diverse processes such as trauma recovery, immune modulation, energy expenditure, obesity, and longevity.

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#### SUPPLEMENTARY MATERIALS

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**The TRPM2 channel is a hypothalamic heat sensor that limits fever and can drive hypothermia**

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## Supplementary Materials for

### **The TRPM2 channel is a hypothalamic heat sensor that limits fever and can drive hypothermia**

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#### **This PDF file includes:**

Materials and Methods  
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Caption for Table S1  
Captions for Movie S1

#### **Other Supplementary Material for this manuscript includes the following:**

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Table S1  
Movie S1

## Supplementary Materials

### Material and Methods

#### Animals

Animal housing and experimentation was in accordance with the local animal welfare and use guidelines (Regierungspräsidium Karlsruhe, Germany).

We generated *Trpm2-Cre* mice using recombineering techniques as described previously (66). In brief, an 8.3 kbp region flanking *Trpm2* exon 34 was retrieved from BAC clone RP23-13818. A selection cassette containing a T2A sequence (67) linked to iCre-recombinase (68) and a FRT-flanked neomycin resistance gene was inserted upstream of the stop codon of the *Trpm2* gene (fig. S3A left panel). The final targeting vector was linearized by NotI, electroporated into A9 (129/C57Bl6-derived hybrid-) ES cells (EMBL Mouse Biology Unit, Monterotondo). Individual ES cell clones were identified by southern blot (fig. S3A, middle panel). A positive clone was injected into 8-cell stage embryos to generate founder mice heterozygous for the targeted allele. Founder males were crossed with FLPeR-expressing transgenic females (69) to remove the FRT flanked neomycin cassette. *Trpm2-Cre* mice were kept on a C57BL/6N background and heterozygous animals were used for experiments. The genotype of *Trpm2-Cre* animals was verified by PCR with primers flanking the 2A-Cre cassette (P1 5'-GATCCCTCTGTATGCGAACC-3', P2 5'-AAAGCATGGCACTTGAGGACATA-3') (FigS3A right panel).

*Nestin-Cre*, *Vglut2-Cre*, *Vgat-Cre*, *Trpm2<sup>-/-</sup>*, *Ai9* and *Z/AP* mouse lines have been derived previously (41, 45, 54, 55, 70, 71).

For all *in vivo* experiments examining autonomous functions ( $T_{\text{core}}$  measurements, IR recordings, mouse activity), both male and female mice were used. The animals were randomly assigned for experiments. The experimenters were aware of the animals' genotype when conducting experiments.

## **1. Primary neuronal culture and Calcium-imaging**

### **1.1 Primary preoptic neuronal cultures**

Preoptic neurons were isolated from brains of 1-3 day old postnatal C57Bl/6N mouse pups purchased from Charles River Laboratories. In brief, brains were isolated quickly, and 200  $\mu\text{m}$  thick brain slices containing the preoptic area of the hypothalamus were cut on a vibratome (HM650V, Thermo Scientific, USA) in artificial cerebral spinal fluid (ACSF; in mM: 135.5 NaCl, 1 KCl, 2 CaCl<sub>2</sub>, 1.3 MgCl<sub>2</sub>, 1.26 K<sub>2</sub>HPO<sub>4</sub>, 26 NaHCO<sub>3</sub> and 9 glucose oxygenized in carbogen (95% O<sub>2</sub> and 5% CO<sub>2</sub>). The preoptic area was micro-dissected from each brain slice in HBSS solution (Invitrogen, USA). Small POA tissue pieces were collected under microscopic control and incubated in HBSS solution containing 0.5 mg/ml papain at 37 °C for 20 min. After the digestion procedure, the tissue was mechanically dissociated into a single-cell suspension by trituration in neurobasal medium (Invitrogen, USA) with a glass pasteur pipette. After centrifugation (300 g, 5 min), the cell pellet was resuspended in neurobasal medium and plated on coverslips sequentially coated with poly-D-lysine (Sigma, USA) and laminin (Sigma, USA). The isolated POA neurons were cultured in a humidified incubator at 37 °C, 5% CO<sub>2</sub> for two days before imaging.

## 1.2 Calcium-imaging of primary preoptic neuronal cells

Cells in culture were first washed once with Ringer's solution (in mM) 140 NaCl, 5 KCl, 2 CaCl<sub>2</sub>, 2 MgCl<sub>2</sub>, 10 glucose, and 10 HEPES pH=7.4) and then incubated with 10 μM of the Ca<sup>2+</sup>-sensitive ratiometric fluorescent dye Fura-2 AM (F1201; Molecular Probes, USA) for 90 min at RT in Ringer's solution. After the loading procedure, coverslips were mounted in a perfusion chamber (RC-22; Warner Instruments, USA) and perfused by an 8-channel modular perfusion system (ValveBank-8; AutoMate Scientific, USA). Calcium-imaging was performed using an inverted fluorescent microscope (Axio Vert.A1; Carl Zeiss, Germany) equipped with a 10X objective (Fluar 10X/0.5 M27, Carl Zeiss, Germany) and a cooled CCD camera (CoolSNAP-HQ2; Photometrics, USA). Fura-2 AM loaded cells were excited at the wavelengths of 340 nm and 380 nm (Chroma Fura-2 filter set), and the fluorescence was collected at 510 nm. The obtained images were analyzed for changes in the fluorescence intensity within regions of interest (ROIs) using MetaFluor 7.1 (Molecular Devices, USA). The intracellular calcium level [Ca<sup>2+</sup>]<sub>i</sub> was represented as the fluorescence ratio F340/F380. The calcium-free Ringer's solution was prepared by substituting MgCl<sub>2</sub> for CaCl<sub>2</sub> at the same concentration and adding 2 mM EGTA. To change the temperature of the perfusion solution for the application of heating or cooling stimuli, Ringer's solution flowed through a glass coil with a custom-made thermal jacket surrounding it. A circulating water bath with heating/cooling function (MultiTemp III; Pharmacia Biotech, Sweden) pumped water into the jacket, allowing rapid and highly reproducible heat exchange with the perfusate (Ringer's solution). The bath temperature in the perfusion chamber was monitored by a thermocouple probe (IT-18; Physitemp Instruments, USA) connected with a digital

thermometer (BAT-12;Physitemp Instruments, USA). The bath temperature was simultaneously recorded along with the calcium signals.

For the statistical analysis of the heat responses using calcium-imaging data and for defining the cut-off value for warm-sensitive neurons (WSNs), calcium responses of individual cultured mouse POA neurons or HEK293 cells were calculated according to the difference between the basal ratio at room temperature ( $\sim 24^{\circ}\text{C}$ ) and the peak ratio during a thermal stimulus of up to  $45^{\circ}\text{C}$  using Clampfit software (Ver.10.3). In brief, the initial fluorescence ratio F340/F380 for each cell at time point 0 was taken as the relative basal calcium level ( $R_0$ ) at room temperature. Next, the peak ratio F340/F380 during the thermal stimulus ( $R_T$ ) was determined for each cell. The response magnitude during the thermal stimulus was calculated by the formula  $R_{\Delta T}=R_T-R_0$ . Unspecific temperature dependent fluorescent changes of Fura-2 were accounted for by averaging  $R_{\Delta T}$  Fura-2 signals of thermally unresponsive HEK293 cells and determining the mean + 5-times Standard Deviation (5SD). This value corresponded to a relative Fura-2 Ratio of  $R_{\Delta T}=0.18$  (see Table S1 and fig. S1E, F for details).

### **1.3 Electrophysiological recordings of cultured primary preoptic area neurons**

Cultures of preoptic area neurons were prepared as described above (see section 1.1). Differential interference contrast images were acquired with a 20X objective, F340/F380 calcium imaging was performed with a 10X objective (Fig. 1E). Using an inverted microscope, a diamond-shaped recording chamber was constantly perfused ( $\sim 2$  ml/min) with a gravity-driven ValveLink 8.2-system (AutoMate Scientific, USA) with (in mM)

136 NaCl, 4 KCl, 2 CaCl<sub>2</sub>, 1 MgCl<sub>2</sub>, 10 glucose and 10 HEPES, pH=7.4 with NaOH at room temperature, ~295 mOsm. Recordings were performed at room temperature. One to three heat stimuli of 45 °C and ~30 s duration were applied with pre-heated channels of the perfusion system. Robust Ca<sup>2+</sup> responders were patched in whole-cell mode with an Axopatch 200B amplifier, sampled at 20 kHz, low pass filtered at 5 or 10 kHz and digitized with a Digidata 1440A using Clampex 10.6. Patch pipette solution contained (in mM) 0.1 or 10 EGTA, 10 HEPES, ~120 CsCl, 2 NaCl, 2 TEA-Cl, 1 MgCl<sub>2</sub>, 6 Na<sub>2</sub>-phosphocreatine, 3 Mg-ATP, 0.3 Na-GTP and 0 or 0.1 ADP-ribose (Sigma-Aldrich, A0752), pH 7.35 with CsOH at room temperature, ~275 mOsm. Open patch pipette resistance was ~4 MΩ, subsequent gigaseal ~5 GΩ. Neurons were held at -65 mV in voltage clamp. In order to inactivate voltage-gated sodium and calcium currents, the cells were depolarized to +50 mV for 500 ms before a ramp to -100 mV was applied within 750 ms to record current-voltage curves, (inset in fig. 1F). After 500 ms at -100 mV, the neuron was held again at -65 mV and holding current was recorded. The ramp protocol was repeated every 5 s.

The neurons were either patched with or without 0.1 mM ADP-ribose in the patch pipette solution. Cell capacity was measured directly after establishing whole-cell mode by applying voltage steps and averaging  $\geq 5$  current responses. If a current of at least -1 nA developed at -100 mV within 10 min in whole-cell mode, the neurons were counted as responders to ADP-ribose and ~10-20 ramps of the stable peak current were averaged. Cells patched without ADP-ribose were recorded for at least 10 min. For non-responders and control cells, the ramps of the last two minutes of the 10 min interval were averaged (fig. S1R and S).

Sodium-free extracellular solution consisted of (in mM) 136 NMDG, 4 KCl, 2 CaCl<sub>2</sub>, 1 MgCl<sub>2</sub>, 12 glucose and 10 HEPES, pH=7.4 with HCl at room temperature, ~295 mOsm. Analysis was carried out in Igor Pro 6.37 (WaveMetrics, USA), after importing Axon binary files with the DataAccess (Bruxton Corporation, USA). For the example trace in fig. S1P, the holding current was resampled with 1 kHz, plotted over time and completed by linear splines. No corrections were made for junction potentials. Conductance was measured by the slope of a linear fit to the respective I-V-curve.

## **2. Viral injections and calcium-imaging in brain slices**

### **2.1 Stereotactic viral injections**

Stereotactic injections were performed as previously described (72). Surgical procedures were performed aseptically. Mice were deeply anesthetized by an intraperitoneal (i.p.) injection of “sleep-mix“ containing metedomidin (0.5 mg/kg), midazolam (5 mg/kg) and fentanyl (0.05 mg/kg). Fur on the head was removed by depilatory cream and eyes were protected by an eye ointment (Bepanthen; Bayer, Germany). Mice were placed on a stereotaxic apparatus (Kopf Model 1900; Tujunga, USA). After exposing a minimal area of the skull, a small hole of 0.2 mm diameter was drilled with a frame-mounted drill (Kopf 1911; Tujunga, USA) for injection. A pulled glass pipette with 20–40 µm tip diameter was inserted into the brain and virus was injected by an air pressure system. 100-200 nl of the recombinant adeno-associated virus encoding the Cre-dependent calcium indicator GCaMP6s (44) under control of the human synapsin-1 promoter (AAV1.Syn.Flex.GCaMP6s.WPRE.SV40, purchased from the University of

Pennsylvania Gene Therapy Program Vector Core) was injected unilaterally into the POA with the coordinates, Bregma: ML: 0.300 mm, AP: 0.000 mm and DV: -5.300 mm. After viral injection, the scalp was carefully sutured with sterile absorbable-needled sutures (Marlin® 17241041; Catgut, Germany). During the whole operation procedure, mice were kept on a 37 °C heating pad. Once the surgery was finished, the anesthesia was antagonized using a subcutaneous injection of “wakeup-mix” (antipamezol 2.5 mg/kg, flumazin 0.5 mg/kg and naloxon 1.2 mg/kg). After the surgery, mice were transferred to their home cages and exposed to infrared warming overnight to help recovery from the surgery. For postoperative care, mice were injected subcutaneously with carprofen (5 mg/kg, Rimadyl; Zoetis, USA). Brain slices were prepared for calcium-imaging two weeks after viral injection.

## **2.2 Calcium-imaging of acute brain slices**

*Nestin-Cre<sup>+/-</sup>;Trpm2-WT* and *Nestin-Cre<sup>+/-</sup>;Trpm2<sup>-/-</sup>* mice aged between P47 and P60 were used in this experiment. On the day of the experiment, mice were euthanized using an overdose of ketamine (Ketavet; Zoetis, USA) and xylazine (Rompun; Bayer, Germany). After cervical dislocation, animals were briefly perfused transcardially with a freshly-made ice-cold carbogen-bubbled NMDG-HEPES solution containing (mM): 93 NMDG, 2.5 KCl, 1.2 NaH<sub>2</sub>PO<sub>4</sub>, 5 L(+)-ascorbic acid, 2 thiourea, 3 sodium pyruvate, 10 MgSO<sub>4</sub>, 0.5 CaCl<sub>2</sub>, 20 HEPES, 30 NaHCO<sub>3</sub>, 25 glucose and 10 N-acetyl-L-cysteine, titrated to pH 7.3 with HCl (73). After dissection, brains were sliced at 300 µm thickness in chilled carbogen-bubbled NMDG-HEPES solution using a vibration microtome (HM 650V; Thermo Scientific, USA). Three coronal slices covering the POA region

(approximately at Bregma AP: +0.24 to -0.16 mm) were recovered per brain. Thereafter, brain slices were allowed to recover for 15 min in carbogen-bubbled NMDG-HEPES solution at 32 °C. Afterwards, tissue slices were transferred to a chamber at room temperature filled with carbogen-bubbled ACSF containing (mM): 125 NaCl, 2.5 KCl, 25 NaHCO<sub>3</sub>, 1.25 NaH<sub>2</sub>PO<sub>4</sub>, 1 MgCl<sub>2</sub>, 2 CaCl<sub>2</sub> and 25 glucose until used for imaging within a period of four hours.

For GCaMP6 calcium-imaging, experiments were performed using an upright microscope (BX51W; Olympus, Japan) attached to a motorized base plate (SliceScope SS-1000-00; Scientifica, UK), a 20X objective (XLUMPLFLN 20XW Plan Fluorit; Olympus, Japan), 470 nm excitation illumination (pE-2; CoolLED, UK) and a digital CCD camera (ORCA-R2 C10600-10B; Hamamatsu, Japan). Camera frequency acquisition was set typically at 10 Hz. During experiments, slices were continually perfused with ACSF, gassed with 95% O<sub>2</sub> and 5% CO<sub>2</sub> at 3 ml/min and bath temperature was monitored using a thermistor (TA-29; Warner Instrument, USA). Changes in bath temperature were performed and recorded using an in-line temperature controller (CL-200; Warner Instruments, USA) commanded externally using an A/D board (Digidata 1440A; Molecular Devices, USA) and pClamp10.3 software (Molecular Devices, USA). Image acquisition and regions of interest (ROI) analysis were performed using MetaFluor 7.1 (Molecular devices, USA). Specifically, ROIs were placed over the cell bodies to record somatic calcium responses (74). The fluorescence time course of each cell was measured by averaging all pixels within the ROI to estimate the relative calcium responses  $=\Delta F/F_0=(F_i-F_{bgi})-F_0/F_0$ , where  $F_{bgi}$  was the mean fluorescence of a background ROI placed at the darkest region of the field of view at time  $i$  and  $F_0$  was calculated as

the mean fluorescence of the 10 lowest values during the first 100 seconds of imaging. The heat stimulation was performed by increasing bath temperature three times every 5 minutes from the baseline temperature ( $33\pm 1$  °C) to  $38\pm 1$  °C for two minutes. After the temperature protocol was applied, slices were perfused with room-temperature high potassium ACSF solution (HiK), with composition similar to the aforementioned ACSF solution but containing KCl 40 mM (osmolarity was preserved by reducing NaCl concentration to 85 mM). Only cells that did respond to the HiK stimulation were further analyzed.

The relative heat response of each cell was calculated as the maximum amplitude of the calcium signal during the first heat stimulation (Fig 2B). The warm response was calculated as the amplitude of the peak of the calcium response during the first stimulation relative to the amplitude of the peak of the calcium response during HiK stimulation (fig. S2). For this experiment, 79 and 78 cells from *Nestin-Cre; Trpm2-WT* and *Nestin-Cre; Trpm2<sup>-/-</sup>* mice, respectively (three slices per mouse, and two animals per genotype) were analyzed.

There were no significant differences within each genetically-defined (*Trpm2-WT* or *Trpm2<sup>-/-</sup>*) group, either when calculating the relative warm responses (mean relative warm response $\pm$ SEM: *Nestin-Cre; Trpm2-WT*(mouse #1)= $0.305\pm 0.043$ , N=45; *Nestin-Cre; Trpm2-WT*(mouse #2)= $0.389\pm 0.072$ , N=34, P=0.3571; *Nestin-Cre; Trpm2<sup>-/-</sup>*(mouse #1)= $0.112\pm 0.041$ , N=47; *Nestin-Cre; Trpm2<sup>-/-</sup>*(mouse #2)= $0.110\pm 0.058$ , N=31, P=0.7132; two-tailed Mann-Whitney test) or when calculating normalized warm responses (mean normalized warm response $\pm$ SEM: *Nestin-Cre; Trpm2-WT*(mouse #1)= $0.152\pm 0.021$ , N=45; *Nestin-Cre; Trpm2-WT*(mouse #2)= $0.180\pm 0.041$ , N=34,

$P=0.6311$ ; *Nestin-Cre; Trpm2<sup>-/-</sup>* (mouse #1)= $0.028\pm 0.014$ ,  $N=47$ ; *Nestin-Cre; Trpm2<sup>-/-</sup>* (mouse #2)= $0.029\pm 0.073$ ,  $N=31$ ,  $P=0.9674$ ; two-tailed Mann Whitney test). The only significant difference was detected when comparing warm responses of neurons from *Trpm2-WT* and *Trpm2<sup>-/-</sup>* mice (mean relative calcium responses upon heat stimuli  $\pm$  SEM: *Nestin-Cre; Trpm2<sup>-/-</sup>*:  $0.111\pm 0.033$ ,  $N=78$ , *Nestin-Cre; Trpm2-WT*:  $0.341\pm 0.040$ ,  $N=79$ ,  $P<0.0001$ , two-tailed Mann-Whitney test).

### **3. PGE<sub>2</sub>, IL-1 $\beta$ and IL-6 microinjections and telemetric T<sub>core</sub> measurement**

#### **3.1 Cannula and telemetry transmitter implantation**

*Trpm2<sup>+/+</sup>* and *Trpm2<sup>-/-</sup>* littermate adult (3-4 month old) mice of both genders were used in these experiments. General surgical procedures were performed as mentioned in section 3.1. A small hole on the skull was drilled at coordinate Bregma: ML 0.30 mm, AP 0.00 mm. A 5 mm long guide cannula (0.46 mm OD and 0.24 mm ID) was implanted (model C315GRL/Spc; Plastic One, USA) and secured using dental cement (Cyano Veneer; Hager & Werken, Germany). Right after implantation, a dummy cannula of 5 mm in length was screwed onto the guide cannula to prevent contaminants from entering the brain (model C315DC/Spc; Plastic One, USA). A sterile telemetric temperature transmitter (TA11TA-F10; Data Sciences International, USA) was implanted in the abdominal cavity. Then muscle and skin layers were separately sutured with absorbable surgical threads (Marlin®; Catgut, Germany).

#### **3.2 Core body temperature measurement by telemetry**

Mice were allowed to recovery for at least one week after temperature transmitter implantation into the body cavity and cannula implantation in the POA. The circadian temperature cycle was used to assess whether animals had recovered from surgery. For the PGE<sub>2</sub>-induce fever models, 2 µl of a solution containing 0.2 nmol/µl PGE<sub>2</sub> or 2 nmol/µl PGE<sub>2</sub> (P6532; Sigma, USA) or vehicle (50% DMSO in saline) were injected into the POA at 1 µl/min using an injection cannula of 6 mm below pedestal (model C315I/Spc; Plastic One, USA) and an automatic microinjector (UMP3 with SYS-micro4 controller; WPI, USA) under isofluran anesthesia (isofluran 1.5%, N<sub>2</sub>O 70% and O<sub>2</sub> 30%). For the interleukin-induced fever models, 2 µl of a solution containing 10 ng/µl IL-1β or a mixture containing 50 ng/µl IL-1β and 100 ng/µl IL-6 nmol/µl (SRP8033 and SRP3330 respectively; Sigma, USA) or vehicle (saline) were injected into the POA at 1 µl/min in restrained animals using the above-mentioned cannula model and microinjector. The pyrogen-induced fever was monitored at 22°C ambient temperature for the PGE<sub>2</sub> experiments and at 30°C for the interleukin experiments.

The injection sites were verified *postmortem* subsequent to injection of 1µl of a solution of 1 mg/ml alcian blue (A5268; Sigma, USA) through the injection cannula and by tracking the scar left by the guide cannula on the tissue. For maximal body temperature analysis, the highest three T<sub>core</sub> values reached by each mouse during febrile response were averaged and the mean value was used to calculate maximal T<sub>core</sub> per mouse.

For time-course analysis of the low-dose (0.4 nmol) PGE<sub>2</sub>-induced fever responses, due to the skipped/missing telemetric data points (18 missing values), non-repeated measures (non-RM) two-way ANOVA was chosen to analyze the febrile responses. T<sub>core</sub> in PGE<sub>2</sub>-

injected groups (N=5 and 4 mice for *TrpM2*<sup>+/+</sup> and *TrpM2*<sup>-/-</sup>, respectively) was independent of genotype (F(1,122)=1.353, P>0.05, non-RM two-way ANOVA).

For time-course analysis of the high-dose (4 nmol) PGE<sub>2</sub>-induced fever responses, due to the skipped/missing telemetric data points (11 missing values in total for both, vehicle and PGE<sub>2</sub>-injected groups), non-repeated measures (non-RM) two-way ANOVA followed by Fisher's LSD *post hoc* test was applied. T<sub>core</sub> in PGE<sub>2</sub>-injected groups (N=10 and 9 mice for *TrpM2*<sup>+/+</sup> and *TrpM2*<sup>-/-</sup>, respectively) was dependent on genotype (F(1,318)=40.78, P<0.0001, non-RM two-way ANOVA) and time (F(18,318)=52.51, P<0.0001, non-RM two-way ANOVA). T<sub>core</sub> in vehicle-injected groups was not dependent on genotype (F(1,316)=1.16, P=0.2824, non-RM two-way ANOVA) but it was dependent on time (F(18,316)=19.67, P<0.0001, non-RM two-way ANOVA). Interaction between factors (time and genotype) was detected neither in PGE<sub>2</sub>-injected nor vehicle-injected groups (P(interaction) in PGE<sub>2</sub>-injected groups=0.9271 and P(interaction) in vehicle-injected groups=0.9973, non-repeated measures two-way ANOVA).

For time-course analysis of the IL-1 $\beta$ -induced fever responses, because of the skipped/missing telemetric data points (8 missing values), non-repeated measures (non-RM) two-way ANOVA followed by Fisher's LSD *post hoc* test was applied. T<sub>core</sub> in IL-1 $\beta$ -injected groups (N=5 mice for *TrpM2*<sup>+/+</sup> and *TrpM2*<sup>-/-</sup>, respectively) was dependent on genotype (F(1,736)=38.40, P<0.0001, non-RM two-way ANOVA) and time (F(92,736)=5.63, P<0.0001, non-RM two-way ANOVA). Interaction between factors (time and genotype) was not detected (P(interaction)=0.15, P>0.05, non-repeated measures two-way ANOVA). However, *post-hoc* Fisher's LSD test showed no significant differences between any informative paired-comparisons.

For time-course analysis of the IL-1 $\beta$  and IL-6 combination-induced fever responses, due to the skipped/missing telemetric data points (2 missing values, non-repeated measures (non-RM) two-way ANOVA followed by Fisher's LSD *post hoc* test was applied.  $T_{\text{core}}$  in IL-1 $\beta$  and IL-6-coinjected groups (N=4 and 5 mice for *TrpM2*<sup>+/+</sup> and *TrpM2*<sup>-/-</sup>, respectively) was dependent on genotype (F(1,677)=195.2, P<0.0001, non-RM two-way ANOVA) and time (F(96,677)=3.60, P<0.0001, non-RM two-way ANOVA). Interaction between factors (time and genotype) was detected (P(interaction)=1.291, P<0.05, non-repeated measures two-way ANOVA).

#### **4. DREADD viral infection and telemetric measurement**

##### **4.1 Stereotactic DREADD viral infection and telemetric implantation**

The stereotactic injections of viral particles followed the same procedure as detailed in section 3.1. 8-week-old mice were bilaterally injected with AAV particles into the POA (serotype 8, 120nl, coordinates, Bregma: AP:0.800 mm, DV:-5.000 mm, M L:  $\pm$ 0.500 mm). Gq-DREADD (AAV8-DIO-hM3Dq-mCherry), Gi-DREADD (AAV8-DIO-hM4Di-mCherry) and cherry control (AAV8-DIO-mCherry) virus were all purchased from Vector Core (University of North Carolina). Telemetric transmitters were implanted according to section 3.1.

The animals were allowed to recover for two weeks before telemetric measurements. The circadian temperature cycle was used to assess whether animals had recovered from surgery and viral injections. All stereotactic injection sites were verified by immunohistochemistry (see section 6.4. below).

## 4.2 Telemetric measurement

Similar to section 3.2, animals were housed in the home cage for telemetric recordings. *Trpm2-Cre* (N=12), *Vglut2-Cre* (N=4) and *Vgat-Cre* (N=3) mice were handled daily for one week before the experiment to habituate the animals. On day 1, day 3 and day 5, mice were injected with saline (i.p.) at 9:00 am. On day 2, day 4 and day 6, 0.3 mg/kg CNO (BML-NS105-0025, Enzo, diluted in saline) was administered (i.p.) at 9:00 am. Equivalent volumes of saline or CNO solution were delivered.

$T_{\text{core}}$  was measured telemetrically as described in 3.2. Saline and CNO responses were averaged over three trials per animal. The relative change of  $T_{\text{core}}$  compared to the baseline was calculated and plotted ( $\Delta T_{\text{core}}$ ).

In *Trpm2-Cre* DREADD experiments, repeated-measures two-way ANOVA detected no significant differences between saline-treated Gq-DREADD, Gi-DREADD and Cherry virus-infected animals ( $F(2,9)=0.86$ ,  $P=0.45$ ). Similarly, the same test found no significant difference of the treatment (saline vs. CNO) in Cherry control virus-infected animals ( $F(1,6)=1.55$ ,  $P=0.26$ ). However, a significant effect of treatment (saline v.s CNO) in both Gq-DREADD and Gi-DREADD virus-infected animals was detected by repeated-measures two-way ANOVA testing ( $F(1,6)=75.64$ ,  $P=0.0001$  and  $F(1,6)=38.6$ ,  $P=0.0008$ , respectively).

In *Vglut2-Cre* experiments, repeated-measures two-way ANOVA detected significant differences between saline- and CNO-treated Gq-DREADD infected animals ( $F(1,6)=86.08$ ,  $P<0.0001$ ). In *Vgat-Cre* experiments, repeated-measures two-way ANOVA detected no significant differences between saline- and CNO-treated Gq-DREADD virus-infected animals ( $F(1,4)=0.44$ ,  $P=0.55$ ).

When comparing Gq-DREADD infected *Trpm2-Cre*, *Vglut2-Cre* and *Vgat-Cre* animals, repeated-measures two-way ANOVA detected no significant differences when animals were treated with saline ( $F(2, 8)=2.27$ ,  $P=0.17$ ), suggesting that the genetic background of the animals does not significantly contribute to the overall  $\Delta T_{\text{core}}$  changes. However, by applying the same test upon CNO-treatment, significant differences were discovered ( $F(2, 8)=27.49$ ,  $P=0.0003$ ). Subsequently, the *post hoc* Fisher's LSD test detected a significant difference between CNO-treated *Trpm2-Cre* and *Vgat-Cre* ( $P=0.0003$ ), but no difference between CNO-treated *Trpm2-Cre* and *Vglut2-Cre* animals ( $P=0.40$ ). Therefore, we concluded that the extent of CNO-induced hypothermia reached a comparable extent in Gq-DREADD infected *Trpm2-Cre* and *Vglut2-Cre* animals.

### 4.3 Infrared imaging

Mice were imaged in their home cages with a highly sensitive infrared camera (VarioCAM; InfraTec, Germany) positioned 25 cm above the cage floor. Thermograms were collected at 1 Hz and analyzed by IRBIS 3 software (InfraTec). In brief, the tails were outlined manually on individual images and the average surface temperatures ( $T_{\text{surface}}$ ) of three images were calculated for any given time point. For each animal and time point after CNO/saline treatment, the relative change of temperature ( $\Delta T_{\text{surface}}$ ) is calculated by  $T_{\text{surface}}$  minus the baseline  $T_{\text{surface}}$  at time point 0 min.

Repeated two-way ANOVA measures detected significant difference between Gq-DREADD and Cherry control virus infected mice ( $N=4$ , per viral type) upon CNO treatment ( $F(1, 5)=9.3$   $P=0.03$ ). *Post hoc* Fisher's LSD test was applied to compare differences between each time point. Using this test, the difference was found to be

borderline non-significant between Gi-DREADD and Cherry control virus infected mice upon CNO treatment ( $F(1,4)=5.2$ ,  $P=0.085$ ).

## **5. Histology**

### **5.1 Immunocytochemistry**

Mice were transcardially perfused with 10 ml PBS followed by 40 ml 4% PFA. Brains were dissected out and postfixed overnight in 4% PFA at 4 °C with gentle agitation. After several washes with PBS 50  $\mu$ m vibratome brain sections were collected into 24-well plates. Floating sections were blocked with 1% goat serum in PBS containing 0.3% TritonX-100 (PBX0.3) for at least 1 h. Sections were then incubated with rabbit anti *Trpm2* (1:1000) (*10*) and mouse anti NFM (to label fiber tracts used as landmarks, 1:100; 2H3, DSHB) diluted in blocking solution overnight and washed extensively with PBX0.3. Sections were then incubated with secondary antibodies (HRP anti rabbit 1:2000, Alexa488 anti mouse 1:1000) for a minimum of 3 hours and washed with PBX0.3, PBS and PBS/10 mM imidazole. Tyramid signal amplification (TSA) was used for *Trpm2* visualization using self-made rhodamine tyramide (75) in PBS containing 10 mM imidazole and 0.001% H<sub>2</sub>O<sub>2</sub> for 30 min at room temperature. Sections were washed extensively with PBS containing 0.1% Tween-20 and mounted using Immu-Mount (Shandon).

## 5.2 *In situ* hybridization

For *in situ* hybridizations we used either 20  $\mu\text{m}$  cryo-sections or 50  $\mu\text{m}$  free-floating vibratome sections. For cryo-sections brains were freshly dissected, directly embedded into OCT and frozen on dry ice/ethanol bath. 20  $\mu\text{m}$  cryo-sections were cut on a Leica CM3050S cryostat, dried and stored at  $-80\text{ }^{\circ}\text{C}$  until use. For floating sections, mice were transcardially perfused with 10 ml PBS followed by 40 ml 4% PFA. Brains were dissected out and postfixed overnight in 4% PFA at  $4\text{ }^{\circ}\text{C}$  with gentle agitation. 50  $\mu\text{m}$  vibratome sections were cut on a Leica VT1200S vibratome and collected and stored in 1% PFA at  $4\text{ }^{\circ}\text{C}$  until use. DREADD virus infected animals were first i.p. injected with 0.3 mg/kg CNO. Thirty minutes afterwards, the animals were sacrificed for *in situ* staining of *Fos*.

*In situ* hybridization was performed using standard procedures (76). In brief, DIG- and/or FITC-labeled cRNA probes were used for hybridization on either free floating- or cryosections. Hybridization was performed overnight at  $65\text{ }^{\circ}\text{C}$ . Sections were washed at  $60\text{ }^{\circ}\text{C}$  twice in 2xSSC/50% Formamide/0.1% N-lauroylsarcosine, treated with 20  $\mu\text{g/ml}$  RNase A for 15 min at  $37\text{ }^{\circ}\text{C}$ , washed twice in 2xSSC/0.1% N-lauroylsarcosine at  $37\text{ }^{\circ}\text{C}$  for 20 min and twice in 0.2xSSC/0.1% N-lauroylsarcosine at  $37\text{ }^{\circ}\text{C}$  for 20 min. Sections were blocked in MABT/10% goat serum/1% Blocking reagent (Roche, cat# 11096176001). For NBT/BCIP staining sections were incubated overnight with sheep anti-DIG-AP (1:1000, Roche 11093274910). After washing staining was performed using NBT/BCIP in NTMT until satisfactory intensity. Double fluorescent *in situs* were stained by two consecutive rounds of TSA amplification with intermediate peroxidase inactivation. Sections were incubated with sheep anti-FITC-POD (1:2000, Roche cat#

11426346910) or sheep anti DIG-POD (1:1000, Roche cat# 11207733910) and TSA was performed using biotin-tyramide or rhodamine-tyramide (15). Subsequently sections were incubated with Streptavidin-Cy2 and DAPI in blocking solution, washed and mounted in Immu-Mount (Shandon).

Trpm2, Vglut2, tdTomato, Fos, Gad1 and Cre probes covered the ORF of the respective gene. Oxt, Avp, Trh , Crh and Trpm3 probes share identical sequences with the probes used by Allen brain atlas.

### **5.3 AP staining**

*Trpm2-Cre;Z/AP* animals were transcardially perfused with 10 ml PBS followed by 40 ml 4% PFA. Brains were dissected out and cryo-protected with 30% sucrose/PBS at 4 °C. Gelatin embedded brains were cut on a Leica SM2010 sliding microtome. 50 µm free-floating sections were first incubated in PBS containing 2 mM Mg<sup>2+</sup> at 70 °C for 1 hour to inactivate the endogenous alkaline phosphatase activity. Subsequently the sections were stained with NBT/BCIP in NTMT at room temperature until satisfactory signals intensity has been achieved. Sections were mounted on glass slides in Immu-Mount.

### **5.4 Immunohistochemistry**

DREADD virus-infected animals were transcardially perfused with PBS followed by 4% PFA. Brains were dissected and cryo-protected in PBS containing 30% sucrose at 4 °C overnight. Gelatin embedded brains were cut on a Leica SM2010 sliding microtome. 40 µm brain sections were mounted on glass slides in Immu-Mount. Virally encoded

mCherry signals were inspected for labeling of the POA. Only correctly targeted animals were included in the study.

### **5.5 Cell counting**

Confocal images of the POA were first manually assigned by area/nuclei according to the Paxinos Atlas. The appropriate brain area was determined based on landmark structures such as the shape and expansion of the ventricles, anterior commissure and optic tract. Labeled cells were counted manually in ImageJ/FIJI.

### **5.6 Image acquisition and analysis**

Fluorescence images were taken on an A1R or C2+ confocal microscope or on a Ni-E wide field microscope, bright field image acquisition was performed on a Ni-E wide field microscope (Nikon imaging center, Heidelberg). Image analysis, conversion and assembly were done using NIS-Elements AR (Nikon), ImageJ/Fiji, Adobe Photoshop and Illustrator. Brain schemes shown in the manuscript (Fig.1D, Fig.3D, Fig.S1A and Fig.S2E) were modified from Paxinos and Franklin, 2001.

## **6. Data and Statistical analysis**

Data was analyzed using Excel (Microsoft Office 2010 for Windows, Microsoft software) and traces from calcium-imaging experiments were analyzed using Clampfit (Ver.10.3; Molecular Devices, USA). Statistical tests were performed using GraphPad 6 (GraphPad Prism 6.0, GraphPad software, USA). Heat-maps showing calcium responses were

generated using the heatmap.2 function of the R package gplots (<https://cran.r-project.org/web/packages/gplots/index.html>).

Results are presented as mean  $\pm$  standard error of the mean (SEM) unless indicated otherwise. The non-repeated-measures two-way analysis of variance (ANOVA) was applied to analyze time-course differences in  $T_{\text{core}}$  between  $Trpm2^{+/+}$  and  $Trpm2^{-/-}$  mice, followed by Fisher's LSD *post hoc* testing of individual time points. For telemetric DREADD experiments, statistical significance was evaluated by repeated-measures two-way ANOVA, and Fisher's LSD *post hoc* testing. Two-tailed *t*-test for normally distributed data or two-tailed Mann-Whitney test for non-normally distributed data were used for comparisons. Unpaired tests were used for comparisons between  $Trpm2^{+/+}$  and  $Trpm2^{-/-}$  mice and paired tests for evaluating CNO- and saline-treated animals. Distribution of data was assayed using the KS normality test, the D'agostino and Pearson omnibus normality test and the Shapiro-Wilk normality test. '\*' signifies  $P < 0.05$ , '\*\*'  $P < 0.01$ , and '\*\*\*'  $P < 0.001$ .

## Supplementary Figures S1-S4

**Fig. S1 Characterization of heat responses in POA neuron cultures by calcium-imaging.** (A) Schematic depicting a coronal tissue section of the mouse brain. The area dissected for preparation of dispersed POA neuronal cultures is indicated in red. (B) Examples of a representative bright field (BF) image and an fluorescent image of primary POA neuron cultures loaded with the intracellular calcium indicator Fura-2 (scale bar, 50  $\mu\text{m}$ ). (C) Representative pseudocolor image series of POA cultures loaded with Fura-2 and challenged sequentially with a 45 °C stimulus and high potassium (100 mM, HiK) as indicated. Neuronal activation was assessed by ratiometric (F340/F380) calcium-imaging. The color bar (right) indicates relative intracellular calcium levels. (D, E) Cultured mouse POA neurons (D) or HEK293 cells (E) were examined for calcium responses (non-normalized Fura-2 ratios, black traces) to repetitive heat stimuli (45°C, red bars) and high potassium (100 mM, HiK, grey bar) in the presence or absence of the extracellular calcium chelator EGTA (2 mM in nominal  $\text{Ca}^{2+}$ -free Ringer's solution) or 2-APB (30  $\mu\text{M}$ ) as indicated. (F) Population analysis of heat responses in cultured POA neurons and HEK293 cells (corresponding to data represented in D and E). Shown are relative peak responses ( $R_t - R_0$ ) of individual cells to the heat stimulus (for details see Table S1). To err on the side of caution, only POA neurons with a peak temperature response  $\geq$  mean + 5SD of the HEK293-cell response were categorized as warm-sensitive neurons (WSNs) as demarcated by the red line. Note that upon a heat stimulus also HEK293 cells showed a subtle increase in the Fura-2 (F340/F380) ratio. This slight increase likely reflects temperature sensitivity of the Fura-2 dye (20) (HEK293: 246 cells, POA: 491 neurons,

POA+EGTA (WSNs only): 88 neurons). Please refer to Table S1 for details. **(G-K)** Cultured mouse POA neurons were examined for calcium responses (black traces) to repetitive heat stimuli (45 °C, red bars) and high potassium (100 mM, HiK, not shown). The effect of the indicated pharmacological agents (grey bars) on warm sensitivity was assessed: **(G)** Serca-Pump blocker cyclopiazonic acid (CPA, 10  $\mu$ M); **(H)** L-type voltage-gated calcium channel blocker nifedipine (Nif, 10  $\mu$ M); **(I)** broad-spectrum ion channel blocker ruthenium red (RR, 10  $\mu$ M); **(J, K)** voltage-gated sodium channel blocker tetrodotoxin (TTX, 1  $\mu$ M), note that a partially TTX-sensitive population (**J**) and a TTX-insensitive population of WSNs (**K**) could be identified. **(L)** Summary of the results shown in Fig 1A and S1D, and S1G-K expressed as percent heat response compared to control (Ringer's) conditions; Student's *t*-test, \*\* $P < 0.01$ . **(M)** Schematic depicting overlap of 2-APB-sensitive and thermosensitive ion channels. **(N)** Cultured POA neurons (red trace) or sensory dorsal root ganglion neurons (DRG, black trace) were sequentially challenged with the TRPM3 agonist pregnenolone sulfate (PS, 10  $\mu$ M), capsaicin (Cap, 1  $\mu$ M) and high potassium (HiK, 100 mM) as indicated and responses assessed by calcium-imaging. Note that neither PS nor Cap was able to elicit any response in POA neurons under these conditions, but only triggered calcium responses in control DRG cultures. **(O)** *In situ* hybridization of a coronal brain section from an adult mouse at the level of the POA, showing pronounced expression of *Trpm3* transcripts in the lateral septal nucleus and more subtle expression in the POA (scale bar, 1 mm). **(P-S)** Electrophysiological whole-cell recordings in voltage clamp mode of adenosine diphosphate ribose- (ADPR-) induced currents in POA neurons identified as WSNs by preceding calcium imaging that was performed before patch-clamping. **(P)** Dialysis of 100  $\mu$ M ADPR induced a rapidly

evolving inward current at a holding potential of -65 mV. No such current was observed in the absence of ADPR. The ADPR-induced current was partially reduced when extracellular sodium ions ( $\text{Na}^+$ -free; current reduction  $49 \pm 8\%$ ,  $n=5$ , SEM) or extracellular sodium and calcium ions ( $\text{Na}^+$ -,  $\text{Ca}^{2+}$ -free) were excluded from the extracellular solution and replaced with N-Methyl-D-Glucamine (NMDG). The remaining current under these NMDG-replacement conditions is likely carried by extracellular potassium ions that are able to pass through the pore of the non-selective cationic TRPM2 ion channel. **(Q)** Current-Voltage (I-V) curves of voltage-clamped WSNs. Shown are recordings of the same cells as in Fig. 1F but at earlier time points after dialysis of the cell with ADPR (grey) and of a cell without ADPR (shades of blue). Note that on average between 1-3 minutes after ADPR exposure, WSNs started to develop a small current that over the next few minutes fully developed into a large current with a characteristic linear current-voltage relation as shown in (R) and Fig. 1F. No such currents were observed in the absence of ADPR. **(R)** Current-Voltage (I-V) curves of individual POA WSNs responding to ADPR (blue). 4 out of 16 WSNs failed to respond to ADPR over the course of a 10-minute recording period (grey). **(S)** Left panel: Average ADPR-induced current density of all 16 analyzed WSNs shown in (R) and compared to the average current density recorded for WSNs in the absence (-) of ADPR ( $n=9$ ). Current densities were determined at -40 mV for comparison with other cell types expressing TRPM2, see for example (40). Right panel: Conductance (i.e. linear slope) of respective I-V curves. All ADPR responding cells have linear I-V curves. Bar plots are mean + SEM. Two-tailed Mann-Whitney test,  $**P < 0.01$ ,  $***P < 0.001$ .

**Fig. S2 TRPM2 mediates responses to high physiological temperatures and fever but is not required for normothermia in mice.** (A) Cultured mouse POA neurons were examined for calcium responses (black trace) to thermal stimuli of increasing magnitude (red trace). Robust 2-APB-sensitive (black bar, 30  $\mu$ M) calcium responses of dispersed POA neurons were observed at  $45.5 \pm 0.5$  °C. (B, C) Population analysis of cellular responses to a temperature stimulus of up to 40 °C recorded in acute POA slices, similar to the analysis shown in Fig. 2C but normalized to high potassium responses (B) and population analysis of high potassium (HiK) responses (C) demonstrate a reduction of the normalized warm responses in acute slices from *Nestin-Cre; Trpm2<sup>-/-</sup>* (mean normalized heat responses  $\pm$  SEM: *Nestin-Cre; Trpm2<sup>-/-</sup>*:  $0.028 \pm 0.030$ , N=78; *Nestin-Cre; Trpm2-WT*:  $0.164 \pm 0.021$ , N=79, two-tailed Mann-Whitney test,  $P < 0.0001$ ) but no alteration of general excitability when compared to wild type controls (mean calcium response during HiK stimulation  $\pm$  SEM: *Nestin-Cre; Trpm2<sup>-/-</sup>*:  $1.981 \pm 0.158$ , N=78, *Nestin-Cre; Trpm2-WT*:  $2.001 \pm 0.112$ , N=79, two-tailed Mann-Whitney test,  $P = 0.36$ ). (D) Cartoon depicting stereotactic injection into the POA (left panel). Injection of Alcian Blue was used to retrospectively ascertain correct positioning of the injection needle (right panel), subsequent to PGE<sub>2</sub> application and fever measurements shown in Fig. 2E. Scale bar 1 mm. (E) Telemetric T<sub>core</sub> measurements show no difference in diurnal body temperatures between *Trpm2<sup>-/-</sup>* and *Trpm2<sup>+/+</sup>* mice, indicating that TRPM2 is not required for maintaining normothermia at basal ambient temperatures. (F) Fever induced by injection of a low dose of PGE<sub>2</sub> (0.4 nmol/mouse) into the POA of *Trpm2<sup>-/-</sup>* mice (black trace) compared to littermate *Trpm2<sup>+/+</sup>* controls (red trace); N=4 and 5 mice per group, respectively; no significant differences were detected between the fever curves of *Trpm2<sup>-/-</sup>*

<sup>-/-</sup> and *Trpm2*<sup>+/+</sup> mice (two way ANOVA  $F(1,122)=1.353$ ,  $P>0.05$ ). Mice were briefly anaesthetized before microinjection of PGE<sub>2</sub>, resulting in a transient drop of T<sub>core</sub> at all conditions and genotypes. **(G)** Fever induced by injection of Interleukin 1 $\beta$  (IL-1 $\beta$ , 20 ng) into the POA of *Trpm2*<sup>-/-</sup> mice (black trace) compared to littermate *Trpm2*<sup>+/+</sup> controls (red trace); N=5 mice per group per genotype; no significant differences were detected between the fever curves of *Trpm2*<sup>-/-</sup> and *Trpm2*<sup>+/+</sup> mice (two way ANOVA  $F(1,736)=38.40$ ,  $P<0.0001$  followed by *post hoc* Fisher's LSD test,  $P>0.05$ ). Mice were restrained but not anaesthetized for interleukin injections. **(H)** Fever induced by co-injection of Interleukin 1 $\beta$  (IL-1 $\beta$ , 100 ng) and Interleukin 6 (IL-6, 200 ng) into the POA of *Trpm2*<sup>-/-</sup> mice (black trace) compared to littermate *Trpm2*<sup>+/+</sup> controls (red trace); N=5 and 4 mice per group, respectively. *Trpm2*<sup>-/-</sup> mice showed a significantly higher fever response compared to *Trpm2*<sup>+/+</sup> littermate controls at the indicated time points (two way ANOVA  $F(1,677)=195.2$ ,  $P<0.0001$ , followed by *post hoc* Fisher's LSD test,  $*P<0.05$ ). Mice were restrained but not anaesthetized for co-injections. **(I)** Maximum body temperatures reached during the fever experiments shown in Fig. 2E and fig. S2F-H. Two-tailed unpaired Student's *t*-test for PGE<sub>2</sub> 4 nmol condition and two-tailed Mann-Whitney test for all additional conditions.  $*P<0.05$ ,  $**P<0.01$ .

### **Fig. S3 Generation and characterization of *Trpm2-Cre* mice**

**(A)** Mice expressing iCre recombinase under the control of the endogenous *Trpm2* promoter were generated by inserting a cassette encoding a cleavable 2A peptide followed by iCre, before the *Trpm2* stop codon. The wild type allele, the targeting construct, and the targeted allele are shown as well as the final recombined allele that

results after removing the neomycin selection cassette by crossing the animals with the Flpe expressing deleter strain. Restriction sites and the positions of 3'/5' DNA probes are indicated that were used for screening targeted ES cells by Southern blotting (middle panel) as well as primers (P1 and P2) used for PCR genotyping of *Trpm2-Cre* mice (right panel). **(B)** Immunohistochemistry using antibodies against TRPM2 (cyan) and neurofilament (NFM, magenta) were used to label mouse brain sections covering the POA at different rostral to caudal (r → c) levels. **(C-D)** *In situ* hybridization of brain sections from wild type (B) or *Trpm2-Cre* (C) mice with probes detecting *Trpm2* or *iCre* messages, respectively. Note that both probes robustly label the POA and a weaker signal can also be detected in the cortex, a region that is not labeled by the TRPM2 antibody (B). **(E)**, Alkaline phosphatase staining of brain sections from *Trpm2-Cre* mice crossed with *Z/AP* reporter mice recapitulates the strong labeling of the POA similar to antibody labeling and *in situ* hybridization (B-D) and shows a somewhat patchy pattern in the cortex, likely reflecting lower expression of *iCre* in this brain region. Scale bar, 1 mm. **(F)** Representative two-color *in situ* hybridizations of a brain section covering the POA from *Trpm2-Cre* mice crossed to a reporter expressing *tdTomato* upon Cre recombination (*Trpm2-Cre; Ai9*). *In situ* probes detecting *tdTomato* (red) and *Trpm2* (green) were used. Note that there is essentially complete cellular overlap of the respective preoptic expression pattern. Scale bars, 100 μm. **(G)** Acute POA slice preparations of *Trpm2-Cre* mice co-expressing the calcium indicator GCaMP6 and either Gq-DREADD (left panel) or Gi-DREADD (right panel) were challenged with clozapine-N-oxide (CNO, 0.5 μM and 5 μM respectively) and a high potassium stimulus (HiK, 40 mM) as indicated. Neuronal activity was recorded by GCaMP6-mediated calcium imaging at a constant bath

temperature of 33 °C. Cre-inducible expression cassettes for GCaMP6 and the respective DREADDs were virally introduced by microinjection into the mouse POA two weeks prior to slice preparation. A non-linear pseudo-color scale indicates relative calcium responses. **(H)** Telemetric  $T_{\text{core}}$  measurements show normal diurnal body temperature regulation in *Trpm2-Cre* mice in the presence or absence of virally encoded Gq-DREADD in POA neurons, indicating that DREADD expression *per se* (and in the absence of the agonist CNO) does not perturb thermoregulation. **(I)**  $T_{\text{core}}$  was measured telemetrically in *Trpm2-Cre* mice expressing either Gq-DREADD, Gi-DREADD or Cherry control and challenged with CNO (0.3 mg/kg) or vehicle (saline) as indicated and  $T_{\text{core}}$  curves were recorded as shown in Fig. 3 C, D and H. The area under the curve (AUC - integration of the  $T_{\text{core}}$  temperature curves over time) is plotted for the indicated conditions. Time interval used for integrations: t=0 min to 2 hour after CNO application; N=4 for each group, two-tailed paired-test. Gq-CNO vs Gq-saline, \*\*P<0.01, Gi-CNO vs. Gi-saline, \*\*P<0.01. Cherry-CNO vs Cherry-saline P=0.12. **(J, K)** Quantification of whole body (J) and interscapular BAT (K) surface temperature difference ( $\Delta T$ ) compared to a time point (t=0 min) before CNO injection. Shown are average surface temperature changes of Gq-DREADD (red), control (black) or Gi-DREADD (green) infected *Trpm2-Cre* mice upon CNO treatment. N=4 mice per group, two way ANOVA followed by *post hoc* Fisher's Least Significant difference (LSD) test. IR imaging of body: Gq-DREADD F(1,6)=13.54, P=0.0103. Gi-DREADD F (1,6)=0.116, P=0.74. IR imaging of BAT: Gq-DREADD F(1,6)=9.3, P=0.003. Gi-DREADD F (1,6)=0.18, P=0.68.

**Fig. S4 Activation of PVH-projecting  $Vglut2^+$  but not  $Vgat^+$  POA neurons recapitulate  $Trpm2^+$ -driven hypothermia**

(A) Illustration depicting the fraction of all preoptic inhibitory ( $Gad1^+$ , red) and all preoptic excitatory ( $Vglut2^+$ , blue) neurons that are  $Trpm2$  positive (shades of green). Note that in *Vglut2-Cre* mice only a small fraction (~20%) of  $Vglut2^+$  POA neurons are expressing Cre recombinase (shown in D, below). (B and C) Representative brain slices demonstrating targeting of the virally encoded, Cre-dependent Gq-DREADD-cherry fusion proteins in the POA of *Vglut2-Cre* (B) and *Vgat-Cre* (C) mice used in experiments shown in Fig. 4B-D and visualized *post hoc* by fluorescent microscopy. Scale bars, 1 mm. (D) Representative images of a brain section covering the POA of *Vglut2-Cre* mice and visualizing transcripts for Cre recombinase (green) and native *Vglut2* (red) by dual-color *in situ* hybridization. Note that  $Vglut2-Cre^+$  neurons only represent a small fraction (~20%) of all  $Vglut2^+$  neurons as illustrated in (A). Scale bars, 100  $\mu$ m. (E) *In situ* hybridizations of brain sections from *Trpm2-Cre* and *Vglut2-Cre* mice using a probe to detect *Fos* (green). Two weeks prior to preparation of tissue sections, the mice were either injected or not injected with Cre-dependent viral Gq-DREADD expression constructs as indicated. Thirty minutes before preparation of the tissue sections, all animals were injected with the DREADD agonist CNO. *Fos* labeling revealed Gq-DREADD-dependent neuronal activation in the paraventricular nucleus of the hypothalamus (PVH). Scale bars: 1 mm for main graphs and 200  $\mu$ m for insets. (F) Depicted are two-color *in situ* hybridizations of *Vglut2-Cre* mice preoptically expressing Cre-dependent Gq-DREADD receptors and using probes to detect *Fos* (green), *Oxytocin* (*Oxt*, red) and *Vasopressin* (*Avp*, red) as indicated. Animals were sacrificed and brain

sections covering the PVH were prepared 30 min after CNO stimulation. Scale bar, 100  $\mu\text{m}$ .

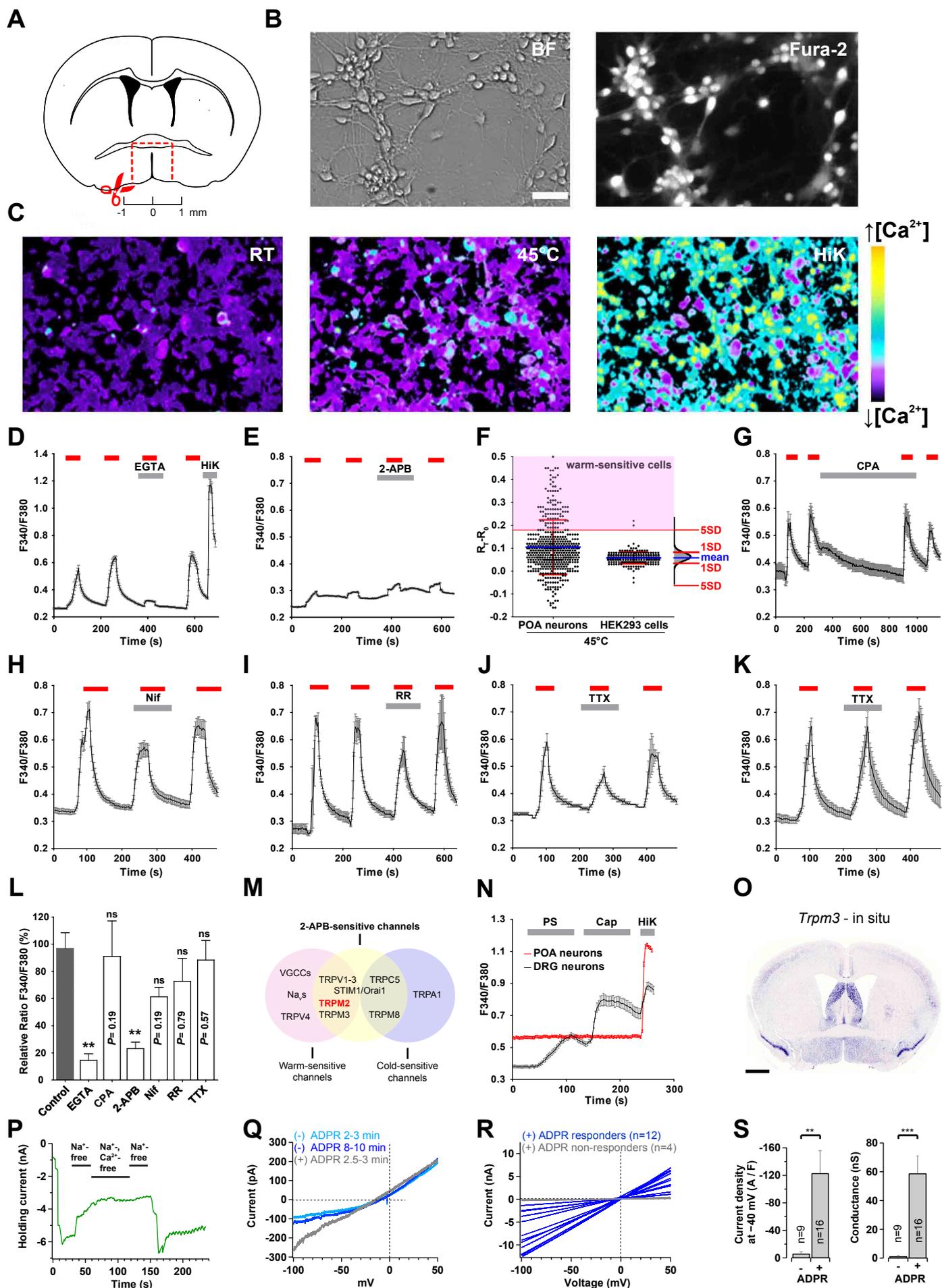
### **Table S1**

#### **Calcium imaging analysis of warm-sensitive POA neurons**

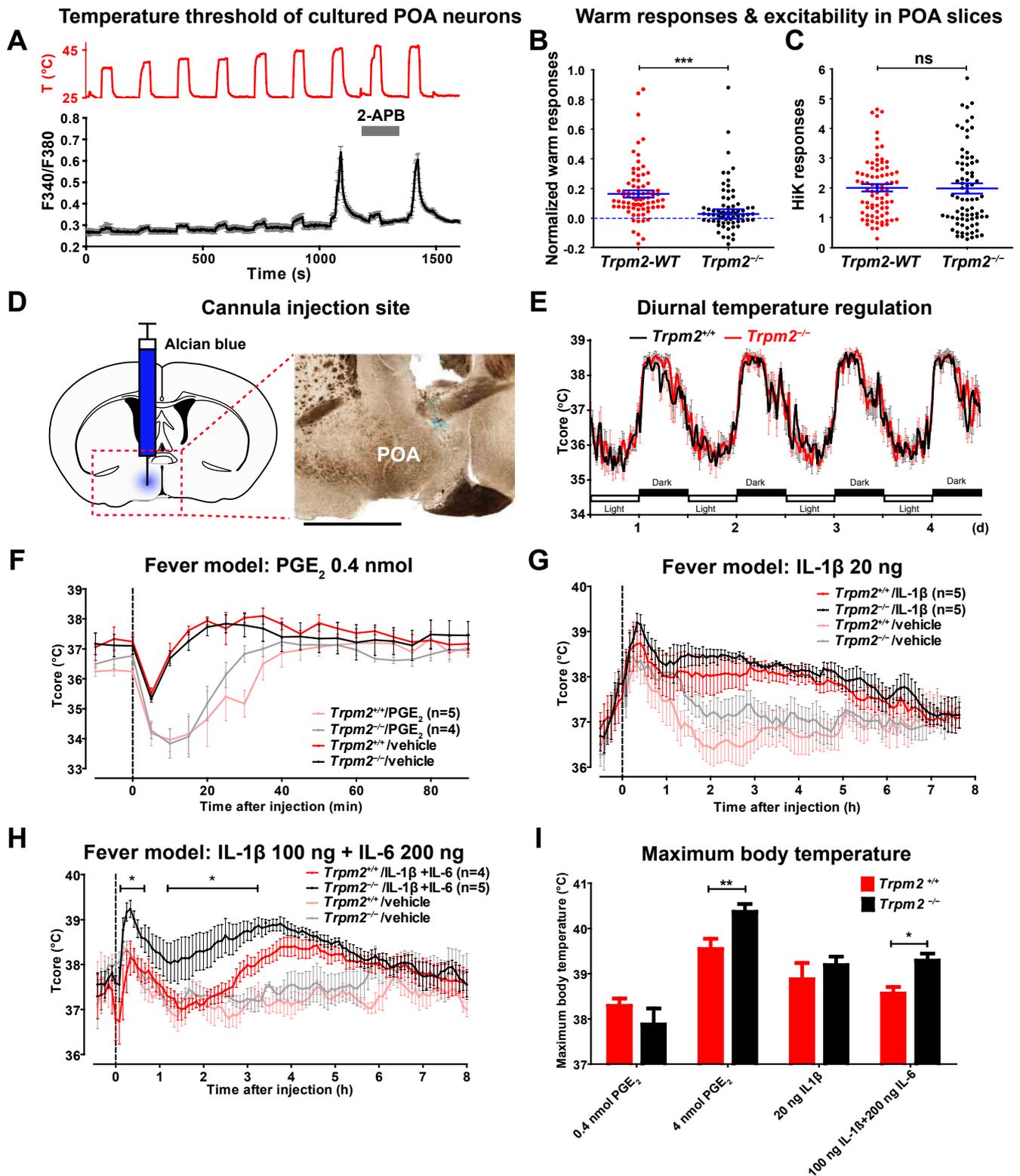
The table displays the raw calcium imaging data (time course experiment of temperature stimulations) of HEK293 cells, and cultured POA neurons of *Trpm2*<sup>+/+</sup> and *Trpm2*<sup>-/-</sup> mice. Additionally, the “Notes and Statistics” sheet describes the step-by-step analysis of warm-responsive neurons. Table S1 corresponds to Fig. 1G-I and fig. S1F.

### **Movie S1**

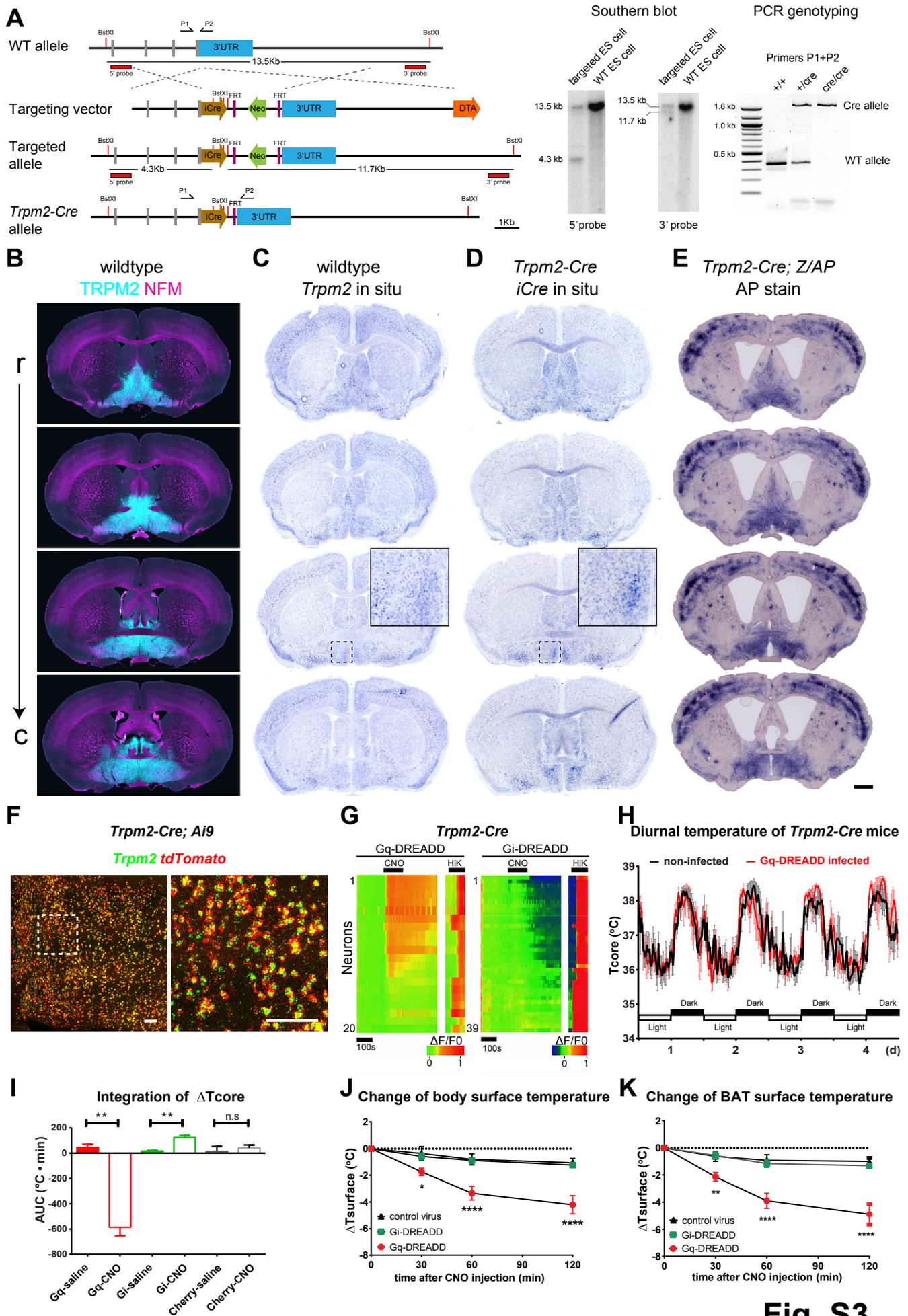
**Infrared thermal imaging of a *Trpm2-Cre* mouse preoptically expressing Gq-DREADD receptors.** Shown is a time-laps movie of a representative Gq-DREADD infected *Trpm2-Cre* mouse at different times before and after CNO injection and imaged with an IR camera from above. The movie is about 1 minute long and covers a total time span of 2 hours in real-time. The movie is broken up into 4 episodes as indicated. Episode 1: before CNO injection; episode 2: 12 to 19 minutes after CNO injection; episode 3: 1 hour after CNO injection; episode 4: 2 hours after CNO injection.



**Fig. S1**

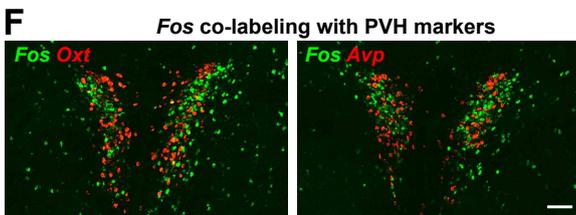
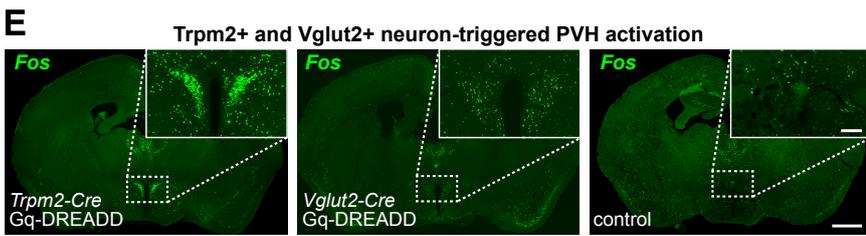
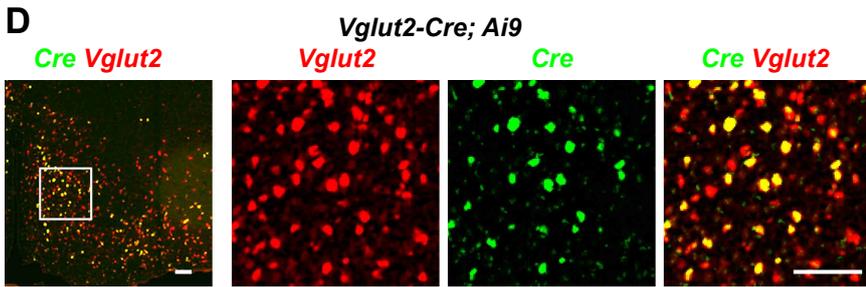
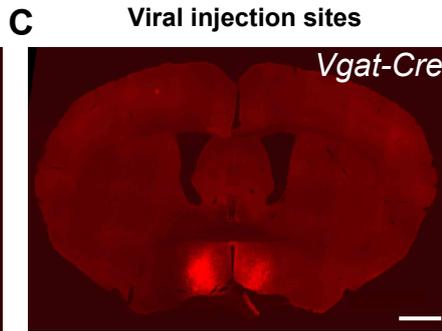
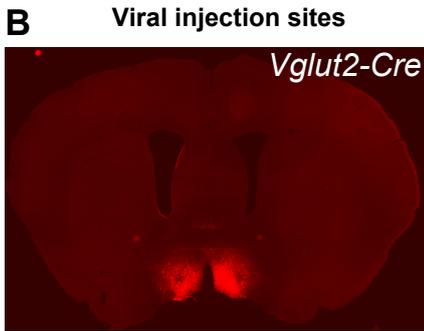
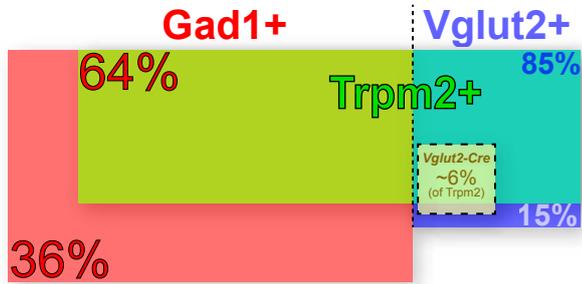


**Fig. S2**



**Fig. S3**

**A** Percent inhibitory and excitatory Trpm2+ neurons



**Fig. S4**

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