Olfactory cortical outputs recruit and shape distinct brain-wide spatiotemporal networks

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Abstract

Odor information is transmitted from the olfactory bulb to several primary olfactory cortical regions in parallel, including the anterior olfactory nucleus (AON) and piriform cortex (Pir). However, the specific roles of the olfactory bulb and cortical outputs in wider interactions with other interconnected regions throughout the brain remain unclear due to the lack of suitable in vivo techniques. Furthermore, emerging associations between olfactory-related dysfunctions and neurological disorders underscore the need for examining olfactory networks at the systems level. Using optogenetics, fMRI, and computational modeling, we interrogated the spatiotemporal properties of brain-wide neural interactions in olfactory networks. We observed distinct downstream recruitment patterns. Specifically, stimulation of excitatory projection neurons in OB predominantly activates primary olfactory network regions, while stimulation of OB afferents in AON and Pir primarily orthodromically activates hippocampal/striatal and limbic networks, respectively. Temporally, repeated OB or AON stimulation diminishes neural activity propagation brain-wide in contrast to Pir stimulation. Dynamic causal modeling analysis reveals a robust inhibitory effect of AON outputs on striatal and limbic network regions. In addition, experiments in aged rat models show decreased brain-wide activation following OB stimulation, particularly in the primary olfactory and limbic networks. Modeling analysis identifies a dysfunctional AON to Pir connection, indicating the impairment of this primary olfactory cortical circuit that disrupts the downstream long-range propagation. Our study for the first time delineates the spatiotemporal properties of olfactory neural activity propagation in brain-wide networks and uncovers the roles of primary olfactory cortical, AON and Pir, outputs in shaping neural interactions at the systems level.

Introduction

The olfactory system plays a crucial role in driving behavioral responses that are essential for daily functions. This system facilitates social behaviors by integrating olfactory stimuli with other sensory stimuli and prior experiences[1,2]. Olfactory systems can discriminate among hundreds to thousands of different odorants given the diversity of odorant receptors that specifically bind to the corresponding odorant molecules[3,4]. While olfaction may be more critical for lower-level mammalian species, it remains important for higher primates, as olfactory stimuli can influence emotions, moods, and behaviors[5]. One unique feature of the anatomical organization of the olfactory system is the absence of thalamic processing, indicating that the brain networks pivotal for olfactory processing are distinctly different from other sensory systems. Odor information detected in the olfactory epithelium is first processed at the olfactory bulb (OB), before being routed in parallel to several primary olfactory cortical regions, including the anterior olfactory nucleus (AON) and piriform cortex (Pir). Present studies have primarily examined anatomical projections and targets of local olfactory microcircuits, such as the OB, AON and Pir, including the synaptic interactions between these three connected regions[6-8]. However, the neural interplay between small numbers of anatomically connected regions (i.e., local micro-circuits) is likely insufficient to describe the dynamic properties of information processing that underlie specific brain functions[9-11]. Recent views have advocated for examining and/or modeling polysynaptic interactions across complex networks spanning multiple circuits distributed throughout the brain[11,12]. Moreover, the associations between olfactory-related dysfunctions and neurological disorders such as aging[13], neurodegenerative diseases[14], and recently COVID-19[15-17], highlight the need for examining the olfactory network at the systems level.

Interrogating brain-wide, long-range olfactory networks and their properties require a large-view brain-wide imaging technique that is sensitive to neural activity. However, mapping the brain-wide extent of olfactory regions via traditional functional MRI (fMRI) mapping approaches is technically challenging due to prominent olfactory response habituation[18,19]. Another complicating factor is the need to present a multitude of odors, which affects experimental efficiency and robustness in examining long-range olfactory networks and their properties. Previous attempts to study brain-wide regions involved in olfactory processing in rodents[20,21] and humans[22,23] using task-based fMRI have only identified some known primary olfactory regions, such as the OB, AON, Pir, olfactory tubercle (Tu), and amygdala (Amg). There is a significant gap in the visualization of long-range olfactory networks and the

characterization of their functional properties. Notably, the specific roles of the bulbar and AON

or Pir cortical outputs in wider interactions with other interconnected regions throughout the

brain in olfactory processing remain unclear. In addition, several higher-order regions suspected

to mediate more complex olfactory processing remain unidentified and/or to be

characterized[5,24,25]. Thus, we designed alternative neuromodulation strategies that can stimulate

olfactory-specific neurons in the OB, AON, and Pir with fMRI to characterize brain-wide

downstream activation targets and the dynamic properties of long-range olfactory networks.

We also aim to employ dynamic causal modeling (DCM)[26], a computational framework that

can be used to infer causal interactions among different regions in the olfactory network under

different stimulation conditions.

In this study, we deployed optogenetic fMRI to achieve reversible, millisecond

precision, and cell-type specific stimulation[12,27-31] of excitatory projection neurons in OB (i.e.,

mitral and tufted cells), and the olfactory-specific neurons of AON and Pir with computational

modeling to investigate the dynamics of large-scale olfactory spatiotemporal networks. Our

findings revealed distinct recruitment of downstream targets by olfactory-specific neural

populations in AON and Pir. AON-driven neural activities strongly activated hippocampal and

striatal networks, while Pir-driven activities preferentially recruited the limbic network.

Repeated excitations of AON or OB across multiple fMRI sessions decreased brain-wide

activations, while Pir excitations did not alter orthodromic neural activity propagation in long-

range olfactory networks. DCM of optogenetic fMRI data showed consistent negative effective

connectivity from AON to various downstream targets in the striatal, and limbic networks,

indicating a robust inhibitory effect of AON outputs on neural activity propagation in long-

range olfactory networks. Further, our systematic examination of long-range olfactory

networks in an aged rat model demonstrated an overall decrease in brain-wide activations upon

OB excitations, particularly in the primary olfactory and limbic networks. DCM showed that

the effective connectivity from AON to Pir switched from positive (healthy) to negative (aged),

indicating impairment of this cortical circuit that disrupted neural activity propagation to

primary olfactory and limbic networks. This study provides the initial delineation of the

spatiotemporal properties of olfactory neural activity propagation in brain-wide networks. We

show that primary olfactory cortices (AON and Pir) recruit distinct higher-order brain networks.

We also uncover the role of the inhibitory effect of AON and the excitatory effect of Pir outputs

in shaping the dynamic properties of neural activity propagation in long-range olfactory

networks.

Results

Olfactory bulb and primary olfactory cortices recruit distinct brain-wide long-range networks

To identify activated neural networks of the olfactory system, we fused Ca^{2+}/calmodulin-dependent protein kinase II alpha (CaMKIIα)-dependent ChR2(H134R) with mCherry to transfect excitatory projection neurons in OB of healthy adult Sprague Dawley rats. Colocalization of mCherry and CaMKIIα confirmed the specific expression of CaMKIIα in mitral and tufted cells within the mitral cell layer and the external plexiform layer of OB (**Fig. 1A**). Optogenetic stimulation was delivered at 1 Hz (10% duty cycle, 40 mW/mm^2), 5, 10, 20 and 40 Hz (30% duty cycle, 40 mW/mm^2) in a block-design paradigm covering the frequency range of delta (1 Hz, low frequency) to gamma (40 Hz, high frequency) oscillations. Different frequencies were presented in a randomized manner within each fMRI session (**Fig. 1B**).

First, we stimulated the OB, the first stage of olfactory information processing in the central nervous system. We detected robust BOLD fMRI activation bilaterally within and beyond primary olfactory networks during 1 Hz stimulation in ipsilateral OB excitatory (**Fig. 2A**). We identified robust and widespread BOLD responses across both hemispheres in known primary olfactory (AON, Pir, tenia tecta, TT, entorhinal cortex, Ent and olfactory tubercle, Tu), limbic (cingulate, Cg, orbitofrontal, OFC and insular, Ins, cortices, and amygdala, Amg), hippocampal (ventral hippocampus, vHP), striatal (nucleus accumbens, NAc and ventral caudate putamen, vCPu), and sensorimotor (motor, MC, somatosensory, S1, auditory, AC and visual, V1, cortices) regions (**Fig. 2A**). These activations indicate the presence of interactions between the primary olfactory network and other networks associated with high-order functions beyond olfactory processing, such as cognition, reward, and cross-modal sensory processing.

Although the first stage of olfactory processing occurs at the OB, primary olfactory cortices (e.g., AON and Pir), which receive parallel direct input from the OB, are also critical for olfaction, as they participate in olfactory memory functions and learning in odor representations within local micro-circuits[32,33] and interactions with the hippocampus[32,34]. As such, we sought to directly examine AON/Pir in our olfactory brain-wide neural networks. We note that direct optogenetic targeting of AON/Pir excitatory neurons may target additional neuronal populations, as not all excitatory neuronal populations in AON and Pir receive olfactory inputs from OB[35,36]. Here, we designed an optogenetic stimulation protocol based on projection targeting (**Supplementary Fig. 1**), whereby the OB afferent terminals were

stimulated at AON/Pir, thereby enabling selective stimulation of neurons in AON/Pir that

exclusively receive olfactory inputs. Upon 1 Hz stimulation of OB afferents at AON, we

observed robust widespread BOLD activations in bilateral primary olfactory, limbic, striatal,

sensorimotor cortical and hippocampal regions (**Fig. 2B**), similar to the spatial patterns in our

findings for OB stimulation. Meanwhile, 1 Hz optogenetic stimulation of OB afferents at Pir

also evoked bilateral BOLD responses in primary olfactory, limbic, and striatal regions albeit

with significantly weaker and sparser BOLD activations in bilateral limbic, striatal, and

sensorimotor cortical regions (**Fig. 2C**).

In addition to the qualitative description of BOLD activation distributions, we further

sought to quantify the strength of neural activity at each downstream activation target upon

stimulation of the three targets (i.e., OB, AON and Pir). We first calculated the area under the

BOLD signal profile (AUC) during the four 20-s 1 Hz optogenetic stimulation blocks for each

downstream activation target (**Fig. 2A-C**). However, direct comparisons between AUCs of the

three separate optogenetic fMRI experiments are erroneous given the differences in absolute

BOLD signal amplitudes, which are likely caused by differences in ChR2 transfection volumes

in OB excitatory neurons and OB afferents at AON/Pir, and subsequently the population size

of neurons/axonal terminals stimulated. The vast majority of OB neurons are interneurons and

not projection neurons that terminate in AON/Pir[37]. Hence, we normalized the AUCs to the

sum of BOLD activations of their respective optogenetic fMRI experiments (i.e., defined as

BOLD activation strength here; **Fig. 2D**). To distinguish the brain networks that are

predominantly recruited by AON- or Pir-driven neural activities compared to OB stimulation,

we found that the bulk of OB-driven neural activities were concentrated within the primary

olfactory network (i.e., OB, AON, Pir, TT, Ent and Tu; $40.5 \pm 1.6\%$ vs. $27.2 \pm 2.7\%$, $P < 0.01$

for OB vs. Pir; and $40.5 \pm 1.6\%$ vs. $35.6 \pm 2.0\%$, $P < 0.1$ for OB vs. AON). The strongest

BOLD activations within the primary olfactory network (i.e., AON, Pir, TT, Ent, Tu) for OB

stimulation experiments aligned with the direct anatomical projections from OB to these

targets[37], indicating the successful excitation of OB projection neurons. Interestingly, we

discovered that AON-driven activities preferentially recruited the striatal (i.e., NAc and vCPu;

$13.8 \pm 0.6\%$ vs. $9.8 \pm 0.8\%$, $P < 0.01$ for AON vs. OB) and hippocampal networks ($2.4 \pm 0.1\%$

vs. $1.2 \pm 0.2\%$, $P < 0.01$ for AON vs. OB), while Pir-driven activities strongly targeted the

limbic network (i.e., Cg, PrL, OFC, Ins and Amg; $39.5 \pm 2.8\%$ vs. $30.3 \pm 1.4\%$, $P < 0.05$ for

Pir vs. OB). Meanwhile, all three stimulated regions showed relatively equal recruitment of

sensorimotor networks. The specificity afforded by perturbing the OB afferents at AON and

Pir indicated that these olfactory inputs recruit distinct downstream targets in primary olfactory,

limbic, hippocampal and striatal networks, despite sharing extensive overlaps in anatomical

projections[25,37-39].

We also examined BOLD activations upon optogenetic stimulation of the three primary

olfactory regions (i.e., OB, AON and Pir) at various frequencies. We observed robust bilateral

activations during 5 Hz stimulation of OB excitatory neurons (**Supplementary Fig. 2A**) and

OB afferents in AON (**Supplementary Fig. 3A**), although the activations at the contralateral

hemisphere were relatively weaker. The subsequent increase in stimulation frequencies to 10

Hz, 20 Hz and 40 Hz resulted in robust activations that were predominantly localized in the

ipsilateral primary olfactory and limbic networks (**Supplementary Figs. 2B-D & 3B-D**).

Meanwhile, BOLD activations were only detected in ipsilateral primary olfactory, limbic, and

striatal networks when higher frequency optogenetic stimulation was presented to OB afferents

at Pir (**Supplementary Fig. 4**). To examine long-range olfactory networks, we focused on the

1 Hz optogenetic stimulation experiments.

Altogether, our approach to stimulating the three primary olfactory regions (i.e., OB,

AON and Pir) provides valuable insights into the networks that are pivotal for olfactory

processing. Notably, the distinction in brain networks that are recruited by olfactory-specific

neural populations in AON vs. Pir such as hippocampal and striatal vs. limbic reveal functional

insight into the AON and Pir in olfactory processing.

AON and Pir circuit outputs shape the dynamic properties of long-range olfactory

networks

The distinct spatiotemporal propagation properties and the recruitment of different long-range

olfactory-related networks suggest varied modulatory roles for OB, AON and Pir when

interacting with their downstream targets. Here, we first examined the dynamics of BOLD

activations across multiple 1 Hz stimulation sessions. We observed distinct changes to the

spatiotemporal properties of BOLD fMRI activations across the three optogenetic fMRI

sessions. Overall, the brain-wide activations at primary olfactory, limbic, striatal, and

sensorimotor cortical networks in our first fMRI session showed an appreciable decreasing

trend in subsequent stimulation sessions. For OB 1 Hz stimulation in the 2nd fMRI session,

there was a decrease in the BOLD activations at most of the downstream targets across multiple

networks. These include the ipsilateral OB (i.e., the stimulation site), contralateral TT, and

bilateral AON, Pir, Ent and Tu of the primary olfactory network; bilateral OFC, Ins and Amg

of the limbic network; bilateral NAc and ipsilateral vCPu of the striatal network; and bilateral

MC and ipsilateral AC of the sensorimotor network (**Fig. 3A** and **Supplementary Fig. 5A**).

BOLD activations during the 3rd fMRI session also showed decreased activations (i.e., 3rd vs 1st fMRI session; **Fig. 3A** and **Supplementary Fig. 5A**). Comparatively, we observed a dramatic decrease in activations upon 1 Hz stimulation of OB afferents at AON during the later fMRI sessions. We found diminished or even absent bilateral activations in downstream targets. Specifically, activations evoked by AON stimulation showed a significant decrease in the 2nd and 3rd sessions at all regions except contralateral OB (**Fig. 3B** and **Supplementary Fig. 5B**). Notably, activations at bilateral vHP also significantly decreased despite having the weakest activation among downstream targets during the 1st fMRI session. However, the activations upon 1 Hz stimulation of OB afferent inputs at Pir remained relatively stable across fMRI sessions in the primary olfactory, limbic, striatal and sensory cortical networks except for a slight decrease in Amg (**Fig. 3C** and **Supplementary Fig. 5C**).

We then sought to identify the driver of these distinct influences of OB, AON and Pir stimulation on the dynamic response properties of long-range olfactory-related networks. Conventional analyses of BOLD fMRI data such as general linear model (GLM) and comparisons of the energy of BOLD signal profiles are insufficient and unable to infer the causal influences of OB-, AON- and Pir-driven neural activity on their downstream targets. Here, we applied DCM[26] on our optogenetic fMRI data to parametrize the downstream effects on selected targets in the right brain hemisphere (i.e., the same hemispheric side to the optogenetic stimulation site). Specifically, we used DCM to computationally model the effective connectivity strength of selected downstream connections in OB, AON and Pir stimulation data. We defined a six-node brain network and its connections that encompassed regions from primary olfactory (OB, AON, Pir and Ent), striatal (vCPu) and limbic (Amg) networks (**Fig. 4A**). These connections were primarily based on the well-defined anatomical feedforward and feedback projections involving the three major primary olfactory regions (i.e., OB, AON and Pir). For simplicity, we did not model other downstream regions to avoid data overfitting issues[40]. Further, we only modeled the optogenetic fMRI data from the 1st session, because BOLD signal profiles for the 2nd and 3rd fMRI sessions had lower SNR due to decreased and in some cases absent activations in long-range olfactory networks. Overall, we observed that the connectivities were primarily positive (i.e., excitatory effects) between the nodes in the primary olfactory network (i.e., OB, AON and Pir; **Fig. 4B**) for all three different stimulations, corroborating documented reciprocal glutamatergic excitatory projections between these three regions[25,38,41,42]. Notably, we observed a significant negative AON and positive Pir connectivity to downstream targets (i.e., Ent, vCPu and Amg; **Fig. 4B**). Quantitative comparisons of the effective connectivity strengths from AON/Pir to Ent, vCPu

and Amg confirmed the robust negative AON vs. positive Pir connectivity to downstream targets (*AON-Ent vs. Pir-Ent*: -0.21 ± 0.14 vs. 0.39 ± 0.08 Hz, $P < 0.05$ for OB stimulation and -0.34 ± 0.09 vs. 0.35 ± 0.06 Hz, $P < 0.01$ for AON stimulation; *AON-vCPu vs. Pir-vCPu*: -0.03 ± 0.06 vs. 0.31 ± 0.10 Hz, $P = 0.053$ for OB stimulation, -0.37 ± 0.08 vs. 0.40 ± 0.06 Hz, $P < 0.01$ for AON stimulation and -0.14 ± 0.06 vs. 0.28 ± 0.07 Hz, $P < 0.05$ for Pir stimulation; *AON-Amg vs. Pir-Amg*: -0.10 ± 0.10 vs. 0.26 ± 0.04 Hz, $P < 0.05$ for OB stimulation, -0.25 ± 0.12 vs. 0.33 ± 0.06 Hz, $P < 0.05$ for AON stimulation and -0.16 ± 0.06 vs 0.30 ± 0.06 Hz, $P < 0.05$ for Pir stimulation; **Fig. 4C**). These results demonstrate distinct inhibitory effect of AON and excitatory effect of Pir outputs in long-range olfactory networks. We speculate that the significant decrease in BOLD activations during repeated OB or AON stimulations, but not during Pir stimulations, arises from the strong inhibitory effect of AON outputs to various targets beyond the local olfactory microcircuits (i.e., OB, AON and Pir). Taken together, our results indicate that the outputs of primary olfactory cortices, AON and Pir, shape the dynamic properties of neural activity propagation in long-range olfactory networks.

Repeated AON stimulations diminish orthodromic neural activity propagation

We recognize that our optogenetic stimulation protocols, particularly the stimulations of ChR2-transfected OB afferents at AON and Pir compared to the stimulation of OB excitatory neurons (**Supplementary Fig. 1**), can be sensitive to antidromic-driven neural activities due to activation of the axon terminals instead of the soma[43,44]. To verify the neural activities underlying our fMRI findings and delineate features of neural activity propagation, we conducted local field potential (LFP) recordings at seven regions in the right brain hemisphere. These regions include the ipsilateral OB, AON, Pir, vCPu, V1, Amg, vHP and dHP. The stimulation paradigm (i.e., block-designed and three recording sessions for each animal) matches that from the fMRI experiments. The averaged four 60-s-block LFP traces (**Fig. 5A-C**) and the subsequent power analysis of these LFP traces for the first recording session (**Fig. 6A-C**) largely concurred with the magnitude and temporal profiles of BOLD fMRI activations (**Fig. 2A-C**), with evoked LFPs observed in all recorded regions during the first experimental session. Parallel to our fMRI findings, the strength of neural activities, which were quantified through LFP power analysis (i.e., area under absolute LFP trace normalized to 30-s baseline LFP power), was also decreased during repetitive 1 Hz stimulation of OB or OB afferents at AON but was relatively unchanged under Pir stimulation (1st vs 2nd and 3rd recording sessions; **Fig. 6**).

We next characterized the latency of neural activity at each recorded region by using

the first appreciable evoked LFP peak to represent the initial neuronal population activity[28,29,45]

(**Fig. 5**). Negative LFP peaks occurred first in each of the three stimulation targets, while

positive peaks were observed for their corresponding downstream targets. During somatic

activation of ipsilateral OB excitatory neurons, there is an evoked LFP response latency of 6.1

$\pm$ 0.7 ms (**Fig. 5A**), which is within the 5-6 ms range documented for response latency of OB

excitatory neurons in the presence of natural odors[46,47]. We measured delays of 8.7 $\pm$ 0.9 ms

and 8.2 $\pm$ 1.0 ms between OB and AON, and OB and Pir, respectively, consistent with the

reported 5-10 ms neural activity propagation along monosynaptic projections of OB to AON[48]

and Pir[49-51]. Neural activities from ipsilateral OB arrived at ipsilateral vCPu and V1 in 9.7 $\pm$

2.6 ms and 11.7 $\pm$ 3.0 ms in line with the axonal conduction speed between OB and

striatal[52,53]/sensorimotor cortical[54,55] regions. Lastly, the long delays of 37.9 $\pm$ 0.9 ms and 39.2

$\pm$ 2.1 ms from ipsilateral OB to vHP and dHP respectively, indicate that neural activities

propagated through polysynaptic projections of OB-primary olfactory cortices-Ent-HP[51,56].

When stimulating the OB afferents in AON, we found a response latency of 7.7 $\pm$ 0.7 ms locally

at AON (**Fig. 5B**). A subsequent delay of 7.2 $\pm$ 1.8 ms and 5.0 $\pm$ 0.9 ms was then measured

between AON and its downstream targets in the primary olfactory network (i.e., OB and Pir),

corroborating reported values of 4-10 ms[57]. The presence of a significant delay between AON

and OB neuronal population activities with appreciable jitter demonstrates that neural activity

propagation through the activation of OB axonal terminals at AON is largely orthodromic in

our experiment. Neural activity from olfactory-specific AON neurons also propagated in an

orthodromic manner to targets in the striatal (vCPu), sensorimotor cortical (V1) and limbic

(Amg) networks, 7.1 $\pm$ 1.0 ms, 7.5 $\pm$ 1.6 ms, and 7.0 $\pm$ 0.4 ms, respectively, in line with axonal

conduction speed[58]. Meanwhile, excitation of the OB afferents at Pir evoked a response latency

of 5.9 $\pm$ 0.6 ms (**Fig. 5C**), which closely matched the latency of ~5 ms upon optogenetic

stimulation of afferents to Pir in a previous study[41]. Subsequently, the neural activity

propagated from Pir to OB, AON, vCPu, V1, and Amg at a delay of 8.1 $\pm$ 0.5 ms, 8.2 $\pm$ 0.3 ms,

7.9 $\pm$ 0.8 ms, 8.6 $\pm$ 0.5 ms, and 8.9 $\pm$ 0.8 ms, respectively. We conclude that the observed

neural activity propagation is predominantly orthodromic given the significant delay and jitter

of excitatory inputs from Pir to OB. Notably, the neural activity propagated in 28.2 $\pm$ 3.0 ms

from Pir to vHP, while the propagation speed from the AON to vHP was faster at 22.2 $\pm$ 1.7

ms. The longer delay between Pir and the hippocampus suggests the presence of an

intermediary region between the Pir and Ent.

Taken together, our electrophysiological recordings showed the direct involvement of

both local and long-range projections that mediate the propagation of neural activity in

olfactory networks. Moreover, orthodromic propagations when stimulating OB afferents at

AON and Pir indicate that the long-range inhibitory effect of AON and excitatory effect of Pir

outputs, which shaped the dynamic properties of BOLD activations in olfactory networks, are

predominantly driven by the feedforward projections of olfactory-specific neurons in the

primary olfactory cortices.

323 **Long-range olfactory networks of aged brains exhibit decreased neural activations and**

324 **impaired AON to Pir connectivity**

Informed by the insights gained from characterizing the spatiotemporal properties of long-

range olfactory networks in normal healthy brains, we deployed our optogenetic fMRI platform

to examine the systemic dysfunction of olfactory networks in a diseased brain. Here, we

employed a D-galactose model of accelerated aging in rodents. This model induces higher

levels of oxidative stress, mitochondrial dysfunction, and cellular apoptosis. These factors can

lead to the decline of cognitive functions, which is a key symptomatic hallmark of age-related

brain senescence[59,60]. Spatially, we observed that 1 Hz stimulation of the OB excitatory neurons

evoked bilateral activations in all brain regions like primary olfactory, limbic, hippocampal,

striatal and sensorimotor networks (**Fig. 7A** vs. **Fig. 2A**). These spatially preserved activations

indicate that neural activity propagation in long-range olfactory networks remains intact in

aged brains. However, the strength of neural activities at downstream activation targets

exhibited a general decrease in both brain hemispheres when compared to healthy animals. A

significant decline in BOLD activations was mostly identified in ipsilateral primary olfactory

(aged vs. healthy; AONR: 16.4 ± 3.8 vs. 33.4 ± 6.5, $P < 0.05$; PirR: 33.5 ± 6.0 vs. 58.0 ± 8.4,

$P < 0.05$; TTR: 15.1 ± 4.8 vs. 41.3 ± 7.0, $P < 0.01$) and ipsilateral limbic networks (CgR: 7.6

± 1.9 vs. 15.5 ± 1.9, $P < 0.01$; OFCR: 19.4 ± 3.0 vs. 36.7 ± 3.8, $P < 0.01$; InsR: 29.2 ± 3.8 vs.

40.6 ± 5.0, $P < 0.05$). We also observed a general trend of decline in the other regions of the

primary olfactory network (PirL: 23.7 ± 5.4 vs. 37.8 ± 7.1, $P = 0.07$ and EntR: 26.0 ± 4.8 vs.

38.3 ± 5.7, P = 0.06), and the ipsilateral sensorimotor network (MCR: 16.5 ± 2.4 vs. 24.5 ± 4.3,

$P = 0.059$). Surprisingly, there was a general trend of increased BOLD activations in the striatal

network (NAcR: 27.7 ± 2.8 vs. 20.3 ± 3.2, $P = 0.055$). DCM of the optogenetic fMRI data in

aged animals revealed a switch in the effect of ipsilateral AON outputs to ipsilateral Pir from

excitatory to inhibitory (healthy vs. aged; 0.23 ± 0.09 vs. -0.12 ± 0.10 Hz, $P < 0.05$), indicating

dysfunction of a key primary olfactory cortical circuit in the olfactory network disrupting

neural activity propagation to primary olfactory and limbic networks. Note, that in healthy animals (**Fig. 2D**), the primary olfactory network was preferentially recruited by AON-driven neural activities, while the limbic network was recruited mainly by Pir-driven neural activities. The predominant effect observed in the right hemisphere was likely due to the stimulation of the right/ipsilateral OB. We expect both hemispheres to be affected equally in aged brains.

Discussion

As a major sensory system, the olfactory system has been linked to numerous non-olfactory regions pivotal for sensory perception and higher-order cognition[5], yet recent views highlight the significant gap in understanding where and how olfactory information is distributed downstream beyond these known primary regions[5,61-64]. In this study, we deployed optogenetic stimulation, fMRI, and electrophysiological recordings to systematically dissect the spatiotemporal properties of long-range olfactory networks. We discovered that olfactory-specific neural populations in AON and Pir distinctly recruited downstream targets such as the hippocampal and striatal networks for AON-driven neural activities, and the limbic network for Pir-driven neural activities. Further analysis revealed robust adaptation of brain-wide BOLD fMRI activations upon repeated optogenetic stimulations of OB afferents in AON or OB excitatory neurons, which were absent when stimulating OB afferents at Pir. Computational modeling of our optogenetic fMRI data through DCM indicated a significant inhibitory effect of AON outputs to downstream targets in the olfactory network compared to the excitatory effect from Pir outputs. These results highlight a likely dominant role of olfactory-specific AON neural populations in mediating sensory adaptation in the presence of olfactory inputs. Electrophysiological recordings demonstrated predominant orthodromic neural activity propagation to downstream targets in long-range olfactory networks for all optogenetic stimulations and verified the adaptation of optogenetically-evoked neural activities observed in the fMRI experiments. In addition, our systematic comparisons of the olfactory networks between aged and healthy brains revealed an overall decrease in the amplitudes of BOLD activations, particularly in regions of the primary olfactory and limbic networks, along with increased inhibitory effect of AON outputs to the Pir, indicating impairment of this key olfactory cortical circuit. Our optogenetic fMRI approach constitutes the initial characterization of 1) the spatiotemporal properties of long-range olfactory networks and 2) the roles of AON and Pir outputs in shaping the dynamic properties of olfactory neural activity propagation in healthy and diseased brains.

Habituation or adaptation shows decreased responses to repeated or continuous
stimuli[65,66]. Olfactory adaptation refers to the ability of the network to adjust its sensitivity at
different stimulus intensities, which is essential for retaining high sensitivity to odd/another
stimulus under repetitive or prolonged odor stimulation[67]. This fundamental property of all
sensory systems prevents overstimulation and retains alertness to the environment[66]. Olfactory
adaption occurs at two levels: peripheral – at the level of olfactory receptors – and central – at
the level of the central nervous system[66]. At present, the underlying neural mechanism of
peripheral adaptation that occurs at the molecular level[67], and central adaptation within the
primary olfactory local network (i.e., OB[68,69], AON[25], Pir[70,71] and Tu[72]) has been shown to
require glutamate release in the OB[73]. However, the significant decreases in BOLD activation
across all downstream targets indicate that olfactory adaptation may occur more broadly across
various targets of the olfactory network rather than being localized at the OB. Specifically, our
results reveal a previously undefined role of the AON in mediating olfactory adaptation at the
systems level, as we showed strong adaptation of optogenetically-evoked neural activities (**Figs.**
3B and **6B**) upon 1 Hz stimulation of OB projection neurons and OB afferent inputs at AON,
but not OB afferents at Pir, along with robust inhibitory effects of AON outputs to Ent, vCPu
and Amg (**Fig. 4C**).
AON sends robust centrifugal inputs to OB that can inhibit bulbar outputs[25,57] and
robust glutamatergic feedforward projections that bilaterally innervate the Pir[41,74], to enhance
activation of piriform neurons in response to odor[41]. Further, recent studies on AON functions
revealed its involvement in interhemispheric communication[25,75,76], sensory gating[77,78] and
odor memory[32,35], indicating a predominant role of AON as a hub to process olfactory
information (i.e., odor identity decoding and gain control of OB outputs[42]). In our study, only
stimulation of the ipsilateral AON evoked bilateral brain-wide BOLD responses in primary
olfactory (i.e., Pir, Tu), limbic (i.e., Ins, OFC) and striatal regions (i.e., NAc, vCPu) at all
frequencies (**Fig. 2B and Supplementary Fig. 2**), indicating the pivotal role of AON in
olfactory processing between both hemispheres. AON receives centrifugal projections from
HP, which represents its ability to process odor memory-related contextual
information[32,35,77,79,80]. Notably, our fMRI analysis also showed that AON-driven neural
activities preferentially recruited the hippocampal network. Further, we observed robust
bilateral brain-wide activations initiated by 1 Hz stimulation at AON during the first fMRI
session, which were followed by dramatically decreased downstream activations under
repetitive stimulations (**Fig. 3**). We speculate that AON exerts robust feedforward neural
activity propagation to long-range downstream targets after the early stage of olfactory

processing at OB. However, in the presence of repeated strong activations of olfactory-specific

neural populations in AON (i.e., through OB or AON stimulations), centrifugal feedback from

the AON, which preferentially controls the gain of OB neuronal responses[42], likely inhibited

OB outputs[25,57], resulting in decreased brain-wide activations in later fMRI sessions. Further,

we identified robust inhibitory AON outputs to non-primary olfactory brain regions (i.e., Ent,

vCPu, Amg; **Fig. 4C**), regardless of the stimulated region, indicating that neural adaptation

processes could also be distributed beyond local olfactory micro-circuits (i.e., centrifugal AON

outputs to OB). We note that olfactory adaptation at the systems level is quite under-studied,

and future studies may examine the circuit properties at these long-range targets when

processing odor information.

Pir sends ipsilateral inputs to OB and AON[39,81] and functions to amplify the inhibition

of odor-related signals and finely tune them in OB[42,82], in order to process and identify critical

features of an odor. Although, studies have noted a significant overlap in the functions of Pir

during olfactory processing with those of AON[37,83,84], the recurrent neural architecture of

Pir[85,86] stands in contrast to that of AON. These structural differences indicate how Pir

processes olfactory inputs from OB distinctly. We found that Pir-driven neural activities

preferentially recruited regions in the limbic network, which differed from those primarily

recruited by AON. Anatomically, Pir projects extensively to regions of the limbic network such

as OFC and Amg[87,88], which demonstrates the specificity in our optogenetic fMRI approach

when dissecting the spatiotemporal properties of long-range olfactory networks. Overall, we

also observed that exciting olfactory-specific neurons in Pir produced weaker brain-wide

BOLD activations than OB or AON stimulation (**Fig. 2C** and **Supplementary Fig. 4**). We

posit this occurs from the recurrent neural architecture of Pir (i.e., dense reciprocal synaptic

connections between olfactory-specific neurons in Pir and inhibitory interneurons), which can

recruit sustained inhibition that subsequently diminishes overall Pir cortical neural activity in

the presence of olfactory inputs[85].

The ability to detect and discriminate odors declines during aging, leading to a decrease

in quality of life and diminished appetite with a concomitant impact on nutrition[89]. Decreased

olfactory function with aging parallels generalized age-related deficits in sensory functions and

cognition that occur in the absence of obvious disease states[14]. However, studies in rodents

demonstrate that the decreased ability to perceive odors was not associated with a general

decline in the number of receptors/cells in the olfactory epithelium or a decline in neuronal

populations (i.e., mitral and tufted cells, and interneurons) in the OB[13]. Instead, the synaptic

density in the glomeruli of the OB appears diminished[13], suggesting impaired downstream

olfactory information processing from the OB to targets in the cortical and subcortical regions.
At present, there is no consensus on the most representative animal model of aging that best
mimics the human condition. D-galactose has been shown to induce advanced glycation end
products formation, a process that contributes to early phases of age-related diseases in both
humans and rodents[90,91]. With the discovery of diminished brain-wide neural activity
propagation, our results in a D-galactose rat model indicate that the impairment of OB
downstream olfactory information processing is likely true in aged brains. We also found the
switch from positive effective connectivity of AON to Pir and AON to vCPu in healthy animals
to significantly negative ones in aged animals, indicating abnormal AON output to downstream
targets in long-range olfactory networks during olfactory information processing. Notably, age-
related neurodegenerative diseases (e.g., Alzheimer's and Parkinson's disease) can exhibit
olfactory impairments in the early stages with cell loss and axonal degradation in AON[92-94].

In summary, our study demonstrates the distinct recruitment of higher-order brain
networks by primary olfactory cortices (AON and Pir), including the identification of AON
and Pir outputs that can shape the dynamic properties of neural activity propagation in long-
range olfactory networks in healthy and aged brains. Such documentation of long-range
olfactory networks and their properties will be of great value to both neuroscientists and
clinicians. By gaining an integrated understanding of olfactory networks and their fundamental
properties in both healthy and aged brains, we can develop therapeutic strategies for
neurological diseases associated with olfactory system dysfunctions.

Methods

Animal subjects. Adult male Sprague Dawley rats were used in all experiments. Animals were individually housed under a 12-h light/dark cycle with access to food and water ad libitum. All animal experiments were approved by the Committee on the Use of Live Animals in Teaching and Research (CULATR) of the University of Hong Kong. Seven groups of animals were used: Group I (n = 11) for optogenetic fMRI experiments upon olfactory bulb (OB) stimulation; Group II (n = 11) for optogenetic fMRI experiments upon anterior olfactory nucleus (AON) stimulation; Group III (n = 9) for optogenetic fMRI experiments upon piriform cortex (Pir) stimulation; Group IV (n = 7) for electrophysiological recordings upon OB optogenetic stimulation; Group V (n = 7) for electrophysiological recordings upon AON optogenetic stimulation; and Group VI (n = 5) for electrophysiological recordings upon Pir optogenetic stimulation; and Group VII (n = 13) for optogenetic fMRI experiments upon OB stimulation in aged rats.

Note that for the electrophysiological experiments, not all animals resulted in successful recordings in all electrode locations. For Group IV, only OB, AON, Pir, vCPu and V1 were recorded in all 7 animals, whereas only 2/7 in vHP and 3/7 animals in dHP. For Group V, AON in 7 animals, OB in 6 animals, whereas only 5/7 in Pir, 3/7 in vCPu, V1 and vHP, and 4/7 animals in Amg. For Group VI, OB, Pir and vHP in 5 animals, followed by 4/5 in AON, 3/5 in vCPu and V1, and 2/5 animals in Amg.

Virus packaging. Recombinant AAV vectors were serotyped with AAV5-coated proteins and produced by the Vector Core of the University of North Carolina. Viral titer in particles/mL was 4×10^{12} for AAV5CaMKII α-ChR2(H134R)-mCherry. Maps are available online from http://www.stanford.edu/group/dlab/optogenetics.

Accelerated aging animal model induction protocol. D-galactose (50 mg/kg, subcutaneous) was administered daily to 10-week-old male adult SD rats for a total duration of 8 weeks. D-galactose administration in rodents has been shown to cause increased oxidative stress, mitochondrial dysfunction and cellular apoptosis, which leads to the deterioration of cognitive functions, a primary symptomatic hallmark of senescence of the brain[59,60].

Stereotactic surgery for viral injection. Stereotactic surgeries were performed with rats at 6-7 weeks of age for group I to VI, and at 15 weeks (i.e., 5 weeks in on the daily D-galactose administration) for group VII. Rats were anesthetized with an intraperitoneal bolus injection of a ketamine (90 mg/kg) and xylazine (40 mg/kg) mixture. After shaving the scalp, rats were

secured in a stereotactic apparatus with non-rupturing ear bars and craniotomy was conducted.

Buprenorphine (0.05 mg/kg, subcutaneous) was administered to minimize pain. A heating pad

was used to prevent hypothermia. Viral injection was performed at two depths in OB (+7.5 mm

anterior to Bregma, +1.8 mm medial-lateral right hemisphere, -1.8 mm and -2.1 mm from the

surface of dura), with subsequent verification by high-resolution anatomic MRI. Viral

constructs were delivered at 1.5 µL volume at each depth through a 5 µL syringe and 33-gauge

beveled needle injected at a rate of 150 nL/min. The injection needle was held in place for 10

min and retracted from the brain slowly. Scalp incision was sutured, and animals were kept on

a heating pad until recovery from anesthesia. Buprenorphine (0.05 mg/kg, subcutaneous) was

administered post-injection twice daily for 72 hours to minimize discomfort. Enrofloxacin was

also administered orally for 72 hours to minimize infection and inflammation post-surgery.

Animals recovered for 4-5 weeks before conducting MRI and electrophysiology experiments.

Animal preparation for optogenetic fMRI experiments. Before surgery, animals were

anesthetized with isoflurane (induction 3% and maintenance 2%) and secured on a stereotactic

frame. Before animals were placed in-magnet, surgery was performed under 1.2-2.0%

isoflurane to implant a custom-made plastic optical fiber cannula (core diameter = 450 µm,

outer diameter = 500 µm). Implantation of the cannula was made at three different locations in

the ipsilateral/right brain hemisphere depending on the experiments, namely at OB (+7.5 mm

anterior to Bregma, +1.8 mm medial-lateral, -2.1 mm from the surface of dura), AON (+5.2

mm anterior to Bregma, +1.8 mm medial-lateral, -4.3 mm from the surface of dura) and Pir

(+4.2 mm anterior to Bregma, +3.1 mm medial-lateral right hemisphere, -5.4 mm from the

surface of dura). Following a midline incision, a craniotomy was performed at these located

coordinates. The fiber tip was scored to create a beveled edge to facilitate insertion of the fiber

and minimize injury to brain tissue. The fiber cannula was then made opaque using heat

shrinkable sleeves to prevent light leakage during stimulation and the eyes of the animals were

blindfolded to prevent undesired visual stimulation, as in our previous studies[28-31,45,95]. Dental

cement was applied to the fiber cannula to affix it to the skull. The optical fiber tip was typically

situated in the mitral cell layer and external plexiform layer in OB stimulation experiments,

dorsal pars principalis of the AON to target OB afferents at AON, and anterior Pir for OB

afferents at Pir with verification by high-resolution anatomical MRI (**Fig. 1A**). The spatial

spread from the fiber tip for 473 nm blue light is rather small (200 µm and 350 µm at 50% and

10% of initial light intensity, respectively) and has little or no in the backward and lateral

directions[28,96], confining the optogenetic stimulation to the excitatory neurons in OB, and the

axonal terminals of OB at AON and Pir, respectively. Buprenorphine (0.05 mg/kg,
subcutaneous) was administered post-implantation.

The animals were mechanically ventilated at a rate of 60 breaths per minute with 1.0%
isoflurane in room-temperature air using a ventilator (TOPO, Kent Scientific, Torrington, CT).
One drop of 2% lidocaine was applied to the chords to minimize discomfort before
endotracheal intubation. During all fMRI experiments, animals were placed on a plastic holder,
and their heads were fixed with a tooth bar and plastic ear bars with screws. Continuous
physiological monitoring was performed using sensors from an MRI-compatible system (SA
Instruments, Stony Brook, NY). Rectal temperature was maintained at ~37.0 °C using a water
circulation system. Vital signs were within normal physiological ranges (rectal temperature:
36.5-37.5 °C, heart rate: 350-420 beats/min, breathing: ~60 breaths/min but not synchronized
to optogenetic stimulation, oxygen saturation level: >95%) throughout the experiments.

MRI scanner-synchronized optogenetic stimulation. An Arduino programming board
synchronized the scanner trigger and the lasers for optogenetic stimulation. Computers and
light delivery systems were kept outside the magnet, and optical patch cables (5-10 m)
delivered light into the bore of the scanner. For optogenetic stimulation, blue light was
delivered using a 473 nm Diode-pumped solid-state laser, measured before scanning as 8 mW
at the fiber-tip (450 μm, NA = 0.5) corresponding to a light intensity of 40 mW/mm^2. For all
fMRI experiments (i.e., OB, AON or Pir stimulation), five frequencies were used (1 Hz with a
10% duty cycle; 5, 10, 20 and 40 Hz with a 30% duty cycle). OB, AON or Pir was stimulated
with a block design paradigm that consisted of four 20s-on and 60s-off periods with the order
of stimulation frequencies pseudorandomized (**Fig. 1B**). Typically, three fMRI sessions were
performed for each animal (i.e., five scans per session covering the five different frequencies).

MRI acquisition procedure. All MRI experiments were carried out on a 7T MRI scanner
(ParaVision v5.1, PharmaScan 70/16, Bruker Biospin GmbH, Ettlingen, Germany) using a
transmit-only birdcage coil in combination with an actively decoupled single-channel receive-
only surface coil. Scout T2-weighted (T2W) images were first obtained using a rapid
acquisition with relaxation enhancement (RARE) sequence for accurate positioning and as an
anatomical reference with field of view (FOV) = 32 × 32 mm^2, matrix = 256 × 256, RARE
factor = 8, echo time (TE) = 36 ms, and repetition time (TR) = 4200 ms. Seventeen contiguous
1 mm slices were then positioned in the transverse orientation according to the rat brain atlas
to cover most of the brain from the OB to the cerebellum. All fMRI measurements were made

using a single-shot gradient-echo echo-planar imaging (GE-EPI) sequence with FOV = 32 ×

32 mm^2, matrix = 64 × 64, flip angle = 56°, and TE/TR = 20/1000 ms.

fMRI data analysis. fMRI analysis was performed using the procedure established in our

previous studies[28-31,45,95]. For each fMRI session, all GE-EPI images were first corrected for

slice timing differences and then realigned to the mean image of the first fMRI scan using the

SPM12 fMRI toolbox (Wellcome Department of Imaging Neuroscience, University College

London, UK). fMRI scans suffering from motion artifacts (i.e., voxel shifts > 0.05 mm detected

by realignment) and sudden physiological changes (i.e., abrupt changes in respiration pattern,

heart rate, and oxygen saturation level) were discarded. After resampling EPI and T2 images

to 128 × 128 × 17, the EPI images from each animal were first registered to their respective

T2W images. T2W images from each animal were then co-registered to a custom-made brain

template acquired from separate age-matched rats with identical T2 scanning parameters using

affine transformation and Gaussian smoothing to maximize normalized mutual information

(SPM12). The transformation matrix was then applied to co-register the fMRI EPI volumes.

Voxel-wise linear detrending with least-squares estimation was subsequently performed

temporally for the EPI volumes to eliminate any baseline drift caused by physiological noise

and/or system instability. For individual animals, data from the three fMRI sessions were

smoothed (full width at half maximum, FWHM = 1 pixel with no out-of-plane smoothing), and

high pass filtered (128 s). Then, a general linear model (GLM) was applied to calculate the

response coefficient (β) maps for each stimulus. Student's t-test was performed to identify

activated voxels using the threshold $P < 0.001$ (t value > 3.1). BOLD signal profiles were then

extracted from regions of interest (ROIs) delineated from the rat brain atlas for further

evaluation. These steps performed at the individual animal level were done to ensure that visual

inspection and comparison of the data quality across animals could be conducted before group-

level analysis. At the group level, resliced, realigned and co-registered EPI images

corresponding to the same fMRI session and stimulation frequency were averaged across

animals. Similar smoothing, filtering, and GLM processes were then applied to generate group-

averaged BOLD activation maps with a threshold of $P < 0.001$. These thresholded activation

maps were further corrected for multiple comparisons (i.e., masked) with masks generated from

one-sample group level t-tests using nonparametric inference with threshold-free cluster

enhancement multiple comparison correction of family wise-error rate (TFCE-FWE, $P < 0.05$).

For the third fMRI session in experiments stimulating the AON and Pir, TFCE-FWE was not

applied in the masking of thresholded BOLD activation maps were only masked with the

group-level masks generated from one-sample group-level t-tests without TFCE-FWE corrections (no activations after t-tests and TFCE-FWE corrections, t > 3.1).BOLD signal profiles for each ROI were generated by averaging those extracted earlier from individual animals.

In the fMRI experiments upon 1 Hz stimulations of OB, AON and Pir, quantification of the strength of neural activity at downstream activation targets was performed separately for the three consecutive fMRI sessions in each animal. First, the cumulative area under the BOLD signal profile (AUC) for each atlas-defined ROI was computed. Here, only the area during the four 20-s-on period of optogenetic stimulation was considered. Given that the BOLD response onset in our experiments was ~3 s after the presentation of the stimulus, we systematically delayed our area under BOLD signal profile measurements by 3 s. Second, the computed AUCs were grouped into five networks such as the primary olfactory (OB, AON, Pir, tenia tecta, TT, entorhinal cortex, Ent and olfactory tubercle, Tu), limbic (cingulate, Cg, orbitofrontal cortex, OFC, insular, Ins and amygdala, Amg), striatal (nucleus accumbens, NAc and ventral caudate putamen, vCPu), hippocampal (ventral hippocampus, vHP) and sensorimotor (motor, MC, somatosensory, S1, auditory, AC and visual, V1, cortices). Third, BOLD activation strength for each defined network was then measured. Here, the AUC of each network was normalized to the AUC of all five networks. Lastly, the BOLD activation strengths for each network were statistically compared between the three different fMRI stimulation experiments targeting OB excitatory neurons, and OB afferents at AON and Pir, respectively (outlier detection with nonlinear regression & false discovery rate at 1 %, Tukey's multiple comparisons test). Note that direct comparisons between AUCs of each downstream target from the three separate optogenetic fMRI experiments were not considered as they are likely erroneous given the differences in absolute BOLD signal amplitudes. For example, BOLD signal amplitudes in Pir stimulation experiments were generally weaker than OB and AON stimulations. The variations here are likely caused by differences in ChR2 transfection volumes in OB excitatory neurons and OB afferents at AON/Pir, including the population size of neurons/axonal terminals stimulated. Note that the vast majority of OB neurons are interneurons, and not projection neurons to OB and AON[37].

Comparison of BOLD power between fMRI sessions (i.e., 1st vs. 2nd, 1st vs. 3rd fMRI session) upon 1 Hz stimulation was done using Dunnett's multiple comparisons test to evaluate olfactory adaptation under repetitive OB/AON/Pir stimulation. Normalized BOLD power was only shown for display (not all data points were shown, because some values were extremely

huge) with statistical results calculated based on true BOLD power comparison (**Fig. S5**).

When comparing the BOLD power between healthy vs. aged rats upon first 1 Hz OB

stimulation, we conducted an uncorrected Fisher's LSD test (**Fig. 7B**).

Spectral dynamic causal modeling (sDCM), and statistical analyses. To parametrize the

downstream effects of OB-, AON- and Pir-driven neural activities on selected targets in the

right brain hemisphere (i.e., same hemispheric side to the optogenetically-stimulated region)

in healthy and aged animals, we further analyzed the BOLD fMRI data upon 1 Hz stimulation

using sDCM[26]. A six-node brain network model was defined, encompassing connections

between regions in the primary olfactory (OB, AON and Pir), striatal (vCPu) and limbic (Amg)

networks (**Fig. 4A**). Specifically, sDCM was utilized to computationally model the effective

connectivity strength of these downstream connections in OB, AON and Pir stimulation of

healthy animals, and OB stimulation data of aged animals. To avoid over-fitting of the data,

the BOLD signals were band-pass filtered with low cutoff frequency at 1/128 Hz and a range

of high cutoff frequencies at 0.1-0.3 Hz. We then evaluated the connectivity parameters from

the choice of 21 high cutoff frequencies with a step size of 0.01 Hz. We selected the optimal

high cutoff frequency for each stimulation target (i.e., OB, AON, Pir) that maximized the

weighted sum of scores for all modeled connections (significant, $P < 0.05$, score = 1; close to

significant, $0.05 < P < 0.1$, score = 0.5; FDR corrected). The optimal high cutoff frequencies

were 0.1, 0.18 and 0.14 Hz for OB, AON and Pir stimulations, respectively, in healthy animals,

and 0.1 Hz for OB stimulation in aged animals. Stochastic sDCM has been demonstrated to

perform better compared to deterministic modeling[26]. Here, we utilized Bayesian model

selection to determine the autoregressive model order (e.g., autoregressive models of order one,

AR1; order two, AR2), which was used to parameterize neural fluctuations and observed noise

in stochastic models. We then applied Bayesian model averaging (BMA) to generate the mean

effective connectivity for each defined connection since we did not obtain a clear winning

autoregressive model order. Following BMA, the mean effective connectivity was statistically

compared across animals and summarized in the form of connectivity matrices (**Fig. 4B**, n =

11, n = 11 and n = 9 for OB, AON and Pir stimulation in healthy animals; n = 13 for OB

stimulation in aged animals; one-sample t-test with FDR correction). We further quantified the

differences in the effective connectivity of AON-related vs. Pir-related connections to

downstream targets such as Ent, vCPu and Amg by conducting paired t-test with FDR

correction (**Fig. 4C**). Unpaired t-test was used to compare the effective connectivity of AON-

related and Pir-related connections between healthy and aged animals (**Fig. 7D**).

Multi-site local field potential (LFP) recordings and analyses. Electrophysiological recordings were performed under the same anesthesia protocols and similar physiological conditions as in the fMRI experiments with body temperature maintained using a heating pad. Guided by the fMRI findings, craniotomies were conducted and dura matter was removed with reference to the ROIs of ipsilateral OB (+7.5 mm anterior to Bregma, +1.8 mm ML), AON (+5.2 mm anterior to Bregma, +1.8 mm ML), Pir (+4.2 mm anterior to Bregma, +3.1 mm ML), vCPu (+2 mm anterior to Bregma, +2.7 mm ML), Amg (-3 mm posterior to Bregma, +3.8 mm ML)/ V1 (-6 mm posterior to Bregma, +4.2 mm ML), vHP (-6.4 mm posterior to Bregma, +6.4 mm ML) / dHP (-3.6 mm posterior to Bregma, +2.4 mm ML). For coverage of the majority of layers in the dentate gyrus of ipsilateral vHP/dHP, a linear microelectrode silicon array (16 recording channels equally spaced at 100 μm; 1.5 MΩ impedance; NeuroNexus Technologies, Ann Arbor, MI) was inserted perpendicularly to the brain surface with the tip of the linear array approximately 4.5 mm/3.7 mm below the dura matter via micromanipulators (Narishige, Amityville, NY). For other regions, single tungsten microelectrodes with 2.0 MΩ and 10 μm tip diameter were utilized (OB: -1.8 mm DV; AON: -4.3 mm DV; Pir: -5.4 mm DV; vCPu: -5.4 mm DV; V1: -1.1 mm DV; Amg: -8.4 mm DV). Fiber and a single channel electrode were constructed into an optrode[97] and were implanted following identical procedures as described for optogenetic fMRI experiments. LFP data were acquired using a multi-channel neurophysiology recording system (sampling frequency: 24 kHz; notch-filter: 50 Hz, 100 Hz, and 150 Hz; Tucker Davis Technologies/TDT, Alachua, FL). Synchronized laser stimulation was controlled by the same system, and the trigger signal for blue light pulse trains was recorded simultaneously with the extracellular recording data. Experimental paradigms were identical to those used in fMRI experiments.

After recordings, raw data were bandpass-filtered (0.01-150 Hz) and down-sampled to 1 kHz using MATLAB. Individual LFP and group-averaged LFP traces were generated. Averaged LFP traces were presented with standard error mean to reflect the reproducibility across animals as shown by our previous analyses and literature[28,29,45,98-100]. The latency of evoked LFPs was measured from the onset of light pulse to the first appreciable peak location (i.e., either negative or positive). To quantify the optogenetically evoked neural activity power during the 20-s optogenetic stimulation block, the area under absolute LFP traces was measured as LFP power during the optogenetic stimulation. Given that the LFP response onset was a few milliseconds after the presentation of the stimulus, no time delay for area under the LFP power measurements (**Fig. 6**).

Histology, immunohistochemistry, and confocal imaging. To confirm the specific expression of ChR2-mCherry in the OB excitatory neurons and the axonal inputs to AON/Pir, histology was performed according to the previously published procedure[28-31,45,95]. Upon completion of experiments, a selected number of optogenetically transfected animals (n = 3 for OB; n = 2 for AON/Pir) were anesthetized with pentobarbital and then transcardially perfused with ice-cold 4% paraformaldehyde (PFA) in phosphate-buffered saline (PBS). The brains were equilibrated in 20% sucrose in PBS at 4 °C overnight. Axial sections (40 μm) were prepared on a freezing microtome (model 860, AO Scientific Instruments). Consecutive sections were mounted and examined with a laser confocal microscope (Carl Zeiss LSM780/LSM980). For immunohistochemistry, free-floating sections were processed with 5% normal goat serum and 0.3% Triton X-100 in PBS with primary antibodies against rabbit polyclonal to CaMKIIα (1:400; Abcam) at 4 °C for 24 h. After washing with PBS, sections were then incubated for 2 h at room temperature with secondary antibodies Alexa Fluor 647 conjugate goat anti-rabbit IgG and Alexa Fluor 488 conjugate goat anti-guinea pig IgG (both 1:500; Molecular Probe). Slices were then washed and mounted using FluoroShield mounting medium with DAPI (Abcam). Double or triple immunofluorescence was assessed with a laser confocal microscope (Carl Zeiss LSM780/LSM980).

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Author Contributions

Study design and conceptualization: T.M., X.W., A.T.L.L. and E.X.W.. Animal fMRI experiments: T.M.. Animal fMRI data analysis: T.M., X.W., A.T.L.L. and X.L.. Electrophysiology experiments: T.M. and J.W.. Electrophysiology data analysis: T.M., X.W. and A.T.L.L.. Visualization: T.M., A.T.L.L., X.W., X.L. and L.X.. Suggestions: X.W., A.T.L.L. and L.X.. Supervision: A.T.L.L., E.X.W., X.W. and P.C.. Original draft writing: T.M. and A.T.L.L.. Writing-review and editing: A.T.L.L., E.X.W., T.M. and X.W..

Competing Interests

The authors declare no competing interests.

Additional Information

Supplementary information accompanies this paper.

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Figure Legends

A **Immunohistochemistry of CaMKIIα::ChR2(H1234R) Viral Expression in OB, AON & Pir**

B **Illustration of Optogenetic fMRI Stimulation Setup and Paradigm**

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Figure 1. Histological characterization of ChR2::CaMKIIα viral expression in the
olfactory bulb (OB) excitatory neurons and OB afferents at the anterior olfactory nucleus
(AON) and the piriform cortex (Pir), and optogenetic fMRI stimulation setup. **(A)**
Confocal images, 10x magnification (Left) and 40x magnification (Right), of ChR2-mCherry
expression in histological slices covering OB, AON and Pir. Overlay of images co-stained for
the nuclear marker DAPI and mCherry in OB (Top) revealed ChR2 expression in the soma of
the mitral cell layer & external plexiform layer in OB (indicated by green arrows). Meanwhile,

ChR2 expressions were found in OB afferents at AON and Pir (Middle & Bottom), indicated

by no colocalization between mCherry and DAPI. **(B)** Illustration of three optogenetic

stimulation targets, namely the OB excitatory neurons, and OB afferents in AON and Pir (Left),

which were conducted in separate animal groups. T2-weighted anatomical MRI image showing

the location of the implanted optical fiber (asterisk: stimulation target). Optogenetic fMRI

block-designed stimulation paradigm (Right). Five different frequencies (1, 5, 10, 20, and 40

Hz) were presented in a pseudorandomized order within each session. We repeated the

stimulations for a total of three sessions in each animal fMRI experiment. All frequencies were

presented at a 30% duty cycle except for 1 Hz, which was presented at a 10% duty cycle.

Figure 2. Optogenetic excitation of OB excitatory neurons and OB afferents at AON and Pir recruit distinct brain-wide long-range olfactory networks. Averaged BOLD activation

maps and signal profiles (Left) upon 1 Hz stimulation of (**A**) OB excitatory neurons, (**B**) OB afferents in AON, and (**C**) OB afferents in Pir during the first fMRI session (n = 11 for fMRI experiments in A & B; n = 9 for C; t > 3.1 corresponding to $P < 0.001$; error bars indicate $\pm$ SEM), and the corresponding area under the BOLD signal profiles (AUC) for each atlas-defined region-of-interests (ROIs; Right). Note that the BOLD activation maps displayed in A-C were further corrected for multiple comparisons with threshold-free cluster enhancement with family-wise error rate (TFCE-FWE) at $P < 0.05$. (**D**) Summary of BOLD activation strength to quantify the extent of various brain networks engaged upon respective optogenetic stimulation of the three different primary olfactory stimulation targets. BOLD activation strength was computed as the ratio of AUC for each ROI over AUC of all ROIs (individual animal data points are shown; error bars indicate $\pm$ SEM; Tukey's multiple comparisons test; **$P < 0.01$, and *$P < 0.05$). OB-driven neural activities mainly activated the primary olfactory network (AON, Pir, TT, Ent and Tu). AON-driven neural activities were found to preferentially recruit the striatal (NAc and vCPu) and hippocampal networks (vHP), while Pir-driven activities strongly targeted the limbic network (Cg, PrL, OFC, Ins and Amg).

Abbreviations. Regions in primary olfactory network: olfactory bulb (OB), anterior olfactory nucleus (AON), piriform cortex (Pir), tenia tecta (TT), entorhinal cortex (Ent), olfactory tubercle (Tu); limbic network: cingulate (Cg), prelimbic cortex (PrL), orbitofrontal cortex (OFC), insular (Ins), amygdala (Amg); hippocampal network: ventral hippocampus (vHP); striatal network: nucleus accumbens (NAc), ventral caudate putamen (vCPu); sensorimotor network: motor (MC), somatosensory (S1), auditory (AC), visual (V1) cortex.

Figure 3. Repetitive optogenetic excitation of OB excitatory neurons and OB afferents at AON and Pir reveal varied neural activity adaptation properties in olfactory networks. Averaged BOLD activation maps (Top) upon 1 Hz stimulation of (**A**) OB excitatory neurons,

(**B**) OB afferents in AON, and (**C**) OB afferents in Pir during the second (n = 11 for fMRI

experiments in A & B; n = 9 for C; t > 3.1 corresponding to *P* < 0.001) and third (n = 8 for

fMRI experiments in A; n = 9 for B; n = 6 for C; t > 3.1 corresponding to *P* < 0.001) fMRI

sessions, respectively. We found a dramatic decrease or even absent bilateral activations in

long-range olfactory networks upon stimulation of OB excitatory neurons or OB afferents at

AON, but not Pir, indicating strong neural activity adaptation mediated by AON-driven neural

activity to downstream targets. Note that similar to Fig. 2, the BOLD activation maps displayed

in A-C were further corrected for multiple comparisons with threshold-free cluster

enhancement with family-wise error rate (TFCE-FWE) at *P* < 0.05, except for that the maps

during the third session in B and C were generated directly after one-sample t-test. The extent

of neural activity adaptation in brain-wide olfactory networks was quantified with the true

BOLD power (AUC of the BOLD signal profiles) for each atlas-defined ROI (Bottom;

individual animal data points are shown; error bars indicate ± SEM; Dunnett's multiple

comparisons test of BOLD power during 2nd vs. 1st fMRI session, 3rd vs. 1st session; ***P <

0.001; **P < 0.01; *P < 0.05). Note that the AUC for each ROI was normalized to their

respective counterpart in the first fMRI session for display.

Figure 4. Inhibitory effect of AON outputs and excitatory effect of Pir outputs to downstream targets in long-range olfactory networks. (**A**) Graphical representation of matrix A which defines the a priori anatomical connections of the olfactory network model utilized in dynamic causal modeling (DCM). (**B**) Effective connectivity matrices of the olfactory network model upon stimulation of OB excitatory neurons and their afferents at AON and Pir. (**C**) Quantitative summary of the significant differences in connectivity strength of downstream connections identified from the statistical comparison of matrices in B (Left; n = 11 for OB and AON stimulations; n = 9 for Pir stimulation; individual animal data points are shown; error bars indicate ± SEM; paired t-test with FDR correction; **$P < 0.01$, and *$P < 0.05$) and the corresponding graphical representation of inhibitory effect of AON and excitatory effect of Pir outputs to downstream targets (Right). We identified robust AON negative vs. Pir positive connectivity to downstream targets such as Ent, vCPu and Amg, indicating distinct inhibitory vs. excitatory effect of the two primary olfactory cortical outputs that can shape the dynamic properties (i.e., BOLD activations across repeated stimulation sessions) of long-range olfactory networks.

A — Evoked Neural Activity & Response Latency upon 1 Hz OB Stimulation

B — Evoked Neural Activity & Response Latency upon 1 Hz AON Stimulation

C — Evoked Neural Activity & Response Latency upon 1 Hz Pir Stimulation

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1070 **Figure 5. Local field potential (LFP) recordings confirm neuronal activity underlying**

1071 **BOLD activations and reveal orthodromic neural activity propagation to downstream**

targets in primary olfactory, striatal, visual, limbic and hippocampal networks. Illustration of electrophysiological recording regions (Top-Left), averaged evoked LFPs from the seven recorded regions (Bottom-Left & Middle), and summary of orthodromic neural activity propagation latencies measured from the stimulation onset over mono- and poly-synaptic projections (Right) upon 1 Hz optogenetic stimulation of (**A**) OB excitatory neurons, (**B**) OB afferents at AON and (**C**) OB afferents at Pir (n = 7, n = 7 and n = 5, respectively; 10% duty cycle; error bars indicate ± SEM; crosshairs denote LFP peaks that represent initial neuronal population activity for latency measurements). Significant delay between AON/Pir and OB neuronal population activity with appreciable jitter showed that neural activity propagation through the excitations of OB axonal terminals at AON or Pir, respectively, were largely orthodromic in our experiment.

Figure 6. LFP recordings confirm varied neural activity adaptation in olfactory networks upon repetitive optogenetic excitations of OB excitatory neurons and OB afferents at AON and Pir. LFP power of evoked neural activities upon 1 Hz optogenetic stimulation of (**A**) OB excitatory neurons (n = 7; 10% duty cycle; error bars indicate ± SEM), (**B**) OB afferents at AON (n = 7; 10% duty cycle; error bars indicate ± SEM) and (**C**) OB afferents at Pir (n = 5; 10% duty cycle; error bars indicate ± SEM) across three electrophysiological recordings sessions. (**D**) Summary of the area under LFP power to quantify the extent of neural activity adaptation in the recorded regions of olfactory networks (individual animal data points are

shown). In accord with our fMRI findings, the strength of neural activities, which were

quantified through LFP power analysis here (i.e., area under absolute LFP trace normalized to

30-s baseline LFP power), was also decreased under repetitive 1 Hz stimulation of OB and OB

afferents at AON but was relatively unchanged under Pir stimulation (1st vs 2nd and 3rd

recording sessions). The LFP recordings demonstrate strong neural activity adaptation

mediated by AON-driven neural activities to downstream targets.

Figure 7. Decreased BOLD activations in primary olfactory and limbic networks, and impaired AON to Pir connectivity in aged animals. (A) Averaged BOLD activation maps and signal profiles from the 1st fMRI session upon 1 Hz stimulation of OB excitatory neurons in D-galactose aged rat model (n = 13; t > 3.1 corresponding to $P < 0.001$; error bars indicate ± SEM). Note that the BOLD activation maps displayed in A were further corrected for multiple comparisons with TFCE-FWE at $P < 0.05$. **(B)** Comparison of the corresponding area under the BOLD signal profiles for each atlas-defined region-of-interests between healthy and aged animals (individual animal data points are shown; Fisher's LSD test; **$P < 0.01$, and *$P < 0.05$; error bars indicate ± SEM). **(C)** Effective connectivity matrices of the olfactory network model upon stimulation of OB excitatory neurons in aged brains. **(D)** Quantitative summary of

the significant differences in connectivity strength of downstream connections identified from

the statistical comparison of effective connectivity matrices between healthy and aged brains

(individual animal data points are shown; unpaired t-test; *$P < 0.05$; error bars indicate $\pm$ SEM).

We uncovered a switch in the polarity from positive to negative in the connectivity of AON

neural activity outputs to Pir in aged animals, indicating the dysfunction of a key primary

olfactory cortical circuit that likely caused the significantly decreased BOLD activations in the

primary olfactory and limbic networks.

A Immunohistochemistry of CaMKIIα::ChR2(H1234R) Viral Expression in OB, AON & Pir

B Illustration of Optogenetic fMRI Stimulation Setup and Paradigm

A **1 Hz Stimulation of Excitatory Neurons at the Olfactory Bulb (OB)**

B **1 Hz Stimulation of OB Afferents at the Anterior Olfactory Nucleus (AON)**

C **1 Hz Stimulation of OB Afferents at the Piriform Cortex (Pir)**

D **BOLD Activation Strength Across Various Networks upon Stimulation of Distinct Olfactory Targets**

A BOLD Activations for 2nd & 3rd fMRI Sessions upon 1 Hz Stimulation of OB Excitatory Neurons

B BOLD Activations for 2nd & 3rd fMRI Sessions upon 1 Hz Stimulation of OB Afferents at AON

C BOLD Activations for 2nd & 3rd fMRI Sessions upon 1 Hz Stimulation of OB Afferents at Pir

A Connection Matrix

Source

B Effective Connectivity Matrices

OB Stimulation   AON Stimulation   Pir Stimulation

Effective Connectivity Strength (Hz)

-0.5   0   0.5

Significant (P < 0.05)   Close to Significant (0.05 < P < 0.10)

C Quantitative Comparison of Connectivity Strength between AON- vs. Pir-Related Connections

Schematic Illustration of Significant AON Inhibitory & Pir Excitatory Outputs to Downstream Olfactory Network Targets

A **Evoked Neural Activity & Response Latency upon 1 Hz OB Stimulation**

B **Evoked Neural Activity & Response Latency upon 1 Hz AON Stimulation**

C **Evoked Neural Activity & Response Latency upon 1 Hz Pir Stimulation**

A **LFP Power upon Repetitive 1 Hz Stimulation at OB Excitatory Neurons**

B **LFP Power upon Repetitive 1 Hz Stimulation of OB Afferents at AON**

C **LFP Power upon Repetitive 1 Hz Stimulation of OB Afferents at Pir**

D **Underlying Neural Activity Dynamics upon Repetitive 1 Hz Stimulation**

A

1 Hz Stimulation of OB Excitatory Neurons in Aged Animals

Bregma +7 mm   +5 mm   +4 mm   +2 mm   0 mm   -2 mm   -4 mm   -6 mm

t   3.1   15

B

Comparison of BOLD Activations in Healthy vs. Aged Animals upon 1 Hz OB Stimulation

Primary Olfactory Network   Limbic Network   vHP   Striatal Network   Sensorimotor Network

■ Healthy ■ Aged

C

Effective Connectivity Matrix in Aged Animals

Source

Target

□ Significant (P < 0.05)   □ Close to Significant (0.05 < P < 0.10)

D

Comparison of Connectivity Strength in Healthy vs. Aged Animals

■ Healthy ■ Aged

P values shown are uncorrected

AON-Pir   AON-vCPu   Ent-OB   Amg-Pir