1 A prefrontal motor circuit initiates persistent movement

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9 Abstract

Persistence ensures continuity in a course of action. It is crucial to human success and animal survival.
However, how the brain applies persistence to movements is unknown. Here we demonstrate that
persistence arises during the initial phase of movement and is maintained until terminal signaling. Using
single-unit extracellular recordings in awake mice, we found that neural encoding of persistent
movement phases is independent of the valuation (i.e., valence) of the external stimuli. Next, using
opto-tagging, we identified a group of dorsal medial prefrontal cortex (dmPFC) motor cortex projecting
(MP) neurons, which encodes the initial phase of a persistent movement rather than valence.
Inactivation of dmPFC MP neurons impairs the initiation of persistence and reduces neuronal activity in
the insular and motor cortex. Finally, a computational model suggests that a successive sensory stimulus
acts as a triggering signal for the initiation of persistent movements. These results reveal a neural
mechanism that switches the brain state from neutral to persistent during movement.

21 Introduction

Persistence is a fundamental property that can help humans succeed in many areas, such as athletic and
social competition [1, 2]. It has a long-lasting effect on the maintenance of motivation [3]. It also contributes
to the animal's survival, such as dominating in social hierarchy and escaping from predators [4]. It can
drive actions successively from seconds to minutes and even outlast the eliciting stimulus. Socially
dominant animals exhibit effortful behaviors toward their rivals [5]. Repeated visual threat stimuli lead a
persistent state in *Drosophila* [6]. Although persistence is influential in both humans and animals, there
are few studies on how the brain applies it to movements. Moreover, the psychological definition of
persistence is not appropriate to solve certain behavioral problems, including how long it should last and
how to quantify its continuity, as it can vary from behavior to behavior. To examine persistence, we
focus on a motivated movement [7] that can persist for a certain period of time and stabilize at a range of
high frequency. This type of motivated movement is triggered by internal states from deep brain regions
and is regulated by cortex neurons [7, 8, 9]. It is also accompanied with extensive valence coding in the
brain [10, 11, 12, 13]. Here, we attempt to identify the neural coding in the different phases of motivated,
persistent movement and to test whether it is the same with the coding of valence.

36 Quantification of persistent movements

First, we defined a persistent movement as the continuous repetition of a single movement (e.g., a cycle
of tongue or limb movements) and the maintenance of that continuity for at least 5 seconds. To impose
a persistent movement on mice, they were deprived of water for 16 to 36 hours until their body weight
decreased by about 22% (Subject details, **Methods**). The mice were then head-fixed to a custom made
set-up and trained to lick various types of liquid after delivery onset (DO) (**Fig. 1a**, **Supplementary Fig.**
1b, and Behavioral details, **Methods**), but received no other artificial stimuli. Licking signals, facial and
locomotor activities were measured. After training, we observed that the mice showed licking
movements sustained for approximately 15-30 seconds during water delivery (**Supplementary Fig. 1c-f**).
Consistent with the standard pattern of affective dynamics performed previously [14], licking frequency
was maximized during the initial phase (**Supplementary Fig. 1c-f**, right column, peaks of licking
frequency are indicated by black arrows) and then stabilized at 6 to 7 Hz until the end of delivery
(**Supplementary Fig. 1d, e**) or when the liquid was switched to quinine (5mM, **Supplementary Fig. 1c, f**).
Higher hedonic stimuli, 20% sucrose, did not increase this frequency (**Supplementary Fig. 1e, f**). As an
aversive stimulus, quinine administration was more likely than the interruption of water or sucrose
administration to cause termination of persistent licking (p<0.05 with respect to termination bias,
Supplementary Fig. 1i). In addition, we found no significant difference between male and female mice in
the initiation and termination of persistent licking movement (**Supplementary Fig. 1r, s**). Therefore,
quinine was used to terminate the persistent licking movement and to evaluate negative valence; by
contrast, water or sucrose was used to trigger licking movement and to evaluate positive valence.

Next, we determined the temporal window for the study of valence and movement phases. For valence,
the assessment of whether water or quinine is hedonic or aversive should be based on the single
contact (approximately 180ms around the licking onset (LO), **Supplementary Fig. 1c-f**) to them. For
movement phase, we divided it into an initial and a terminal phase. To our observation, the maximum
licking frequency was confined within the initial phase and a sharp decrease of licking frequency was in
the terminal phase (**Supplementary Fig. 1c-f**), so we suspected that there was additional neural coding
to increase and decrease licking frequency besides the coding of licking command. Based on the period
of peak licking frequency (approximately 3s, as indicated by the arrow, **Supplementary Fig. 1c**) in the
initial phase and the sharp drop of licking frequency (approximately 3s from 6-7 Hz to less than 1 Hz, as
indicated by the arrow, **Supplementary Fig. 1c**) in the terminal phase, we designed a time window of 5-
second long (including 2s baseline before the licking onset) as a temporal window for data analysis.
Together, we considered the neuronal firings of the valence and movement phases were confined
within a small (S, LO-100ms to LO+80ms) and a large (L, 1st LO-2s to 1st LO+3s) scale window,
respectively (**Fig. 1c**).

Dissociable neural coding of movement phases and valence

To characterize the single-units that statistically represent valence and movement phases, we used *in*
vivo silicon probe to collect neuronal activity data from all three brain regions, including the insular
cortex (IC), which is known to encode valence in the gustatory system [11, 15], the primary motor cortex
(M1), which represents the motor commands of voluntary movement [16], and the dorsal medial
prefrontal cortex (dmPFC), which has been shown to connect these two brain regions [17]. The single-units
that were generated from the neural activity data were then classified into different groups of neural
representations according to the extent to which they can discriminate liquid types (for valence) or
movement phases (**Methods**).

We then asked whether the neuronal coding of movement phases and valence are associable or not by
examining the proportion of neural representations of valence and movement phases in three brain
regions (IC, M1, and dmPFC). As population, we found that 27% of single-units represented movement
phases (initial or terminal phase, ≥65% in movement phase correlation) but showed weak or no taste
tuning (**Fig. 1d-f**). 14% of single-units represented valence (positive or negative valence (PV or NV),
z>1.64 in taste correlation) but exhibited poor specification on movement phases (**Fig. 1d-f**). By contrast,
only small fraction (8%) of single units displayed the preference to both valence and movement phases
(**Fig. 1e, f**), though this number varied trivially from region to region (**Supplementary Fig. 4**).

To test whether the neural networks representing movement phases and valence interact with each
other, we examined the connectivity between the neurons solely tuned to initial phase and PV using
Total Spiking Probability Edges (TSPE) [18] and compared it with shuffled connectivity. Our results showed
that there is no overall excitatory impact from the initial phase to PV or from PV to initial phase tuned
neurons (mean of real TSPE < 99 percentile of shuffled TSPE, **Supplementary Fig. 5d**). Note that
connectivity between the terminal phase and NV tuned neurons was not available because their spike
data were unable to construct a cross-correlation in a 50ms-time window, suggesting that the coding of
terminal phase and NV are not connected. Together, these results suggest that the movement phases
and valence are encoded separately.

Coding of the initial phase in dmPFC MP neurons

Next, we investigated the neural basis representing the initial phase of persistent movement. We first
compared the fraction of clustered neural representations in the IC, M1, and dmPFC. Second, we
examined decoding performance in these brain regions by training a linear discriminant decoder on
firing rate data. Using cell-specific recordings enabled by the opto-tagging approach [19] (**Fig. 2a-c**), we
found that motor cortex projecting (MP) neurons [17] in the dmPFC exhibited the best representation of
the initial phase among the three mapped brain regions (**Fig. 2, Supplementary Fig. 4, 6,** and **7**).
Specifically, the neuronal clustering results showed that 32% of dmPFC MP neurons exhibited a degree
(≥65% shuffled activity at initial phase & ≤35% shuffled activity at terminal phase) of initial phase
representation (**Fig. 2h**), whereas only 3% of them exhibited a degree (≥65% shuffled activity at terminal
phase & ≤35% shuffled activity at initial phase) of terminal phase representation (**Fig. 2h**). No more than
20% of dmPFC MP neurons showed a degree (z>1.64 in valence correlations compared to shuffled
activity) of valence representation (PV+NV, **Fig. 2e**). The decoding performance results showed that
dmPFC MP neurons had a low ability to discriminate positive and negative valence (p<0.0001 lower than
shuffled cumulative decoding accuracy, **Fig. 2k left**), while had a high representation of movement
phases (p<0.0001 higher than IC and shuffled cumulative decoding accuracy, **Fig. 2k right**). To confirm
the discriminability of dmPFC MP neuron between valence and movement phases, we first embedded
neuronal population activity in the S- and L-window of dmPFC MP neurons into trajectories using
principal component analysis (PCA) and then measured the mean Euclidean distances in all PC
dimensions. The results showed that the discrimination of neural activity was significant higher in the L-
window than it in the S-window (p<0.05, **Fig. 2j**), indicating that dmPFC MP neurons discriminate better
between movement phases than between valence values.

Bodily arousal is usually accompanied by valence changes [4]. It is, therefore, necessary to investigate
whether or not dmPFC MP neuron also encode bodily arousal. In rodents, facial activity can reflect

bodily arousal [20]. In the persistent licking task, the facial activity showed a reliable increase immediately
after water DO (p<0.05 compared to baseline in all sessions, **Supplementary Fig. 1k-n**). Thus, we used
facial activity to assess bodily arousal in the initial phase. To investigate whether the firing changes of
dmPFC MP neuron in the initial phase were due to bodily arousal, we trained a Hammerstein-Wiener
model and tested its prediction accuracy using facial activity data. Although some dmPFC neurons
exhibited a degree of facial activity representation in the initial phase (p<0.05 initial phase (median=-
4.5671; 21/194 neurons with accuracy>30%) vs terminal phase (median=-36.9341; no neuron with
accuracy>30%), **Supplementary Fig. 8d2**), all dmPFC MP neurons showed poor prediction performance
at both initial (median=-14.5639; no neuron with accuracy>30%) and terminal phase (median=-27.4588;
no neuron with accuracy>30%) on facial activity and no significant difference between these two phases
(p>0.05, **Fig. 2l, m**). These results excluded the possible coding of bodily arousal in dmPFC MP neurons.

Effects of dmPFC MP neuron silencing on persistent movement

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To test the role of dmPFC MP neuron on persistent licking movement in different phases, we examined
the initiation and termination bias, as well as licking frequency (**Methods**). dmPFC MP neurons were
optogenetically manipulated by expressing stGtACR2 [21] and shining laser during the different phases of
the persistent licking task (**Fig. 3a**). Our results showed that optogenetic silencing of dmPFC MP neurons
impaired the initiation of licking (p<0.001 compared to the sham trials, **Fig. 3b, c**), but had no effect on
the termination of lick (p>0.05 compared to the sham trials, **Fig. 3d, e**) or the licking frequency in the
middle phase (**Supplementary Fig. 10b2**) in thirsty mice, suggesting that dmPFC MP neurons are not
involved in motor control. The similarity of thirst level was confirmed by comparing body weight loss in
sham and laser trials (**Supplementary Fig. 10b4**, **c3**, and **d3**). To further confirm the function of dmPFC
MP neuron in initiating persistent licking movement, dmPFC MP neurons, expressing hm4D(Gi), were
chemogenetically silenced by administrating with CNO in mice and the licking chances were tested
during the persistent licking task. As expected, thirsty mice with chemogenetically silenced dmPFC MP
neuron had lower chance of drinking water (p<0.05 compared to the saline trials, **Supplementary Fig.
9c**). To investigate whether the dampened licking initiation was due to a decrease in bodily arousal, we
tested facial and locomotor activity with or without optogenetic silencing. The mice in the sham and
laser trials displayed the similar levels of facial and locomotor activity in all initial, middle, and terminal
phase (**Fig. 3g** and **Supplementary Fig. 10b-d**), suggesting that the attenuated licking initiation was not
due to the decrease of bodily arousal.

We next asked whether the effect of dmPFC MP neuron silencing was specific to the initiation of
persistent licking or also general to other types of initiation of persistent behavior. We took advantage
that mice showed a phase of persistent running after a mild electric shock (**Supplementary Fig. 9h**). To
test whether inactivation of dmPFC MP neurons also affected this behavior, we examined body activity
after a 1s electrical tail shock. Indeed, chemogenetic silencing of dmPFC MP neurons suppressed
escaping behavior (**Supplementary Fig. 9h, i**), suggesting that dmPFC MP neurons are generally involved
in the initiation of persistent movements. This result is consistent with previously reported general
population of dmPFC neurons, silencing of which has been shown to delay the initiation of avoidance
movements [9]. Together with the evidence that dmPFC MP neurons are required for the persistent licking
movement, this also supports the theory of mixed selectivity in the prefrontal cortex [22].

Next, we hypothesized that the increase in the brain state of positive valence and tongue movement in
the initial phase was due to the coding of movement phases. IC is thought to encode taste valence [11, 15]
(38/42 in w-q vs 11/42 in w-s vs 33/42 in w-w with mean decoding accuracy higher than chance,
Supplementary Fig. 6a), whereas M1 was involved in voluntary control of tongue movement during
licking [23, 24] (12% neurons with z>1.64 in lick correlations compared with shuffled activity,
Supplementary Fig. 4b1). Therefore, the activities in IC and M1 should be affected by the optogenetic
silencing of dmPFC MP neurons. To test this hypothesis, we measured the neural activity in these two
brain regions with or without shining laser on dmPFC. As expected, the neuronal activities of M1 and IC
were decreased (p<0.05) after dmPFC MP neurons were optogenetically silenced (**Fig. 4f, g**). We further
confirmed this initial phase specificity by excluding the effect of silencing of dmPFC MP neuron on
valence at middle phase (no significant difference of sucrose-licking frequency between sham and laser
trials, **Supplementary Fig. 10a**). Overall, our results suggest that the dmPFC MP neuron is required to
initiate continuous licking, and further promoting taste valence during the initial phase, but loses its
necessity upon the persistent movement already initiated.

A MP network-based computational model

Finally, we asked what causes the MP network to initiate a persistent movement. To answer this
question, we built a neural network-based model (**Fig. 5a**) and examined how the output of licking
performance changes in response to different types of inputs. The design of this model was mainly
based on two criteria: (1) the inter-spike interval of a single neuron in the model is matched to the MP
neuron in mPFC (**Supplementary Fig. 11a**); (2) the neuronal population of modeled network performs
rotational dynamics [25] because we assumed that tongue movement follows a rhythmic pattern
(**Supplementary Fig. 11b, c**). To verify if the output of this model matched the performance of thirsty
mice in the actual experiment, we manipulated the firing rate of the simulated network by inserting the
inputs with different amplitudes and examined the output licking frequency and initiation bias. We
found no linear relationship between the mean neuronal firing rate and above two parameters
(**Supplementary Fig. 11d**, red), which is consistent with the experimental observations (**Supplementary
Fig. 11d**, black). To further ensure the feasibility of the MP network-based model, we simulated the
optogenetic silencing of dmPFC MP neurons through decreasing the number of all neurons in the
modeled network. Consistent with the experimental data, the network with reduced neural population
inhibited the initiation of persistent lick (**Fig. 5b**). We then examined the effect of temporally continuous
input on the output of the network. We found that even a single short-term interruption (200ms)
disrupted the initiation of continuous lick (average 57% decrease on initiation bias, **Fig. 5c**). The
percentage of bias with two interruptions decreased to 5.7±0.08% (compared with 21±1.17% with no
interruption, **Fig. 5c**). This temporally continuous input of the model suggests that the triggering signal
for the MP network should be intact, continuous sensory stimulation. For the persistent licking task, we
suspected this sensory stimulation may come from the continuous sensation of the liquid delivery.

Summary and Discussion

Our study identified a neural circuit, responsible for initiating persistent movements. After receiving a
sensory signal, dmPFC MP neurons can send the command signals to the primary motor cortex and
striatum, which in turn initiates the downstream machineries for a persistent movement (**Fig. 6**). Our
results excluded the possibility of the neural coding of valence, bodily arousal, and motor command by

dmPFC MP neurons. Our findings suggest that the engagement in persistent movement is not based
solely on a snap judgment of the current situation.

Summary of neural principles. Based on the membrane properties and network connectivity of MP
neurons, we summarized two neural principles that govern the initiation of persistent movement. First,
the circuits should consist of slow spiking neurons. mPFC layer V MP neurons are the main functional
units linking deep brain regions and motor cortex, and they fire more slowly than the pyramidal neurons
of layer II [17, 26]. In line with the observation of maximal licking frequency (about 8 Hz) during the initial
phase (**Supplementary Fig. 1c-f**), the firing rate of neurons modulating the tongue should not exceed
this value. Second, the circuit should be constructed with monotonic outputs and divergent inputs.
Within the mPFC, the projection of the MP network is relatively sparse, projecting only to the
contralateral dmPFC and motor regions, whereas the inputs to the MP network are originated from
different deep brain regions [17]. This connection configuration may help to ensure that the MP network
not only simultaneously receive the signals from the neurons that tuned multiple aspects of sensory
cues, but also regulate downstream motor output independent of other neural networks.

Formation of a generalized command. Although we have uncovered a neural basis for encoding the
initial phase of persistent movements, it is unclear how the generalized sensory command is formed or
how a specific movement is executed under a persistent pattern: First, how are the valuable aspects of
sensory cues recognized? This unsupervised learning process requires a generalizing neural network [27]
that extracts only a few optimal aspects of stimulus while maintaining specificity. The question is why
some of the aspects are optimal but others are not. We suspect that these optimized aspects are the
ones that are most similar or closest in time to the reward or punishment itself, as they can efficiently
and directly link to a specific behavioral pattern. Second, how can the sensory cue be associated with a
particular movement pattern? Based on the reciprocal link between mPFC and motor cortex [17], we
reason that a feedback loop between the generalization network and motor regions may be required.
Meanwhile, a valence input may be also necessary to construct the association and sensitize the
response to sensory cue.

Explanation of the dynamics of valence coding. To explain the gradually fading firings (**Supplementary**
Fig. 4d, e) of PV neural representations during water licks, we postulate two possibilities: (1)
Transformation of neural representations. Some neurons may initially encode PV but abolish this
encoding property in subsequent phases. Other neurons may not encode PV in the initial phase but
adopt this encoding later. This possibility can explain the persistent valence encoding in the IC
(**Supplementary Fig. 6a3**); (2) Valence decreases. Since the thirsty mice may become satiated as the lick
progresses, the value of water to the mice will decrease accordingly. Similarly, the gradually increasing
activity of the neural representations of NV (**Supplementary Fig. 4d, e**) could also be caused by the
transformation of their encoding properties or by an increasing negative valuation of water.

The function of MP neuron in the IC-MP-motor circuit. Based on the structure of the circuit, the MP
neuron acts as a node connecting IC and motor region [17]. Since IC encodes valence, it is possible that the
MP neuron integrates valence signal and sends it to motor region. However, our result showed that MP
neuron does not encode valence (**Fig. 2k**). Because some of the neurons in IC also encode movement
phases (**Fig. 2k** and **Supplementary Fig. 4a**), we speculate the function of this circuit is to transmit
movement phase signals, instead of valence, during persistent movement control.

Relationships to persistent spiking activity. Persistent spiking activity in prefrontal cortex has been
observed in both rodents and humans during the maintenance of working memory [28, 29]. However, only a
few neurons in the prefrontal cortex showed persistent activity (small fraction of neural representations
of lick, **Supplementary Fig. 4c**, and gradually decreasing firing of neural representations of PV and initial
phase, **Supplementary Fig. 4d, e**) during persistent licking. Rather than maintain the neural activity,
dmPFC MP neurons are responsible for triggering such activity in the motor cortex. Because this
triggering process also requires a successive, short-term (about 5s) external stimulus, we summarize the
function of the dmPFC MP network as converting a relatively short-term, external, persistent input into
a relatively long-term, internal, persistent activity.

METHODS

Subject details

All experimental procedures were approved by the Institutional Animal Care and Use Committee
(IACUC) and the Biosafety Committee of the University of Wyoming. 14 male and 6 female
immunocompetent mice of specified age (indicated in **Surgeries** and **Behavioral details**) were used for
various experimental purposes. All mice were bred on a C57/BL6J background. Mice older than 30 days
were housed with same-sex littermates or alone in a vivarium at 21-23 ^{o}C and a 12-h light/dark cycle.
Mice implanted with an electrode or head bar were housed alone. For chemo-genetic experiments, 2
male and 2 female mice received AAV injection and head bar implant. For optogenetic experiments, 6
male mice and 2 female mice received AAV injection, an opto-electrode, and head bar implantation. 3
male mice and 1 female mouse were implanted with electrodes and head bar. 1 male mouse and 1
female mouse were implanted with optical fibers and electrodes. Optical fibers were implanted in 2
male mice. For all behavioral experiments, mice were deprived of water for 16-36 hours before the
experimental day until they reached 75-80% of their free-feeding body weight. After each behavioral
experiment, mice were returned to their home cage where they had unlimited access to water for at
least five days until the next behavioral test day.

Surgeries

The preparatory procedures are similar for both implantation and injection. Mice were anesthetized
using oxygenated (2 LPM for induction and 0.4 LPM for maintenance) 2% isoflurane (v/v). Mice heads
were fixed to the stereotactic device (NARISHIGE SG -4N) and maintained at 37 ^{o}C with a heating pad
(K&H No. 1060). Seventy percent isopropyl alcohol and iodine were placed on the incision site. The skull
was exposed by cutting the skin and removing the dura and connective tissue. The coordinates used for
positioning the injection and implantation sites were relative to Bregma (antero-posterior A-P, medio-
lateral M-L, dorsal-ventral D-V) in mm. After the surgeries, the mice were administered intraperitoneal
ibuprofen (50 mg kg^{-1}), and they were kept at 37°C for 30-60 minutes before returning to the home
cage.

For viral injection, P14-30 mice were used. A small craniotomy (approximately 0.2 mm in diameter) was
made over the injection site. Glass filaments (Drummond Scientific Co.) with a tip diameter of
approximately 5 μm were filled with 2 μL of virus solution. By pressure injection with a custom-made
device driven by a single-axis hydraulic manipulator (NARISHIGE mmo-220A), the viral solution

(undiluted, 100 nL at each injection site) was delivered to the desired regions at a rate of 30-50 nL min^{-1}.
Opto-labeled dmPFC MP neurons were labeled with pAAV- CAG -hChR2-mcherry (Addgene_28017-
AAVrg) injected into motor cortex (A-P -0.6, M-L 1.0, D-V 0.2 0.5 0.8). For chemogenetic inhibition,
pAAV-Ef1a-mCherry- IRES -Cre (Addgene_55632-AAVrg) was injected into bilateral motor cortex (A-P -
0.6, M-L ±1.0, D-V 0.2 0.5 0.8), followed by injection of pAAV-hsyn-DIO-hm4D(Gi) (Addgene_44362-
AAV5) into the bilateral dmPFC (A-P 1.35, M-L ±0.2, D-V 0.2 0.5 0.8 at an angle of 30° to the upright
position). For optogenetic silencing, pAAV-CKIIa-stGtACR2-FusionRed (Addgene_105669-AAVrg) was
injected into the bilateral motor cortex (A-P -0.6, M-L ±1.0, D-V 0.2 0.5 0.8). Mice were returned to the
home cage until at least three weeks before implantation.

For implantation, silicon probes (A4x8-Edge-2mm-100-200-177-CM32 or A1x32-Edge-5mm-25-177-
CM32, NeuroNexus) or 32-tetrode bundles (Bio-Signal technologies) were implanted followed by optic
fibers and head bar. To build opto-electrode, optic fibers (MFC_200/245-0.37_2.0mm_MF1.25_FLT)
were fixed around 0.5 mm above the electrodes using crazy glue and dental cement (Lang Dental). In
opto-inhibition experiments, two optic fibers (MFC_600/710-0.37_1.0mm_MF1.25_FLT) were implanted
to the bilateral prefrontal cortices (A-P 1.7, M-L ±0.5, D-V 0.5). To fit the shape of prefrontal, motor, and
insular cortex, customized 32-tetrode bundles were split into one or two clusters. To record the single
unit, electrodes were implanted in the left hemisphere with the following designs and coordinates:
silicon probe (A1x32-Edge -5mm-25-177-CM32) was implanted at the pIC (A-P −0.5-(-1.5), M-L 3-4, D-V
3); silicon probe (A1x32-5mm-25-177-CM32) was implanted at the aIC (A-P 1.5-1.7, M-L 2.5-3.5, D-V
2.5); silicon probe (A4x8-Edge-2mm-100-200-177-CM32) based opto-electrode was implanted at the
dmPFC (A-P 1.0-1.7, M-L 0.1-1.5, D-V 1); 32-tetrode bundles (one-cluster) based opto-electrode was
implanted at the dmPFC (A-P 1.0-1.7, M-L 0.1-1.5, D-V 1); 32-tetrode bundles (two-cluster) were
implanted at the pIC (A-P −0.5-(-1.5), M-L 3-4, D-V 3) and motor cortex (A-P 1-2, M-L 1-2, D-V 0.5).
During the surgery, the skull was horizontally aligned through a fixing apparatus (Stoelting Co.). An
anchor screw was placed on the right cerebellum to connect ground wires of the electrodes. After
placing the anchor screw and electrodes, silicone sealant (kwik-cast, world precision instrument) was
applied above the exposed brain tissue. A customized head bar
(github.com/ywang2822/Multi_Lick_ports_behavioral_setup) was then positioned over the skull. To
affix the implant, Metabond (C&B Metabond, Parkell) and dental cement (Lang Dental) were applied.
The behavioral experiments started at least one week after the surgery.

Behavioral details

The head-fix setup was connected to a construction rod (Throlabs) by a 3d printed connector
(github.com/ywang2822/Multi_Lick_ports_behavioral_setup). Multi-lick-ports
(github.com/ywang2822/Multi_Lick_ports_behavioral_setup) were placed in front of the head fix and
connected to the Dual Lick Port Detector (www.janelia.org/open-science/dual-lick-port-detector). Three
Clearlink sets (Baxter) were used for liquid delivery. The delivery speed was manually calibrated to 0.15 -
0.2mL min^{-1} every time before the behavioral test. The delivery switch was controlled by three solenoid
valves (LFVA1220210H, THE LEE CO.) in a noise-reducing box. The switch timing was programmed
through the Bpod (Sanworks). The signal of mice locomotor activity was collected through an optical
shaft encoder (H5-360-IE-S, US digital). For facial videography set-up, the camera (S3-U3-91S6C-C,
Teledyne FLIR) was positioned at the right side of the mouse's lateral face surface, which illuminated by
two infrared arrays. For laser delivery, a solid-state laser (Shanghai Laser& Optics Century Co., 473 nm)
was connected to fiber optic patch cord (Doric Lenses), which attached to the implanted optic fibers

using ceramic mating sleeves. To conditionally control the laser delivery by water, sucrose, or quinine
onset, we used a 4-way data switch box (BNC, Kentek) to bridge the laser and solenoid valves. A
programmable stimulator (A-M system, model 4100) was used to control laser delivery and a voltage
pulse for tail shock experiment. All signals, including frame timing, wheel speed, liquid delivery timing,
lick timing, shock timing, and laser delivery timing, were sent to an USB interface board (Intan
Technologies, RHD).

For licking task, the deprivation was conducted until the body weight decreased approximately 22% (the
deprivation time was 16 to 36 hours, varied from mouse to mouse). During training phase, mice learned
to sense the water through their whiskers. We considered mice to become proficient at the task when
licking happened within 3s after the delivery onset (DO) in all repeated trials. During the test phase, we
first delivered water and 20% sucrose in a random sequence for a total of 30s. After at least 5 min, we
then orderly delivered water, 20% sucrose, and 5 mM quinine for 10s each or water and 5 mM quinine
for 15s each.

For tail shocking task, 16-23 volts electrical shocks were administered to the tail by a customized
shocker (electric shock box machine kit, STEREN). Two conductive adhesive copper tapes were
connected to the shocker and positioned 2 cm apart at the tail by sticking on customized heat shrink
tube (various on the circumference of mouse tail). During the first time of training, the voltage of
electrical shocks were adjusted until escaping behavior was observed (speed>10 cm s^{-1} right after the
shock). This voltage was recorded and used for the following tests. Those who did not perform escaping
behavior were excluded from the test.

Spike sorting and firing rate estimation

Before spike sorting, single unit data were acquired from 32-channel RHD head stage, which connected
with a signal acquisition system (USB board, Intan Technologies) with sampling rate at 20 kHz. All spike
sorting procedures were performed with an offline software Spikesorter [30, 31, 32]
(swindale.ecc.ubc.ca/home-page/software/) under following parameters: (0.5 kHz and 4 poles high pass
Butterworth filter) for signal filtering, (noise calculation: median; threshold: 80µV, 5x noise, 0.75ms
window width) for even detection, and (pca dimensions = 2; template window: -0.8 to 0.8; starting
sigma = 5; threshold = 9) for clustering. We used Bayesian adaptive kernel smoother [33] with following
parameters, α = 4 and β = (number of spike events) ^ (4/5), to estimate the firing rate of sorted spikes.
For small scale temporal window (180ms), we used a bandwidth of 5ms. While for large scale temporal
window (5s), we used a bandwidth of 200ms.

Optogenetic silencing

We illuminated bilateral prefrontal cortices using 473 nm laser to activate stGtACR2 [21]. Laser pulses
(40ms width at 20Hz) were delivered in a 5s duration. The onset of laser pulses was triggered based on
either water DO or quinine DO. The optogenetic silencing experiments were only performed after mice
reached stable behavioral level (after at least two test phases and (lick onset (LO) – DO < 3s) in all test
phases). The trials, of which licking frequency > 0.5 Hz in the time course 3s before DO, were excluded.
Histological characterizations were used to identify the viral infection.

Opto-tagging

We applied laser pulses (1ms width at 20 Hz, 3s duration) on the unilaterally prefrontal cortex of viral
(AAV-ChR2) injected mice. Laser and network-evoked spikes (see also **Spike sorting and firing rate**
estimation) were identified using the Stimulus Associated spike Latency Test (SALT[19]). Specifically, laser
and network-evoked spikes were assessed in a 0-5ms and a 6-10ms temporal window after laser onset,
respectively. For those units with significant correlation (correlation coefficient > 0.85) of average
waveform and significantly different distribution (P < 0.05) of spike latency with baseline units were
identified as laser or network-evoked units.

Chemogenetic inhibition

Viral pAAV-hsyn-DIO-hm4D(Gi) (Addgene_44362-AAV5) injected mice were administered
intraperitoneally with Clozapine N-oxide dihydrochloride (CNO, 2mg kg^{-1}, Tocris) ten minutes before the
licking or tail shocking task. Only the mice reached stable behavioral level (after at least two test phases
and (LO – DO < 3s) in all test phases) were used for chemogenetic experiments. In the licking task, mice
were re-trained to lick the water one to two times after recovery from CNO administration. The re-
trained phases were not included in test phases.

Analysis of facial and locomotor activity

We collected the frames during the licking or tail shocking task. We then converted these frames into
histogram of oriented gradients (HOG) vectors by using 8 orientations, 32 pixels per cell and 1 cell per
block. To extract the most variant facial part, we cropped the ear part with 364x296 pixels fixed size and
manually selected position of each transformed HOG vector [20]. Temporally adjacent HOG vectors were
paired, the facial activity at each time point was calculated as follows: *1-ΔR*, where *ΔR* is the correlation
coefficient between two temporally adjacent HOG vectors.

The signals that collected from the encoder were digital pulses. The locomotor activity was calculated as
speed (cm s^{-1}): $\frac{circumf}{CPR \cdot dt}$, where *circumf* is the circumference (cm) of the wheel, *CPR* (cycles per
revolution) is 360, and *dt* is the time interval between two digital pulses.

Analysis of licking initiation/termination bias

With the feeling of extremely thirsty, the mice will start a non-stop licking behavior when water is
available until feeling satiated or the delivery stopped [7] (**Supplementary Fig. 1d**). To evaluate if the mice
start or stop the continuous, but not discrete, lick, we calculated the initiation and termination bias. We
first calculated simple moving averages (SMAs) after water or quinine DO as following: $SMA = \frac{\sum_{i=1}^{n} l_i}{n}$,
where *l* is the lick times during a 200ms time window and *n* = 5 . For the initiation bias, all values of
SMAs were ignored if there was a zero value after DO. We created a vector that contained SMAs
sampled at 200ms intervals. The SMA value was counted from the last non-zero value. The initiation bias
(ibias) was calculated as: $ibias = \frac{1}{idx}$, where *idx* is the first time point of the SMA > 1.2 (6Hz). If all SMA
values equal zeros in 6 seconds, *ibias* was set as zero. For the termination bias, the SMA value was
counted from the first time point after the quinine DO (for the water-quinine and water-sucrose-quinine
session) or the end time point of water delivery (for the session water-water session) or sucrose delivery
(for the session water-sucrose session). The termination bias (tbias) was calculated as: $tbias = \frac{1}{idx}$,
where *idx* is the first time point of the SMA < 1 (5Hz). If all SMA values ≥ 1, *tbias* was set as zero.

Cell classification

For the cell classification to discriminate water- and quinine-licks, we categorized single-units into four separate groups of neural representation (lick, positive valence (PV), negative valence (NV), and mixed valence (MV)) based on firing rate estimation at the lick window (LO-100ms: LO+80ms). To determine if the firing rate is significantly higher than normal condition, we created pseudo-trials that have the same lick interval with the corresponding real licks during the 10s baseline. Individual time bins of each pseudo lick trials were concatenated horizontally and shuffled. This procedure was repeated 1000 times and the pseudo-trial matrix was calculated as the mean among shuffled datasets. For real lick trials, we only selected the first four trials for encoding analysis. At each individual time bin of pseudo and real trials, we calculated Euclidean norm of two temporally adjacent firing rate estimations (5ms each). When absolute z-scores of two distributions ($z_{12} = \frac{\mu_1 - \mu_2}{\sqrt{\sigma_1^2 + \sigma_2^2}}$, where μ and σ represent mean and standard deviation, respectively, of the distribution1 and 2) exceeded 1.29, they were considered significantly different. We selected single-units with significantly high firing rate in water or quinine lick trials for further analysis. To evaluate the time bias of firing rate across lick trials, we mean centered the whole firing rate matrix. The Frobenius norms were calculated as follows: $norm = \sqrt{\sum_{i=1}^{n}(t_i - c)^2}$, where t_i is the mean column value of firing rate matrix with 10ms time bin across lick trials, c is centered mean, and n equals the number of time bins. We categorized a single-unit with the time bias of lick trials when its real norm greater than 95% of 1000 shuffled norms. To estimate if the peak of firing rate of two distributions is different, we compared the times of the maximum firing rate across lick trials between two distributions using two sample t-test. Their firing rate peaks were considered different when p-value less than 0.05 (z-score>1.64). Single-units were categorized into the group of lick, positive or negative valence, or mixed response when their firing rates met following conditions: lick, quinine trials > pseudo trials & water trials > pseudo trials & without time bias of the firing peak between quinine trials and water trials & with the time bias of water and quinine trials; positive valence (PV), water trials > pseudo trials & water trials > quinine trials & quinine trials ≤ pseudo trials; negative valence (NV), quinine trials > pseudo trials & quinine trials > water trials & water trials ≤ pseudo trials; mixed valence (MV), water trials > pseudo trials & quinine trials > pseudo trials & with time bias of the firing peak between quinine trials and water trials & with the time bias of water and quinine trials; others were grouped into unrelated valence (UV).

To examine if there is a trial bias of firing rate during the water-licks, we compared the spike times at the small scale time window of water-licks. The matrices were binned (8 trials per group) and the mean values of each group were calculated. New generated binned firing rate matrices were then used. We next calculated Frobenius norm: $\sqrt{\sum_{i=1}^{n}(t_i - c)^2}$, where t_i is the mean row value of the binned firing rate matrix, c is centered mean, while n is the number of trials. When the real Frobenius norm was greater than 95% of 1000 shuffled norm, the single-unit was considered with trial bias in water-lick trials.

For the cell classification to discriminate initial and terminal phase of persistent movement, we assessed the firing rates at the 2s temporal window before or after the first water LO or quinine DO. The Five time point (0s, 0.5s, 1s, 1.5s, and 2s) were used for classification analysis. To construct pseudo data, the over 70s spike train data were used to extract five-time point random temporal window. We defined a single-unit with estimated firing rate higher than 65% pseudo data at the temporal window (1st water LO-2s: 1st

water LO) and lower than 35% pseudo data at the temporal window (quinine DO: quinine DO+2s) as
initial phase neural representation before LO; with estimated firing rate higher than 65% pseudo data at
the temporal window (1st water LO: 1st water LO+2s) and lower than 35% pseudo data at the temporal
window (quinine DO: quinine DO+2s) as initial phase neural representation after LO; with estimated
firing rate higher than 65% pseudo data at the temporal window (quinine DO: quinine DO+2s) and lower
than 35% pseudo data at the temporal window (1st water LO-2s: 1st water LO+2s) as terminal phase
neural representation. Others were classified as neural representations of unrelated movement phases.

Connectivity estimation

To estimate the connections among different neural representations, we used Total Spiking Probability
Edges (TSPE [18]). This method allows us to calculate the cross-correlation between pairs of spike trains
and to evaluate excitatory connections. Furthermore, it gives high accuracy estimation in a short
recording period. However, this method can only be used for the comparison of two individual single-
units. To apply this method on two networks, we first selected spike trains of photo-tagged and network
evoked single-units. We assumed the connection from photo-tagged to network evoked single-units is
positive. For the individual single-unit, 3s time duration of spike times data was cropped and sent to
TSPE calculation. We next selected spike trains of the neural representations of NV, PV, and initial phase
in three brain regions (IC, M1, and dmPFC). Individual time windows of 180ms across the lick trials (LO-
100ms: LO+80ms) were extracted and recombined. The connectivity among these neural
representations was calculated using TSPE. To perform a statistical test, we compared the real TSPE and
the pseudo TSPE, in which 3s spike times data were shuffled 200 times and created pseudo spike times.
70, 80, 90, 95, and 99 percentile of pseudo TSPE were used for the comparison based on the neural
representation number of real data. Specifically, if the real data number is less than 10, they will not be
used for the comparison; if the real data number is between 10 and 15, 80 percentile of pseudo TSPE
will be used for the comparison; if the real data number is between 15 and 20, 90 percentile of pseudo
TSPE will be used for the comparison; if the real data number is between 20 and 32, 95 percentile of
pseudo TSPE will be used for the comparison; if the real data number is larger than 32, 99 percentile of
pseudo TSPE will be used for the comparison.

Decoding analysis

For both small and large scale temporal window decoding, we employed a multiclass linear discriminant
analysis. We first estimated spike firing rate (see **Spike sorting and firing rate estimation)** in water and
quinine lick window (LO-100ms to LO+80ms for small scale window and first LO-2s to first LO+3s for
large scale window). The firing rate was normalized by dividing the maximum firing rate value of water
and quinine lick window. To create firing rate pseudo data, pseudo LOs were randomly selected for the
entire time duration of the spike train. This procedure was repeated 50 times. Same with real data
normalization, the pseudo data were divided by the max firing rate value in each repeat. To improve the
subsequent decoding performance, we reduced the dimensionality of the neural activity by PCA. The
dimensions that explained over 85% of the data variance were selected to train a decoder, which is
based on an error-correcting output codes (ECOC) classifier using binary support vector machine (SVM)
learner (MATLAB 'fitcecoc' function). 50 percent of real and pseudo data were used to train the
decoder, and the rest of them were used to test the decoder's performance. This procedure was
repeated 10 times to get an average accuracy and standard deviation.

Facial activity prediction

We trained Hammerstein-Wiener model to predict behavioral activity using estimated firing rate [34].
Before the modeling, the firing rates were estimated in a large scale temporal window (first LO-2s to
first LO+3s) and normalized into a range from 0 to 1 by dividing the maximum firing rate value of water
and quinine lick window. Empty and zero firing rates vectors were removed. The facial dynamics (see
Facial activity analysis) were smoothed using Gaussian filter. The parameters that can best simulate
facial dynamics were used to construct Hammerstein-Wiener model. Specifically, the number of zeros
was set in a range from 0 to 2 (nb-1); the number of poles was set in a range from 1 to 3 (nf); and the
degree of input nonlinearity estimator (one-dimensional polynomial) was set from 2 to 5. We then used
MATLAB function 'predict' to obtain decoding accuracies of facial dynamics from the test spike data.
Specifically, the data (four trials in total) were split 50/50. First two trials were used for training the
model and last two trials were used for testing. To avoid and overflow error, the values which lower
than negative 1500 were excluded.

Histology

To check the position of implanted electrodes and site of injection, mice were anesthetized with 2%
isoflurane (v/v) and perfused intracardially with 0.9% saline followed by 4% paraformaldehyde (PFA).
Fixed brains were washed three times before dehydration in 30% sucrose for 24 hr. Slices were cut on a
cryostat (MICROM, HM505E) at 70μm thickness after embedding with an optimal cutting temperature
compound (Tissue tek). Fluorescent images were acquired by an LSM 980 microscope (Zeiss), with a 10 ×
0.45NA objective or a 2.5 × 0.085NA objective.

MP network-based model

The purpose of this modeling was to create lick raster readouts through giving a type of current input,
simulating a network of neural activities, and transforming the simulated spike timing to the lick timing.

For the input current, we set the starting time t_s with a flexible (1-200ms) or a fixed (50ms) delay after
the DO. The amplitudes of input current were varied from 50-200pA according to the different
simulations. To test the effect of input continuity, the current was cut using one or twice 200ms zero
amplitude.

We simulated neural activity using Brian2 simulator package in a customized python code based on a
sparsely connected spiking neuron network [35]. The network consisted of 1000 excitatory and 800
inhibitory neurons as default (we decreased this number in the simulation of optogenetic inhibition).
The membrane potential of each neuron was modeled according to the MP neuron membrane
properties [17] based on the Hodgkin-Huxley model as following:

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$$\frac{dV}{dt} = \frac{gl \cdot (El-V) - g_{Na} \cdot m^3 \cdot h \cdot (V-E_{Na}) - g_K \cdot n^4 \cdot (V-E_K) + g_e \cdot (E_e-V) + g_i \cdot (E_i-V) + I_{ex} \cdot t_s}{C_m},$$

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$$\frac{dg_e}{dt} = -\frac{g_e}{\tau_e},$$

519
$$\frac{dg_i}{dt} = -\frac{g_i}{\tau_i},$$

where the excitatory and inhibitory synaptic time constants τ_e and τ_i were set as 5 and 10ms,
respectively. Other metrics were adjusted to adapt the inter-spike-intervals of L5 mPFC MP neuron,
which proposed to be a functional and dominate interneurons that bridge the gap between deep brain

regions and motor cortex [17]. The membrane potential V was initiated randomly at a range from -65mV

to -63mV. The excitatory conductance was set at a range from 0 to 0.06ns and the inhibitory

counterpart was in 0 to 1.5ns.

The synapse action was dependent on the usage u and availability x of released neurotransmitter

before and after an action potential as following: $\frac{du}{dt} = -\omega_f \cdot u, \frac{dx}{dt} = \omega_d - \omega_d \cdot x$, where the

facilitation rate was set as 3.33 s^{-1} and the depression rate was set as 2 s^{-1}. The rest synaptic release

probability was set as 0.6. The probability of excitatory connection of the network was given as 0.1 and

the inhibitory counterpart was 0.2.

To generate lick raster data from the network, we first assumed that one single lick cycle is governed by

a rotational neural dynamic [25]. Then we divided one cycle of the lick into nine phases. The triggering

probability P of a lick signal was calculated from the network through a decision algorithm:

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$$P(t) = \begin{cases} 1, & \prod_{\theta=0}^{2\pi}(\sum_{i=1}^{N} \varphi_{\theta i}(t - 280)) > 0 \ \& \ S\left(\sum_{i=1}^{N} \varphi_{\theta i}(t - 280)\right) < 30 \\ 0, & \prod_{\theta=0}^{2\pi}(\sum_{i=1}^{N} \varphi_{\theta i}(t - 280)) = 0 \ or \ S\left(\sum_{i=1}^{N} \varphi_{\theta i}(t - 280)\right) \geq 30 \end{cases}'$$

where S represents the standard deviation of the spike counts in whole phases.

Statistical test

For comparison of the mean of facial and locomotor activity (**Supplementary Fig. 1k-n** and

Supplementary Fig. 9e, h), comparison of the mean of licking frequency at different sessions

(**Supplementary Fig. 1c-f**), and comparison of TSPE (**Supplementary Fig. 5**), we used two-tailed Wilcoxon

signed rank test. For evaluation of the slope of spike counts trend line, we used one sample t test

(**Supplementary Fig. 4e**). The rest of statistical tests were two sample t test.

DATA AVAILIBILITY

The raw data that support these findings are available from corresponding author upon reasonable

request. Source data are provided with this paper.

CODE AVAILIBILITY

Materials and setup instruction for head fix and triple lick ports are freely available

(https://github.com/ywang2822/Multi_Lick_ports_behavioral_setup). Spike extraction used SpikeSorter

(swindale.ecc.ubc.ca/home-page/software). Connectivity estimation used TSPE toolbox

(https://github.com/biomemsLAB/TSPE). Estimated firing rate using Bayesian adaptive kernel smoother

(https://github.com/nurahmadi/BAKS). Identified opto-tag single-units using stimulus-associated spike

latency test (http://kepecslab.cshl.edu/software/). Cartoonish mice were created using Biorender

(biorender.com). Venn diagrams were plot using a customized Matlab script

(https://www.mathworks.com/matlabcentral/fileexchange/22282-venn). Other customized codes are

available from https://doi.org/10.5281/zenodo.7953792.

Figure Legends

Fig 1. Dissociable neural representations of movement phases and valence

a. Schematic of the experimental setup. Water or quinine was continuously delivered through a multi-lick-ports with a time delay of zero at the delivery switches. Mice were trained to respond the liquid delivery without conditioning. **b.** Schematic of recording sites. IC, insular cortex; dmPFC, dorsal medial prefrontal cortex; M1, primary motor cortex. **c.** A representative task of persistent lick. Two recording windows were selected near the water or quinine delivery onset (DO). For the large scale window (L-window), the recording epoch began 2 seconds before the 1st water or quinine lick onset (LO) until 3 seconds after the first water or quinine LO. For the small scale window (S- window), the recording epoch began from 100 ms before the LO to 80 ms after the LO. **d.** Spike raster and firing rate estimation of four representative single- units classified to represent the initial phase and terminal phase of persistent movement, positive valence (PV), and negative valence (NV) in the L- window and S- window, respectively. Spike raster and firing rate under water and quinine licks are colored blue and red, respectively. **e.** Pie chart showing the percentage of the classified neural representations. **f.** Venn diagram of the neural representations of valence and movement phases. Percentages of each category in all recorded neurons are shown in the diagram.

Fig 2. Coding of initial phase in dmPFC MP neurons

a. Schematics showing the labeling and recording of MP neuron in dmPFC. **b.** Representative image showing channelrhodopsin-2 (ChR2) expression in MP neuron. **c.** Identification of labeled MP neuron. We identified the unit as MP neuron when there was a significant probability of evoked spikes, appeared from 0 to 5ms after laser onset (light blue), and when there was a high correlation (R>0.85) between evoked spikes (light blue waveform) and other spikes (black waveform). **d & g.** Color-coded plot showing MP neural response in S-window (d) and in L-window (g) from one representative trial. **e & h.** Fraction of valence (e) and movement phases (h) classified neural representations in MP and IC neurons. **f & i.** Left, PCA trajectories of dmPFC MP neuron. Right, bar plot showing cumulative variance explained percentage of first two PCs. **j** Comparison of mean Euclidean distance from all PC dimensions of dmPFC MP neurons, between S-window and L-window. Values are mean ± s.e.m. **k.** Decoding of taste signals (water or quinine) in S-window (left) phase (initial or terminal) in L-window (right). **l.** Examples of facial activities (black traces) overlaid with MP firing predictions (Left, initial phase; right, terminal phase). **m.** Summary of facial activity predictions by recorded MP neurons. *P < 0.05, **P < 0.01, ****P < 0.0001, n.s. p > 0.05.

Fig 3. dmPFC MP neuron is required for the initiation of persistent licking movement

a. Left: schematic of bilateral silencing MP neurons in dmPFC. Right: experimental design for optogenetic silencing. Mice received laser or sham stimulation (5s, 20Hz) concurrent with water or quinine DO. **b & d.** Raster plot showing licking movement relative to water DO (b) or quinine DO (e). Laser (left) or sham (right) was triggered by DO. Cyan background and dash box represent laser (b) and sham (d) delivery period, respectively. **c & e.** Percentage of bias that started (c) or stopped (e) persistent lick (see Methods). **f.** Color-coded plot showing facial activity relative to water DO. **g.** Z scored facial activity relative to water DO. Values are mean ± s.e.m. *p<0.05, n.s. p>0.05.

Fig 4. Silencing of dmPFC MP neuron impairs the neural activity in motor and insular cortex

a. Schematic of bilateral silencing MP neurons in the dmPFC. **b & c.** Schematic of recording sites in IC (b) and M1 (c). **d & e.** Baseline subtracted, z scored firing rate, relative to water DO, for the recorded neurons in IC (d) and in M1 (e) from one representative trial. Cyan background and dash box represent laser (top) and sham (bottom) delivery period, respectively. **f & g.** Mean baseline subtracted, z scored firing rate, relative to water DO, of IC (f) and M1 (g) neurons. Values are mean ± s.e.m. *p<0.05.

Fig 5. Modulation of neuron number and input in the MP network-based computational model

a. The MP network-based model. A spiking network receives an input of current after the DO (delivery onset) with a flexible (1 to 200ms, for panel B) or fixed (50ms, for panel C) delay. 56% connectivity is excitatory and 44% of it is inhibitory. The lick raster outputs are calculated by a decision algorithm (see Methods) using the simulated spike data. **b.** Simulation of optogenetic inhibition of dmPFC MP neurons by reducing the neuron number (both excitatory and inhibitory neurons) in the MP network based model. Left 1-3: lick raster produced by the model under indicated neuron number. Right: percentage of initiation bias that calculated from left lick raster data. **c** Performance of licking behavior under the different continuities of input current. Left 1-3: top: lick raster plots under, bottom: inputs with indicated interruption times. Right: percentages of initiation bias with indicated interruption times. *p < 0.05, ****p < 0.0001

Fig 6. Summarized model for how dmPFC MP neurons initiate persistent movement.

a. Pattern of neuronal activation during the persistent licking movement. After the onset of stimulation, there is a delay of approximately 1.3 seconds until the first lick onset. In the initial phase (the licking frequency is driven to the maximum), the indicated three types of neurons are activated. In the persistent phase, dmPFC MP neurons are inactivated, while M1 and IC neurons show persistent activity. **b.** During the initial phase, when the sensory signals are received, dmPFC MP neurons activate themselves and the downstream motor neurons. The latter drive the licking movements that bring the tongue into contact with the water. The taste signal from the tongue is further transmitted to the valence-encoding neurons in the IC. During the persistent phase, no signal is transmitted from the dmPFC MP neurons to the motor neurons, but the motor neurons is continuously activated. This results in persistent licking movement and persistent valence encoding in the IC.

Supplementary Fig 1. Behavioral performance in persistent licking tasks

a. Schematic of timeline for licking tasks per trial. **b.** Schematic of behavioral setup and equipment. **c-f.** Lick behavior relative to delivery onset (DO, left) and to 1st lick onset (LO, right) after DO in four sessions. Top, raster plots showing lick behavior in the session water (w, 15s)-quinine (q, 15s) (c), water (w, 15s)-water (w, 15s) (d), water (w, 15s)-sucrose (s, 15s) (e), and water (w, 10s)-sucrose (s, 10s)-quinine (q, 10s). Bottom, lick frequency plot. The 'peak' indicates the maximum lick frequency. Wilcoxon signed-rank test, in comparison with baseline: ****p<0.0001, n.s. not significant. Values are mean ± s.e.m. **g.** Comparison of the percentage of body weight decrease among four sessions using one-way anova. **h & i.** Percentage of bias that started (h) or stopped (i) persistent lick (Methods). One-way anova: p=0.1147 for initiation bias; p<0.0001 for termination bias. Bonferroni multi-compare: **p<0.01 ****p<0.0001. **j.** Illustration of calculating facial activity as 1-Δcorrelation (see Methods). **k-n.** Color-coded plot and traces showing facial and locomotor activity relative to water DO. We compared the epochs from DO to LO, from switch onsets to switch onsets+5s, from end onsets to end onsets+5s with the baseline. Wilcoxon signed-rank test: *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001, n.s. not significant. Values are mean ±

s.e.m. **o-s.** Comparison of licking performance (o), facial and locomotor activity (p and q), initiation bias

(r), and termination bias (s) between male and female mice in w-q session. *$p<0.05$, **$p<0.01$,

***$p<0.001$, ****$p<0.0001$, n.s. not significant. Values are mean ± s.e.m.

Supplementary Fig 2. Histological and electrophysiological verification

a. Histological verification of IC implant sites. From left to right: representative images of anterior IC

implant is shown in 1st to 2nd columns and posterior IC implant is shown in 3rd to 4th columns. The

magnified images are shown2^{nd} and 4th columns. **b.** Histological verification of M1 implant sites. From

left to right: representative anterior M1implant is shown in 1st to 2nd columns and posterior M1 implant

is shown in 3rd to 4th columns. The magnified images are shown in 2nd and 4th columns. **c.** Histological

verification of optic fiber implans for optogenetic silencing in bilateral dmPFC. **d.** Histological verification

of viral injections for chemogenetic inhibition of hM4D labeled neuron in bilateral dmPFC. **e.** Histological

verification of viral injections and electrodes and optic fiber implants for optogenetic identification of

ChR2 labeled neuron in unilateral dmPFC. **f.** Electrophysiological verification of photo-tagged and

network-excited neurons (see Methods).

Supplementary Fig 3. Analysis of single-unit classification

a. Procedure for shuffling baseline data. Left: color coded firing rates in the real and pseudo lick trials

(the pseudo lick number equals to the real lick number after the baseline). Right: distributions of mean

adjacent Euclidean norm obtained from the 1000 pseudo datasets and 1 real dataset. **b.** Procedure for

categorizing the single-units with or without time bias. Top: example of a single-unit with time bias.

Bottom: example of a single-unit without time bias. Left: color-coded firing rates showing shuffled data

during lick trials. Middle: real data firing rates. Right: distributions of Frobenius norms of shuffled data.

The single-unit was categorized as time bias when its Frobenius norm higher than 95 percentile of

shuffled data. **c.** Spike raster and firing rates plots showing representative classified single-units that

encode lick, mix valence (MV), negative valence (NV), and positive valence (PV) in the session water-

quinine (wq, top) and the session water-sucrose-quinine (wsq, bottom). Black dots denote the peak

firing rate at the each lick trial. **d.** Procedure for categorizing the neural representations with or without

water-lick trial bias. Example of a single-unit with (top) or without (bottom) lick trial bias (matrix was

binned with each 8 trials). Left: color-coded firing rates showing shuffled data during water-lick trials.

Middle: real data firing rates. Right: distributions of Frobenius norms of shuffled data. The single-unit

was categorized as water-lick bias when its Frobenius norm higher than 95 percentile of shuffled data. **e.**

Color-coded firing rates showing the two examples of PV neural representations with (right) or without

(left) water-lick trial bias. **f.** Procedure for generating pseudo data. Left: illustration of pseudo data

selecting. Right: distributions of firing rates in 5 pseudo time points. **g.** Example of a single-unit firing

rate in three recording epochs. **h.** Spike raster and firing rates plots showing representative classified

single-units that represented initial phase-after lick onset (ALO), initial phase-before lick onset (BLO),

terminal phase, and mixed phase (MG).

Supplementary Fig 4. The percentage and firing trend of specific neural representations in the brain

regions IC, M1, and dmPFC

a-c. Percentage of the classified neural representations in IC (a), M1 (b), and dmPFC (c). **a1** & **b1** & **c1**:

percentage of PV, NV, MV, lick, and UV (unrelated valence) neural representations. **a2** & **b2** & **c2**:

percentage of initial phase (including ALO and BLO), terminal phase, mixed phase, and unrelated

movement phase neural representations. **a3** & **b3** & **c3**: venn diagram showing the overlap and non-
overlap percentage of indicated neural representations. **d.** The spike counts of indicated neural
representations during water-licks. A blue dot denotes the spike times in the small scale window (LO-
100ms to LO+80ms) of a water-lick trial. Blue lines represent the trend line of blue dots. **e.** Slopes of
trend line of overall non-overlapping neural representation groups in IC, M1, and dmPFC as indicated.
The symbol + and − represent the significant positive and negative value of its labeled trend line slope,
respectively. + or - P<0.05, one sample t-test.

Supplementary Fig 5. The connectivity among specific neural representation groups

a-b. Connectivity of neural pairs between photo-tagged and network-excited single-units. A: Cross-
correlograms of individual neural pairs. B: TSPE comparison of real data and the shuffled data at
different percentile. P values are measured by Wilcoxon signed-rank test. **c-d.** Connectivity among
classified neural representations. **Left:** Cross-correlograms of individual neural pairs. **Right:** TSPE
comparison of real data and the shuffled data at 99 percentile. All mean of real TSPEs are less than mean
of 99% shuffled data.

Supplementary Fig 6. The decoding of liquid types in IC, M1, and dmPFC along with lick proceeding

a1 & b1 & c1. Decoding of the water-licks in 1st recording epoch (water DO to water DO+15s) and in 2nd
recording epoch (water DO+15s to delivery end). **a2 & b2 & c2.** Decoding of the water-licks in 1st
recording epoch (water DO to water DO+15s) and the 1st sucrose-lick in 2nd recording epoch (sucrose DO
to delivery end). **a3 & b3 & c3.** Decoding of the water-licks in 1st recording epoch (water DO to water
DO+15s) and the 1st quinine-lick in 2nd recording epoch (quinine DO to delivery end). **From left to right:**
1st column, color-coded plot showing z scored neural response in the first lick window of 1st recording
epoch and 2nd recording epoch; 2nd column, decoding of the first water-lick in 1st recording epoch and in
2nd recording epoch; 3rd column, color-coded plot showing neural response in the 42th lick window of 1st
recording epoch and in the 1st lick window of 2nd recording epoch; 4th column, decoding of the 42th
water-lick in 1st recording epoch and the first lick in 2nd recording epoch. 5th column, different brain
regions showing different liquid type discrimination and changing the discrimination level (decoding
performance) along with lick trials evolving. Decoding of water-licks (from 1st to 42th lick trial) in 1st
recording epoch and first lick in 2nd recording epoch. Values are mean ± s.e.m.

Supplementary Fig 7. The decoding of initial versus terminal phase in IC, M1, and dmPFC

a1 & b1 & c1. Decoding of the initial phase in 1st recording epoch (1st water LO-2s to 1st water LO+3s)
and the terminal phase in 2nd recording epoch (water delivery end-2s to water delivery end+3s). **a2 & b2**
& c2. Decoding of the initial phase in 1st recording epoch (1st water LO-2s to 1st water LO+3s) and the
terminal phase in 2nd recording epoch (sucrose delivery end-2s to sucrose deliver end+3s). **a3 & b3 & c3.**
Decoding of the initial phase in 1st recording epoch (1st water LO-2s to 1st water LO+3s) and the terminal
phase in 2nd recording epoch (1st quinine LO-2s to 1st quinine LO+3s). **Top:** color-coded plot showing z
scored neural response in the 1st recording epoch and 2nd recording epoch. **Bottom:** decoding
performance. Values are mean ± s.e.m.

Supplementary Fig 8. Prediction of facial activity from spike activity in regions IC, M1, and dmPFC

718 **a.** Examples of facial activities (black traces) overlaid with Hammerstein-Wiener model using indicated
719 parameters. **b-d.** Firing rate predictions of facial activity using Hammerstein-Wiener model. B1 & C1 &

720 D1: examples of facial activities (black traces) overlaid with firing rate predictions (right, water-lick; left,
721 quinine-lick). B2 & C2 & D2: summary of facial activity predictions. *p<0.05

Supplementary Fig 9. Behavioral effect of dmPFC MP neuron chemogenetically silencing

a. Schematic of bilateral silencing MP neurons in dmPFC. **b.** Lick behavior relative to DO with or without
CNO administration. From left to right: 1st column and 3rd column, raster plots showing lick behavior in
the session w (15s)-w or s (15s). 2nd column and 4th column, lick frequency plot. **c.** Comparison of lick
chance between saline and CNO administrated mice after water delivery. **d.** Comparison of the
percentage of body weight decrease between saline and CNO administrated mice. **e.** Color-coded plot
and traces showing z scored facial and locomotor activity relative to water DO. We compared mean
facial and locomotor activity in the epochs from DO to LO, from switch onsets to switch onsets+5s, from
end onsets to end onsets+5s with that in the baseline. Wilcoxon signed-rank test: *p<0.05, **p<0.01,
n.s. not significant. **f-g.** Comparison of z scored facial activity (f) and locomotor activity (g) between
saline and CNO administrated mice at various epochs as indicated. Two sample t-test: **p<0.01, n.s. not
significant. **h.** Color-coded plot and traces showing facial and locomotor activity relative to shock onset
(SO). The mean facial and locomotor activity in the epoch from SO to SO+5s were compared with that in
the baseline. Wilcoxon signed-rank test: *p<0.05, ****p<0.0001, n.s. not significant. **i.** Comparison of z
scored facial activity (top) and locomotor activity (bottom) between saline and CNO administrated mice
at the epoch SO to SO+5s. Two sample t-test: *p<0.05, ****p<0.0001.

Supplementary Fig 10. Behavioral effect of dmPFC MP neuron optogenetically silencing

a1 & b1. Top, raster plots showing Lick behavior relative to DO shining with or without laser. Bottom,
Color-coded plot and traces showing facial and locomotor activity relative to DO. Laser (left) or sham
(right) was triggered by DO. **c1 & d1.** Color-coded plot and traces showing facial and locomotor activity
relative to DO. Laser (left) or sham (right) was triggered by DO. Cyan background and dash box represent
laser and sham delivery period, respectively. **a2 & b2.** Left, lick frequency relative to DO. Right,
Comparison of lick times at laser or sham treatment periods. **a3 & b3 & c2 & d2.** Comparison of facial
(left) and locomotor (right) activity between laser and sham treatment periods. **a4 & b4 & c3 & d3.**
Comparison of the percentage of body weight decrease between laser and sham treatment groups. **c4.**
lick frequency relative to DO. The analyzing window was extended to full time period. No significant
difference between laser and sham treatment was found after laser off. Two sample t-test: n.s. not
significant **p<0.01

Supplementary Fig 11. Comparison of the MP network based model and the experimental data

a. Voltage responses of a representative MP neuron in the experiment (data from our previous work[17],
black traces) and the model (red traces) to step current injections. **b.** Tongue movement and MP
network is quantified by rotational phases. Left: representative image showing tongue movement.
Right: one cycle of MP network and tongue movement was set as 125ms and there was a 280ms delay
before the signal from MP network arrived to tongue. **c.** Polar histogram of experimental (top) and
simulated (bottom) network phases for indicated LO numbers. The blue area of each phase indicates the
relative spike counts. **d.** Neural activity of real (left, black) and simulated (right, red) MP network related
to lick frequency and initiation bias. Dash lines denote the 95% intervals. R-squares and P values of the
linear regression are labeled at each panel. In both **c** and **d**, the experimental data were only chosen

from the neurons that the normalized firing rate lower than 50 (65% of the total neuron number) given
that slow spiking MP neurons are mainly functional linking deep brain regions and motor cortex[17].

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AUTHOR CONTRIBUTIONS

Q.S. supervised the project, acquired the funding, and edited the manuscript. Y.W. built the behavioral
set-up, collected and analyzed data, performed the simulation, and wrote the original manuscript.

880 COMPETING INTERESTS

881 The authors declare no competing interests.

Fig 1

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Supplementary Fig 1

Supplementary Fig 2

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Supplementary Fig 8

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Supplementary Fig 11