# Natural and pathological aging distinctively impact the vomeronasal system and social behavior

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**Abstract**

Normal aging and many age-related disorders such as Alzheimer´s disease cause deficits in olfaction, however it is currently unknown how natural and pathological aging impact the detection of social odors which might contribute to the impoverishment of social behavior at old age further worsening overall health. Here, we investigated the effect of aging in the recognition of social cues and the display of social behavior. Our findings indicate that aging distinctively disrupts the processing of social olfactory cues decreasing social odor exploration, discrimination and habituation in both wild type senescent (2-year-old) mice and in 1-year-old double mutant model of Alzheimer´s disease (APP/PS1[Het]). Furthermore, social novelty was diminished in 1-year-old APP/PS1 mice, indicating that alterations in the processing of social cues are accelerated during pathological aging. Analysis of the vomeronasal organ, the main gateway to pheromone-encoded information, indicated that natural and pathological aging distinctively reduce the neurogenic ability of the vomeronasal sensory epithelium. Cell proliferation remained majorly preserved in 1-year model of Alzheimer´s disease (APP/PS1[Het]), whereas naturally aged animals exhibited significant deficiencies in the number of mature, proliferative and progenitor cells. This study reveals fundamental differences in the cellular processes by which natural and pathological aging disrupt the exploration of social cues and social behavior.

**Introduction**

Social recognition is essential for survival allowing to appropriately adapt behavior across a variety of contexts (Adolphs, 2001; Brennan and Kendrick, 2006). As aging progresses, social odors have been shown to elicit attenuated responses (Hlinák and Krejcí, 1991; Garrat et al., 2011), a phenomenon interpreted as a natural consequence of age-related decline in sensory perception (Doty et al., 1984; Murphy et al., 2002; Doty and Kamath, 2014), that might negatively impact the social behavior of aged animals, whose survival might become increasingly dependent on the social environment. In humans, social impoverishment has been identified as a major aggravating factor for decreased life expectancy, and an indicator of the appearance of dementia and neurodegenerative disorders such as Alzheimer´s disease (AD) (Wilson et al., 2007; Heinrichs et al., 2003; Donovan et al., 2016). Despite the central role of social interaction in maintaining overall well-being, the mechanisms by which aging, natural or pathological, alter social information processing remain unclear.

Mouse social communication strongly depends on chemosignals comprised by a heterogeneous mix of volatile and non-volatile molecules (pheromones) (Petrulis and Johnston, 1997; Beynon and Hurst, 2004; Yoon et al., 2005; Baum and Kelliher, 2009; Chamero et al., 2012; Noack et al., 2010; Tobin et al., 2010; Li and Dulac, 2018). Volatile odors are primarily detected at the main olfactory epithelium (MOE) which projects to the main olfactory bulb (MOB), whereas pheromones are mainly detected by

46　the vomeronasal sensory epithelium (VSE) (Jacobson, 1813; Halpern and Martinez-Marcos, 2003;
47　Munger et al., 2009) which connects to the accessory olfactory bulb (AOB) (Shipley et al., 2008).
48　However, both systems have overlapping and complementary functions in relation to the stimuli they
49　are able to detect and the behavioral or physiological output they can generate (Brennan and Zufall,
50　2006; Zufall and Leinders -Zufall, 2007).
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A remarkable property of the VSE is that contains proliferative niches capable of generating new cells
majorly becoming functional neurons during adulthood (Weiler, 2005; Weiler et al., 1999; Whitman
and Greer, 2009; Graziadei and Monti Graziadei, 1983; Halpern and Martínez-Marcos, 2003; Brann
and Firestein, 2014; De la Rosa-Prieto et al., 2009; Oboti et al., 2015). The neurogenic properties of
olfactory structures are known to decline with age (Bailey et al., 2004; Ahlenius et al., 2009; Wagner
et al., 2009), a phenomenon proposed to underlie the olfactory deficits associated to both natural aging
and neurodegenerative disorders (Doty et al., 1984; Murphy et al., 2002; Doty and Kamath, 2014;
Devanand, 2016; Mobley et al., 2014; Lazarov and Marr, 2010). However, how natural and pathological
aging affect the regenerative capacity of the VSE has been scarcely studied despite being central to
socio-sexual cue detection. Previous studies have shed some light into VSE anatomical alterations
induced by natural aging. An elegant study by Brann and Firestein (2010) showed that the proliferative
capacity of the VSE of 2-year-old animals was attenuated in comparison to young animals. More
recently, Mechin et al., (2021) explored the effects of natural aging on VSE cell proliferation, finding a
significant reduction in mature Olfactory Marker Protein (OMP)-expressing cells (OMP$^+$), indicating
structural modifications of the aged VNO.  However, whether these alterations in the aged VSE have
any impact in odor detection and subsequent display of social behaviors, particularly in animal models
of neurodegeneration, is currently unclear.

To gain insight into these questions, we investigated the impact of natural and pathological aging in
social odor-evoked sniffing behavior, social-odor habituation/dishabituation and social behavior during
the aging process of wild type C57/BL6 mice and amyloid precursor protein (APP) and presinilin-1
(PS1) (APPswe/PS1ΔE9) double transgenic mice (APP/PS1$^{Het}$ mice, therein), a widely used animal
model of AD (Jankowsky et al., 2001; Jankowsky et al., 2004). Analysis of the number of progenitors,
proliferative and mature neurons revealed that naturally aged animals show reduced neuronal
proliferation and decreased levels of sensory mature (OMP$^+$) neurons. In contrast, the VSE of middle
age (1-year-old) APP/PS1$^{Het}$ mice exhibited normal levels of mature neurons and cell regeneration,
suggesting that natural and pathological aging distinctively impact the neurogenic capacity of the VSE.

We explored the functional consequences of these cellular alterations observing reduced exploration
time of social odors and decreased performance in a social odor habituation-dishabituation test in both
senescent and middle age APP/PS1$^{Het}$ mice, suggesting that despite the overall normal VSE structure
and cell composition, APP/PS1$^{Het}$ animals exhibit an accelerated decay of the processing of social cues.
Moreover, 1-year-old APP/PS$^{Het}$ mice exhibited a significant reduction of social novelty which was no
apparent in naturally aged animals, exposing a fundamental difference in how natural and pathological
aging impact the regulation of social behavior.

**Materials and Methods**

*Animals*

Experiments were performed in either control C57/BL6 mice (young: 2-4 months-old; adult: 6-8
months-old; middle age: 12-14 months-old; old (senescent): 20-24 months-old) or APP/PS1 on a
C57/BL6 genetic background (young: 2-4 months-old; middle-age: 12-14 months-old) (Jackson Labs,
Stock No. 004462, MMRRC Stock No. 34829). APP/PS1$^{Het}$ mice express a chimeric mouse/human
amyloid precursor protein (Mo/HuAPP695swe) and a mutant human presenilin 1 (PS1-dE9). These
animals develop beta-amyloid deposits at ~ 6 - 7 months of age and exhibit early-onset cognitive
impairments (Jankowsky et al., 2001, 2004; Reiserer et al., 2007). Animals were kept group-housed and

experimentally-naïve during all their lifetime under pathogen-free conditions. Animals were housed in
ventilated cages with free access to food and water on a 12 h light/dark cycle.

All experiments were performed according to Spanish and European Union regulations regarding
animal research (2010/63/EU), and the experimental procedures were approved by the Bioethical
Committee at the Instituto de Neurociencias and the Consejo Superior de Investigaciones Científicas
(CSIC).

### *VNO dissection*

Mice were perfused transcardially with PBS, pH 7.4, followed by 4% paraformaldehyde in 0.1 M
phosphate buffer (PB), pH 7.4. A fixed mouse head was placed under a scope and the lower jaw was
removed to get a palate view. To facilitate VNO extraction, the palate and the nasal septum were
removed and the bilateral VNOs were split in two parts with a gentle twisting motion. Finally, the
cartilaginous tissue that covers each portion was carefully removed to extract the VNOs. Tissue was
incubated in a 30 % sucrose solution for cryoprotection and kept at -4 ºC until sectioning. Samples were
embedded in OCT and frozen at -80 ºC for cryosectioning.

### *Immunohistochemistry*

VNOs embedded in OCT were sliced in 16 μm thick sections in a cryostat apparatus (Leica CM 3050S).
Slices were incubated in blocking solution containing 0.5 % Triton X-100 and 5 % normal horse serum
in 0.1 M TBS for 2 h at room temperature (RT). Primary antibody incubation was performed overnight
(o/n) at 4 ºC against OMP (Wako goat polyclonal; 1:2000), PCNA (Sigma, rabbit monoclonal; 1:2000);
Sox2 (R&D Systems, goat polyclonal; 1:300). For PCNA immunostaining, slices were incubated in 10
mM citrate buffer (100 ºC; pH, 6) for 5 minutes prior staining. Sections were incubated with Alexa
Fluor 488- or 594- conjugated secondary antibodies (Jackson Laboratories, 1:500) for 2 h at RT.
Hoechst (Sigma, 1:10000) was added during 5 minutes after the secondary antibody incubation for
nuclei visualization. Imaging was performed using a vertical confocal Microscope Leica SPEII. Final
images were assembled with Adobe Illustrator.

### *Sox2 fluorescence intensity at the VNO supporting cell layer*

120 For determining the Sox2 fluorescent labeling area in the VNO supporting cell layer (SCL) where
121 single-cell quantification was limited by the densely packed disposition of the cells, we calculated the
122 corrected total fluorescence (CTF) of the area of interest employing the Freehand ROI tool of Image J.
123 CTF was calculated by subtracting the background fluorescence from a minimum of 3 sections, 16 μm
124 thick from at least 4 animals per condition.

### *Stereology, cell quantification and AOB volume estimation*

The total number of olfactory mature neurons (OMP$^+$ neurons) in the entire VSE was estimated using
the optical fractionator method (Wong et al., 2018) included in the Stereo Investigator software (MBF
Bioscience). Two different tools are combined in this unbiased quantification method: a 3D optical
dissector for cell counting and a fractionator, which is based on a systematic, uniform and random
sampling over a known area of the tissue (West and Gundersen, 1990; Dorph-Petersen and Lewis, 2011;
Parrish-Aungst et al., 2007). The number of neurons is estimated as:

$$N = (1/h \times 1/ssf \times 1/asf) \times \sum Q - t$$

In which $\sum Q$ is the total cell number in the region of interest acquired using the optical dissector
method; **t** stands for the mean mounted section thickness; **h** is the optical dissector height; **asf** is the
area sampling fraction and **ssf** is the frequency of sampling (section sampling fraction). For the VSE
stereological analysis, sampling was performed with a 20x objective (Leica, NA 0.6), the counting
frame area was 2500 μm$^2$ and the sampling grid area was 22500 μm$^2$; h for VSE stereology was 8 μm

with 1 μm as upper and lower guard zones and t was set at 10 μm. Quantifications were performed for
the central, intermediate and marginal regions of the VSE dividing the area in segments of equal length
as previously described (Giacobini et al., 2000; Brann and Firestein, 2010). For the cell quantification
in the anterior-posterior regions, the VSE was divided in 10 sections per animal and the results were
divided by the area to obtain the number of cells/mm$^2$ in each subdivision. Quantification of the number
of cells in the marginal and central areas was performed in three slices of each animal delineated with
a 50x50 dissector employing Stereo Investigator (MBF Bioscience).

VSE volume was calculated by multiplying the sampled area by the slice thickness (16 μm) and the
number of series (10 slices per animal). Ten sections per series were analyzed to obtain an estimation
of the VSE area and volume in young (4 months-old), aged (24 months-old), middle aged (12-months
old) APP/PS1$^{WT}$ and middle aged APP/PS1$^{Het}$ mice.

The volume of the AOB was estimated by measuring the AOB area and multiplying the sampled area
by the section thickness (50 μm) and the number of analyzed sections (5 series per animal).

### *Odorants*

Social and non-social scents were employed to discern the impact of aging in social odor perception.
As non-social scents, we used **i)** the banana-like odor isoamyl acetate (Sigma) shown to be primarily
detected at the MOE level (Xu et al., 2005) with neutral valence within a broad dilution range (Fortés-
Marco et al., 2015; Root et al., 2014; Saraiva et al., 2016), and **ii)** food pellets to perform a food finding
test (FFT, see experimental details below). As a social scent, we used mouse urine as urine from
conspecifics has been shown to elicit robust VNO activity (Chamero et al., 2017). Urine samples were
collected according to standard procedures (Kurien et al., 2004). For habituation-dishabituation tests,
frozen urine samples from different young subjects and cage mates were used to test fine odor
discrimination. For sensitivity tests, urine samples were combined in a stock sample used for each round
of odor presentations until completing the experiments. For female urine, samples obtained at different
points of the estrous cycle were pooled to generate a uniform stock.

### *Behavioral assessment*

All the behavioral experiments using social and non-social stimuli were performed in a dedicated room
with continuous air reposition under dim indirect light (20 lx). Odor dilutions were prepared in a room
outside the animal house. Odor presentation was performed in a homemade methacrylate box with
removable walls for cleaning. The chamber dimensions were 15 cm width, 15 cm length and 30 cm
high. A small hole (1cm diameter) was performed in the middle of one of the sides located 5 cm above
from the box ground to fit standard cotton sticks impregnated with 1 μL of the odorant (**Figure 3A**)
allowing direct contact with the stick. Habituation to the testing conditions was performed before
odorant presentations consisting of: handling (5 min), free exploratory activity in the box and
habituation to the cotton stick movement (5 min). Odors were presented embedded in a cotton stick in
order to preserve non-volatile pheromones and mimic the natural detection method of physical contact
between conspecifics (Leypold et al., 2002; Stowers et al., 2002; Kimchi et al., 2007). To avoid
variability due to the intensity of the volatile components of the urine, only direct nose contact with the
tip of the cotton stick were considered as explorative behavior (sniffing/exploration time). Experiments
were monitored by a video recording camera fixed 15 cm above the box. Videos were collected and
analyzed off-line using SMART video-tracking software (PanLab S.L.).

a) *Odor exploration test*: After a period of habituation (10 min), mice were exposed to two control trials
(mineral oil for isoamyl acetate assays or water for urine tests) during 1 min separated by intervals of 1
min to avoid lack of novelty and odor habituation (Breton-Provencher et al., 2009; Sanderson and
Bannerman, 2011). Subsequent presentations consisted on serial dilutions of either urine samples
(diluted in water: 1:1.000, 1:500, 1:250, 1:100, 1:50, 1:10 and non-diluted, nd) or isoamyl-acetate were

tested in ascending order (diluted in mineral oil: $1:5 \times 10^5$, 1:100.000, 1:10.000; 1:1.000, 1:100). Animals
were considered to detect the olfactory stimulus when spent more time investigating the odors than the
vehicles (water or mineral oil).

b) *Food finding test (FFT)*: Food finding test (FFT) was performed following standard procedures
(Deacon et al., 2009). The mice were food deprived for 24 h before testing. Five food pellets (~ 35 g)
were placed a corner and covered by 5 cm of litter bedding. Animals were considered to detect the food
pellet when spent digging, touching and holding the food pellet for more than 5 s.

c) *Social odor habituation-dishabituation test:* The effect of aging in social odor discrimination and
habituation was explored using an adapted habituation-dishabituation test (Gheusi et al., 2000). After a
period of cage habituation (10 min), two control trials were performed employing cotton sticks soaked
with 1μl of water (vehicle). Three successive presentations of urine from an animal of the opposite sex
(S1a-c) were followed by three presentations of urine from a different subject of the opposite-sex (S2a-
c) to evaluate fine chemo-olfactory discrimination and habituation. Urine samples came from young
animals of the opposite sex to maximize approaching behavior (Xu et al., 2005; Garratt et al., 2011).
Each sample presentation lasted 1 min separated by 1 min intervals. Habituation was estimated as a
decrease in the exploration time (sniffing) over the cotton stick tip after the first exposure of urine from
the same animal (S1a). Dishabituation (discrimination) occurs in response to a new odor presentation
(S2a) as a measurable increase in the exploration time. The test allows to explore two consecutive
phases of discrimination ($H_2O^B$-S1a and S1c-S2a) and habituation (S1a-S1b and S2a-S2b). Trend lines
between $H_2O^B$-S1a, S1c-S2a, S1a-S1b and S2a-S2b were fitted to obtain the slope values indicating the
amplitude of the discrimination and habituation effect in each tested condition. Higher positive values
indicated more pronounced social discrimination and higher negative values corresponded to more
robust habituation (**Supplementary Figures 8 and 9**).

d) *Long-term social habituation-dishabituation test*: The aforementioned social odor habituation task
was adapted to assay social cue memory by presenting the urine sample employed in S1a 24 hours after
the first test was performed. Intact social odor memory manifested as a reduction in the sniffing time
during the presentation of S1a during the second trial, an effect which was clearly apparent in young
animals (**Figure 5F-G**).

e) *Three chamber test*: Social testing was performed in a cage ($60 \times 40 \times 22$ cm) following standard
procedures (Nadler et al., 2004). Dividing walls were made from clear Plexiglas, with openings
allowing access into each chamber. The test mouse was first placed in the middle chamber and allowed
to explore for 10 minutes. After the habituation period, an unfamiliar subject of the same sex (mouse 1,
M1), was placed in one of the side chambers. The stranger mouse was enclosed in a small, round wire
cage, which allowed nose contact between the bars. In this first session (sociability), the test mouse had
a choice of spending time in either the empty chamber (E) or the chamber occupied by M1. At the end
of the sociability session, each mouse was tested in a second 10-minute session to evaluate social
preference for a new subject. A second, unfamiliar mouse (mouse 2, M2) of the same sex was placed
in the chamber that had been empty during the first session. This second stranger was enclosed in an
identical wire cage than M1. The test mouse had a choice between the first, already-investigated mouse
(M1), and the novel unfamiliar mouse (M2) which indicates their social preference or social novelty
(Nadler et al., 2004). Continuous video recordings were collected and analyzed off-line using BORIS
(Friard and Gamba, 2016) SMART video-tracking software (PanLab S.L.). Measures of time spent
sniffing E, M1, and M2 were quantified for each session.

*Statistical analysis*

All data were tested for statistical significance using GraphPad Prism 8. Shapiro-Wilk test was used to
determine data normality. One-way ANOVA with Tukey´s test for multiple comparisons with a single
variable was implemented for cell quantifications and anatomical data. For the behavioral analysis, a

two-way ANOVA with Tukey´s test was used to test multiple comparisons with more than one variable.
* $P \leq 0.05$; ** $P \leq 0.01$; *** $P \leq 0.001$.

## 233 Results

### 234 *Structural modifications of the mouse VSE during natural and pathological aging*

We sought to investigate the structure of the VSE during natural and pathological aging, as the main
gateway of pheromone (non-volatile)-encoded information (Cheetham et al., 2007; Sharrow et al., 2002;
Ferrero et al., 2013). Stereological analysis revealed significantly smaller VSE volumes in 2 year-old
(senescent) mice (**Figure 1A**: VSE volume mm$^3$: 4 month-old: $0.190 \pm 0.008$; 1-year-old APP/PS1$^{WT}$:
$0.170 \pm 0.007$; 24 month-old: $0.160 \pm 0.007$; 1-year-old APP/PS1$^{Het}$: $0.160 \pm 0.008$). This observation
was supported by the reduced number of olfactory marker protein positive cells (OMP$^+$) along the
anterior-posterior axis of the VSE and the analysis of the marginal (M), intermediate (I) and central (C)
regions of the medial VSE (Halpern and Martinez-Marcos, 2003) in the VSE of 2-year old mice
(**Supplementary Figure 1** and **Figure 1C, E**: number of OMP$^+$ cells x10$^3$ /$\mu$m$^2$: **Anterior**: 4 month-
old: $10.69 \pm 0.56$; 12 month-old APP/PS1$^{WT}$: $9.50 \pm 0.45$; 24 month-old: $7.80 \pm 0.53$; 12 month-old
APP/PS1$^{Het}$: $8.60 \pm 0.53$; **Medial**: 4 month-old: $9.92 \pm 0.23$; 12 month-old APP/PS1$^{WT}$: $9.15 \pm 0.78$; 24
month-old: $7.70 \pm 0.70$; 12 month-old APP/PS1$^{Het}$: $8.40 \pm 0.40$; **Posterior**: 4 month-old: $9.15 \pm 0.70$;
12 month-old APP/PS1$^{WT}$: $9.53 \pm 0.77$; 24 month-old: $7.90 \pm 0.75$; 12 month-old APP/PS1$^{Het}$: $10.00 \pm$
$0.84$).

We hypothesize that the drop in the number of OMP$^+$ cells could translate into a reduced axonal
projection to the AOB, the main VNO target area, resulting in smaller AOB volumes. We observed
reduced AOB size in 2 year-old mice (**Figure 1B:** AOB volume mm$^3$: 4 month-old: $0.48 \pm 0.03$; 12
month-old APP/PS1$^{WT}$: $0.48 \pm 0.04$; 24 month-old: $0.41 \pm 0.03$; 12 month-old APP/PS1$^{Het}$: $0.44 \pm 0.03$),
indicating that aging alters basic structural features of the VNO-AOB axis.

We expanded our analysis to the SRY-box transcription factor 2 (Sox2)-expressing cells (Sox2$^+$) in the
supporting cell layer (VSE SCL), a neuronal stem cell-marker which also stain mature differentiated
sustentacular cells (Guo et al., 2010; Taroc et al., 2020; Katreddi et al., 2021). The estimation of the
labeling intensity of Sox2 in the SCL showed a drastic reduction in senescent mice but not in
APP/PS1$^{Het}$ mice (**Figure 1D, F**: Sox2 fluorescence intensity / area ($\mu$m$^2$): 4 month-old: $33.24 \pm 2.43$;
12 month-old: $26.23 \pm 3.95$; 24 month-old: $10.74 \pm 1.32$; 12 month-old APP/PS1$^{Het}$: $27.37 \pm 0.96$),
indicating natural aging may disrupt the VSE structure by reducing the number of both sensory neurons
and sustentacular cells (OMP$^+$ and Sox2$^+$).

### 262 *Natural and pathological aging differentially impact cell proliferation in the VSE*

The observed decrease in the number of mature OMP$^+$ and Sox2$^+$ cells in the VSE of senescent mice
could be due to the reduction of the VSE´s regenerative ability. Furthermore, there is currently no
available data on the status of the VSE neurogenic niche in neurodegenerative animal models. Thus, we
analyzed the number of proliferative cells identified by the expression of proliferative cell nuclear
antigen (PCNA) in the anterior, medial and posterior VNO. The marginal zone of the anterior and
medial VSE proliferation was found more abundant consistently with previous reports (Giacobini et al.,
2000; Brann and Firestein, 2010; **Figure 2A-B**). Senescent mice exhibited a significant reduction in
PCNA$^+$ cells (**Figure 2A-B**, number of PCNA$^+$ cells x 10$^3$/mm$^3$: **Anterior**: 4 month-old: $2.26 \pm 0.80$;
12 month-old APP/PS1$^{WT}$: $0.73 \pm 0.3$; 24 month-old: $0.75 \pm 0.35$; 12 month-old APP/PS1$^{Het}$: $3.42 \pm$
$0.82$; **Medial**: 4 month-old: $2.15 \pm 0.52$; 12 month-old APP/PS1$^{WT}$: $1.61 \pm 0.32$; 24 month-old: $0.23 \pm$
$0.13$; 12 month-old APP/PS1$^{Het}$: $1.57 \pm 0.20$; **Posterior**: 4 month-old: $3.04 \pm 0.90$; 12 month-old APP$^{WT}$:
$2.42 \pm 0.60$; 24 month-old: $0.17 \pm 0.08$; 12 month-old APP/PS1$^{Het}$: $1.93 \pm 0.57$), indicating reduced
proliferation. Surprisingly, the number of PCNA$^+$ cells in APP/PS1$^{Het}$ mice was significantly higher that
of aged-matched controls in the anterior VNO (**Figure 2A-B**), suggesting an increased cell turnover in

these animals. Importantly, no significant overlap between OMP and PCNA staining was observed indicating that these two markers recognize cell populations at different maturation stages (**Supplementary Figure 2**).

We then sought to explore whether the reduced number of proliferative PCNA$^+$ cells in senescent mice could be due to a decrease of stem cell generation by analyzing the number of Sox2$^+$ neural precursor cells in the VSE (Tucker et al., 2010; Panaliappan et al., 2018). We observed a significant reduction in the number of Sox2$^+$ cells in 2 year-old mice but not in APP/PS1$^{Het}$ animals, indicating a reduced ability for progenitor generation in the aged VSE (**Figure 2C**: number of Sox2$^+$ cells in the VSE, number of cells x 10$^4$ cells/$\mu m^2$: 4 month-old: 3.08 ± 0.40; 12 month-old APP/PS1$^{WT}$: 2.92 ± 0.38; 24 month-old: 2.37 ± 0.36; 12 month-old APP/PS1$^{Het}$: 2.15 ± 0.24). Altogether, these findings revealed a fundamental difference in how natural and diseased aging impact the VSE structure and proliferative capacity.

### *Late onset of deficits in social exploration during natural aging*

Our data indicated that natural aging induced significant alterations in the structure and proliferative ability of the VSE, thus we investigated whether these adaptations might translate into impairments in the processing of socio-sexual information. Cumulative evidence points to olfactory decline as a common symptom of natural and pathological aging (Doty et al., 1984; Murphy et al., 2002; Doty and Kamath, 2014; Rawson et al., 2012; Roberts et al., 2016). However, most of these studies and diagnostic tests employed synthetic odors with reduced emotional valence, therefore, the temporal course and severity of the age-related involution of the recognition of social cues, largely processed by the VNO-AOB system in the mouse, remains unclear.

To explore the impact of natural aging on social odor detection, we conducted an odor-evoked sniffing test in which serial dilutions of urine from young conspecifics of the opposite sex were presented to either male or female subjects over a range of different ages: young (2-4 mo.), adult (6-8 mo.), middle-aged (12-14 mo.) and old/senescent mice (20-24 mo.) (**Figure 3A**, see Material and Methods for details on urine sample preparation). Our results indicated a significant reduction in the exploration time throughout the various urine dilutions in comparison to adult wild type mice (**Figure 3B-C** sniffing time normalized to water control 2, **undiluted sample**: aged: 3.83 ± 0.71; adult: 6.06 ± 0.88; **1:10 dilution**: aged: 3.04 ± 0.50; adult: 4.90 ± 1.10; **1:50 dilution**: aged: 1.90 ± 0.33; adult: 3.14 ± 0.55; **1:100 dilution**: aged: 1.56 ± 0.37; adult: 2.95 ± 0.43; **1:250 dilution**: aged: 1.37 ± 0.36; adult: 2.36 ± 0.70; **1:500 dilution**: aged: 0.86 ± 0.20; adult: 2.50 ± 0.35; **1:1000 dilution**: aged: 0.60 ± 0.20; adult: 2.80 ± 0.43). Raw sniffing time data showed a similar decrease in the exploration of social odors in senescent animals across different dilutions (see details in **Supplementary Figures 3 and 4**).

Furthermore, senescent mice progressively increased the exploration time in parallel with odor concentration, whereas young, adult and middle-age animals showed a habituation phase at intermediate dilutions (1:500, 1:250, 1:100 and 1:50), indicating effective odor detection and recognition capabilities at low urine concentrations (Yang and Crawley, 2009). Data analysis disaggregated by sex showed that the reduction in exploration time was similar in male and female animals (**Supplementary Figure 5**; but see Martínez et al., 2017 for an alternative view).

### *Reduction in the exploration of social information is accelerated in an animal model of neurodegeneration*

Next, we investigated whether pathological aging would affect the exploration of social olfactory cues despite no obvious effect on VSE structure or proliferative capacity. Social odor sensitivity tests in 1 year-old APP/PS1$^{Het}$ mice revealed reduced sniffing times when compared to age-matched APP/PS1$^{WT}$ control mice for low urine dilutions, suggesting that pathological aging accelerates the decline of exploration time of social odors (**Figure 3D-E** sniffing time normalized to water control 2, **undiluted sample**: 1 year-old APP/PS1$^{Het}$: 2.58 ± 0.43; 1 year-old APP/PS1$^{WT}$: 5.65 ± 1.51; **1:10 dilution**: 1 year-old APP/PS1$^{Het}$: 1.82 ± 0.27; 1 year-old APP/PS1$^{WT}$: 4.94 ± 1.21; **1:50 dilution**: 1 year-old APP/PS1$^{Het}$:

1.43 ± 0.46; 1 year-old APP/PS1$^{WT}$: 1.40 ± 0.26; **1:100 dilution**: 1 year-old APP/PS1$^{Het}$: 1.37 ± 0.43; 1 year-old APP/PS1$^{WT}$: 2.03 ± 0.55; **1:250 dilution**: 1 year-old APP/PS1$^{Het}$: 0.90 ± 0.17; 1 year-old APP/PS1$^{WT}$: 0.70 ± 0.14; **1:500 dilution**: 1 year-old APP/PS1$^{Het}$: 0.52 ± 0.10; 1 year-old APP/PS1$^{WT}$: 0.97 ± 0.20; **1:1000 dilution**: 1 year-old APP/PS1$^{Het}$: 1.00 ± 0.20; 1 year-old APP/PS1$^{WT}$: 0.84 ± 0.31). Similar to senescent mice, these results were reproduced when raw sniffing time data were compared (**Supplementary Figures 6 and 7**).

*Aging and neurodegeneration impact social odor recognition more severely than other odor modalities*

Next, we asked whether non-social odor modalities were also affected in senescent and APP/PS1$^{Het}$ animals. We investigated this question by analyzing the exploration time to both food and synthetic neutral odors. First, we exposed naturally aged and APP/PS1$^{Het}$ mice to serial dilutions of isoamyl acetate (IA), a synthetic banana like odor of neutral valence when used at high dilutions (Saraiva et al., 2016; Fortes-Marco, 2015). Consistent with previous studies, mice of all conditions showed significantly reduced responses to the neutral odorant than to urine (**Figure 4A-D**) according to the higher emotional valence of urine in comparison to a synthetic odor (Jagetia et al. 2018; Kobayakawa et al. 2007; Mandairon et al. 2009; Saraiva et al. 2016). Our results showed no significant differences on the exploration times of IA in senescent mice and a modest but significant increase in the sniffing time of the 1:10$^4$ and 1:100 IA dilutions in APP/PS1$^{Het}$ animals (**Figures 4A-D,** normalized sniffing time, **1:100 dilution**: 1 year-old APP/PS1$^{Het}$: 1.40 ± 0.12; 1 year-old APP/PS1$^{WT}$: 0.93 ± 0.16; young: 1.76 ± 0.26; aged: 1.24 ± 0.20;  1:1000 dilution: 1 year-old APP/PS1$^{Het}$: 1.40 ± 0.22; 1 year-old APP/PS1$^{WT}$: 1.42 ± 0.30; young: 1.90 ± 0.35; aged: 1.56 ± 0.50; **1:10$^4$ dilution**: 1 year-old APP/PS1$^{Het}$: 1.50 ± 0.25; 1 year-old APP/PS1$^{WT}$: 0.91 ± 0.13; young: 1.44 ± 0.32; aged: 1.20 ± 0.26; **1:10$^5$ dilution**: 1 year-old APP/PS1$^{Het}$: 0.75 ± 0.11; 1 year-old APP/PS1$^{WT}$: 0.75 ± 0.15; young: 1.41 ± 0.50; aged: 0.90 ± 0.20; **1:5x10$^5$ dilution**: 1 year-old APP/PS1$^{Het}$: 0.85 ± 0.18; 1 year-old APP/PS1$^{WT}$: 0.82 ± 0.11; young: 1.65 ± 0.42; aged: 1.23 ± 0.35). Similarly, the food finding test showed no significant differences in the latency to find food pellets after 24 hours of food deprivation in either naturally aged or middle age APP/PS1$^{Het}$ mice (**Figure 4E** latency of finding food pellet (min), young: 5.22 ± 1.08; aged: 5.15 ± 0.95; 1-year-old APP/PS1$^{Het}$: 4.27 ± 0.58; 1-year-old APP/PS1$^{WT}$: 3.87 ± 0.32). These results suggest that deficits in social exploration time at advanced stages of natural aging and in an animal model of AD are more pronounced than other odor modalities.

*Social odor discrimination and habituation are reduced in naturally aged and APP/PS1$^{Het}$ mice*

To further investigate the impact of aging and neurodegeneration in the detection of social information, we performed a habituation – dishabituation test, which relies on the animal's ability to discriminate novel smells (Yang and Crawley, 2009). For these experiments, young (2-4 mo.) and aged (20-24 mo.) wild type animals were presented three consecutive replicates of urine samples from two different animals (S1a-c and S2a-c) (**Figure 5A**). Aged animals were able to discriminate between urine sources, but showed reduced sniffing times during both the first and second discrimination and habituation phases (**Figure 5B-C**). Similarly, middle age APP/PS1$^{Het}$ mice also exhibited reduced sniffing times in comparison to aged-matched APP/PS1$^{WT}$ controls (**Figures 5B-E** sniffing time in seconds, **water control 1**: aged: 0.56 ± 0.09; young: 0.42 ± 0.05; APP/PS1$^{WT}$: 0.72 ± 0.08; APP/PS1$^{Het}$: 0.57 ± 0.06; **water control 2**: aged: 0.56 ± 0.11; young: 0.67 ± 0.10; APP/PS1$^{WT}$: 0.50 ± 0.07; APP/PS1$^{Het}$: 0.71 ± 0.09; **S1a**: aged: 1.36 ± 0.27; young: 2.45 ± 0.40; APP/PS1$^{WT}$: 2.81 ± 0.40; APP/PS1$^{Het}$: 1.75 ± 0.22; **S1b**: aged: 0.70 ± 0.14; young: 0.87 ± 0.20; APP/PS1$^{WT}$: 0.95 ± 0.10; APP/PS1$^{Het}$: 0.68 ± 0.10; **S1c**: aged: 0.70 ± 0.21; young: 0.51 ± 0.10; APP/PS1$^{WT}$: 0.84 ± 0.22; APP/PS1$^{Het}$: 0.71 ± 0.13; **S2a**: aged: 1.51 ± 0.27; young: 2.44 ± 0.43; APP/PS1$^{WT}$: 1.85 ± 0.42; APP/PS1$^{Het}$: 1.00 ± 0.17; **S2b**: aged: 0.82 ± 0.14; young: 0.81 ± 0.15; APP/PS1$^{WT}$: 0.80 ± 0.22; APP/PS1$^{Het}$: 0.62 ± 0.08; **S2c**: aged: 0.96 ± 0.18; young: 0.55 ± 0.11); APP/PS1$^{WT}$: 0.60 ± 0.13; APP/PS1$^{Het}$: 0.50 ± 0.09). Analysis of the slope values of the first and second discrimination and habituation phases revealed that senescent mice exhibited

significant deficits during both rounds of social habituation – dishabituation, and APP/PS1[Het] mice only
during the second phase of discrimination (**Supplementary Figures 8 and 9**).

Then, we adapted the habituation-dishabituation test to assay potential changes in long-term social odor
memory (**Figure 5F**). To this aim, animals were exposed to the same S1a urine sample 24 hours after
the first presentation. A reduction of the sniffing time during the first discrimination phase was
interpreted as an indicator of memory. This reduction was clearly observed in young animals, but absent
in naturally aged and 1-year-old APP/PS1[Het] mice (**Figure 5G:** sniffing time of S1a after 24 hours in
seconds, young: $1.20 \pm 0.20$; aged: $1.12 \pm 0.20$; APP/PS1[WT]: $1.40 \pm 0.50$; APP/PS1[Het]: $1.20 \pm 0.33$),
suggesting long-term social odor memory impairments during both natural and pathological aging.

*Age-related deficits in social discrimination and habituation are not influenced by previous experience*

Next, we compared animals exposed to either familiar (littermate urine, L) or novel social odors (novel
urine, N) in the habituation-dishabituation test (**Figure 6A**). Aged animals performed poorly in the L-
N dishabituation task (no significant increase in sniffing time), suggesting that the reduction in social
odor discrimination and habituation is independent of previous experience.

Since impairments in social odor discrimination were more prominent in naturally aged mice, we asked
whether these deficits also extended to the recognition of animal's own odors, as the loss of self-
awareness is a disrupting symptom of common occurrence in senescent subjects (La Joie et al., 2016;
Valech et al., 2018). To this aim, animals were presented to samples of their own urine (O) during the
second discrimination phase of the habituation-dishabituation test (**Figure 6, C-D**). Our data indicated
that animal´s own urine was effective to elicit a typical discrimination (dishabituation) response,
(**Figure 6, C-D**: sniffing time in seconds, **water control 1**: aged: $0.70 \pm 0.10$; young: $0.81 \pm 0.10$; **water
control 2**: aged: $0.31 \pm 0.09$; young: $0.48 \pm 0.11$; **La**: aged: $2.07 \pm 0.43$; young: $3.06 \pm 0.35$; **Lb**: aged:
$0.64 \pm 0.16$; young: $0.74 \pm 0.17$; **Lc**: aged: $0.48 \pm 0.10$; young: $0.45 \pm 0.10$; **Oa**: aged: $1.23 \pm 0.18$;
young: $1.22 \pm 0.33$; **Ob**: aged: $0.48 \pm 0.13$; young: $0.84 \pm 0.13$; **Oc**: aged: $0.49 \pm 0.11$; young: $0.55 \pm
0.15$), suggesting that self-recognition is overall preserved in senescent mice.

*Social novelty is disrupted during pathological aging*

Last, we explored whether the observed impairments in social odor exploration and discrimination may
negatively impact social interaction in naturally aged and APP/PS1[Het] mice. To this aim, we performed
a three-chamber test (Nadler et al., 2004; Yang et al., 2011; **Figure 7A**) to assess general sociability
and social novelty in senescent and APP/PS1[Het] animals. Our data indicated a reduction in social novelty
in middle age APP1/PS1[Het] mice, which was not detected in 2 year-old animals (**Figure 7B-E**,
**Sociability**, sniffing time in seconds, **E**: aged: $14.50 \pm 1.90$; young: $25.06 \pm 2.96$; APP/PS1[WT]: $39.15 \pm
3.63$; APP/PS1[Het]: $41.00 \pm 4.07$; **M1[A]**: aged: $47.24 \pm 4.60$; young: $64.35 \pm 5.23$; APP/PS1[WT]: $69.40 \pm
6.05$; APP/PS1[Het]: $69.73 \pm 7.10$; **Social novelty**, sniffing time in seconds, **M1[B]**: aged: $49.15 \pm 4.20$;
young: $62.37 \pm 5.01$; APP/PS1[WT]: $71.30 \pm 5.75$; APP/PS1[Het]: $65.12 \pm 6.70$; **M2**: aged: $63.70 \pm 5.30$;
young: $99.93 \pm 9.83$; APP/PS1[WT]: $93.07 \pm 11.70$; APP/PS1[Het]: $77.02 \pm 8.43$). Although senescent mice
did not show significant impairments in either sociability or social novelty, they exhibited a mild
increase in the latency to approach M1 during the sociability phase (**Supplementary Figure 10**),
consistent with the overall decrease in the exploration time of social odors (**Figures 3, Supplementary
Figures 3 and 4**). These findings exposed exacerbated social deficits in an animal model of AD,
suggesting a distinct impact of natural and pathological aging on social behavior.

**Discussion**

The quality of social life has been proposed as a predictive factor for developing dementia or mental
illness (Cohen et al., 1985; Kawachi and Berkman, 2001; Stockdale et al., 2007; Moak and Agrawal,
2010; Wilson et al., 2007). However, the impairment of social functions with age is poorly understood

with some authors suggesting that social deficits might be the consequence of a more generalized brain
impairment (Bottiroli et al., 2016; Johansson et al., 2018), but also an early manifestation of AD (Moran
et al., 2012; Cavallini et al., 2013) which might be developed independently of cognitive deficits (Mohs
et al., 2000; Spalletta et al., 2004).

Our results revealed that both natural and pathological aging affect several key aspects of social
information processing in mice including the exploration of social odors, social odor discrimination and
habituation. To gain insight into the mechanisms underlying these deficits, we explored the age-related
adaptions of the VSE, a central gateway for pheromone-encoded information in mammals (Jacobson,
1813; Halpern and Martínez-Marcos, 2003; Munger et al., 2009). The accessory olfactory system has
been indeed largely overlooked in contrast to other aspects of olfaction, whose functional decline has
been suggested as a potential biomarker of neurodegenerative disorders like AD (recently reviewed in
Murphy, 2019).

Our data showed VSE anatomical alterations during natural aging but not in APP/PS1$^{Het}$ mice, a
standardized animal model of AD. Whereas the VSE of APP/PS1$^{Het}$ mice maintained fairly stable
neurogenic capabilities, senescent wild type animals showed a significant reduction in proliferative and
stem cells (PCNA$^+$ and Sox2$^+$ VSE cells) (**Figure 2**) which may explain the reduction of mature OMP$^+$
neurons, sustentacular Sox2$^+$ cells and VSE volume (**Figure 1** and **Supplementary Figure 1**). Our
findings are consistent with two previous studies directly exploring the effects of natural aging in the
VSE (Brann and Firestein, 2010; Mechin et al., 2021), which reported an overall thinning of the
vomeronasal sensory epithelium (Mechin et al., 2021) and reduced VSE neurogenesis in 2-year-old
animals (Brann and Firestein, 2010). Here, we expand this current knowledge by providing novel
information on the properties of the VSE in a commonly used animal model of AD (APP/PS1$^{Het}$),
revealing an unexpected preservation of the organ and its proliferative capabilities in middle age
APP/PS1$^{Het}$ mice. These results contrast with the reduction in SVZ neurogenesis reported in similar
animal models (Scopa et al., 2020; Zeng et al., 2016; Verret et al., 2007; Zhang et al., 2007), suggesting
that VSE neurogenesis might be less susceptible to lesions (Brann and Firestein, 2010) and pathological
conditions than other proliferative areas. However, we cannot rule out the possibility that APP/PS1$^{Het}$
animals may develop potential alterations in the VSE structure and cell composition at a later age (> 1
year-old). An aspect that we were prevented to investigate due to the life expectancy (~ 14-16 months-
old) of this mouse model.

A relevant aspect of our work is that most previous studies addressing olfactory decline during natural
and pathological aging employed synthetic and neutral odors, thus remaining uncertain the effect of
healthy and diseased aging in the recognition of social cues. To gain insight into this question, we
investigated the exploration time and the habituation-dishabituation response to social odors (urine)
(**Figures 3, 5**). Our findings revealed that despite the distinctive effects of natural and pathological
aging in VSE structure and neurogenic capability, both processes affected the exploration of socio-
sexual information, social discrimination-habituation, and social behavior, suggesting fundamental
differences in how these processes impact social information encoding.

Further experiments are needed to establish causality, but our results suggest that the observed VSE
alterations in 2-year-old mice may elicit deficits in urine exploration time and the lack of habituation to
intermediate urine dilutions (**Figure 3; Supplementary Figures 3 and 4**). VSE volume data from
middle age APP/PS1$^{WT}$ control animals may be suggestive that VSE structural changes might start to
appear at a younger age (**Figure 1A**: VES volume mm$^3$: 4 month-old: 0.190 ± 0.008; 1-year-old
APP/PS1$^{WT}$: 0.170 ± 0.007; p = 0.03; two-way ANOVA with Tuckey´s test), although their functional
consequences might not become apparent until advanced stages of aging (2-year-old). This scenario
suggests an aged-related parsimonious decay of the VNO that matches the reported effect of natural
aging in other olfactory areas like the OB. Several studies have shown that despite the reduced
regeneration rate of olfactory sensory neurons (OSNs), OB alterations involving a reduced number of

synaptic contacts (Richard et al., 2010), expression defects of odorant receptor genes (Lee et al., 2009;
Khan et al., 2013) and changes in the OSNs dynamic range (Kass et al., 2018) are of late occurrence
and just becoming obvious in 2-year-old mice. Although the olfactory system may undergo age-related
changes from middle age (Mobley et al., 2014), functional deficits during natural aging might exhibit a
late onset, suggesting compensatory mechanisms to preserve the processing of olfactory cues like social
information relevant for the survival of aged animals (Tobiansky et al., 2012; Tikhonova et al., 2015).
In contrast to the steadily decline of natural aging, pathological conditions may accelerate functional
deficits even in the absence of obvious structural changes of peripheral organs like the VNO, suggesting
impairments in the central processing of social information. In this regard, defects in the exploration of
social odors were found exacerbated in APP/PS1[Het] mice, which exhibited significant impairments
already at middle age (**Figure 3, D-E**), a condition likely to aggravate frailty and reduce life expectancy
in this animals (as observed in this mouse strain in our experimental conditions).

Furthermore, a detailed analysis of the first discrimination phase of the modified social habituation-
dishabituation (presentation of S1a urine sample after 24 hours; **Figure 5, F-G**) revealed significant
impairments in both 2-year old wild type animals and middle age APP/PS1[Het] mice, suggesting that in
parallel to peripheral sensory decline, the downstream pathways involved in social cue recognition may
be affected during both natural and pathological aging. To control for a potential contribution of novelty
in the social habituation-dishabituation test, we presented urine samples from littermate and novel
animals finding similar deficits in discrimination (**Figure 6**). Interestingly, although senescent animals
presented discrimination deficits they preserved the ability to differentiate between urine from a novel
subject and their own, preventing to explore further the underlying mechanisms of subjective perception
loss, a disrupting symptom of senescence and dementia which currently lacks appropriate animal
models for preclinical studies (Jessen et al., 2020).

Age-related decline of olfactory detection is likely to occur due to alterations of both the main olfactory
epithelium and the VNO (Apfelbach et al., 1991; Lee et al., 2009; Ueha et al., 2018), thus potentially
affecting other odor modalities. Here, we sought to explore this question by performing odor-evoked
sniffing tests employing social and neutral synthetic odors (**Figure 4**). Quantification of the exploration
time across various IA dilutions exposed equivalent measurements in senescent and young mice, and
even enhanced responses in the case of APP/PS1[Het] mice. Furthermore, results from a food seeking test,
exposed no significant differences in the latency time to find the hidden food pellets (**Figure 5E**)
indicating that defects in social odor detection related to natural and pathological aging may be more
severe than other odor modalities. A plausible explanation for the specific impairment in the exploration
of social odors of aged and APP/PS1[Het] animals may involve the reduction of reproductive activity in
old animals, but not in foraging and feeding behaviors (Harb et al., 2014). The decline to detect specific
odors may rise from alterations on the number of sensory neurons and synaptic connections specifically
related to the discrimination of conspecifics (Wacker et al., 2010; Bielsky et al., 2005; Tobin et al.,
2010) and heterospecific social cues (e.g. predator) (Kang et al., 2009; Wacker et al., 2010). In
particular, impairments in social odor exploration and discrimination may rise from maladaptations
specific to the VNO-AOB axis, consistent with the observed decrease in AOB volume in senescent
animals (**Figure 1B**). Future studies should deepen into the anatomical and functional properties of the
VNO-AOB circuit in order to obtain a complete picture on how healthy and pathological aging affect
social information processing.

Finally, we sought to determine how the age-related defects in social odor sensitivity, discrimination
and memory shaped social behavior. Results from a three chamber test (**Figure 7**) indicated that
sociability was overall preserved in senescent and APP/PS1[Het] animals. In contrast, social novelty was
severely impaired in APP/PS1[Het] mice consistently with previous studies (Cheng et al., 2014; Olesen et
al., 2016; Locci et al., 2021; Shoji and Miyakawa, 2019). Deficits in social novelty in APP/PS1[Het] mice
could be linked to the deficits in social discrimination observed in the habituation-dishabituation test
(**Figure 5D, E** and **Supplementary Figure 8;** Enwere et al., 2004; Patel and Larson, 2009; Moreno et

al., 2014) which could impair the recognition of M2 as a novel subject. Consistent with the overall
decrease in the exploration time of social odors (**Figures 3, Supplementary Figures 3 and 4**), aged
animals exhibited a mild increase in the latency to approach the novel mouse (M1) during the first
sociability phase (**Supplementary Figure 10**). These findings indicate that despite the reduction in
social odor exploration and discrimination, overall sociability and novelty are majorly preserved in
naturally aged mice. In contrast, the APP/PS1 neurodegenerative model shows a measurable
impairment in social preference. This evidence opens the intriguing possibility that neuronal circuits
underlying specific social functions (sociability vs social memory) may be particularly susceptible to
pathological aging.

The present study offers new evidence on how natural and pathological aging distinctively impact the
pheromone detection system and different aspects of social information processing, a necessary first
step towards developing novel strategies to reduce the deterioration of olfactory deficits and social
behavior in aged subjects.

**Figure legends**

**Figure 1. Structural modifications of the mouse VSE during natural and pathological aging. A)**
Dispersion plot of the VSE volume indicates a significant reduction during natural but not pathological
aging. **B)** Dispersion plot of AOB volume. Thick lines indicate the mean ± SEM. Data were obtained
from 4 animals per condition. **C)** Representative images of OMP staining in the VSE during natural and
pathological aging. Scale bar indicates 100 μm. **D)** Representative images of Sox2 staining in the VSE
and the supporting cell layer. Scale bar indicates 100 μm. VL: vomeronasal lumen; SCL: supporting
cell layer; NSE: non-sensory epithelium; M: marginal zone; I: intermediate zone; C: central zone. **E, F)**
Dispersion plots represent the number of OMP$^+$ cells in the VSE (number of cells x $10^3$/μm$^3$) and the
intensity of Sox2 fluorescence in the SCL (Sox2 CTF /μm$^2$). Thick lines indicate the mean ± SEM. Data
were obtained from 10-16 slices from at least three animals per condition. Statistical comparisons were
calculated by two-way ANOVA with Tukey´s test. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001.

**Figure 2. Natural and pathological aging differentially impact cell proliferation in the VSE. A)**
Representative confocal images of PCNA staining in the VSE proliferative niche during natural and
pathological aging. To note there is no overlapping signal between the two markers indicating the
identification of cells at different maturation stages (**Supplementary Figure 2**). Circles indicate the
proliferative marginal region of the medial VNO. Scale bar represents 20 μm. **B, C)** Dispersion plots
represent the number of PCNA$^+$ cells in the VSE (number of cells x $10^3$/mm$^3$) and the number of Sox2$^+$
cells in the VSE (number of cell x $10^4$/μm$^2$). Thick lines indicate the mean ± SEM. Data were obtained
from ten slices from animal of each condition. Statistical analysis was calculated by two-way ANOVA
with Tukey´s test for multiple comparisons. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001.

**Figure 3. Natural and pathological aging reduce exploration to social odors. A)** Schematics of the
olfactory test used in this study in which urine dilutions are presented as a social signal. **B)** Average of
the exploration (sniffing) time of urine serial dilutions normalized to the exploration time of the vehicle
(water) of adult and aged wild type mice. A typical habituation was observed in adult mice at
intermediate dilutions (1:500, 1:250, 1:100; 1:50). **C)** Dispersion plot of the normalized sniffing time
of each urine dilution (nd, non-diluted; 1:10; 1:50; 1:100; 1:250; 1:500; 1:1000) for adult and aged wild
type mice. **D)** Average of the sniffing time of urine dilutions normalized to the exploration time of the
vehicle (water) of middle age APP/PS1 controls (APP$^{WT}$) and middle age APP/PS1 heterozygous mice
(APP/PS1$^{Het}$). **E)** Dispersion plots of normalized sniffing time of middle age APP$^{WT}$ control and
APP/PS1$^{Het}$ mice. Grey lines in the dispersion plots indicate mean ± SEM. Data were analyzed by a one-
way ANOVA with Tukey´s test to test multiple comparisons with more than one variable. * P ≤ 0.05;
** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.

**Figure 4. Aging and neurodegeneration impact social odor recognition more severely than other odor modalities. A)** Average of the sniffing time in response to a neutral synthetic odor (isoamyl acetate, IA) normalized to the vehicle (mineral oil) of young and aged wild type mice. **B)** Dispersion plots of normalized sniffing time in response to various dilutions of IA in young and aged wild type mice. **C)** Average of the sniffing time in response to a neutral synthetic odor (isoamyl acetate, IA) normalized to the vehicle (mineral oil) of middle age APP/PS1 controls (APP^WT) and middle age APP/PS1 heterozygous mice (APP/PS1^Het). **D)** Dispersion plots of normalized sniffing time in response to various dilutions of IA in APP/PS1^WT and APP/PS1^Het mice. Note that APP/PS1^Het mice exhibit significantly longer exploration times for the $1:10^4$ and $1:100$ IA dilutions. **E)** Food deprived animals performed a food seeking test (schematics on the right) which revealed equivalent latencies to find hidden food pellets as shown in the dispersion data plot (left panel). Thick lines in dispersion plots indicate mean ± SEM. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more than one variable. Number of mice: *Neutral odor sensitivity*: young: n = 19; aged: n = 27; middle age control: n = 10; middle age APP/PS1^Het: n = 13. *Food finding test*: young: n = 21; aged: n = 27; APP^WT: n = 20; APP^Het: n = 16. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.

**Figure 5. Social odor discrimination and habituation are reduced in naturally aged and APP/PS1^Het mice. A)** Schematics of the social habituation-dishabituation test used in this study: after habituation to the experimental cage, the animal is exposed to the same urine sample three times (S1a-c) which induces a typical increase in exploration time (1st discrimination) to be followed by a reduction in the sniffing time (1st habituation). Dishabituation induced by a urine sample from a new subject (S2a) elicits a second round of discrimination - habituation. **B)** Average sniffing time of the social habituation-dishabituation test performed by young and aged wild type mice. **C)** Dispersion plot of the sniffing time corresponding to the social habituation-dishabituation test of young and aged animals. **D)** Average sniffing time of the social habituation-dishabituation test performed by middle age APP/PS1^WT control and APP/PS1^Het mice. **E)** Dispersion plot of the sniffing time corresponding to the social habituation-dishabituation test of middle age APP/PS1^WT control and APP/PS1^Het mice. **F)** The habituation-dishabituation test was modified to assess potential deficits in long-term discrimination due to natural or pathological aging by presenting the same S1a sample 24 hours after. **G)** Paired data corresponding to the sniffing time during the first and second (24 hours later) S1a presentation is plotted for each condition. Thick lines in dispersion plots and black dots in G indicate mean ± SEM. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more than one variable. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.

**Figure 6. Age-related deficits in social discrimination and habituation are not influenced by previous experience. A)** Average sniffing time of social habituation-dishabituation test in response to odors from novel (N) or littermate (L) subjects. **B)** Dispersion plot of the exploration time of aged and young animals in response to novel or littermate urine samples. **C)** Average sniffing time of social habituation-dishabituation test in response to littermate and the animal´s own urine (O). **D)** Dispersion plot of exploration time of aged and young animals in response to their own´s or littermate urine samples. Thick lines in dispersion plots indicate mean ± SEM. Number of mice: young: n = 21; aged: n = 23. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more than one variable. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.

**Figure 7. Social novelty is disrupted during pathological aging. A)** Schematics of the three chamber test used in this study to test sociability and social novelty. In the sociability phase the sniffing time of E and M1^A are compared. Social novelty is estimated by quantifying the sniffing time of exploring M1^B versus M2. **B)** Dispersion plots of the sniffing times of E-M1^A (sociability) and M1^B-M2 (social novelty) for young wild type mice. **C)** Dispersion plots of the sniffing times of E-M1^A (sociability) and M1^B-M2 (social novelty) for aged wild type mice. **D)** Dispersion plots of the sniffing times of E-M1^A (sociability) and M1^B-M2 (social novelty) for middle age APP^WT mice. **E)** Dispersion plots of the

sniffing times of E-M1^A (sociability) and M1^B-M2 (social novelty) for middle age APP^Het mice. Black
dots in the paired plots indicate mean ± SEM. Number of mice: young: n = 10; aged: n = 14; middle
age control: n = 10; middle age APP/PS1^Het: n = 12. Data were analyzed by a one-way ANOVA with
Tukey´s test to test multiple comparisons with more than one variable. * P ≤ 0.05; ** P ≤ 0.01; *** P
≤ 0.001; **** P ≤ 0.0001.

**Supplementary Figure 1 (related to Figure 1). Structural modifications of the mouse VSE and**
**AOB during natural and pathological aging. A)** Schematics depicting the VNO divisions into
marginal (M), intermediate (I) and central (C). **B-D)** Dispersion plots of the number of OMP^+ cells in
the marginal, intermediate and central zone of the VSE. Thick lines in the dispersion plots indicate
mean ± SEM. Data were obtained from 10-16 slices from at least three animals per condition. Statistical
comparisons were calculated by one-way ANOVA with Tukey´s test for multiple comparisons with a
single variable. P value < 0.05 is considered significant. P values are indicated in the plot.

**Supplementary Figure 2 (related to Figure 2). The proliferative niche of the VSE is reduced**
**during natural, but not pathological aging.** Representative confocal images of both OMP and PCNA
staining in the VSE proliferative niche. To note there is no overlapping signal between the two markers
indicating the identification of cells at different maturation stages. Circles indicate the proliferative
marginal region of the VSE. Scale bar represents 20 μm. Data were obtained from ten slices from animal
of each condition.

**Supplementary Figure 3 (related to Figure 3). Statistical comparison of the exploration time**
**between different ages for each dilution.** Dispersion plots represent the sniffing time of 2, 6, 12 and
24 months-old animals for the various serial dilutions tested in Figure 3. Thick grey lines indicate mean
± SEM. Number of mice for each condition and the P values for each statistical comparison is indicated
in the tables below. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple
comparisons with more than one variable.

**Supplementary Figure 4 (related to Figure 3). Statistical comparison of exploration time between**
**different dilutions for each analyzed age.** Dispersion plots represent the sniffing time and trend curves
of 2, 6, 12 and 24 months-old animals for the various serial dilutions tested in Figure 3. Thick dots
indicate mean ± SEM. P values for each statistical comparison is indicated in the tables on the right.
Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more
than one variable.

**Supplementary Figure 5 (related to Figure 3). Aged-related deficits in social odor exploration**
**exhibit no significant sex differences. A)** Average exploration (sniffing) time of urine serial dilutions
of adult and aged female wild type mice. **B)** Average exploration (sniffing) time of urine serial dilutions
of adult and aged male wild type mice. **C)** Dispersion plots of the sniffing time of urine dilutions that
show significant impairments in female mice (nd, non-diluted; 1:10; 1:500; 1:1000). **D)** Dispersion
plots of the sniffing time of the urine dilutions that show significant changes in male mice (nd, non-
diluted; 1:250; 1:500; 1:1000). Plots indicate mean ± SEM. Number of mice: adult females: n = 13;
adult males: n = 11; aged females: n = 14; aged males: n = 11. Data were analyzed by a one-way
ANOVA with Tukey´s test to test multiple comparisons with more than one variable. * P ≤ 0.05; ** P
≤ 0.01; *** P ≤ 0.001.

**Supplementary Figure 6 (related to Figure 3). Statistical comparison of the exploration time**
**between APP/PS1^WT and APP/PS1^Het for each dilution.** Dispersion plots represent the sniffing time
middle age of APP^WT and APP^Het mice for the various serial dilutions tested in Figure 3. Thick grey
lines indicate mean ± SEM. Number of mice for each condition and the P values for each statistical
comparison is indicated in the tables below. Data were analyzed by a one-way ANOVA with Tukey´s
test to test multiple comparisons with more than one variable.

**Supplementary Figure 7 (related to Figure 3). Statistical comparison of the exploration time between different dilutions for APP/PS1$^{WT}$ and APP/PS1$^{Het}$ mice.** Dispersion plots represent the sniffing time and trend curves of middle age APP$^{WT}$ and APP$^{Het}$ mice for the various serial dilutions tested in Figure 3. Thick dots indicate mean ± SEM. P values for each statistical comparison is indicated in the tables on the right. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more than one variable.

**Supplementary Figure 8 (related to Figure 5). Social discrimination is reduced during natural and pathological aging. A)** Dispersion plots of the sniffing time of the first discrimination phase (H$_2$O-S1a) for young and aged wild type animals. **B)** Dispersion plots of the sniffing time of the second discrimination phase (S1c-S2a) for young and aged wild type animals. **C)** Dispersion plots of the slope values of the first and second discrimination phases of young and aged mice. **D)** Dispersion plots of the sniffing time of the first discrimination phase (H$_2$O-S1a) for of APP$^{WT}$ and APP$^{Het}$ mice. **E)** Dispersion plots of the sniffing time of the second discrimination phase (S1c-S2a) for of APP$^{WT}$ and APP$^{Het}$ mice. **F)** Dispersion plots of the slope values of the first and second discrimination phases of APP$^{WT}$ and APP$^{Het}$ mice. Black dots in **A, B, D**, and **E** and thick lines in **C** and **F** indicate mean ± SEM. Data were analyzed by one-way ANOVA with Tukey´s test. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001.

**Supplementary Figure 9 (related to Figure 5). Social habituation is diminished during natural and pathological aging. A)** Dispersion plots of the sniffing time of the first habituation phase (S1a-S1b) for young and aged wild type animals. **B)** Dispersion plots of the sniffing time of the second habituation phase (S2a-S2b) for young and aged wild type animals. **C)** Dispersion plots of the slope values of the first and second habituation phases of young and aged mice. **D)** Dispersion plots of the sniffing time of the first habituation phase (S1a-S1b) for of APP$^{WT}$ and APP$^{Het}$ mice. **E)** Dispersion plots of the sniffing time of the second habituation phase (S2a-S2b) for of APP$^{WT}$ and APP$^{Het}$ mice. **F)** Dispersion plots of the slope values of the first and second habituation phases of APP$^{WT}$ and APP$^{Het}$ mice. Black dots in **A, B, D**, and **E** and thick lines in **C** and **F** indicate mean ± SEM. Data were analyzed by one-way ANOVA with Tukey´s test. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001.

**Supplementary Figure 10 (related to Figure 7). Latency to approach the first stranger (M1) in the three chamber sociability test is increased in naturally aged animals. A)** The sociability phase of the three chamber test involves exposing the test animal to a stranger mouse (M1) for the first time, which elicits an increase in the exploration time in that side of the chamber. B) Dispersion plots of the latency to approach M1 during the first encounter of the sociability phase. Thick lines indicate mean ± SEM. Data were analyzed by one way-ANOVA with Tukey´s test. * P ≤ 0.05.

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**Author contributions**

The project was led by S.J. Brain sample preparation, data acquisition, and analysis was done by A.P. Result interpretation, manuscript preparation, and editing was done by S.J., P.C. and A.P. with help of members from the Jurado Lab at the Instituto de Neurociencias (CSIC-UMH).

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**Figure 1. Structural modifications of the mouse VSE during natural and pathological aging. A)** Dispersion plot of the VSE volume indicates a significant reduction during natural but not pathological aging. **B)** Dispersion plot of AOB volume. Thick lines indicate the mean ± SEM. **C)** Representative images of OMP staining in the VSE during natural and pathological aging. Scale bar indicates 100 μm. **D)** Representative images of Sox2 staining in the VSE and the supporting cell layer. Scale bar indicates 100 μm. VL: vomeronasal lumen; SCL: supporting cell layer; NSE: non-sensory epithelium; M: marginal zone; I: intermediate zone; C: central zone. **E, F)** Dispersion plots represent the number of OMP+ cells in the VSE (number of cells x $10^3/\mu m^3$) and the intensity of Sox2 fluorescence in the SCL (Sox2 CTF $/\mu m^2$). Thick lines indicate the mean ± SEM. Data were obtained from 10-16 slices from at least three animals per condition. Statistical comparisons were calculated by two-way ANOVA with Tukey´s test. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001.

**Figure 2. Natural and pathological aging differentially impact cell proliferation in the VSE. A)** Representative confocal images of PCNA staining in the VSE proliferative niche during natural and pathological aging. To note there is no overlapping signal between the two markers indicating the identification of cells at different maturation stages (**Supplementary Figure 2**). Circles indicate the proliferative marginal region of the medial VNO. Scale bar represents 20 μm. **B, C)** Dispersion plots represent the number of PCNA⁺ cells in the VSE (number of cells x 10³/mm³) and the number of Sox2⁺ cells in the VSE (number of cell x 10⁴/μm²). Thick lines indicate the mean ± SEM. Data were obtained from ten slices from animal of each condition. Statistical analysis was calculated by two-way ANOVA with Tukey´s test for multiple comparisons. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001.

**Figure 3. Natural and pathological aging reduce exploration to social odors. A)** Schematics of the olfactory test used in this study in which urine dilutions are presented as a social signal. **B)** Average of the exploration (sniffing) time of urine serial dilutions normalized to the exploration time of the vehicle (water) of adult and aged wild type mice. A typical habituation was observed in adult mice at intermediate dilutions (1:500, 1:250, 1:100; 1:50). **C)** Dispersion plot of the normalized sniffing time of each urine dilution (nd, non-diluted; 1:10; 1:50; 1:100; 1:250; 1:500; 1:1000) for adult and aged wild type mice. **D)** Average of the sniffing time of urine dilutions normalized to the exploration time of the vehicle (water) of middle age APP/PS1 controls (APP^WT) and middle age APP/PS1 heterozygous mice (APP/PS1^Het). **E)** Dispersion plots of normalized sniffing time of middle age APP^WT control and APP/PS1^Het mice. Grey lines in the dispersion plots indicate mean ± SEM. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more than one variable. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.

# NEUTRAL ODOR EXPLORATION

# FOOD FINDING TEST

Figure 4. Aging and neurodegeneration impact social odor recognition more severely than other odor modalities. A) Average of the sniffing time in response to a neutral synthetic odor (isoamyl acetate, IA) normalized to the vehicle (mineral oil) of young and aged wild type mice. B) Dispersion plots of normalized sniffing time in response to various dilutions of IA in young and aged wild type mice. C) Average of the sniffing time in response to a neutral synthetic odor (isoamyl acetate, IA) normalized to the vehicle (mineral oil) of middle age APP/PS1 controls (APP[WT]) and middle age APP/PS1 heterozygous mice (APP/PS1[Het]). D) Dispersion plots of normalized sniffing time in response to various dilutions of IA in APP/PS1[WT] and APP/PS1[Het] mice. Note that APP/PS1Het mice exhibit significantly longer exploration times for the 1:104 and 1:100 IA dilutions. E) Food deprived animals performed a food seeking test (schematics on the right) which revealed equivalent latencies to find hidden food pellets as shown in the dispersion data plot (left panel). Thick lines in dispersion plots indicate mean ± SEM. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more than one variable. Number of mice: Neutral odor sensitivity: young: n = 19; aged: n = 27; middle age APP[WT]: n = 10; middle age APP/PS1[Het]: n = 13. Food finding test: young: n = 21; aged: n = 27; APP[WT]: n = 20; APP[Het]: n = 16. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.

**Figure 5. Social odor discrimination and habituation are reduced in naturally aged and APP/PS1[Het] mice. A)** Schematics of the social habituation-dishabituation test used in this study: after habituation to the experimental cage, the animal is exposed to the same urine sample three times (S1a-c) which induces a typical increase in exploration time (1st discrimination) to be followed by a reduction in the sniffing time (1st habituation). Dishabituation induced by a urine sample from a new subject (S2a) elicits a second round of discrimination - habituation. **B)** Average sniffing time of the social habituation-dishabituation test performed by young and aged wild type mice. **C)** Dispersion plot of the sniffing time corresponding to the social habituation-dishabituation test of young and aged animals. **D)** Average sniffing time of the social habituation-dishabituation test performed by middle age APP/PS1[WT] control and APP/PS1[Het] mice. **E)** Dispersion plot of the sniffing time corresponding to the social habituation-dishabituation test of middle age APP/PS1[WT] control and APP/PS1[Het] mice. **F)** The habituation-dishabituation test was modified to assess potential deficits in long-term discrimination due to natural or pathological aging by presenting the same S1a sample 24 hours after. **G)** Paired data corresponding to the sniffing time during the first and second (24 hours later) S1a presentation is plotted for each condition. Thick lines in dispersion plots and black dots in G indicate mean ± SEM. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more than one variable. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.

**Figure 6. Age-related deficits in social discrimination and habituation are not influenced by previous experience. A)** Average sniffing time of social habituation-dishabituation test in response to odors from novel (N) or littermate (L) subjects. **B)** Dispersion plot of the exploration time of aged and young animals in response to novel or littermate urine samples. **C)** Average sniffing time of social habituation-dishabituation test in response to littermate and the animal´s own urine (O). **D)** Dispersion plot of exploration time of aged and young animals in response to their own´s or littermate urine samples. Thick lines in dispersion plots indicate mean ± SEM. Number of mice: young: n = 21; aged: n = 23. Data were analyzed by a one-way ANOVA with Tukey´s test to test multiple comparisons with more than one variable. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.

**Figure 7. Social novelty is disrupted during pathological aging. A)** Schematics of the three chamber test used in this study to test sociability and social novelty. In the sociability phase the sniffing time of E and M1$^A$ are compared. Social novelty is estimated by quantifying the sniffing time of exploring M1$^B$ versus M2. **B)** Dispersion plots of the sniffing times of E-M1$^A$ (sociability) and M1$^B$-M2 (social novelty) for young wild type mice. **C)** Dispersion plots of the sniffing times of E-M1A (sociability) and M1$^B$-M2 (social novelty) for aged wild type mice. **D)** Dispersion plots of the sniffing times of E-M1$^A$ (sociability) and M1$^B$-M2 (social novelty) for middle age APPWT mice. **E)** Dispersion plots of the sniffing times of E-M1$^A$ (sociability) and M1$^B$-M2 (social novelty) for middle age APP$^{Het}$ mice. Black dots in the paired plots indicate mean ± SEM. Number of mice: young: n = 10; aged: n = 14; middle age control: n = 10; middle age APP/PS1$^{Het}$: n = 12. Data were analyzed by a one-way ANOVA with Tukey's test to test multiple comparisons with more than one variable. * P ≤ 0.05; ** P ≤ 0.01; *** P ≤ 0.001; **** P ≤ 0.0001.