

1 **Male-specific vasotocin expression in the medaka tuberal**  
2 **hypothalamus: androgen dependence and probable role in**  
3 **aggression**

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16 **Abstract**

17

18 Terrestrial vertebrates have a population of androgen-dependent vasotocin (VT)-expressing neurons in  
19 the extended amygdala that are more abundant in males and mediate male-typical social behaviors,  
20 including aggression. Teleosts lack these neurons but instead have novel male-specific VT-expressing  
21 neurons in the tuberal hypothalamus. Here we found in medaka that *vt* expression in these neurons is  
22 dependent on post-pubertal gonadal androgens and that androgens can act on these neurons to directly  
23 stimulate *vt* transcription via the androgen receptor subtype Ara. Furthermore, administration of  
24 exogenous VT induced aggression in females and alterations in the androgen milieu led to correlated  
25 changes in the levels of tuberal hypothalamic *vt* expression and aggression in both sexes. However,  
26 genetic ablation of *vt* failed to prevent androgen-induced aggression in females. Collectively, our results  
27 demonstrate a marked androgen dependence of male-specific *vt* expression in the teleost tuberal  
28 hypothalamus, although its relevance to male-typical aggression needs to be further validated.

29

30 **Keywords:** aggression, androgen, hypothalamus, teleost, vasotocin

## 31 Introduction

32

33 Aggression is an adaptive behavioral trait that is crucial for competition for territories, food, and  
34 mating partners and the establishment of social hierarchies. Although the regulation of aggression in  
35 vertebrates involves many different neural mechanisms, particular attention has been given to two  
36 hormonal systems in the brain: sex steroids and nonapeptides (Kelly and Wilson, 2020).

37 Sex steroids are peripherally derived or produced in the brain and act on neural circuits to modulate  
38 behavior, primarily through binding to specific nuclear receptors that serve as ligand-gated transcription  
39 factors (Yang and Shah, 2014). Androgens, among other sex steroids, play a central role in facilitating  
40 aggression, and males are typically more aggressive than females due to their androgen-dominated  
41 steroid milieu (Hashikawa *et al.*, 2018; Lischinsky and Lin, 2020). In rodents, the stimulatory effects of  
42 androgens on aggression are largely mediated by the activation of estrogen receptors (ESRs) after their  
43 conversion to estrogens in the brain (Yang and Shah, 2014). Recent studies have revealed that activation  
44 of the ESR subtype ESR1 in the ventromedial hypothalamus (VMH) is particularly important for the  
45 expression of aggressive behavior (Chen and Hong, 2018; Hashikawa *et al.*, 2018; Lischinsky and Lin,  
46 2020). However, the transcriptional targets of ESR1 that mediate aggressive behavior remain elusive,  
47 and the specific role of sex steroids in the regulation of aggression is unclear. Furthermore, and  
48 importantly, it is unlikely that the findings in rodents apply to other vertebrates including primates and  
49 teleost fish, where androgens act directly on behaviorally relevant neural circuits via the androgen  
50 receptor (AR) without conversion to estrogens (Okubo *et al.*, 2019; 2022).

51 Nonapeptides, namely, vasotocin (VT, also called vasopressin in mammals) and oxytocin (OT),  
52 are evolutionarily conserved neuropeptides that have been associated with a wide range of social  
53 behaviors, including aggression (Theofanopoulou *et al.*, 2021; Mennigen *et al.*, 2022). The largest  
54 population of neurons expressing VT and that expressing OT both lie in the paraventricular nucleus  
55 (PVN), where they project throughout the brain, as well as to the pituitary, to modulate behavior (Rigney  
56 *et al.*, 2022). In many terrestrial vertebrates, additional neuronal populations expressing VT in an  
57 androgen-dependent, and hence male-biased, manner have been identified in the extended amygdala,  
58 specifically in the bed nucleus of the stria terminalis (BNST) and the medial amygdala (MeA) (Kelly  
59 and Goodson, 2014; Aspesi and Choleris, 2022; Rigney *et al.*, 2022; 2023). These neurons serve as the  
60 main regulators of male-typical social behaviors, and elicit pro- or anti-aggressive behavioral responses  
61 in males, depending on species and social context (Aspesi and Choleris, 2022).

62 Notably, however, teleost fish lack VT-expressing neurons in the extended amygdala, even though  
63 their aggression—like that of terrestrial vertebrates—seems to depend on VT and is generally more  
64 prevalent in males (Godwin and Thompson, 2012; Rigney *et al.*, 2023). This suggests that, in teleosts,  
65 VT may function within another neural circuit to elicit high levels of aggression in males. In line with  
66 this idea, teleosts have several populations of VT-expressing neurons in the tuberal hypothalamus, in  
67 addition to the major population that spans the brain nucleus homologous to the PVN and its immediate  
68 surroundings (Godwin and Thompson, 2012; Oldfield *et al.*, 2015). Our previous findings further

69 revealed that, in medaka fish (*Oryzias latipes*), the populations in the posterior tuberal nucleus (NPT)  
70 and the posterior part of the ventral tuberal nucleus (pNVT) are confined to males (Kawabata *et al.*,  
71 2012). The NPT and pNVT are considered homologous to the ventral tegmental area/substantia nigra  
72 and the anterior hypothalamus, respectively (Forlano and Bass, 2011; Loveland and Hu, 2018), both of  
73 which have been implicated in nonapeptide-regulated social behavior (Rigney *et al.*, 2022).

74 To our knowledge, no information is available on the regulation or role of VT-expressing neuronal  
75 populations in the NPT and pNVT, but it has been reported in pupfish (*Cyprinodon nevadensis*  
76 *amargosae*) that VT expression in the hypothalamus is higher in socially dominant, highly aggressive  
77 males (Lema *et al.*, 2015). Taken together, the above observations led us to hypothesize that VT  
78 expression in either or both of these neuronal populations is induced exclusively in males in an  
79 androgen-dependent manner and contributes to the high levels of aggression typical of males. Here we  
80 tested this hypothesis by investigating the regulatory mechanisms and physiological roles of male-  
81 specific VT expression in the tuberal hypothalamus of medaka.

## 82 **Materials and Methods**

83

### 84 **Animals**

85 All experimental procedures involving animals were performed in accordance with the University  
86 of Tokyo Institutional Animal Care and Use Committee guidelines. The committee requests the  
87 submission of an animal-use protocol only for use of mammals, birds, and reptiles, in accordance with  
88 the Fundamental Guidelines for Proper Conduct of Animal Experiment and Related Activities in  
89 Academic Research Institutions under the jurisdiction of the Ministry of Education, Culture, Sports,  
90 Science and Technology of Japan (Ministry of Education, Culture, Sports, Science and Technology,  
91 Notice No. 71; June 1, 2006). Accordingly, we did not submit an animal-use protocol for this study,  
92 which used only teleost fish and thus did not require approval by the committee.

93 Wild-type medaka of the d-rR strain and *vt* knockout medaka produced in this study were kept  
94 under controlled conditions at 28 °C and a photoperiod of 14:10 light/dark, and were fed with live  
95 *Artemia nauplii* and dry food (Otohime; Marubeni Nisshin Feed, Tokyo, Japan) 3–4 times a day.  
96 Sexually mature fish between 3 and 6 months of age were used in all experiments except for the analysis  
97 of age-dependent changes in *vt* expression, for which fish of 1, 2, 3, and 7 months of age were employed.  
98 To control for genetic diversity and environmental variation, siblings raised in the same conditions were  
99 assigned as the comparison group in all experiments, including those with knockout fish. All sampling  
100 was done 1–2.5 hours after initiation of the light period.

101

### 102 **Gonadectomy and drug treatment**

103 A small incision was made in the ventrolateral abdominal wall of anesthetized fish (0.02% tricaine  
104 methane sulfonate). The gonad was removed through the incision, which was then closed with nylon  
105 thread. Following a 3-day recovery period in saline (0.9% sodium chloride), gonadectomized fish were  
106 reared for 6 days in water containing 100 ng/ml of 11-ketotestosterone (KT; the primary androgen in  
107 teleosts that cannot be converted to estrogens) (Cosmo Bio, Tokyo, Japan) or estradiol-17 $\beta$  (E2; the  
108 primary estrogen in teleosts and other vertebrates) (Fujifilm Wako Pure Chemical, Osaka, Japan), or  
109 vehicle alone (ethanol) and then sampled. Sham-operated control fish were subjected to the same  
110 surgical procedure as gonadectomized fish but without removing the gonad and then treated with vehicle  
111 alone.

112 In separate experiments, males and females (including *vt* knockout females) with intact gonads  
113 were reared for 9 days in water containing 250 ng/ml of the AR antagonist cyproterone acetate (CA)  
114 (LKT Laboratories, St. Paul, MN) and 100 ng/ml of KT, respectively. These fish were observed daily  
115 for changes in aggressive behavior and, when necessary, were sampled on day 9. The concentration of  
116 sex steroids used was determined based on steroid levels in medaka serum (Tilton *et al.*, 2003). Ovarian-  
117 intact females were also treated intraperitoneally with 0.02 ng/mg body weight of Vt peptide (Fujifilm  
118 Wako Pure Chemical) or vehicle alone (saline) and observed for changes in aggressive behavior 2 hours  
119 after treatment.

120

### 121 **Single-label *in situ* hybridization**

122 A digoxigenin (DIG)-labeled cRNA probe for *vt* was generated by PCR amplification of a DNA  
123 fragment corresponding to nucleotides 1–845 (845 bp) of the medaka *vt* cDNA followed by *in vitro*  
124 transcription using T7 RNA polymerase and DIG RNA Labeling Mix (Roche Diagnostics, Basel,  
125 Switzerland). The *in situ* hybridization procedure has been outlined in detail elsewhere (Hiraki-  
126 Kajiyama *et al.*, 2019). Briefly, brains were fixed in 4% paraformaldehyde (PFA), paraffin embedded,  
127 and coronally sectioned at 10- $\mu$ m thickness. Hybridization signal was visualized with an anti-DIG  
128 antibody conjugated to alkaline phosphatase (RRID: AB\_514497; Roche Diagnostics) and nitro blue  
129 tetrazolium/5-bromo-4-chloro-3-indolyl phosphate (NBT/BCIP) substrate (Roche Diagnostics). The  
130 color was allowed to develop for 5 hours. All sections to be compared were processed in the same batch.  
131 To obtain quantitative data, all relevant sections were photographed and converted to black and white  
132 binary images by thresholding using Adobe Photoshop (Adobe, San Jose, CA), and the total area of *vt*  
133 expression signal across all the relevant sections was calculated for each brain nuclei using ImageJ  
134 (<https://imagej.nih.gov/ij/>).

135

### 136 **Double-label *in situ* hybridization**

137 The double-label *in situ* hybridization procedure has been described in detail elsewhere (Kawabata-  
138 Sakata *et al.*, 2020). Briefly, brains were fixed in 4% PFA, embedded in 20% sucrose/5% agarose, and  
139 cryosectioned at 20- $\mu$ m thickness in the coronal plane. Sections were simultaneously hybridized with  
140 the *vt* probe, which was labeled with fluorescein using T7 RNA polymerase and Fluorescein RNA  
141 Labeling Mix (Roche Diagnostics), and a DIG-labeled AR (*ara*, NM\_001122911; *arb*, NM\_001104681)  
142 probe (Hiraki *et al.*, 2012). Fluorescein was detected with an anti-fluorescein antibody conjugated to  
143 horseradish peroxidase (RRID: AB\_2737388; PerkinElmer, Waltham, MA) and visualized using the  
144 TSA Plus Fluorescein System (PerkinElmer); DIG was detected with an anti-DIG antibody conjugated  
145 to alkaline phosphatase (RRID: AB\_514497; Roche Diagnostics) and visualized with Fast Red (Roche  
146 Diagnostics). Sections were counterstained with 4',6-diamidino-2-phenylindole (DAPI) to identify cell  
147 nuclei. Fluorescent images were captured using a TCS SP8 confocal laser-scanning microscope (Leica  
148 Microsystems, Wetzlar, Germany) with the following excitation/emission wavelengths: 405/410–480  
149 nm (DAPI), 488/495–545 nm (fluorescein), and 552/565–700 nm (Fast Red).

150

### 151 **Transcriptional activity assay**

152 A bacterial artificial chromosome (BAC) clone containing the medaka *vt* locus (clone ID: ola1-  
153 127B15) was obtained from National BioResource Project (NBRP) Medaka  
154 (<http://www.shigen.nig.ac.jp/medaka/>). The *vt* gene and flanking regions were sequenced and analyzed  
155 for the presence of potential canonical bipartite androgen-responsive element (ARE)-like sequences  
156 using Jaspar (version 5.0\_alpha; <http://jaspar.genereg.net/>). cDNA fragments encoding full-length  
157 medaka Ara and Arb (NM\_001104681 and NM\_001170833, respectively) were amplified by PCR and

158 inserted into the expression vector pcDNA3.1/V5-His-TOPO (Thermo Fisher Scientific, Waltham, MA).  
159 Fragments of genomic DNA upstream of the first methionine codon (2677 bp) and downstream of the  
160 stop codon (1469 bp) of *vt* were amplified by PCR from the BAC clone and inserted into the respective  
161 NheI and XbaI sites of the luciferase reporter vector pGL4.10 (Promega, Madison, WI). The resultant  
162 reporter construct was transiently transfected into COS-7 cells using Lipofectamine LTX (Thermo  
163 Fisher Scientific), together with an internal control vector pGL4.74 (Promega) and either the Ara or Arb  
164 expression construct. Six-hour post-transfected cells were treated with 0,  $10^{-10}$ ,  $10^{-8}$ , or  $10^{-6}$  M KT for  
165 18 hours in Dulbecco's modified Eagle's medium (phenol red-free) containing 5% charcoal-treated fetal  
166 bovine serum (Thermo Fisher Scientific). After cell lysis, luciferase activity was determined using the  
167 Dual-Luciferase Reporter Assay System (Promega) on the GloMax 20/20n Luminometer (Promega).  
168 All assays were conducted in duplicate and repeated independently three times.

169 To determine the ARE responsible for androgen induction of *vt* transcription, assays were also  
170 conducted with luciferase reporter constructs containing only the fragment upstream of the first  
171 methionine codon or carrying point mutations in the identified ARE-like sequences. A construct  
172 containing the upstream fragment was prepared as described above. Constructs containing point-  
173 mutated ARE-like sequences were prepared using the PrimeSTAR Mutagenesis Basal Kit (Takara Bio,  
174 Shiga, Japan). Because an ARE half-site can function alone to confer androgen inducibility (Pihlajamaa  
175 *et al.*, 2015), both half-sites of each ARE-like sequence were mutated (into a HindIII recognition site  
176 sequence [AAGCTT] to facilitate confirmation of the mutation). The procedures for cell transfection,  
177 KT treatment, and luciferase activity measurements were the same as above, except that a single dose  
178 ( $10^{-6}$  M) of KT was used.

179 Additional assays were performed with luciferase reporter constructs containing truncated versions  
180 of the downstream fragment to see the effects of simultaneous loss of multiple ARE-like sequences in  
181 this fragment. Constructs containing the full-length (1469 bp) or serially 3'-truncated (970 bp and 485  
182 bp) downstream fragment were generated as described above and used in these assays. Cell transfection,  
183 KT treatment ( $10^{-6}$  M), and luciferase activity measurements were performed as described above.

184

### 185 **Aggressive behavior test**

186 The aggressive behavior test was conducted as previously described (Yamashita *et al.*, 2020).  
187 Briefly, four fish of the same genotype and sex that were not familiar with one another were housed in  
188 a 2-liter rectangular tank and separated from each other by opaque partitions. After 10-min acclimation  
189 to the tank, the partitions were removed to allow the fish to interact. Their behavior was recorded for 30  
190 min with a digital video camera (Everio GZ-G5; Jvckenwood, Kanagawa, Japan, or iVIS HF S11/S21;  
191 Canon, Tokyo, Japan). All tests were done 1–5 hours after initiation of the light period. Video recordings  
192 were manually analyzed for the total number of each aggressive act (chase, fin display, circle, strike,  
193 and bite).

194

### 195 **Production of knockout medaka**

196 The *vt* knockout medaka line was produced using transcription activator-like effector nuclease  
197 (TALEN) technology, targeting the sequence corresponding to the mature Vt peptide (Supplementary  
198 Fig. 1A), essentially as described (Takahashi et al., 2016). In brief, TALE repeat arrays were assembled  
199 using the Joung Lab REAL Assembly TALEN kit (Addgene 1000000017). Synthesized TALEN  
200 mRNAs were injected into the cytoplasm of one-cell stage embryos. Once these fish reached adulthood,  
201 they were outcrossed to wild-type fish and the resulting progeny were tested for target site mutations by  
202 T7 endonuclease I assay (Kim et al., 2009) and direct sequencing. A founder fish was identified that  
203 reproducibly produced progeny carrying a 10-bp deletion that introduced a frameshift into the mature  
204 Vt peptide. The progeny were intercrossed to obtain homozygous, heterozygous, and wild-type siblings.  
205 Both male and female homozygous fish were viable and displayed no obvious morphological or  
206 developmental defects. Each fish was genotyped by direct sequencing of the targeted locus.

207 Only female knockout fish were used in this study because we considered that the most direct  
208 approach to test our hypothesis would be to confirm that exogenous androgens induce aggression in  
209 wild-type females but not in *vt* knockout females. We considered males to be unsuitable for this  
210 experiment because they have high levels of endogenous androgens and would be less susceptible to  
211 exogenous androgens.

212

### 213 **Statistical analysis**

214 All quantitative data were presented as mean  $\pm$  standard error of the mean. On graphs, individual  
215 data points were also plotted to indicate the underlying distribution. All statistics were analyzed using  
216 GraphPad Prism (GraphPad Software, San Diego, CA). Data between two groups were compared using  
217 unpaired two-tailed Student's *t*-test. When the F-test indicated that the variances between groups were  
218 significantly different, Welch's correction was applied. To compare data among more than two groups,  
219 one-way analysis of variance (ANOVA) was performed, followed by either Dunnett's (comparisons  
220 between control and experimental groups) or Bonferroni's (comparisons among experimental groups)  
221 *post hoc* test. In cases where the Bartlett's and Brown-Forsythe tests revealed significant differences in  
222 variances among groups, the data were subjected to log transformation to correct for heterogeneity of  
223 variance. When variances remained heterogeneous following transformation, Kruskal-Wallis test was  
224 utilized, followed by Dunn's *post hoc* test. For analyses of age-dependent changes in *vt* expression and  
225 aggressive behavior in *vt*-deficient females treated with KT, two-way ANOVA was used to determine  
226 the effects of and interactions between age and sex and between genotype and KT treatment,  
227 respectively. If a significant interaction was detected, differences between groups were further analyzed  
228 by Bonferroni's *post hoc* test. All data points were included in the analyses and no outliers were defined.



## 229 Results

230

### 231 Male-specific *vt* expression in the tuberal hypothalamic nucleus is dependent on post- 232 pubertal gonadal androgens

233 First, we determined the spatiotemporal pattern of *vt* expression and related sex differences in the  
234 medaka brain. *In situ* hybridization analysis of brains from different ages showed that *vt* was expressed  
235 in the PMp/PPa/PMm/PMg (the latter two brain nuclei are homologous to the PVN) in the preoptic area  
236 and the SC/aNVT, NAT, NPT, and pNVT in the tuberal hypothalamus at all ages examined (Fig. 1A  
237 and B; see Supplementary Table 1 for abbreviations of medaka brain nuclei). Signal quantification  
238 revealed that expression in the PMp/PPa/PMm/PMg, shared by males and females, increased with age  
239 similarly in both sexes, but levels were slightly higher in males overall (main effect of age,  $p < 0.0001$ ;  
240 main effect of sex,  $p = 0.0020$ ; interaction between age and sex,  $p = 0.5837$ ) (Fig. 1C). Expression in  
241 both the SC/aNVT and NAT peaked at 2 months of age, when secondary sexual characters were well-  
242 developed but no spawning had yet occurred (spawning began at 3 months); there were no significant  
243 sex differences except for a slight female bias in the SC/aNVT at 2 months of age ( $p = 0.016$  between  
244 sexes) (Fig. 1C). Expression in the NPT and pNVT was male-specific at all ages examined, with  
245 significant sex differences detected after 2 months of age ( $p = 0.0005$ ,  $0.0003$ , and  $0.0078$  for NPT at 2,  
246 3, and 7 months, respectively;  $p < 0.0001$  for pNVT at 2, 3, and 7 months) (Fig. 1C).

247 These results led us to speculate that sex steroid hormones produced by the gonads after puberty  
248 onset influence the pattern of *vt* expression in the medaka brain. We tested this idea by quantitative *in*  
249 *situ* hybridization of *vt* expression in fish that were gonadectomized in adulthood and treated with KT  
250 or E2. In the male pNVT, castration significantly reduced *vt* expression ( $p < 0.0001$ ), which was restored  
251 by KT treatment ( $p = 0.0014$ ) but not by E2 treatment (Fig. 1D and E). In the female pNVT, *vt* expression  
252 was not detected under normal conditions but was induced by KT treatment following ovariectomy ( $p$   
253  $< 0.0001$ ) (Fig. 1F and G). In other brain nuclei, no significant differences between treatments were  
254 observed (Fig. 1D and F). Collectively, these results suggest that high circulating levels of androgens  
255 released from the testis stimulate *vt* expression in the pNVT in post-pubertal males, whereas in females,  
256 the lack of androgen stimulation prevents its induction. In addition, the effect of androgens on *vt*  
257 expression in the pNVT is transient and reversible; thus, *vt* expression is attenuated in androgen-depleted  
258 adult males and, conversely, induced in estrogen-depleted and androgen-supplemented adult females.

259

### 260 Androgens can directly activate the transcription of *vt* through Ara

261 To investigate the possible direct action of androgens on *vt*-expressing neurons in the pNVT, we  
262 first determined if these neurons coexpress ARs. Most teleosts, including medaka, possess two subtypes  
263 of AR, designated Ara and Arb (Okubo *et al.*, 2022). Double-label *in situ* hybridization for *vt* and each  
264 AR subtype revealed that *ara*, but not *arb*, was abundantly expressed in the pNVT of both sexes, and  
265 virtually all of the *vt*-expressing neurons in the male pNVT were positive for *ara* expression (Fig. 2A  
266 and b). Note that the names of *ara* and *arb* used in the present study follow the nomenclature of Ogino

267 *et al.* (2016; 2023), which reflects the orthology/paralogy relationships of teleost ARs. Thus, the gene  
268 referred to as *ara* in our previous publications (*e.g.*, Hiraki *et al.*, 2012; 2014; Kawabata-Sakata *et al.*,  
269 2020; Yamashita *et al.*, 2017; 2020) is denoted as *arb* in this study, and the gene referred to as *arb* as  
270 *ara*.

271 Next, we tested the ability of androgens to directly activate the transcription of *vt*. Our search for  
272 potential AREs in the *vt* locus of medaka identified six bipartite ARE-like sequences within the upstream  
273 flanking region (positions -2600, -2225, -2168, -1620, -817, and -242 relative to the first methionine  
274 codon) and nine within the downstream flanking region (positions +179, +336, +340, +448, +682, +883,  
275 +1250, +1411, and +1435 relative to the stop codon) (Supplementary Fig. 2). No ARE-like sequences  
276 were found in the gene body of *vt*. A transcriptional activity assay using a luciferase-based reporter  
277 construct containing the upstream and downstream flanking fragments of *vt* that carried these ARE-like  
278 sequences revealed that luciferase activity was dose-dependently induced by KT in the presence of either  
279 Ara ( $p = 0.0015$  at  $10^{-8}$  M and  $0.0006$  at  $10^{-6}$  M) or Arb ( $p = 0.0437$  at  $10^{-8}$  M and  $0.0201$  at  $10^{-6}$  M),  
280 with higher induction observed for Ara (Fig. 2C). An additional assay using only the upstream fragment  
281 (the downstream fragment was removed from the reporter construct) resulted in loss of KT-induced  
282 luciferase activity in the presence of Ara ( $p = 0.0379$  and  $0.6724$  with and without the downstream  
283 fragment, respectively), suggesting that the *cis*-element responsible for KT induction is present in the  
284 downstream region (Fig. 2D). To further explore this finding, we introduced point mutations in each of  
285 the nine ARE-like sequences located in the downstream fragment in the reporter construct and studied  
286 the resulting changes in luciferase activity. However, none of the mutations had a significant impact on  
287 KT-induced luciferase activity in the presence of either Ara or Arb (Fig. 2E).

288 We therefore performed additional assays using reporter constructs containing serially 3'-truncated  
289 downstream fragments to examine the effects of simultaneous loss of multiple ARE-like sequences.  
290 However, even after truncating the downstream fragment from 1469 bp to 485 bp, KT-induced luciferase  
291 activity was not abolished in the presence of either Ara ( $p < 0.0001$ ,  $= 0.0015$ , and  $0.0178$  for 1469, 970,  
292 and 485 bp fragments, respectively) or Arb ( $p = 0.0136$ ,  $0.0198$ , and  $0.0487$ , respectively). These results  
293 suggest that the 485-bp region downstream of the stop codon of *vt* contains the *cis*-element responsible  
294 for KT induction. Considering that none of the mutations in the ARE-like sequences found in this region  
295 abolished KT induction, it may be that AR interacts with *cis*-elements other than the canonical ARE to  
296 activate the transcription of *vt*. Overall, these findings indicate that androgens can act on *vt*-expressing  
297 neurons in the pNVT to directly stimulate the transcription of *vt* via Ara, although the *cis*-element  
298 responsible for this process remains to be identified.

299

### 300 **Administration of exogenous Vt elicits aggression in females**

301 If the high levels of aggression typically observed in male medaka result from male-specific  
302 production of Vt in the pNVT, then female medaka should also exhibit aggression when given  
303 exogenous Vt. We tested this idea by treating females with Vt and analyzing the resulting changes in  
304 intrasexual aggression. Aggression in medaka and many other teleosts includes five behavioral acts:

305 chase, fin display, circle, strike, and bite (Oliveira *et al.*, 2011; Kagawa, 2013). Vt treatment led to a  
306 significant increase in the number of chases ( $p = 0.0086$ ), fin displays ( $p = 0.0104$ ), and bites ( $p =$   
307  $0.0011$ ), but no strikes were induced (Fig. 3). Although not statistically significant, the treatment also  
308 evoked circles, which are not typically observed in females (Fig. 3). From these findings, it was evident  
309 that exogenous Vt elicits aggression in female medaka.

310

### 311 **Altering the androgen milieu leads to correlated changes in the levels of aggression** 312 **and vt expression in the tuberal hypothalamus**

313 If androgen-dependent vt expression in the pNVT facilitates aggression, there is likely to be a  
314 correlation among the levels of vt expression in the pNVT, androgen action, and aggression. To explore  
315 this relationship, we administered KT to females with intact ovaries and evaluated changes in intrasexual  
316 aggression and vt expression in the pNVT. There was a significant increase in the number of chases  
317 after 4 days of KT treatment ( $p = 0.0421$ ), and this increase persisted throughout the remaining treatment  
318 period ( $p = 0.0136$  on day 5 and  $0.0054$  on day 7) (Fig. 4A). Although not statistically significant, the  
319 number of fin displays, circles, strikes, and bites also increased (Fig. 4A). *In situ* hybridization analysis  
320 revealed that KT treatment, even without ovariectomy, induced vt expression in the pNVT ( $p = 0.0043$ ),  
321 along with increased aggression (Fig. 4B and C). In other brain nuclei, no significant changes in vt  
322 expression were noted with KT treatment (Fig. 4B).

323 Next, we treated males with intact testes with the AR antagonist CA to inhibit androgen/AR  
324 signaling and conducted similar behavioral and expression analyses. The number of chases, fin displays,  
325 and circles was significantly reduced after 3 days of treatment ( $p = 0.0094$ ,  $0.0044$ ,  $0.0289$ , and  $0.0007$   
326 for chases on days 3, 5, 7, and 8, respectively;  $p = 0.0323$ ,  $0.0275$ , and  $0.0038$  for fin displays on days  
327 3, 4, and 5 onward, respectively;  $p = 0.0024$ ,  $0.0024$ , and  $0.0003$  for circles on days 3, 4, and 5 onward,  
328 respectively), and the number of bites was significantly reduced after 1 day of treatment ( $p = 0.0075$  on  
329 days 1, 4, 5, and 6) (Fig. 4D). *In situ* hybridization analysis revealed that CA treatment substantially  
330 attenuated vt expression in the pNVT ( $p = 0.0005$ ) (Fig. 4E and F). There were no notable changes in vt  
331 expression in other brain nuclei, except for a slight decrease in the PMp/PPa/PMm/PMg ( $p = 0.0402$ )  
332 (Fig. 4E).

333 Taken together, these results demonstrate a clear correlation among vt expression in the pNVT,  
334 androgen action, and aggression, and strengthen our hypothesis that the high levels of aggression typical  
335 of males are attributable to androgen-dependent vt expression in the pNVT.

336

### 337 **vt deficiency fails to prevent androgen-induced aggression in females**

338 Lastly, we further tested our hypothesis by producing a vt knockout medaka line and comparing  
339 the aggression levels of females treated with KT between genotypes. We considered that, if our  
340 hypothesis is correct, KT treatment should elicit aggressive behavior in vt<sup>+/+</sup> and vt<sup>+/-</sup> females, but not  
341 in vt<sup>-/-</sup> females. We employed TALEN-mediated genome editing to produce a medaka line with a  
342 deleterious frameshift mutation in vt (Supplementary Fig. 1A), resulting in an inability to produce the

343 mature Vt peptide (Supplementary Fig. 1B). We then treated  $vt^{+/+}$ ,  $vt^{+/-}$ , and  $vt^{-/-}$  females with KT and  
344 analyzed their aggression levels on days 0 (before treatment), 5, and 9. Contrary to our expectations,  
345 however, aggression was induced even in  $vt^{-/-}$  females, and no significant differences between  
346 genotypes were found for any aggressive act (Supplementary Fig. 1C). Therefore, this analysis did not  
347 support our hypothesis.

## 348 Discussion

349

350 We previously identified male-specific populations of VT-expressing neurons in the tuberal  
351 hypothalamus of medaka (Kawabata *et al.*, 2012). Here we proposed and tested the hypothesis that VT  
352 expression in these neuronal populations is induced exclusively in males in an androgen-dependent  
353 manner and contributes to high, male-typical levels of aggression. Teleost fish, like terrestrial  
354 vertebrates, have a major population of VT-expressing neurons in the brain nucleus homologous to the  
355 PVN and its surrounding areas (PMp/PPa/PMm/PMg) (Godwin and Thompson, 2012). It has been  
356 reported in several teleost species, including medaka, that the number of one or more subpopulations of  
357 these VT-expressing neurons correlates with the level of aggression, and for this reason, male-typical  
358 aggression in teleosts has been attributed to this neuronal population (Godwin and Thompson, 2012;  
359 Kagawa, 2013; Silva and Pandolfi, 2019). On the other hand, no attempt has been made to assess the  
360 contribution of the tuberal hypothalamic populations to aggression.

361 Here we first demonstrated that male-specific VT expression in the tuberal hypothalamus is indeed  
362 dependent on androgens. More specifically, VT expression in the pNVT of the tuberal hypothalamus is  
363 induced exclusively in post-pubertal males by large amounts of androgens secreted by the testes. To our  
364 knowledge, this is the first report showing that teleosts, although lacking the VT-expressing neurons in  
365 the extended amygdala that are androgen-dependent and thus male-biased, have VT-expressing neurons  
366 with comparable properties in a distinct brain nucleus. We observed no sex steroid dependence of VT  
367 expression in other brain nuclei including the PMp/PPa/PMm/PMg, as reported in other teleost species  
368 such as bluehead wrasses (*Thalassoma bifasciatum*) (Semsar and Godwin, 2003). In teleosts, therefore,  
369 the neuronal population in the pNVT is presumably responsible for reproductive cycle-dependent VT  
370 functions, including increased aggression during the mating period.

371 Despite the similarities, there are important differences in androgen-dependent VT expression in  
372 the extended amygdala of terrestrial vertebrates and the tuberal hypothalamus of teleosts. First, in the  
373 rodent extended amygdala, a large fraction of testosterone is converted locally to E2 and then acts  
374 through the ESR (Aspesi and Choleris, 2022), whereas in the medaka tuberal hypothalamus, KT, an  
375 androgen that cannot be converted to E2, acts directly through the AR. Because the effects of androgens  
376 on male behavior rely primarily on an AR-mediated pathway in many vertebrate species (Okubo *et al.*,  
377 2022), our findings in medaka may be broadly applicable to other species. The second difference  
378 concerns whether treating adult females with sex steroids induces VT expression above basal levels. In  
379 adult female rats, VT expression in the BNST is elevated by estrogen treatment but only to levels  
380 observed prior to ovariectomy (De Vries *et al.*, 1994; Turano *et al.*, 2019). This is because, in rodents,  
381 and probably in other terrestrial vertebrates, sex steroids produced in the fetal gonads in a sex-specific  
382 manner act on the developing BNST to shape irreversible and enduring sex differences in VT expression  
383 (Rigney *et al.*, 2022; 2023). Furthermore, in rodents, this sex difference is due in part to the direct  
384 neuronal actions of sex chromosome-linked genes, which are independent of sex steroid effects (De  
385 Vries *et al.*, 2002). In medaka, by contrast, treating females with androgens markedly induced VT

386 expression in the pNVT, which was virtually absent in ovary-intact and ovariectomized conditions.  
387 Taken together with the observation that castration in males severely reduced VT expression in the  
388 pNVT, it seems that the sexually dimorphic pattern of VT expression in medaka can be reversed between  
389 the sexes in response to changes in the adult androgen milieu. It is known that the sexual phenotypes of  
390 teleosts (behavioral or otherwise), unlike those of terrestrial vertebrates, are highly labile across the  
391 lifespan and can be reversed between the sexes, even as adults (Okubo *et al.*, 2019; 2022); the present  
392 study now shows that manipulation of the adult androgen milieu effectively reverses sex-typical  
393 aggressive behavior. The reversibility of sexually dimorphic VT expression may reflect this fact.

394 It may be relevant to note here that KT-induced *vt* expression in the pNVT of females was apparent  
395 but at a relatively low level compared to that of castrated males. A possible explanation for this  
396 observation is that full induction of *vt* expression by KT in females requires a relatively long period of  
397 time because it involves the generation of new neurons or the activation of the counterparts of male *vt*-  
398 expressing neurons that are in a quiescent, non-activated state. It would be worthwhile to determine if  
399 treating females with KT for a longer period of time results in comparable levels of *vt* expression as  
400 males.

401 The question then arises whether androgen/AR signaling induces VT expression in pNVT neurons  
402 directly or indirectly through other target genes or cells. We found that *ara*, one of the two AR genes in  
403 teleosts, is expressed in almost all VT-expressing neurons in the pNVT. We further showed by  
404 transcriptional activity assays that androgens can directly stimulate the transcription of *vt* and that the  
405 magnitude of this stimulation is greater through Ara than through Arb. These results suggest that  
406 androgens directly activate VT expression in the pNVT through binding to Ara. This finding is  
407 consistent with recent work in cichlids (*Astatotilapia burtoni*) showing that *ara* and *arb* are functionally  
408 differentiated, with *ara* responsible for promoting aggression (Alward *et al.*, 2020). Although it is not  
409 known whether VT is a direct transcriptional target of AR signaling (or ESR signaling) in terrestrial  
410 vertebrates, as it is in medaka, androgen-induced *Vt* expression in the BNST of adult rats has been  
411 reported to involve changes in the DNA methylation pattern of the *Vt* promoter (Auger *et al.*, 2011).  
412 Future work will be needed to clarify whether a similar mechanism exists in the medaka pNVT and to  
413 identify functional AREs in the medaka *vt* locus (which eluded detection in the present survey) in order  
414 to clarify the evolutionary conservation and divergence of regulatory mechanisms for sexually  
415 dimorphic VT expression.

416 We also tested the hypothesized role of male-specific VT expression on aggression. Although VT  
417 has been implicated in aggressive behavior in many teleost species (Godwin and Thompson, 2012; Silva  
418 and Pandolfi, 2019), Yokoi *et al.* (2015) observed no anomalies in intrasexual aggression of male  
419 medaka carrying a missense mutation in *vt* (leading to replacement of the first methionine with arginine  
420 and loss of the ability to produce the mature Vt peptide). While this observation may suggest that VT is  
421 not involved in aggression in medaka, we showed here that administration of exogenous Vt peptide  
422 elicited aggressive behavior in female medaka, which typically exhibit little or no aggression. This result  
423 indicates that VT certainly serves to facilitate aggression in medaka as well as in other teleosts (although

424 it is possible that the induced aggression may represent a pharmacological effect rather than a  
425 physiological role of VT). Furthermore, manipulation of the androgen milieu of adult male and female  
426 medaka to inhibit or facilitate aggression revealed a clear correlation among *vt* expression in the pNVT,  
427 androgen action, and aggression. More specifically, treatment of male medaka with an AR antagonist  
428 resulted in a marked reduction in both aggression and *vt* expression in the pNVT, whereas androgen  
429 treatment of females elicited both of them. All of these findings support our hypothesis that male-  
430 specific VT expression contributes to male-typical high levels of aggression, thereby strengthening its  
431 validity.

432 Finally, we further tested this hypothesis using *vt*-deficient female medaka but, contrary to our  
433 expectations, androgen treatment induced comparable levels of aggression in *vt*-deficient females as in  
434 wild-type females. A straightforward interpretation of this result would suggest that our hypothesis is  
435 incorrect and androgen-induced aggression is not mediated by VT; however, the lack of detectable  
436 effects of *vt* deficiency might also be due to functional compensation by other genes or pathways. A  
437 good candidate for this functional compensation is OT, a nonapeptide closely related to VT that can  
438 partially activate VT receptors (Mennigen *et al.*, 2022; Rae *et al.*, 2022). In medaka, *ot*-expressing  
439 neurons are not observed in the pNVT, where male-specific *vt*-expressing neurons reside (Kawabata *et al.*,  
440 2012), but it is possible that *ot*-expressing neurons in another brain nucleus project to the same brain  
441 region as male-specific *vt*-expressing neurons and compensate for *vt* function. Contrary to male medaka  
442 carrying a missense mutation in *vt*, which showed comparable levels of aggression to wild-type males,  
443 as described above (Yokoi *et al.*, 2015), male medaka carrying a missense mutation in the VT receptor  
444 subtype *vla2* (leading to replacement of asparagine at position 68 with isoleucine) were found to be less  
445 aggressive (Yokoi *et al.*, 2015). This discrepancy can also be explained if OT compensates for the loss  
446 of VT function via V1A2. It would be interesting to test this idea in future studies by producing double-  
447 knockout medaka for *vt* and *ot* and analyzing their behavioral phenotypes.

448 In summary, our results have demonstrated that VT expression in the tuberal hypothalamic nucleus  
449 is induced exclusively in males, most likely as a result of direct transcriptional activation of VT by  
450 androgen/Ara signaling, although it remains to be determined whether this VT expression is relevant to  
451 the high levels of aggression typical of males. The anterior hypothalamus of rodents, which is  
452 homologous to the teleost pNVT, contains no VT-expressing neurons; however, it receives heavy  
453 projections from male-biased VT neurons in the BNST and abundantly expresses the behaviorally-  
454 relevant V1A receptor, thus representing a primary site of action of VT for aggression (Rigney *et al.*,  
455 2022; 2023). If the V1A receptor homolog is also expressed in the teleost pNVT, VT produced in this  
456 brain nucleus may contribute to male-typical aggression by acting in an autocrine/paracrine manner.  
457 Further studies, including analyses of receptor expression and axonal projection patterns, will be needed  
458 to test this idea. It would also be worthwhile investigating whether the current findings in medaka apply  
459 to other teleosts. The resulting information on species variation in the spatiotemporal expression patterns  
460 and regulatory mechanisms of VT may provide some insight into the diversity of social structures and  
461 behavioral patterns in teleosts.

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463

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470

471 **Declarations of interest**

472

473 The authors declare no competing interests.



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475

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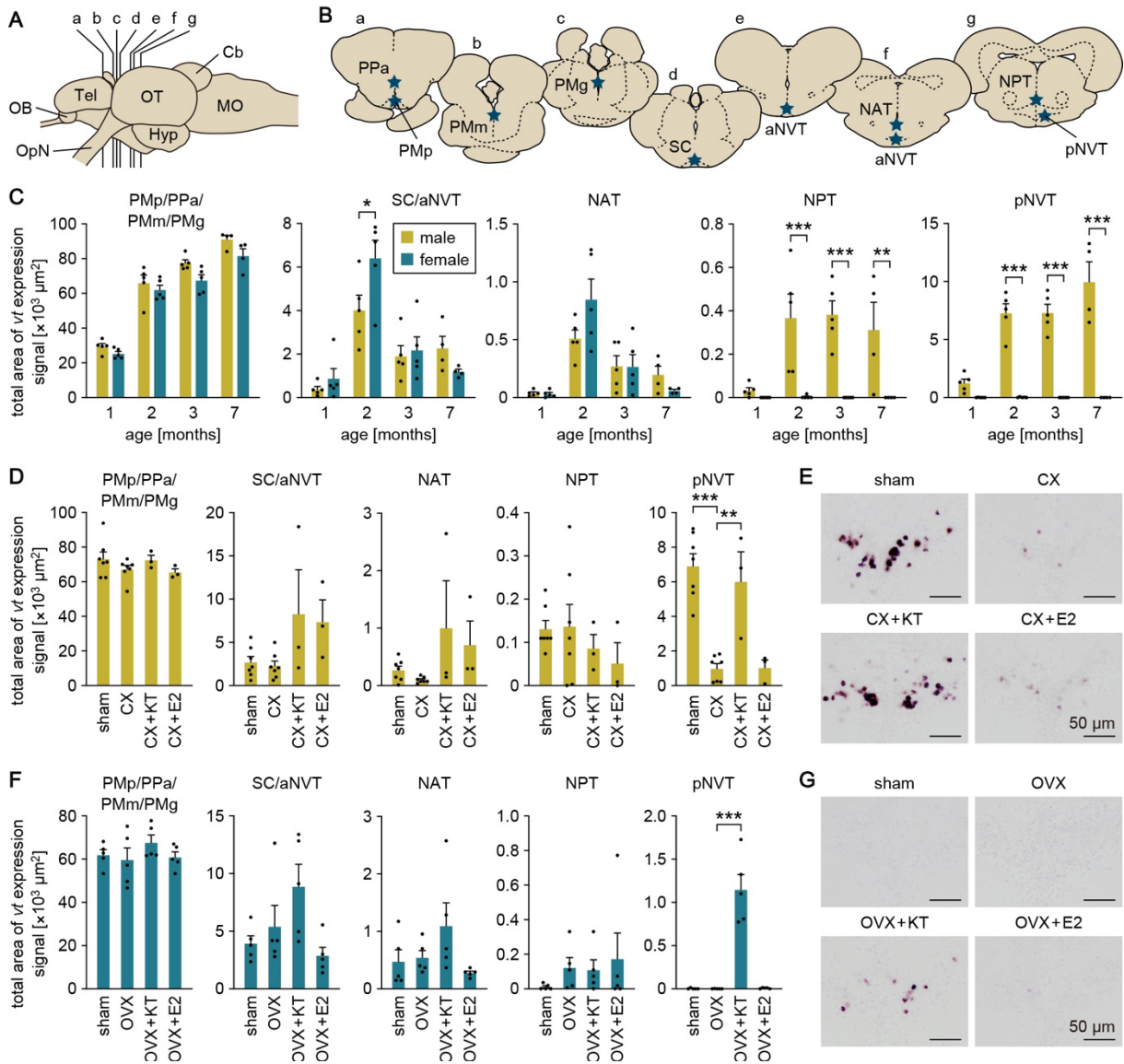
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593 **Figures**

594



595

596

597 **Fig. 1. Male-specific *vt* expression in the tuberal hypothalamic nucleus is dependent on**

598 **post-pubertal gonadal androgens.** (A) Schematic drawing of the medaka brain (lateral view,

599 anterior to left) depicting the approximate levels of sections shown in panel B. (B) Coronal brain sections

600 containing nuclei in which *vt* is expressed (stars). See Supplementary Table 1 for abbreviations of brain

601 regions and nuclei. (C) Total area of *vt* expression signal in each brain nucleus of males (yellow columns)

602 and females (blue columns) at different ages (n = 5 per sex and age, except n = 4 at 7 months). (D) Total

603 area of *vt* expression signal in each brain nucleus of sham-operated males (sham) and castrated males

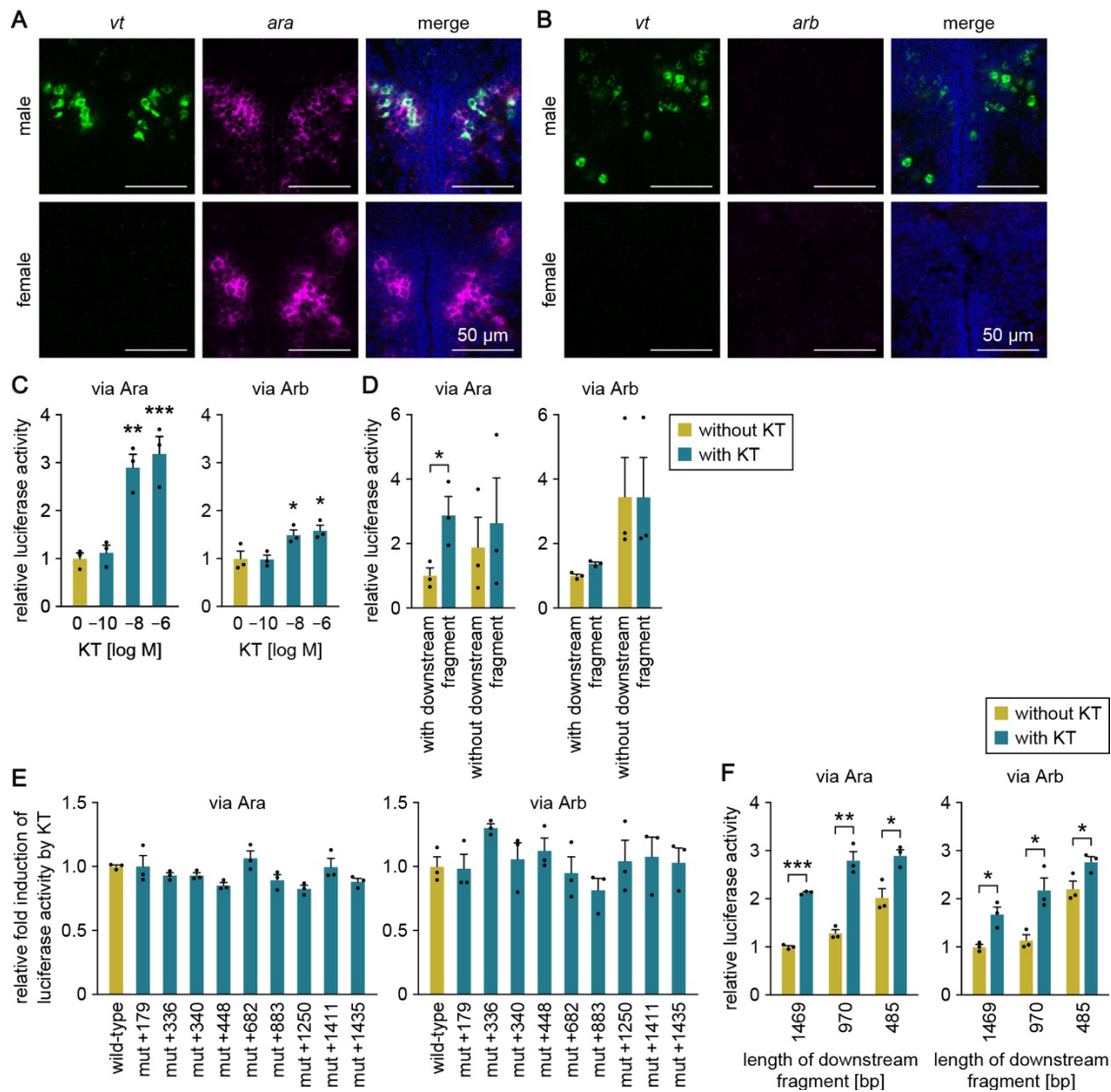
604 treated with vehicle alone (CX), KT (CX+KT), or E2 (CX+E2) (n = 7 for sham and CX; n = 3 for

605 CX+KT and CX+E2). (E) Representative images of *vt* expression in the pNVT of sham, CX, CX+KT,

606 and CX+E2 males. Scale bars are all 50  $\mu\text{m}$ . (F) Total area of *vt* expression signal in each brain nucleus

607 of sham females and ovariectomized females treated with vehicle alone (OVX), KT (OVX+KT), or E2

608 (OVX+E2) (n = 5 for each group). (G) Representative images of *vt* expression in the pNVT of sham,  
609 OVX, OVX+KT, and OVX+E2 females. Scale bars are all 50  $\mu$ m. Statistical differences were  
610 determined by Bonferroni's *post hoc* test (C, D, F). \* $p$  < 0.05; \*\* $p$  < 0.01; \*\*\* $p$  < 0.001.

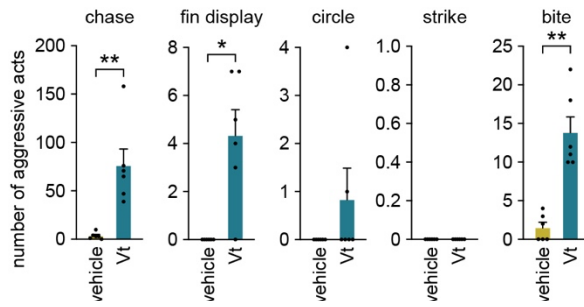


611  
612

613 **Fig. 2. Androgens can directly activate the transcription of *vt* through Ara.** (A, B)  
614 Representative images of the expression of *ara* (A) and *arb* (B) in the pNVT, where *vt* exhibits sexually  
615 dimorphic expression. In each row, left panels show *vt* expression (green), middle panels show *ara/arb*  
616 expression (magenta), and right panels show the merged images with DAPI staining (blue). Scale bars  
617 are all 50  $\mu$ m. (C) Stimulation of *vt* transcriptional activity by KT. A luciferase-based reporter construct  
618 containing genomic fragments upstream of the first methionine codon and downstream of the stop codon  
619 of *vt* was transfected into COS-7 cells, in conjunction with an Ara or Arb expression construct. Cells  
620 were stimulated to varying concentrations of KT, and luciferase activity was measured. Fold induction  
621 was calculated relative to unstimulated cells. (D) Effects of removing the downstream fragment on KT-  
622 induced luciferase activity. A luciferase reporter construct with or without the downstream fragment  
623 was transfected into cells, along with the Ara or Arb expression construct. Cells were stimulated with or  
624 without KT (blue and yellow columns, respectively), and luciferase activity was determined. Fold  
625 induction was calculated relative to the construct with the downstream fragment without KT stimulation.  
626 (E) Effects of mutations in ARE-like sequences on KT-induced luciferase activity. A wild-type

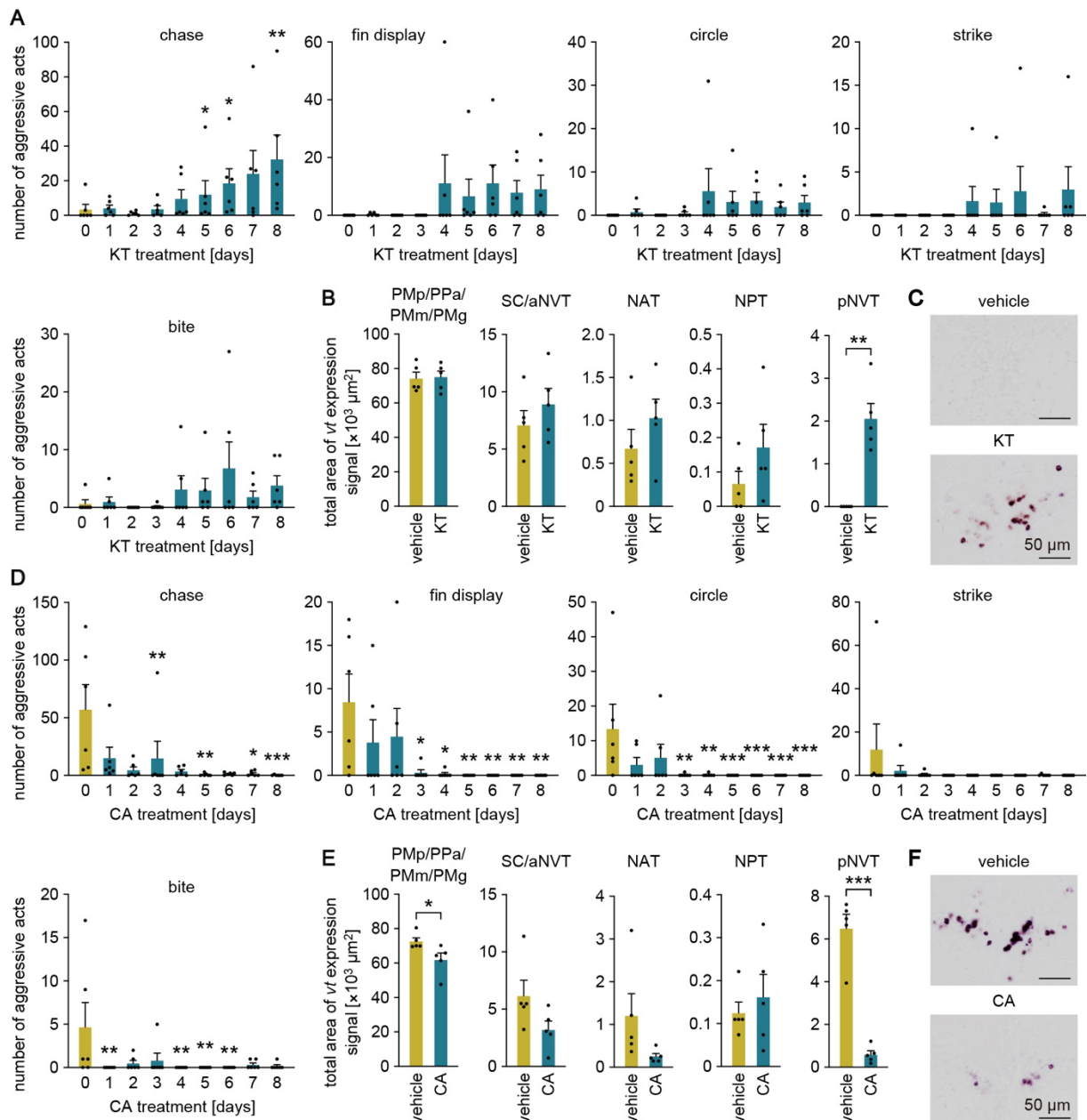
627 luciferase reporter construct (wild-type) or a construct carrying a mutation in the ARE-like sequence at  
628 position +179 (mut+179), +336 (mut+336), +340 (mut+340), +448 (mut+448), +682 (mut+682), +883  
629 (mut+883), +1250 (mut+1250), +1411 (mut+1411), or +1435 (mut+1435) was transfected into cells,  
630 along with the Ara or Arb expression construct. Cells were stimulated with or without KT, and fold  
631 induction of luciferase activity was calculated relative to the wild-type construct. (F) Effects of 3'-  
632 truncation of the downstream fragment on KT-induced luciferase activity. Cells were transfected with a  
633 reporter construct containing 1469-bp, 970-bp, or 485-bp downstream fragment and an Ara or Arb  
634 expression construct. Cells were stimulated with or without KT (blue and yellow columns, respectively),  
635 and fold induction of luciferase activity was calculated relative to the construct containing the 1469 bp  
636 fragment without KT stimulation. Statistical differences were determined by Dunnett's *post hoc* test  
637 (versus unstimulated control (C) or wild-type construct (E)) and unpaired t-test (D and F). \* $p < 0.05$ ;  
638 \*\* $p < 0.01$ ; \*\*\* $p < 0.001$ .





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641 **Fig. 3. Administration of exogenous Vt elicits aggression in females.** Shown is the sum of  
642 each aggressive act (chase, fin display, circle, strike, and bite) exhibited by females receiving vehicle  
643 only or Vt (n = 6 for each treatment). Statistical differences were determined by unpaired *t*-test with  
644 Welch's correction. \**p* < 0.05; \*\**p* < 0.01.



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**Fig. 4. Altering the androgen milieu leads to correlated changes in the levels of aggression and vt expression in the tuberal hypothalamus.** (A) Daily changes in each aggressive act (chase, fin display, circle, strike, and bite) exhibited by KT-treated females (n = 6). (B) Total area of vt expression signal in each brain nucleus of vehicle- or KT-treated females (n = 5 per treatment). (C) Representative images of vt expression in the pNVT of vehicle- or KT-treated females. Scale bars are 50  $\mu\text{m}$ . (D) Daily changes in each aggressive act (chase, fin display, circle, strike, and bite) exhibited by males treated with the androgen receptor antagonist CA (n = 6). (E) Total area of vt expression signal in each brain nucleus of vehicle- or CA-treated males (n = 5 per treatment). (F) Representative images of vt expression in the pNVT of vehicle- or CA-treated males. Scale bars are 50  $\mu\text{m}$ . Statistical differences were determined by Dunnett's *post hoc* test (versus day 0) (A, D), except for the fin display data in panel D, which was determined by Dunn's *post hoc* test, and unpaired *t*-test with

658 or without Welch's correction (B, E).  $*p < 0.05$ ;  $**p < 0.01$ ;  $***p < 0.001$ .