

1 ***fruitless* tunes functional flexibility of courtship circuitry during development**

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14 **ABSTRACT**

15• *Drosophila* male courtship is controlled by the male-specific products of the *fruitless*
16 (*fru^M*) gene and its expressing neuronal circuitry. *fru^M* is considered a master gene that
17 controls all aspects of male courtship. By temporally and spatially manipulating *fru^M*
18 expression, we found that *fru^M* is required during a critical developmental period for
19 innate courtship towards females, and its function during adulthood is relatively
20 trivial. By altering or eliminating *fru^M* expression, we generated males that are
21 innately heterosexual, homosexual, bisexual, or without innate courtship but could
22 acquire such behavior in an experience-dependent manner. These findings show that
23 *fru^M* is not absolutely necessary for courtship but is critical during development to
24 build a sex circuitry with reduced flexibility and enhanced efficiency and provide a
25 new view about how *fru^M* tunes functional flexibility of a sex circuitry.

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27• **Keywords:**

28• *Drosophila*; courtship; innate behavior; fruitless; circuit flexibility; sexual orientation

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30 **Introduction**

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32 *Drosophila* male courtship is one of the best understood innate behaviors in terms of
33 genetic and neuronal mechanisms (Dickson, 2008; Yamamoto and Koganezawa,
34 2013). It has been well established that the *fruitless* (*fru*) gene and its expressing
35 neurons control most aspects of such innate behavior (Ito et al., 1996; Manoli et al.,
36 2005; Ryner et al., 1996; Stockinger et al., 2005). The male-specific products of the
37 P1 promoter of the *fru* gene (*fru^M*) are expressed in ~2000 neurons, which are inter-
38 connected to form a sex circuitry from sensory neurons to motor neurons (Cachero et
39 al., 2010; Lee et al., 2000; Manoli et al., 2005; Stockinger et al., 2005; Usui-Aoki et
40 al., 2000; Yu et al., 2010). *fru^M* function is necessary for the innate courtship behavior
41 and sufficient for at least some aspects of courtship (Baker et al., 2001; Demir and
42 Dickson, 2005; Manoli et al., 2005). Thus, the study of *fru^M* function in controlling
43 male courtship serves as an ideal model to understand how innate complex behaviors
44 are built into the nervous system by regulatory genes (Baker et al., 2001).

45 Although *fru^M* serves as a master gene controlling *Drosophila* male courtship, we
46 recently found that males without *fru^M* function, although do not court if raised in
47 isolation, were able to acquire at least some courtship behaviors if raised in groups
48 (Pan and Baker, 2014). Such *fru^M*-independent but experience-dependent courtship
49 acquisition requires another gene in the sex determination pathway, the *doublesex*
50 (*dsx*) gene (Pan and Baker, 2014). *dsx* encodes male- and female-specific DSX

51 proteins (DSX^M and DSX^F, respectively) (Burtis and Baker, 1989), and DSX^M is
52 expressed in ~700 CNS neurons, the majority of which also express *fru*^M (Rideout et
53 al., 2010; Robinett et al., 2010). It has been found that the *fru*^M and *dsx*^M co-
54 expressing neurons are required for courtship in the absence of *fru*^M function (Pan and
55 Baker, 2014). Thus *fru*^M-expressing neurons, especially those co-expressing *dsx*^M,
56 control the expression of courtship behaviors even in the absence of FRU^M function.
57 Indeed, although the gross neuroanatomical features of the *fru*^M-expressing circuitry
58 are largely unaffected by the loss of *fru*^M, detailed analysis revealed morphological
59 changes of many *fru*^M-expressing neurons (Kimura et al., 2005; Kimura et al., 2008;
60 Mellert et al., 2010). Recent studies further reveal that FRU^M specifies neuronal
61 development by recruiting chromatin factors and changing chromatin states, and also
62 by turning on and off the activity of the transcription repressor complex (Ito et al.,
63 2012; Ito et al., 2016; Sato et al., 2019a; Sato et al., 2019b; Sato and Yamamoto,
64 2020).

65 That FRU^M functions as a transcription factor to specify development and/or
66 physiological roles of certain *fru*^M-expressing neurons, and perhaps the
67 interconnection of different *fru*^M-expressing neurons to form a sex circuitry raises
68 important questions regarding when *fru*^M functions and how it contributes to the sex
69 circuitry (*e.g.*, how the sex circuitry functions differently with different levels of
70 FRU^M), especially in the background that *fru*^M is not absolutely necessary for male
71 courtship (Pan and Baker, 2014). To at least partially answer these questions, we

72 temporally or spatially knocked down *fru^M* expression, and compared courtship
73 behavior in these males with that in wild-type males or *fru^M* null males and revealed
74 crucial roles of *fru^M* during a narrow developmental window for the innate courtship
75 towards females. We also found that the sex circuitry with different *fru^M* expression
76 has distinct function such that males could be innately heterosexual, homosexual,
77 bisexual, or without innate courtship but could acquire such behavior in an
78 experience-dependent manner. Thus, *fru^M* tunes functional flexibility of the sex
79 circuitry instead of switching on its function as conventionally viewed.

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82 **Results**

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84 To specifically knock down *fru^M* expression, we used a microRNA targeting *fru^M*
85 (*UAS-fruMi*) and a scrambled version as a control (*UAS-fruMiScr*) as previously used
86 (Chen et al., 2017). We firstly tested male courtship without food in the behavioral
87 chamber. Knocking down *fru^M* in all the *fru^{GAL4}* labeled neurons eliminated male
88 courtship towards females (courtship index [CI], which is the percentage of
89 observational time that males displayed courtship, is nearly 0) (Figure 1A), consistent
90 with previous findings that *fru^M* is required for innate male-female courtship (Demir
91 and Dickson, 2005; Pan and Baker, 2014). We then added a temperature dependent
92 *tub-GAL80^{ts}* transgene to restrict *UAS-fruMi* expression (e.g., at 30°C) at different

93 developmental stages. We raised *tub-GAL80^{ts/+}; fru^{GAL4}/UAS-fruMi* flies at 18°C
94 (permissive for GAL80^{ts} that inhibits GAL4 activity) and transferred these flies to
95 fresh food vials every two days. In this way we generated *tub-GAL80^{ts/+};*
96 *fru^{GAL4}/UAS-fruMi* flies at 9 different stages from embryos to adults and incubated all
97 flies at 30°C to allow *fru^M* knock-down for 2 days, then placed all flies back to 18°C
98 until courtship test (Figure 1B). We found that males with *fru^M* knocked down from
99 stage 5 to 6, matching the pupation phase, rarely courted (CI < 10%) and none
100 successfully mated, while males with *fru^M* knocked down near this period showed a
101 partial courtship deficit (Figure 1C, D). Knocking down *fru^M* specifically during
102 adulthood for 2 days did not affect male courtship (CI > 80%) and mating success.
103 These results reveal a critical developmental period where *fru^M* is required for adult
104 male courtship towards females.

105 As we did not see an obvious courtship deficit in males with *fru^M* knocked down
106 in adulthood for 2 days, we further tested the role of *fru^M* in adulthood using different
107 approaches. We set out to express the female-specific *transformer (traF)* gene (Baker
108 and Ridge, 1980; McKeown et al., 1988) to feminize all *fru^{GAL4}* labeled neurons, in
109 addition to the *fru^M* RNAi experiments. We express *UAS-traF* or *UAS-fruMi* in all the
110 *fru^{GAL4}* labeled neurons specifically during adulthood for 4 days before test (see
111 procedure above each figure) for single-pair male-female, male-male, and male
112 chaining (in groups of 8 males) behaviors. We found that overexpression of *traF* in all
113 *fru^{GAL4}* labeled neurons during adulthood for 4 days did not affect male-female

114 courtship (Figure 1E), but slightly increased male-male (Figure 1F) and male chaining
115 behaviors (Figure 1G). Furthermore, knocking down *fru^M* in all *fru^{GAL4}* labeled
116 neurons during adulthood for 4 days did not affect male-female (Figure 1E) or male-
117 male courtship (Figure 1F), but slightly increased male chaining behaviors (Figure
118 1G). Together these results indicate that *fru^M* function during pupation is crucial for
119 adult courtship towards females, while its function during adulthood is dispensable
120 for female-directed courtship, though it plays a minor role in inhibiting male-male
121 courtship behaviors.

122 To further reveal the role of *fru^M* in male courtship, we tried to spatially knock
123 down *fru^M* expression using a simple way: *fru^M* in brain and *fru^M* outside brain. We
124 used *Otd-Flp* expressing FLP specifically in the central brain (Asahina et al., 2014) to
125 divide *fru^{GAL4}* expression (Figure 2A) into two parts: *fru^M*- and *Otd*-positive neurons
126 (specifically in brain) in *Otd-Flp/UAS-myrGFP; fru^{GAL4}/tub>GAL80* males (Figure
127 2B), and *fru^M*-positive but *Otd*-negative neurons (theoretically outside brain, but still
128 with few in brain) in *Otd-Flp/UAS-myrGFP; fru^{GAL4}/tub>stop>GAL80* males (Figure
129 2C). We also checked GFP expression in peripheral nervous system in these males,
130 and found a few GFP-positive cells in antennae and forelegs in *Otd-Flp/UAS-*
131 *myrGFP; fru^{GAL4}/tub>stop>GAL80* males, but no expression in *Otd-Flp/UAS-*
132 *myrGFP; fru^{GAL4}/tub>GAL80* or wild-type males (Figure 2D-G). Thus, we
133 successfully divided *fru^{GAL4}* expression into two categories, one with *GAL4* expressed

134 in *fru*⁺*Otd*⁺ neurons in brain, and the other with *GAL4* expressed in *fru*⁺*Otd*⁺ neurons
135 outside brain.

136 We then used the above intersectional strategy to knock down *fru*^M expression
137 in all *fru*^{GAL4} neurons or those neurons in or outside brain, and compared male
138 courtship with that in wild-type males and *fru*^M null males. We tested one-time single-
139 pair male-female and male-male courtship (single-housed before test) as well as male
140 chaining in groups of 8 males over 3 days on food for better comparison of these
141 courtship assays, as courtship by *fru*^M null males largely depends on food presence
142 (Pan and Baker, 2014). We found that male-male courtship in *fru*^M knocked down
143 males is higher if tested on food, consistent with a courtship promoting role by food
144 (Grosjean et al., 2011; Pan and Baker, 2014), while courtship in wild-type males on
145 food or without food is not changed in our assays (Figure 3—figure supplement 1).
146 We found that wild-type males performed intensive courtship behavior towards virgin
147 females (CI > 80%) and rarely courted males (CI ~0) (Figure 3A). Furthermore, these
148 control males did not show any chaining behavior after grouping from 3 hours to 3
149 days (ChI = 0) (Figure 3B). In striking contrast, *fru*^M null mutant (*fru*^{LexA}/*fru*⁴⁻⁴⁰) males
150 rarely courted either females or males (Figure 3C); however, these males developed
151 intensive chaining behavior after grouping for 1-3 days (Figure 3D). These
152 observations replicated previous findings that there exists a *fru*^M-independent
153 experience- and *dsx*^M-dependent courtship pathway (Pan and Baker, 2014) (Figure
154 3E). To compare behavioral differences by *fru*^M null males and *fru*^M RNAi knocked

155 down males, we quantified to how much extent the microRNA against *fru^M* worked.

156 We found that the *fru^M* mRNA level was reduced to ~40% of that in control males

157 (Figure 3F). Interestingly, while males with *fru^M* knocked down in all *fru^M* neurons

158 rarely courted females (CI~5%, Figure 3G), they displayed a high level of male-male

159 courtship behavior (CI > 50%, Figure 3G) and constantly high level of male chaining

160 (Figure 3H), dramatically different from *fru^M* null males. These results reveal distinct

161 roles of low *fru^M* (RNAi) and high *fru^M* (wild-type) in regulating male-male and male-

162 female courtship (Figure 3I). To further dissect the role of *fru^M* in male courtship, we

163 knocked down *fru^M* specifically in brain, and found that such males had a reduced

164 level of courtship towards females (CI = $56.61 \pm 5.86\%$), but their sexual orientation

165 was not changed as they courted males in a much lower level (CI = $15.94 \pm 3.26\%$,

166 Figure 3J). Furthermore, males with *fru^M* knocked down in brain showed low male

167 chaining behavior initially but increasing levels of chaining behavior over 1-3 days

168 (ChI[3h] = $9.35 \pm 5.40\%$, ChI[3d] = $68.82 \pm 5.53\%$, Figure 3K). These results

169 indicate that *fru^M* function in brain promotes male-female courtship and inhibits

170 acquisition or progression of the experience-dependent chaining behavior (Figure 3L).

171 In contrast, males with *fru^M* knocked down outside brain showed equally intensive

172 male-female and male-male courtship (CI[male-female] = $85.62 \pm 1.42\%$, CI[male-

173 male] = $82.89 \pm 2.76\%$, Figure 3M), indicating an inhibitory role of *fru^M* in these

174 neurons for male-male courtship (Figure 3O). These males performed a high level of

175 male chaining behavior initially (ChI[3h] = $92.90 \pm 3.08\%$), but decreased levels of

176 chaining behavior over 1-3 days ($\text{ChI}[3\text{d}] = 20.01 \pm 3.75\%$, Figure 3N), consistent
177 with the above finding that *f^{ru}^M* function in the brain which is intact in these males
178 inhibits acquisition or progression of male chaining behavior (Figure 3L).

179 Taken together, the above results indicate crucial role of *f^{ru}^M* expression level
180 during a critical developmental period for the manifestation of courtship behaviors
181 and reveal functional flexibility of the *f^{ru}^M*-expressing sex circuitry (Figure 4).

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183

184 **Discussion**

185

186 Previous findings show that *f^{ru}^M* expression commences at the wandering third instar
187 larval stage, peaks at the pupal stage, and thereafter declines but does not disappear
188 after eclosion (Lee et al., 2000), which suggests that *f^{ru}^M* may function mainly during
189 development for adult courtship behavior despite of no direct evidence. Here we
190 temporally knocked down *f^{ru}^M* expression in different developmental stages for 2
191 days and found that males with *f^{ru}^M* knocked down during pupation rarely courted,
192 while males with *f^{ru}^M* knocked down during adulthood courted normally towards
193 females. This is the first direct evidence that *f^{ru}^M* is required during development but
194 not adulthood for courtship behavior. However, we also found a minor role of *f^{ru}^M*
195 during adulthood in suppressing male-male courtship, as males with *f^{ru}^M* knocked
196 down or *tra* overexpressed during adulthood displayed slightly enhanced male-male

197 courtship or male chaining behaviors. Note that a previous study found that removal
198 of *transformer 2 (tra2)* specifically during adulthood using a temperature sensitive
199 *tra2* allele induced 8 out of 96 females to show male-type courtship behaviors (Belote
200 and Baker, 1987), which suggests that expression of FRU^M and DSX^M (by removal of
201 *tra2* function in females) during adulthood is sufficient to masculinize CNS to some
202 extent and induce a small fraction of females to display courtship behaviors. A recent
203 study also found that *fru*^M expression in the *Or47b*-expressing olfactory receptor
204 neurons depends on the activity of these neurons during adulthood (Hueston et al.,
205 2016). Based on all these findings, we propose that *fru*^M expression during pupation is
206 crucial for specifying a sex circuitry that allows innate courtship towards females, and
207 its expression during adulthood may be activity-dependent in at least some neurons
208 and modulates some aspects of courtship (e.g., inhibits male-male courtship). Thus,
209 there are at least two separate mechanisms that *fru*^M contributes to the sex circuitry,
210 one during a critical developmental period to build the female-directed innate
211 courtship into that circuitry, and the other during adulthood to modulate neuronal
212 physiology in an experience-dependent manner.

213 Most importantly, we revealed striking flexibility of the fly sex circuitry by
214 manipulating *fru*^M expression. We listed four cases with *fru*^M manipulation here for
215 comparison: (1) males with a sex circuitry having wild-type *fru*^M function have innate
216 heterosexual courtship, as they court readily towards females, but do not court males
217 no matter how long they meet; (2) males with a sex circuitry having no *fru*^M function

218 lose the innate courtship ability, but have the potential to acquire courtship towards
219 males, females, and even other species in an experience-dependent manner; (3) males
220 with a sex circuitry having limited *fru^M* expression (e.g., 40%) have innate
221 homosexual courtship, as they court readily towards other males, but rarely court
222 females; (4) males with a sex circuitry having limited *fru^M* expression outside brain
223 (but intact *fru^M* expression in brain) are innately bisexual, as they court equally
224 towards females or males. Although previous studies found that different *fru^M* alleles
225 (e.g., deletions, inversions or insertions related to *fru*) showed very different courtship
226 abnormalities (Anand et al., 2001; Vilella et al., 1997), it was very hard to link *fru^M*
227 function to the flexibility of sex circuitry, and often seen as allele-specific or
228 background-dependent phenotypes. Our study using relatively simple genetic
229 manipulations that generate dramatical different courtship behaviors promoted us to
230 speculate a different view about the role of *fru^M*: instead of simply being a master
231 gene that controls all aspects of male courtship, *fru^M* is not absolutely necessary for
232 courtship, but changes the wiring of the sex circuitry during development such that
233 the sex circuitry may function in very different ways, ranging from innately
234 heterosexual, homosexual, bisexual to largely experience-dependent acquisition of the
235 behavior. Such flexibility of the sex circuitry is tuned by different *fru^M* expression,
236 such that changes of *fru^M* regulatory regions during evolution would easily select a
237 suitable functional mode of the sex circuitry.

238

239

240 **Materials and methods**

241

242 **Fly Stocks**

243 Flies were maintained at 22 or 25°C in a 12 h:12 h light:dark cycle. Canton-S flies

244 were used as the wild-type strain. Other stocks used in this study include the

245 following: *fru*^{GAL4} (Stockinger et al., 2005), *UAS-fruMi* and *UAS-fruMiScr* (Meissner

246 et al., 2016), *fru*^{LexA} and *fru*⁴⁻⁴⁰ (Pan and Baker, 2014) and *Otd-Flp* (Asahina et al.,

247 2014). *UAS-traF* (BL#4590), *tub-GAL80^{ts}* (BL#7019), *tub>GAL80>* (BL#38881)

248 and *tub>stop>GAL80* (BL#39213) were from Bloomington Drosophila Stock

249 Center.

250

251 **Courtship and Chaining Assays**

252 For the single-pair courtship assay, the tester males and target flies (4-8 days old)

253 were gently aspirated into round 2-layer chambers (diameter: 1 cm; height: 3 mm per

254 layer) and were separated by a plastic transparent barrier that was removed ~30 min

255 later to allow courtship test. Courtship index (CI), which is the percentage of

256 observation time a fly performs any courtship step, was used to measure courtship to

257 female targets or between 2 males. Paired male-male courtship used 2 males of the

258 same genotype but focused on the male fly that first initiated courtship (courtship of

259 the initiator to the other). All tester flies were single housed if not otherwise

260 mentioned. Each test was performed for 10 min.

261 For male chaining assay, tester males (4-8 days old) were loaded into large round
262 chambers (diameter: 4 cm; height: 3 mm) by cold anesthesia. Tests were performed
263 daily for 4 consecutive days (3 hours after grouping as day 0, then days 1–3).

264 Chaining index (ChI), which is the percentage of observation time at least 3 flies
265 engaged in courtship together, was used to measure courtship in groups of 8 males.

266

267 **Immunohistochemistry**

268 We dissected brains and ventral nerve cords of 5-7 days old males in Schneider's
269 insect medium (Thermo Fisher Scientific, Waltham, MA) and fixed in 4%
270 paraformaldehyde (PFA) in phosphate-buffered saline (PBS) for 30 min at room
271 temperature. After washing four times in PAT (0.5% Triton X-100 and 0.5% bovine
272 serum albumin in PBS), tissues were blocked in 3% normal goat serum (NGS) for 60
273 min, then incubated in primary antibodies diluted in 3% NGS for ~24 hr at 4°C,
274 washed (4 × 15-min) in PAT at room temperature, and incubated in secondary
275 antibodies diluted in 3% NGS for ~24 hr at 4°C. Tissues were then washed (4 × 15-
276 min) in PAT and mounted in Vectorshield (Vector Laboratories, Burlingame, CA) for
277 imaging. Primary antibodies used were rabbit anti-GFP (1:1000; A11122, Invitrogen,
278 Waltham, MA) and mouse anti-Bruchpilot (1:50; nc82, Developmental Studies
279 Hybridoma Bank, Iowa City, IA). Secondary antibodies used were goat anti-mouse

280 IgG conjugated to Alexa 555 (1:500, A28180, Invitrogen) and goat anti-rabbit IgG
281 conjugated to Alexa 488 (1:500, A11008, Invitrogen). Samples were imaged at 10× or
282 20× magnification on a Zeiss 700 confocal microscope and processed with ImageJ.

283

284 **Statistics**

285 Experimental flies and genetic controls were tested at the same condition, and data are
286 collected from at least two independent experiments. Statistical analysis is performed
287 using GraphPad Prism and indicated inside each figure legend. Data presented in this
288 study were first verified for normal distribution by D'Agostino–Pearson normality
289 test. If normally distributed, Student's *t* test is used for pairwise comparisons, and
290 one-way ANOVA is used for comparisons among multiple groups, followed by
291 Tukey's multiple comparisons. If not normally distributed, Mann-Whitney U test is
292 used for pairwise comparisons, and Kruskal-Wallis test is used for comparisons
293 among multiple groups, followed by Dunn's multiple comparisons.

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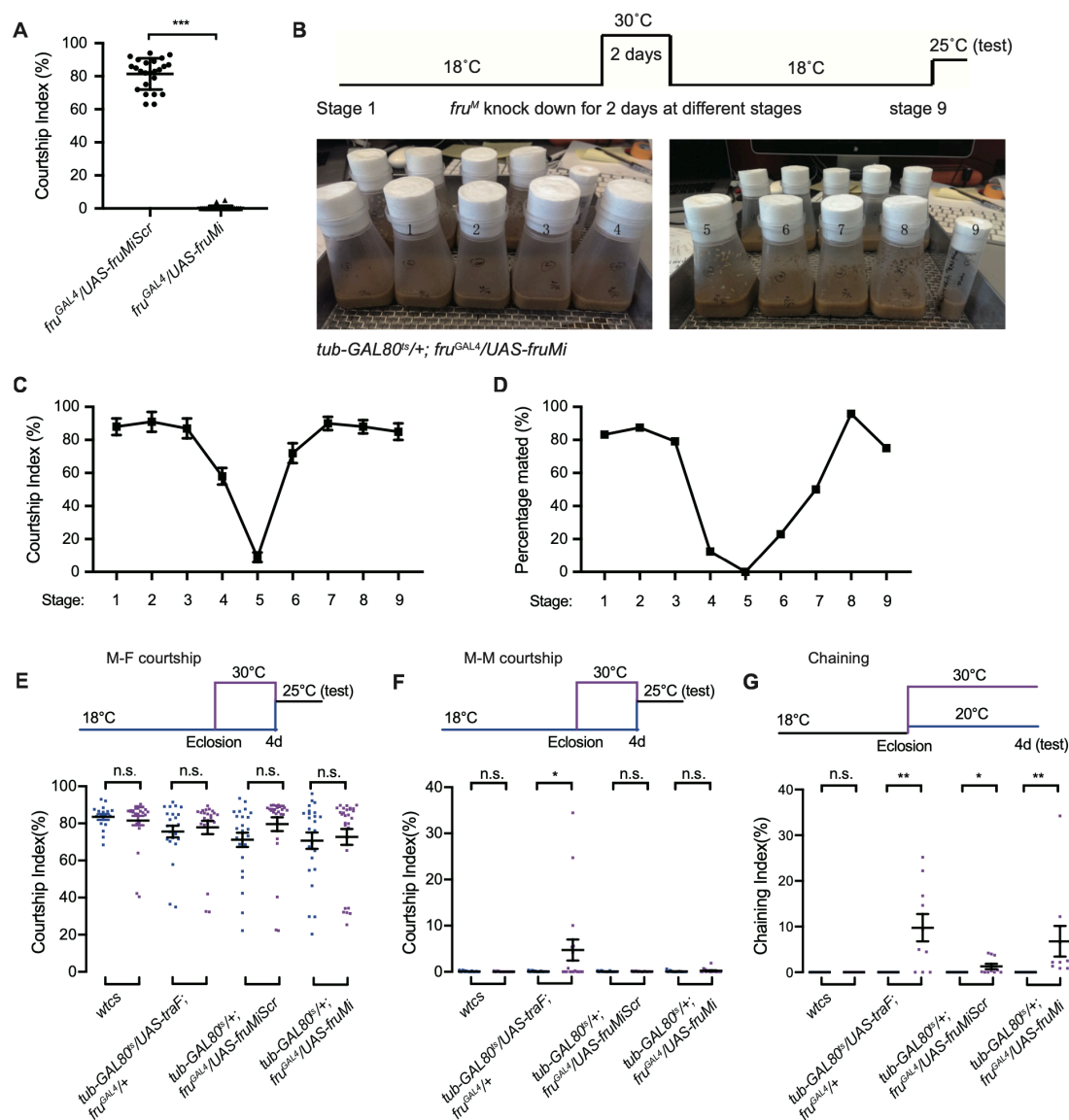
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403 **Figure legends**

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406 **Figure 1.** Distinct roles of *fru^M* during development and adulthood in regulating male

407 courtship. **A:** Knocking down *fru^M* using RNAi throughout development and

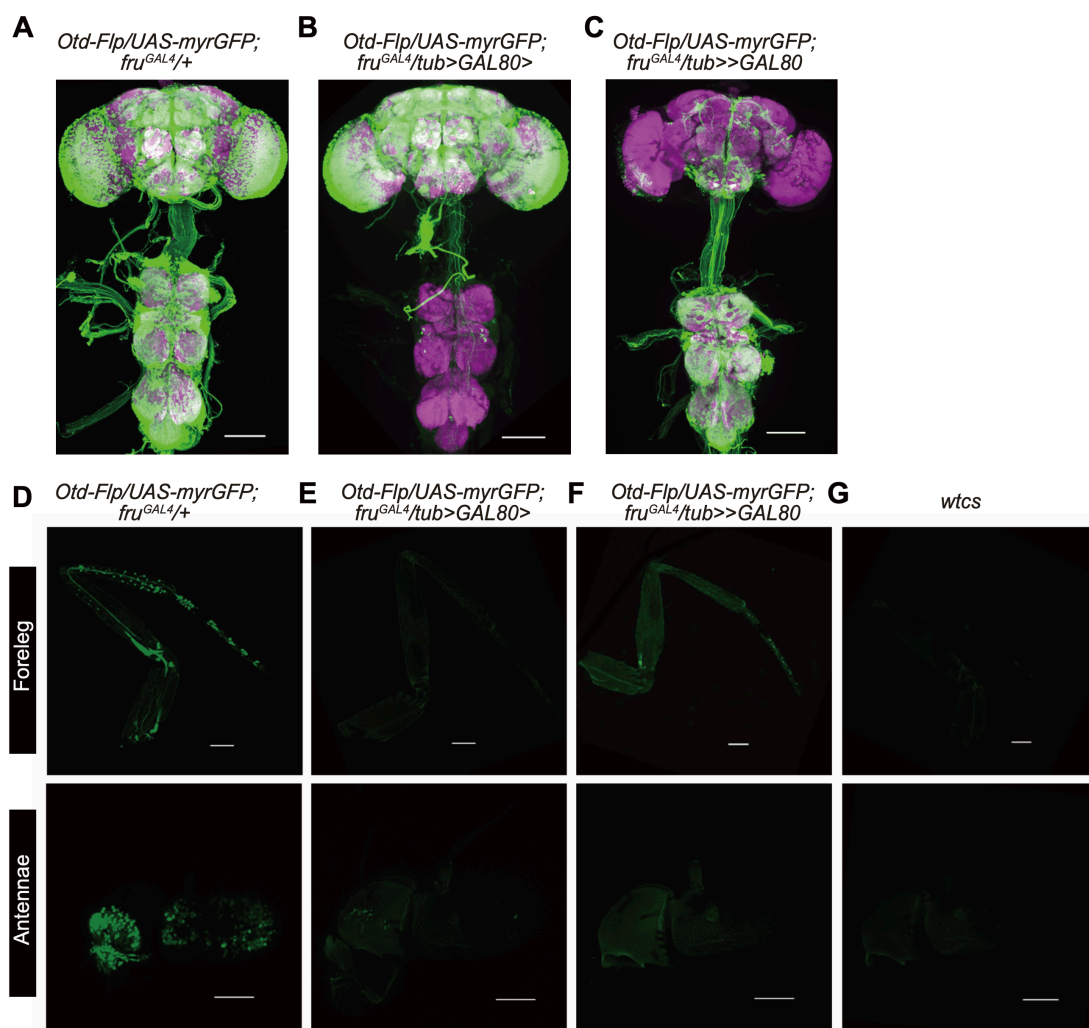
408 adulthood eliminated male courtship towards virgin females. $n = 24$ for each. $***p <$

409 0.001 , unpaired t-test. **B:** A schematic of genetic strategy to knock down *fru^M* at

410 different developmental stages for 2 days. Stages 1 to 9 refer to specific

411 developmental stages from embryos to adults with interval of 2 days. **C and D:**

412 Courtship indices of males with *fru^M* knocked down at specific developmental stages
413 as indicated above towards virgin females. Males with *fru^M* knocked down at stage 5
414 for 2 days (a period of pupation from stage 5 to 6, see above picture) rarely courted
415 virgin females (**C**), and none successfully mated (**D**). Knocking down *fru^M* at stages
416 near 5 (e.g., stage 4 or 6) also partially impairs courtship and mating success.
417 Knocking down *fru^M* at stage 9 (adult) has no obvious effect on courtship and mating.
418 $n = 24$ for each. **E-G**: Knocking down *fru^M* specifically during adulthood slightly
419 increased male-male courtship behaviors. For male-female courtship (**E**), $n = 17, 26,$
420 $23, 23, 24, 27, 24$ and 28 respectively (from left to right), n.s., not significant,
421 unpaired t-test. For single-pair male-male courtship (**F**), $n = 18$ for each. n.s., not
422 significant, $*p < 0.05$, unpaired t-test. For male chaining among 8 males as a group
423 (**G**), $n = 8, 8, 8, 10, 8, 9, 8$ and 9 respectively (from left to right). n.s., not significant,
424 $*p < 0.05$, $**p < 0.01$, Mann-Whitney test. Error bars indicate SEM.
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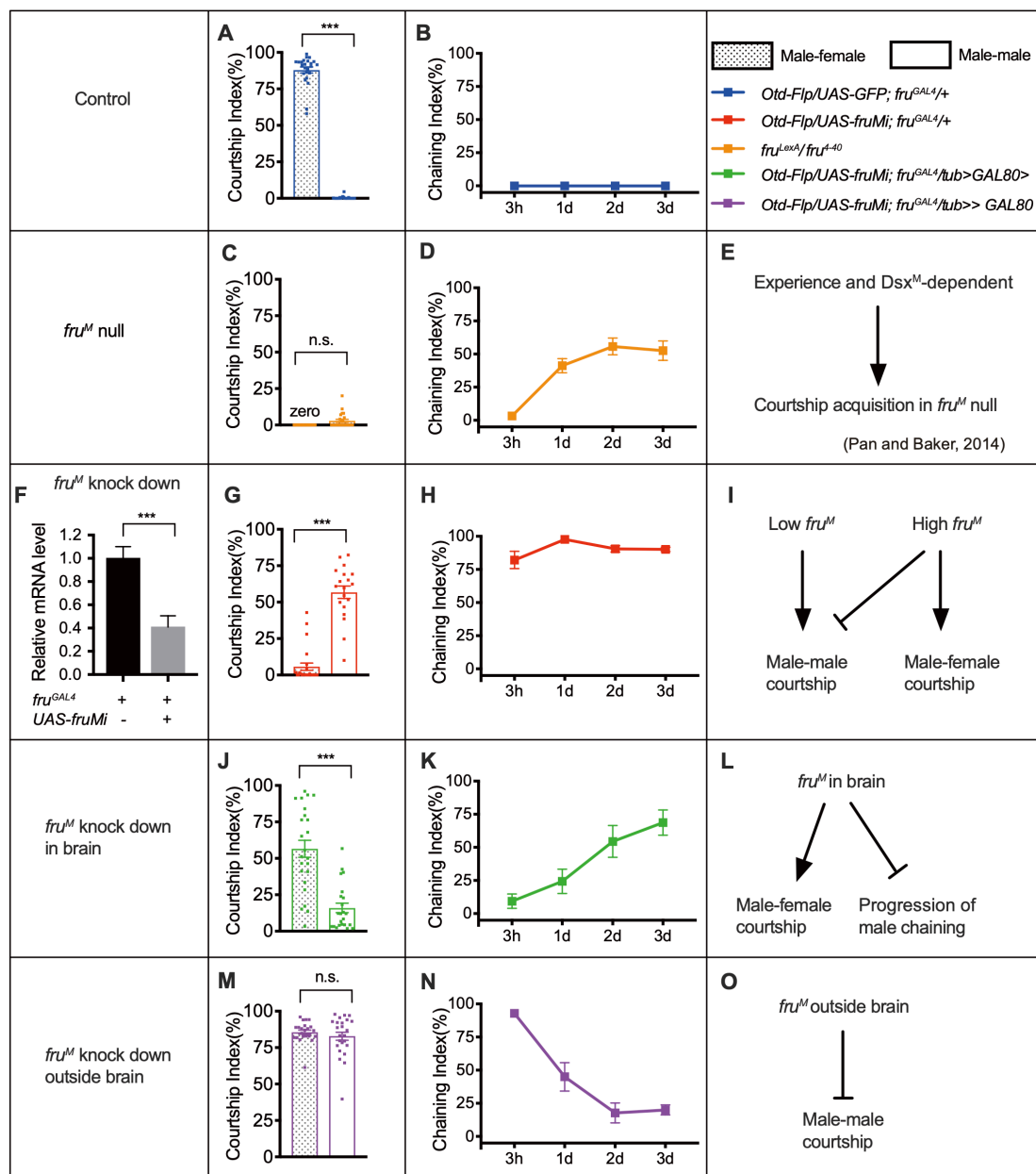
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427 **Figure 2.** Dividing *fru^M* expression into two complementary parts. A: Expression
428 pattern of the *fru^M* circuitry revealed by *fru^{GAL4}* driving *UAS-myrGFP* (green), co-
429 stained with nc82 (magenta). B and C: Genetic strategy to divide *fru^M* neurons (green)
430 into two parts. Scale bars above, 100 μ m. D-G: Expression pattern of *fru^{GAL4}* driving
431 *UAS-myrGFP* (green) in forelegs (scale bars, 100 mm) and antennae (scale bars, 50
432 mm) in males with indicated genotypes. Representative of 5 samples each.

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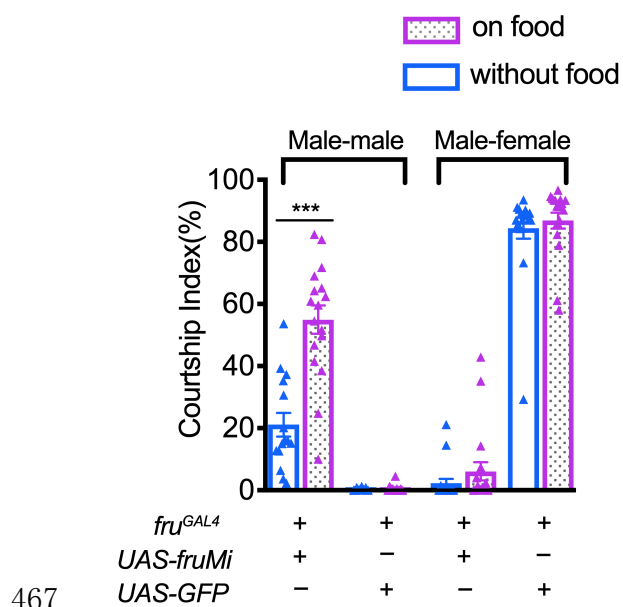
437 **Figure 3. *fru^M* tunes functional flexibility of the *fru^M* circuitry.** **A and B:** Wild-type
 438 males courted intensively towards virgin females (**A**, left bar), but rarely courted
 439 males (**A**, right bar) or displayed chaining behavior in groups of 8 males (**B**). $n = 24$,
 440 24, 8 respectively. *** $p < 0.001$, unpaired t-test. **C:** *fru^{LexA}/fru⁴⁻⁴⁰* (*fru^M* null) males
 441 rarely courted either females or males. $n = 24$ for each. n.s., not significant, unpaired
 442 t-test. **D:** *fru^{LexA}/fru⁴⁻⁴⁰* males did not show chaining behavior after 3-hr group-
 443 housing, but developed intensive chaining behavior after 1-3 days. $n = 8$. **E:** A

444 summary of courtship acquisition independent of *fru^M*. **F:** RNAi against *fru^M*
445 efficiently decreased but not fully eliminated *fru^M* expression. $n = 3$. *** $p < 0.001$,
446 Mann-Whitney U test. **G:** Knocking down *fru^M* in all *fru^{GAL4}* neurons generated males
447 that have reversed sexual orientation such that they rarely courted females but
448 intensively courted males. $n = 24$ and 19 respectively. *** $p < 0.001$, unpaired t-test.
449 **H:** Males with *fru^M* knocked down in all *fru^{GAL4}* neurons showed intensive chaining
450 behavior at all time points (from 3 hours to 3 days upon group-housing). $n = 7$. **I:**
451 Distinct roles of low *fru^M* (RNAi) and high *fru^M* (wild-type) in regulating male-male
452 and male-female courtship. **J:** Males with *fru^M* knocked down in *fru^{GAL4}* neurons in
453 the brain had a lower level of courtship towards females, but their sexual orientation
454 was not changed. $n = 24$ and 23 respectively. *** $p < 0.001$, unpaired t-test. **K:** Males
455 with *fru^M* knocked down in *fru^{GAL4}* neurons in brain showed low male chaining
456 behavior initially but increasing levels of chaining behavior over 1-3 days. $n = 6$. **L:** A
457 summary of the role of *fru^M* in brain in promoting male-female courtship and
458 suppressing the experience-dependent acquisition or progression of male chaining
459 behavior. **M:** Males with *fru^M* knocked down in *fru^{GAL4}* neurons outside brain
460 generated bisexual males that have intensive male-female and male-male courtship. n
461 = 24 for each. n.s., not significant, unpaired t-test. **N:** Males with *fru^M* knocked down
462 in *fru^{GAL4}* neurons outside brain showed high male chaining behavior initially, but
463 decreased levels of chaining behavior over 1-3 days. $n = 8$. **O:** A summary of the role

464 of *fru^M* outside brain in suppressing male-male courtship behavior. Error bars indicate

465 SEM.

466



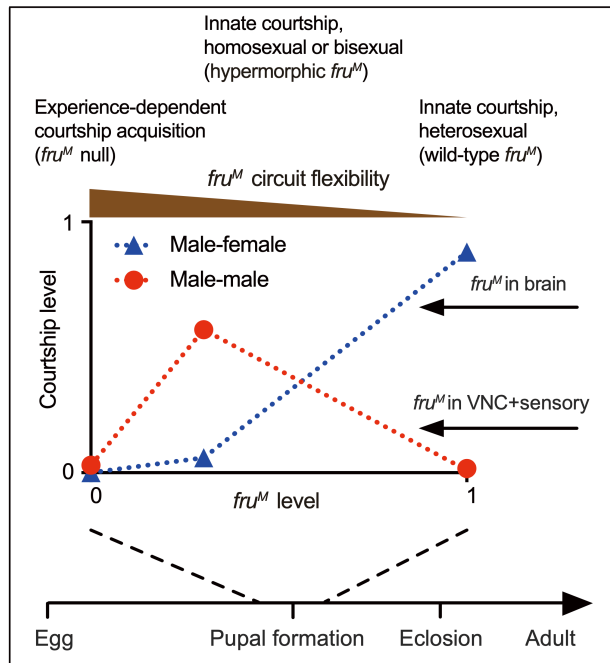
468 **Figure 3—figure supplement 1.** Comparison of male courtship with or without food.

469 Male-male courtship in *fru^M* knocked down males is higher in the presence of food. *n*

470 = 15, 17, 15, 18, 18, 18, 18 and 18 from left to right respectively. ****p* < 0.001,

471 unpaired t-test. Error bars indicate SEM.

472



473

474 **Figure 4.** A summary of *fru^M* function in male courtship. Firstly, *fru^M* is largely
475 required during a specific developmental period for courtship, and plays a minor role
476 during adulthood; secondly, the sex circuitry without *fru^M* or with different levels of
477 *fru^M* has different properties such that males would have experience-dependent
478 courtship acquisition, or innate courtship but with different sexual orientation. Such
479 flexibility of the sex circuitry is tuned by different *fru^M* expression. Red circles and
480 blue triangles represent corresponding *fru^M* levels and courtship levels (red: male-
481 male courtship; blue: male-female courtship).

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