

# *Drosophila* Bruton's Tyrosine Kinase Regulates Habituation Latency and Facilitation in Distinct Mushroom Body Neurons

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Habituation is the adaptive behavioral outcome of processes engaged in timely devaluation of non-reinforced repetitive stimuli, but the neuronal circuits and molecular mechanisms that underlie them are not well understood. To gain insights into these processes we developed and characterized a habituation assay to repetitive footshocks in mixed sex *Drosophila* groups and demonstrated that acute neurotransmission from adult  $\alpha/\beta$  mushroom body (MB) neurons prevents premature stimulus devaluation. Herein we demonstrate that activity of the non-receptor tyrosine kinase dBtk protein is required within these neurons to prevent premature habituation. Significantly, we also demonstrate that the complementary process of timely habituation to the repetitive stimulation is facilitated by  $\alpha'/\beta'$  MB neurons and also requires dBtk activity. Hence our results provide initial insights into molecular mechanisms engaged in footshock habituation within distinct MB neurons. Importantly, dBtk attenuation specifically within  $\alpha'/\beta'$  neurons leads to defective habituation, which is readily reversible by administration of the antipsychotics clozapine and risperidone suggesting that the loss of the kinase may dysregulate monoamine receptors within these neurons, whose activity underlies the failure to habituate.

**Key words:** antipsychotics; Btk; *Drosophila*; habituation; mushroom bodies

## Significance Statement

Habituation refers to processes underlying decisions to attend or ignore stimuli, which are pivotal to brain function as they underlie selective attention and learning, but the circuits involved and the molecular mechanisms engaged by the process therein are poorly understood. We demonstrate that habituation to repetitive footshock involves two phases mediated by distinct neurons of the *Drosophila* mushroom bodies and require the function of the dBtk non-receptor tyrosine kinase. Moreover, habituation failure upon dBtk abrogation in neurons where it is required to facilitate the process is readily reversible by antipsychotics, providing conceptual links to particular symptoms of schizophrenia in humans, also characterized by habituation defects and ameliorated by these pharmaceuticals.

## Introduction

Habituation is a form of adaptive behavioral plasticity that permits animals to ignore repetitive or prolonged non-reinforced

stimuli (Thompson and Spencer, 1966; Rankin et al., 2009). Timely devaluation of such stimuli is the behavioral output of largely undefined molecular processes, apparently engaging multiple cellular systems (Ramaswami, 2014). Because habituation underlies gated selective attention and discrimination between novel and pre-experienced stimuli (Gillberg, 2003), it represents a form of adaptive behavioral flexibility, reported defective in patients with migraines (Siniatchkin et al., 2003; Kalita et al.,

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2014), attention deficit hyperactivity disorder (Jansiewicz et al., 2004; Massa and O'Desky, 2012), schizophrenia (SD; Braff et al., 2001; Meincke et al., 2004) and autism spectrum disorders (ASD; D'Cruz et al., 2013; Tei et al., 2018). Hence, defining molecular mechanisms that govern habituation is likely to contribute toward understanding the molecular etiology of these conditions.

To explore habituation mechanisms of CNS circuits in *Drosophila*, we developed a novel habituation paradigm to repetitive mild electric footshock (Acevedo et al., 2007). Habituation to repetitive footshock requires structurally and functionally intact mushroom bodies (MBs), neurons also essential for associative learning and memory (Guven-Ozkan and Davis, 2014). The MBs are comprised by ~2000 neurons per hemisphere subdivided into  $\alpha/\beta$ ,  $\alpha'/\beta'$  and  $\gamma$  subtypes (Crittenden et al., 1998). Because neurotransmission from  $\alpha/\beta$  MB neurons is required to suppress premature habituation, we proposed that physiologically, it blocks devaluation of the repetitive footshock stimuli, hence required for habituation latency (Acevedo et al., 2007). However, whether other intrinsic or extrinsic MB neurons are required to facilitate habituation remained unclear.

To address this question and to identify proteins governing habituation within the MBs, we have conducted a genetic screen for mutants that present defective habituation. Here we report on the role of one of the proteins identified in the screen, Bruton's tyrosine kinase (dBtk), which belongs to the Src non-receptor kinase superfamily and is associated with agammaglobulinemia in humans (Mattsson et al., 1996). The *Drosophila* dBtk29A (dBtk) gene encodes two proteins by alternative splicing, dBtk type 1 and dBtk type 2, with the latter considered orthologous to the human protein (Gregory et al., 1987). dBtk consists of conserved SH2, SH3 and kinase domains, whereas an N-terminal plekstrin homology (PH) domain characterizes the larger type 2 protein (Tsikala et al., 2014). dBtk is implicated in many essential functions in *Drosophila* (Gregory et al., 1987; Roulier et al., 1998; Baba et al., 1999; Hamada-Kawaguchi and Yamamoto, 2017), including regulation of the actin cytoskeleton (Tsikala et al., 2014). Although dBtk is highly expressed in the fly CNS, there is limited information regarding its functional role(s) therein (Asztalos et al., 2007; Sunouchi et al., 2016). Here we demonstrate acute differential roles for this kinase within distinct MB neuronal populations in the regulation of habituation dynamics to repeated footshock stimuli.

## Materials and Methods

### *Drosophila* culture and strains

*Drosophila* were cultured in standard wheat-flour-sugar food supplemented with soy flour and CaCl<sub>2</sub> (Acevedo et al., 2007) at 18°C or 25°C. All MiMIC insertions were from the Bloomington Stock Center (BDRC; Indiana University; Venken et al., 2011) and they were backcrossed to  $y^1w^1$  for at least seven generations before use in behavioral experiments. MBGal80 (Krashes et al., 2007) was obtained from Ron Davis (Scripps Florida). The Btk-Gal4 (49182), dncGal4 (48571), and Btk RNAi stocks (35159 and 25791) were from BDRC. To generate the driver heterozygote controls for experiments with the RNAi-encoding transgenes, driver-bearing strains were crossed to their  $y^1v^1$  (BDSC, 36303) background. The UAS-Btk lines (109-093 and 109-095) were from the Kyoto Stock Center (Kyoto Institute of Technology). VT44966-Gal4 ( $\gamma$ -driver) was from the Vienna *Drosophila* Resource Center (VDRC; Vienna BioCenter Core Facilities, 203571). The  $\alpha'/\beta'$  Gal4 drivers VT030604 (VDRC, 200228) and c305a were a kind gift from S. Waddell (University of Oxford). The glial driver repo-Gal4, the pan-neuronal drivers elav-Gal4 and Ras2Gal4, and the mushroom body specific drivers 247-Gal4, leo-Gal4, c739-Gal4, c772-Gal4 were described previously (Aso et al., 2009; Gouzi et al., 2018). The Gal80<sup>ts</sup> transgene was added to the driver-bearing chromosomes by recombination or standard crosses as indicated.

### Description of Gal4 expression patterns used in this work.

*Elav*: pan-neuronal expression in all developmental stages (FlyBase ID: FBrf0237128)

*Ras2*: throughout the larval and adult CNS, enriched in the adult MBs (Gouzi et al., 2011)

*repo*: all glia (FlyBase ID: FBrf0237128)

*leo*: adult  $\alpha$ ,  $\alpha'$ ,  $\beta$ ,  $\beta'$ ,  $\gamma$  MB neurons (Messaritou et al., 2009)

*247*: adult  $\alpha$ ,  $\alpha'$ ,  $\beta$ ,  $\beta'$ ,  $\gamma$  MB neurons (RRID:BDSC\_50742)

*dnc*: adult  $\alpha$ ,  $\alpha'$ ,  $\beta$ ,  $\beta'$ ,  $\gamma$  MB neurons, scattered neurons in subesophageal ganglion and ventral optic lobes (Gai et al., 2016)

*c772*: adult  $\alpha$ ,  $\beta$ ,  $\gamma$  MB neurons, antennal lobe, medulla, tritocerebrum (Aso et al., 2009)

*c739*: adult  $\alpha$ ,  $\beta$  MB neurons, antennal lobe, medulla, restricted protocerebral neurons, inferior neuropils (Aso et al., 2009).

*VT44966*:  $\gamma$  MB neurons, wedge neurons, superior lateral protocerebrum, gnathal neurons, medial bundle (Shyu et al., 2017)

*c305a*: adult  $\alpha'$ ,  $\beta'$  MB neurons, antennal nerve, medulla, restricted protocerebral neurons, inferior neuropil, gnathal neurons (Aso et al., 2009)

*VT030604*: adult  $\alpha'$ ,  $\beta'$  MB neurons, gnathal neurons (Shyu et al., 2017)

### Behavioral assays

All flies used in behavioral experiments were tested 3–5 d after emergence. All experiments were performed under dim red light at 25°C and 65–75% relative humidity. To obtain animals for behavioral experiments Gal4 driver homozygotes were crossed *en masse* to strains carrying either UAS-btk, UAS-btk-RNAi, or UAS-shi<sup>ts</sup> transgenes. Animals expressing Gal80<sup>ts</sup> (McGuire et al., 2003) were raised at 18°C until hatching and then placed at 30°C for 2 d before testing. Flies carrying UAS-shi<sup>ts</sup> were reared at 18°C and the dynamin was inactivated by incubation at 32°C for 30 min before the behavioral experiment.

*Electric footshock avoidance*. Experiments were performed as described before (Acevedo et al., 2007). Briefly, ~70 flies were placed at the choice point of a T-maze to choose for 90 s between an electrified and an otherwise identical inert standard copper grid. In the electrified grid, 45 V shocks were delivered every 5.15 s, each lasting 1.2 s. The avoiding fraction (AF) was calculated by dividing the number of flies avoiding the shock by the total number of flies.

*Habituation to electric footshock*. Habituation to electric shock experiments were performed as described before (Acevedo et al., 2007). Briefly, for the training phase ~70 flies were sequestered in the upper arm of a standard T-maze lined with an electrifiable grid. They were exposed to 15 1.2 s electric shocks at 45 V with a 5.15 s interstimulus interval. Air was not drawn through the tube during training to avoid association of the shocks with air. After a 30 s rest and 30 s for transfer to the lower part of the maze, the flies were tested by choosing between an electrified and an inert grid. Therefore, the earliest measures of post-training responses are 1 min after the flies received the last training stimulus. Testing was performed at the same voltage (45 V) as for training. During the 90 s choice period, 17–18 1.2 s stimuli were delivered to the electrified arm of the maze. At the end of the choice period, the flies in each arm were trapped and counted, and the habituation fraction (HF) was calculated by dividing the number of flies preferring the shock by the total number of flies, as above. Finally, the habituation index (HI) was calculated as  $(HF - AF) \times 100\%$  and therefore represents the change in footshock avoidance contingent upon prior footshock experience (habituation). Although the absolute avoidance score is variable, even for the same genotypes (Table 1), as expected for behavioral experiments performed over a significant time period, because the HI measures the relative change in avoidance within each genotype, it is not affected by such variability. In fact, failure to habituate, which is the primary phenotype reported herein, is a manifestation of maintained avoidance relative to that of naive flies of the same genotype.

*Dishabituation*. To distinguish habituation from fatigue or sensory adaptation, flies were dishabituated post-training with an 8 s puff [yeast odor (YO)] of air drawn at 500 ml/min over a 30% (w/v) aqueous solution of Brewer's yeast (Acros Organics) and then were submitted to testing.

**Table 1. Collective avoidance indices and statistics**

Genotype	Mean ± SEM	t Ratio	p
Fig. 1C. ANOVA: $F_{(2,58)} = 1.6020, p = 0.2106$			
c (control)	0.76 ± 0.03		
btk <sup>M1/+</sup>	0.77 ± 0.05	0.106	0.9158
btk <sup>M1/btk<sup>M1</sup></sup>	0.66 ± 0.07	1.5444	0.1281
Fig. 1D. ANOVA: $F_{(5,68)} = 0.8807, p = 0.4993$			
c (control)	0.61 ± 0.02		
btk <sup>M2/M2</sup>	0.62 ± 0.04	−0.007	0.9945
btk <sup>M1/M1</sup>	0.59 ± 0.05	0.4748	0.6365
btk <sup>M1/ML</sup>	0.55 ± 0.05	1.0243	0.310
btk <sup>M2/ML</sup>	0.53 ± 0.05	1.4211	0.1602
btk <sup>ML/+</sup>	0.63 ± 0.03	−0.192	0.848
Fig. 1E. ANOVA: $F_{(3,60)} = 0.8603, p = 0.4670$			
0 μM	0.53 ± 0.03		
0.1 μM	0.49 ± 0.03	1.2032	0.2339
1 μM	0.55 ± 0.03	−0.291	0.7722
10 μM	0.49 ± 0.04	0.9707	0.3358
Fig. 1F. ANOVA: $F_{(3,62)} = 1.9514, p = 0.1312$			
c (control)	0.54 ± 0.04		
btk <sup>M1/M1</sup>	0.67 ± 0.05	−1.737	0.0877
btk <sup>M2/M2</sup>	0.51 ± 0.06	0.3651	0.7163
btk <sup>ML/+</sup>	0.62 ± 0.06	−1.031	0.3066
Fig. 2A. ANOVA: $F_{(2,44)} = 1.3353, p = 0.2740$			
elavGAL4/+; UAS-btk <sup>RNAi1</sup> /GAL80 <sup>ts</sup>	0.59 ± 0.05		
elavGAL4/+; GAL80 <sup>ts</sup> /+	0.61 ± 0.04	−0.237	0.8139
UASbtk <sup>RNAi1</sup> /+	0.49 ± 0.07	1.3101	0.1973
Fig. 2B. ANOVA: $F_{(2,35)} = 2.0785, p = 0.1412$			
elavGAL4/+; UAS-btk <sup>RNAi2</sup> /GAL80 <sup>ts</sup>	0.66 ± 0.04		
elavGAL4/+; GAL80 <sup>ts</sup> /+	0.72 ± 0.03	−1.105	0.2771
UASbtk <sup>RNAi2</sup> /+	0.78 ± 0.03	−2.006	0.0531
Fig. 2C. ANOVA: $F_{(2,30)} = 1.100, p = 0.3468$			
elavGAL4/+; UAS-btk <sup>RNAi1</sup> /GAL80 <sup>ts</sup>	0.73 ± 0.05		
elavGAL4/+; GAL80 <sup>ts</sup> /+	0.63 ± 0.03	1.4796	1.1502
UASbtk <sup>RNAi1</sup> /+	0.68 ± 0.05	0.7939	0.4340
Fig. 2D. ANOVA: $F_{(2,27)} = 0.1719, p = 0.8431$			
Ras2GAL4, GAL80 <sup>ts</sup> /+; UASbtk <sup>RNAi1</sup>	0.67 ± 0.08		
Ras2GAL4, GAL80 <sup>ts</sup> /+	0.65 ± 0.05	0.1811	0.8578
UASbtk <sup>RNAi1</sup> /+	0.70 ± 0.04	−0.386	0.703
Fig. 2E. ANOVA: $F_{(2,46)} = 2.1609, p = 0.1273$			
repoGAL4, GAL80 <sup>ts</sup> /UAS-btk <sup>RNAi1</sup>	0.63 ± 0.05		
repoGAL4, GAL80 <sup>ts</sup> /+	0.73 ± 0.03	−2.075	0.044
UASbtk <sup>RNAi1</sup> /+	0.67 ± 0.03	−0.955	0.3449
Fig. 2F. ANOVA: $F_{(2,47)} = 3.2093, p = 0.0498$			
elavGAL4/+; MBG80, GAL80 <sup>ts</sup> /UASbtk <sup>RNAi1</sup>	0.71 ± 0.02		
elavGAL4/+; MBG80, GAL80 <sup>ts</sup> /+	0.63 ± 0.02	2.2266	0.031
UASbtk <sup>RNAi1</sup> /+	0.71 ± 0.03	−0.085	0.932
Fig. 3A. ANOVA: $F_{(2,34)} = 0.5313, p = 0.5929$			
leoMBGAL4, GAL80 <sup>ts</sup> /UAS-btk <sup>RNAi1</sup> /+	0.46 ± 0.09		
leoMB-GAL4, GAL80 <sup>ts</sup> /+	0.56 ± 0.07	−0.905	0.3723
UAS-btk <sup>RNAi1</sup> /+	0.56 ± 0.07	−0.892	0.3788
Fig. 3B. ANOVA: $F_{(5,66)} = 2.7712, p = 0.0255$			
leoMBGAL4, btk <sup>M1/+</sup> ; GAL80 <sup>ts</sup> /+ UN	0.65 ± 0.03		
leoMBGAL4, btk <sup>M1/btk<sup>M1</sup></sup> ; GAL80 <sup>ts</sup> /+ UN	0.50 ± 0.03	2.205	0.031
leoMBGAL4, btk <sup>M1/btk<sup>M1</sup></sup> ; GAL80 <sup>ts</sup> , UASbtk <sup>5</sup> UN	0.59 ± 0.05	0.9337	0.3541
leoMBGAL4, btk <sup>M1/+</sup> ; GAL80 <sup>ts</sup> /+ IN	0.56 ± 0.04		
leoMBGAL4, btk <sup>M1/btk<sup>M1</sup></sup> ; GAL80 <sup>ts</sup> /+ IN	0.45 ± 0.04	−1.265	0.211
leoMBGAL4, btk <sup>M1/btk<sup>M1</sup></sup> ; GAL80 <sup>ts</sup> , UASbtk <sup>5</sup> IN	0.62 ± 0.04	−0.914	0.364
Fig. 3C. ANOVA: $F_{(2,39)} = 0.5892, p = 0.5599$			
leoMBGAL4/+; UASbtk <sup>RNAi2</sup> /GAL80 <sup>ts</sup>	0.65 ± 0.05		
leoMBGAL4/+; GAL80 <sup>ts</sup> /+	0.68 ± 0.03	−0.471	0.640
UAS-btk <sup>RNAi2</sup> /+	0.61 ± 0.06	0.6403	0.5260
Fig. 3D. ANOVA: $F_{(2,42)} = 0.4747, p = 0.6256$			
247MBGAL4, GAL80 <sup>ts</sup> /UAS-btk <sup>RNAi1</sup>	0.63 ± 0.03		
247MBGAL4, GAL80 <sup>ts</sup> /+	0.60 ± 0.02	0.9688	0.3385
UASbtk <sup>RNAi1</sup> /+	0.61 ± 0.03	0.5817	0.5640

(Table continues.)

**Table 1. Continued**

Genotype	Mean ± SEM	t Ratio	p
Fig. 3E. ANOVA: $F_{(2,30)} = 0.2194, p = 0.8044$			
dncGAL4, GAL80 <sup>ts</sup> /UASbtk <sup>RNAi1</sup>	0.53 ± 0.02		
dncGAL4, GAL80 <sup>ts</sup> /+	0.51 ± 0.02	0.4624	0.6474
UASbtk <sup>RNAi1</sup> /+	0.54 ± 0.03	−0.198	0.8443
Fig. 3F. ANOVA: $F_{(2,44)} = 1.1638, p = 0.3222$			
c772GAL4, GAL80 <sup>ts</sup> /UASbtk <sup>RNAi1</sup>	0.63 ± 0.03		
c772GAL4, GAL80 <sup>ts</sup> /+	0.54 ± 0.06	1.451	0.1542
UASbtk <sup>RNAi1</sup> /+	0.57 ± 0.03	1.1173	0.2702
Fig. 3G. ANOVA: $F_{(2,43)} = 1.891, p = 0.1766$			
UASbtk <sup>RNAi1</sup> /c739GAL4, GAL80 <sup>ts</sup>	0.57 ± 0.05		
c739GAL4, GAL80 <sup>ts</sup> /+	0.46 ± 0.04	1.8657	0.0692
UASbtk <sup>RNAi1</sup> /+	0.53 ± 0.04	0.6998	0.488
Fig. 3H. ANOVA: $F_{(2,47)} = 3.0762, p = 0.0559$			
γlobeGAL4/+; UASbtk <sup>RNAi1</sup> /GAL80 <sup>ts</sup>	0.58 ± 0.03		
γlobeGAL4/+; GAL80 <sup>ts</sup> /+	0.49 ± 0.03	2.0924	0.0421
UASbtk <sup>RNAi1</sup> /+	0.59 ± 0.03	−0.238	0.8133
Fig. 3I. ANOVA: $F_{(2,28)} = 1.0786, p = 0.3548$			
c305aGAL4/+; UASbtk <sup>RNAi1</sup> /GAL80 <sup>ts</sup>	0.77 ± 0.03		
c305aGAL4/+; GAL80 <sup>ts</sup> /+	0.71 ± 0.02	1.3002	0.2049
UASbtk <sup>RNAi1</sup> /+	0.77 ± 0.04	0.0361	0.9715
Fig. 3J. ANOVA: $F_{(2,49)} = 2.5477, p = 0.0156$			
VT030604GAL4/+; UASbtk <sup>RNAi1</sup> 18°C	0.74 ± 0.03		
VT030604GAL4/+; UASbtk <sup>RNAi1</sup> 25°C	0.67 ± 0.02	1.9623	0.0557
VT030604-GAL4/+; UAS-btk <sup>RNAi1</sup> 18 > 30°C	0.78 ± 0.03	−0.972	0.3361
Fig. 3K. ANOVA: $F_{(1,25)} = 9.7589, p = 0.0046$			
c305aGAL4-UAS shi <sup>ts</sup> UN	0.85 ± 0.03		
c305aGAL4-UAS shi <sup>ts</sup> IN	0.75 ± 0.01	−3.124	<b>0.0046</b>
Fig. 4A. ANOVA: $F_{(5,104)} = 4.7413, p = 0.0006$			
control 6 sh	0.79 ± 0.04		
btk <sup>M1/M1</sup> 6 sh	0.63 ± 0.04	2.9535	<b>0.0039</b>
control 10 sh	0.69 ± 0.03		
btk <sup>M1/M1</sup> 10 sh	0.60 ± 0.04	1.6587	0.1003
control 15 sh	0.73 ± 0.02		
btk <sup>M1/M1</sup> 15 sh	0.58 ± 0.04	2.9818	<b>0.0036</b>
Fig. 4B. ANOVA: $F_{(8,156)} = 2.8880, p = 0.0051$			
UASbtk <sup>RNAi1</sup> /elavGAL4; GAL80 <sup>ts</sup> 6 sh	0.82 ± 0.02		
elavGAL4; GAL80 <sup>ts</sup> /+ 6 sh	0.66 ± 0.03	3.7031	<b>0.0003</b>
UAS-btk <sup>RNAi1</sup> /+ 6 sh	0.71 ± 0.04	2.5227	0.0127
UASbtk <sup>RNAi1</sup> /elavGAL4; GAL80 <sup>ts</sup> 10 sh	0.77 ± 0.02		
elavGAL4; GAL80 <sup>ts</sup> /+ 10 sh	0.66 ± 0.03	2.2533	0.0257
UAS btk <sup>RNAi1</sup> /+ 10 sh	0.66 ± 0.06	2.0069	0.0466
UASbtk <sup>RNAi1</sup> /elavGAL4; GAL80 <sup>ts</sup> 15 sh	0.73 ± 0.04		
elavGAL4; GAL80 <sup>ts</sup> /+ 15 sh	0.74 ± 0.03	−0.171	0.8644
UASbtk <sup>RNAi1</sup> /+ 15 sh	0.70 ± 0.03	0.6554	0.5132
Fig. 4C. ANOVA: $F_{(2,28)} = 1.2620, p = 0.2999$			
0 μM	0.58 ± 0.05		
0.1 μM	0.64 ± 0.04	−0.832	0.4129
1 μM	0.68 ± 0.04	−1.588	0.1245
Fig. 4D. ANOVA: $F_{(2,30)} = 0.0147, p = 0.9854$			
Habituation, 15 shocks	0.76 ± 0.02		
YO dishabituation	0.77 ± 0.02	−0.147	0.8839
AVD after yeast puff	0.77 ± 0.02	−0.147	0.8839
Fig. 4E. ANOVA: $F_{(2,30)} = 0.0147, p = 0.9854$			
btk <sup>M1/M1</sup> habituation 6 sh	0.68 ± 0.03		
btk <sup>M1/M1</sup> Dishabituation 6 sh	0.68 ± 0.03	0	1
btk <sup>M1/M1</sup> habituation 10 sh	0.73 ± 0.04		
btk <sup>M1/M1</sup> Dishabituation 10 sh	0.73 ± 0.03	0.0617	0.9511
btk <sup>M1/M1</sup> AVD after yeast puff	0.73 ± 0.04	−1.102	0.2772
Fig. 5A. ANOVA: $F_{(5,119)} = 3.8885, p = 0.0027$			
leoGAL4/+; UASbtk <sup>RNAi1</sup> /GAL80 <sup>ts</sup> UN 6 sh	0.81 ± 0.02		
leoMBGAL4/+; UASbtk <sup>RNAi1</sup> /GAL80 <sup>ts</sup> IN 6 sh	0.78 ± 0.02	1.2528	0.2128
leoMBGAL4/+; UASbtk <sup>RNAi1</sup> /GAL80 <sup>ts</sup> IN 10 sh	0.82 ± 0.01		
leoMBGAL4/+; UASbtk <sup>RNAi1</sup> /GAL80 <sup>ts</sup> UN 10 sh	0.75 ± 0.02	3.053	<b>0.0052</b>
leoMBGAL4/+; UASbtk <sup>RNAi1</sup> /GAL80 <sup>ts</sup> IN 15 sh	0.82 ± 0.01		
leoMB-GAL4/+; UASbtk <sup>RNAi1</sup> /GAL80 <sup>ts</sup> UN 15 sh	0.75 ± 0.02	3.053	<b>0.0028</b>

(Table continues.)

**Table 1. Continued**

Genotype	Mean ± SEM	t Ratio	p
Fig. 5B. ANOVA: $F_{(5,77)} = 0.2798, p = 0.9227$			
leoG4, btk <sup>M1</sup> /btk <sup>M1</sup> , G80ts, UAS-btk <sup>5</sup> UN 6 sh	0.69 ± 0.03		
leoG4, btk <sup>M1</sup> /btk <sup>M1</sup> , G80ts, UAS-btk <sup>5</sup> IN 6 sh	0.71 ± 0.03	−0.391	0.6966
leoG4, btk <sup>M1</sup> /btk <sup>M1</sup> , G80ts, UAS-btk <sup>5</sup> UN 10 sh	0.69 ± 0.03		
leoG4, btk <sup>M1</sup> /btk <sup>M1</sup> , G80ts, UAS-btk <sup>5</sup> IN 10 sh	0.72 ± 0.03	−0.759	0.4505
leoG4, btk <sup>M1</sup> /btk <sup>M1</sup> , G80ts, UAS-btk <sup>5</sup> UN 15 sh	0.69 ± 0.03		
leoG4, btk <sup>M1</sup> /btk <sup>M1</sup> , G80ts, UAS-btk <sup>5</sup> IN 15 sh	0.72 ± 0.03	−0.759	0.4505
Fig. 5C. ANOVA: $F_{(5,77)} = 1.1341, p = 0.3503$			
UAS-btk <sup>RNAi1</sup> /c739-GAL4, GAL80 <sup>TS</sup> UN 6 sh	0.73 ± 0.02		
UAS-btk <sup>RNAi1</sup> /c739-GAL4, GAL80 <sup>TS</sup> IN 6 sh	0.72 ± 0.02	0.4393	0.6617
UAS-btk <sup>RNAi1</sup> /c739-GAL4, GAL80 <sup>TS</sup> UN 10 sh	0.74 ± 1.01		
UAS-btk <sup>RNAi1</sup> /c739-GAL4, GAL80 <sup>TS</sup> IN 10 sh	0.70 ± 0.02	1.2977	0.1985
UAS-btk <sup>RNAi1</sup> /c739-GAL4, GAL80 <sup>TS</sup> UN 15 sh	0.74 ± 1.01		
UAS-btk <sup>RNAi1</sup> /c739-GAL4, GAL80 <sup>TS</sup> IN 15 sh	0.70 ± 0.02	1.2977	0.1985
Fig. 5D. ANOVA: $F_{(5,82)} = 1.3274, p = 0.2614$			
c305aGAL4, GAL80 <sup>TS</sup> /UAS-btk <sup>RNAi1</sup> UN 6 sh	0.78 ± 0.02		
c305aGAL4, GAL80 <sup>TS</sup> /UAS-btk <sup>RNAi1</sup> IN 6 sh	0.74 ± 0.02	1.3687	0.1751
c305aGAL4, GAL80 <sup>TS</sup> /UAS-btk <sup>RNAi1</sup> UN 10 sh	0.78 ± 0.02		
c305aGAL4, GAL80 <sup>TS</sup> /UAS-btk <sup>RNAi1</sup> IN 10 sh	0.74 ± 0.02	1.5464	0.1261
c305aGAL4, GAL80 <sup>TS</sup> /UAS-btk <sup>RNAi1</sup> UN 15 sh	0.78 ± 0.02		
c305aGAL4, GAL80 <sup>TS</sup> /UAS-btk <sup>RNAi1</sup> IN 15 sh	0.74 ± 0.02	1.5222	1.1321
Fig. 6A. ANOVA: $F_{(2,28)} = 1.1989, p = 0.3176$			
0 μM	0.37 ± 0.04		
5 μM	0.45 ± 0.04	−1.45	0.1591
10 μM	0.43 ± 0.03	−1.139	0.2652
Fig. 6B. ANOVA: $F_{(2,30)} = 0.8493, p = 0.4384$			
0 μM	0.50 ± 0.05		
5 μM	0.54 ± 0.05	−0.624	0.5376
10 μM	0.46 ± 0.03	0.5694	0.5736
Fig. 6C. ANOVA: $F_{(3,44)} = 1.3657, p = 0.2667$			
0 μM	0.50 ± 0.03		
0.1 μM	0.57 ± 0.06	−1.432	0.1598
1 μM	0.56 ± 0.02	−1.373	0.1772
10 μM	0.50 ± 0.03	−0.063	0.9502
Fig. 6D. ANOVA: $F_{(2,26)} = 8.8963, p = 0.0013^*$			
0 μM	0.71 ± 0.03		
5 μM	0.53 ± 0.07	2.7508	<b>0.0111</b>
10 μM	0.78 ± 0.03	−1.216	0.0243
Fig. 6E. ANOVA: $F_{(2,27)} = 1.3673, p = 0.2732$			
c305aGAL4/+; UAS-btk <sup>RNAi1</sup> /GAL80 <sup>TS</sup>	0.66 ± 0.04		
c305aGAL4/+; UAS-btk <sup>RNAi1</sup> /GAL80 <sup>TS</sup> Clz	0.73 ± 0.03	−1.54	0.1361
c305aGAL4/+; UAS-btk <sup>RNAi1</sup> /GAL80 <sup>TS</sup> Ris	0.72 ± 0.03	−1.26	0.2193
Fig. 6F. ANOVA: $F_{(2,26)} = 2.2671, p = 0.1254$			
c739GAL4/+; UAS-btk <sup>RNAi1</sup> /GAL80 <sup>TS</sup>	0.66 ± 0.04		
c739GAL4/+; UAS-btk <sup>RNAi1</sup> /GAL80 <sup>TS</sup> Clz	0.73 ± 0.03	1.6255	0.1171
c739GAL4/+; UAS-btk <sup>RNAi1</sup> /GAL80 <sup>TS</sup> Ris	0.72 ± 0.03	−0.378	0.7085
Fig. 6G. ANOVA: $F_{(5,95)} = 1.8751, p = 0.1065$			
btk <sup>M1/M1</sup> 6 sh	0.44 ± 0.05		
btk <sup>M1/M1</sup> Clz 6 sh	0.43 ± 0.02	0.3073	0.7593
btk <sup>M1/M1</sup> 10 sh	0.43 ± 0.05		
btk <sup>M1/M1</sup> Clz 10 sh	0.45 ± 0.03	−0.352	0.726
btk <sup>M1/M1</sup> 15 sh	0.20 ± 0.04		
btk <sup>M1/M1</sup> Clz 15 sh	0.43 ± 0.03	−2.527	0.0133
Fig. 6H. ANOVA: $F_{(5,73)} = 28.6620, p < 0.0001^*$			
btk <sup>M1/M1</sup> 6 sh	0.35 ± 0.03		
btk <sup>M1/M1</sup> Ris 6 sh	0.63 ± 0.03	−6.509	<b>1.1 × 10<sup>−8</sup></b>
btk <sup>M1/M1</sup> 10 sh	0.36 ± 0.02		
btk <sup>M1/M1</sup> Ris 10 sh	0.64 ± 0.04	−6.91	<b>2.1 × 10<sup>−9</sup></b>
btk <sup>M1/M1</sup> 15 sh	0.38 ± 0.03		
btk <sup>M1/M1</sup> Ris 15 sh	0.68 ± 0.04	−6.64	<b>6.3 × 10<sup>−9</sup></b>
Fig. 6I. ANOVA: $F_{(3,52)} = 0.5727, p = 0.6356$			
0 μM 6 sh	0.76 ± 0.03		
10 μM 6 sh	0.72 ± 0.03	0.952	0.3458
0 μM 15 sh	0.78 ± 0.02		
10 μM 15 sh	0.73 ± 0.04	0.8233	0.4143

(Table continues.)

**Table 1. Continued**

Genotype	Mean ± SEM	t Ratio	p
Fig. 6I. ANOVA: $F_{(3,53)} = 65.5164, p < 0.0001^*$			
0 μM 6 sh	0.78 ± 0.01		
10 μM 6 sh	0.85 ± 0.00	−10.26	<b>7 × 10<sup>−14</sup></b>
0 μM 15 sh	0.79 ± 0.01		
10 μM 15 sh	0.84 ± 0.00	−9.555	<b>7 × 10<sup>−13</sup></b>

Avoidance of 45 V footshocks are shown for all genotypes per figure as indicated. Initial ANOVA: values are shown as well as individual statistical comparisons with the respective controls. Significant differences uncovered are highlighted in bold.

**Olfactory habituation.** Olfactory habituation experiments were performed as detailed in Semelidou et al. (2018). Avoidance of the aversive odorant 3-octanol (OCT) carried in an airstream of 500 ml/min in one arm of a standard T-maze versus air was assessed and an index (AF) was calculated. For the “training phase”, ~70 flies were exposed in the upper arm of a standard T-maze to OCT for 4 min. After a 30 s rest period, the flies were lowered to the center of the maze for testing their osmotactic response by a choice of air versus OCT. At the end of the 90 s choice period the flies in each arm were trapped counted and HF and HI were calculated as described.

**Pharmaceutical treatments**

The Btk inhibitor Ibrutinib (Selleck Chemicals) and the antipsychotic drugs clozapine (Sigma-Aldrich) and risperidone (Tocris Bioscience) were diluted in DMSO and mixed at the indicated final concentrations in Brewer’s yeast (Acros Organics) aqueous paste. The concentration ranges used bracketed analogous concentrations as used for humans. Ibrutinib was used at 0.1, 1, and 10 μM; clozapine at 5 and 10 μM; and risperidone at 0.1, 1, and 10 μM. Flies were starved for 5 h in empty vials at 25°C before exposure to drug or vehicle-only containing yeast paste for 14–16 h. The following day, flies were transferred in normal food vials, trained, and tested as detailed for footshock habituation.

**Western blots**

Five adult female heads 1–3 d post-eclosion were homogenized in 1× Laemmli buffer (50 mM Tris, pH 6.8, 100 mM DTT, 5% 2-mercaptoethanol, 2% SDS, 10% glycerol, and 0.01% bromophenol blue). The lysates were boiled for 5 min at 95°C, centrifuged at 10,000 × g for 5 min and separated by SDS-PAGE. Proteins were transferred to a PVDF membrane at 120 V for 80 min and probed with anti-dBtk antibody (Tsilaka et al., 2014) at 1:4000 and anti-syntaxin (Syn) antibody (8C3; Developmental Studies Hybridoma Bank) at 1:3000. Rat and mouse HRP-conjugated antibodies were applied at 1:5000 and proteins were visualized with chemiluminescence (ECL Plus, GE Healthcare). Signals were measured with Bio-Rad Molecular Imager Chemidoc XRS+.

**Confocal microscopy**

BtkGal4 flies were crossed to UAS-mCD8-GFP (Lee and Luo, 1999) and progeny were used to examine the expression pattern of dBtk in the adult brain. Flies were dissected in cold PBS, fixed in 4% paraformaldehyde for 15 min, and imaged by laser-scanning confocal microscopy (Leica, TCS SP8). Images were captured using a 40 ×/1.3 NA oil objective after 488 nm excitation and digital image resolution was set at 1024 × 1024. Image stacks were collected at 0.75 μm intervals to cover the entire brain. The images were converted to grayscale, inverted in Adobe Photoshop 3, and shown as maximum intensity projections derived from confocal stacks.

**Statistical analysis and experimental design**

For all experiments, controls and genetically matched experimental genotypes were tested in the same session in a balanced experimental design. The order of training and testing was randomized. When two genetic controls were used, we required an experimental result to be significantly different from both genetic controls. Untransformed (raw) data were analyzed parametrically with the JMP 7 statistical software package (SAS Institute). If significant, initial ANOVA tests were followed by planned comparisons [least square mean (LSM) contrast analyses] if they indicated significant differences among the genotypes and the level of significance was adjusted for the experiment-wise error rate as suggested by Sokal and Rohlf (1981).



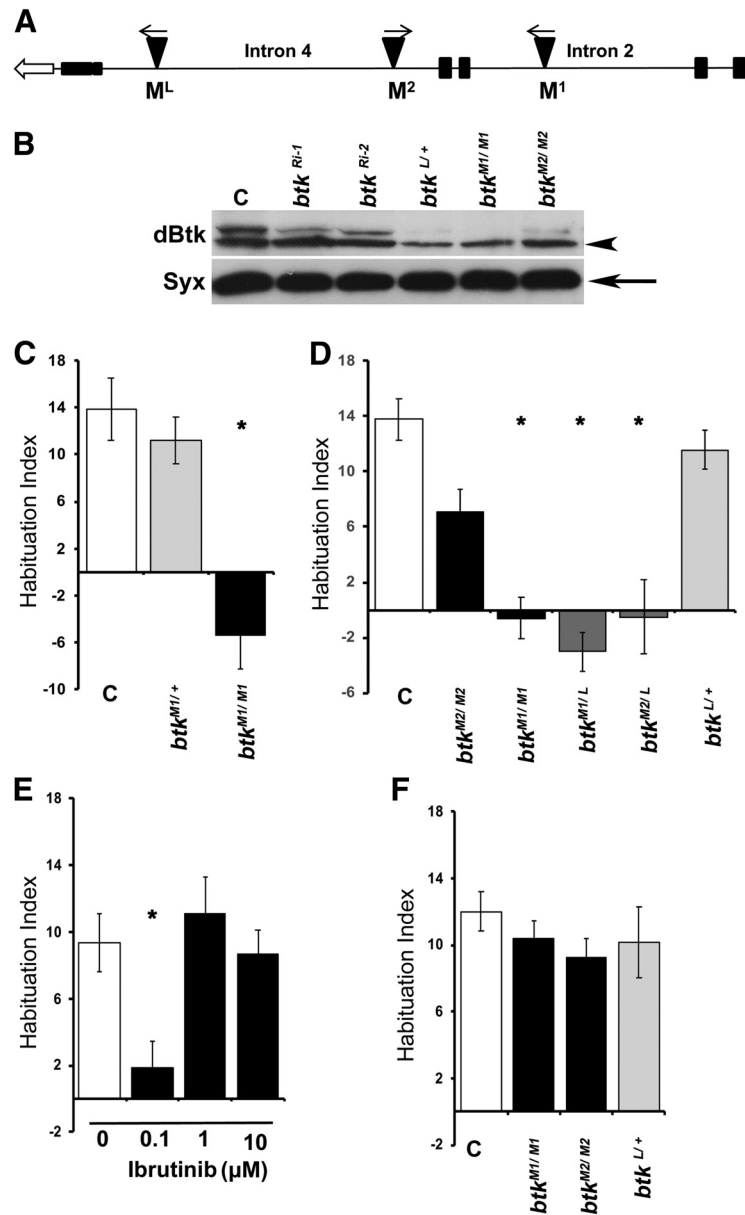
## Results

### dBtk mutants are defective in footshock habituation

In the footshock habituation paradigm (Acevedo et al., 2007), flies avoid the initial 2–8 repetitive stimuli, but their response declines rapidly to an asymptotic baseline as predicted (Rankin et al., 2009), after 10–11 stimuli. This pre-exposure-dependent attenuated avoidance after 15 footshocks relative to that of naive animals is quantified as a positive change in the HI of control flies in Figure 1C (open bar). Failure to establish habituation does not attenuate shock avoidance after shock pre-exposure and therefore yields the zero or negative difference from the naive response reported by the HI. Because the HI measures the relative change in avoidance within each genotype, it is not dependent on absolute 45 V avoidance levels, which can be variable (Table 1).

Viable MiMIC (Venken et al., 2011) insertion mutants (34172-*dBtk*<sup>M1</sup>) in intron 2 of the *dBtk* gene (Fig. 1A), resulting in reduction of both protein isoforms in adult heads (Fig. 1B), did not attenuate shock avoidance following exposure to 15 stimuli, in contrast to controls and heterozygotes, indicating failure to habituate (Fig. 1C, *btk*<sup>M1/M1</sup>; ANOVA:  $F_{(2,58)} = 16.7450$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 1.2 \times 10^{-6}$  vs control). Mutants for a different insertion (34284-*btk*<sup>M2</sup>) in intron 4 (Fig. 1A), which has a milder effect on dBtk levels (Fig. 1B), habituated to the footshocks, albeit not to the same degree as controls (Fig. 1D, *btk*<sup>M2/M2</sup>; ANOVA:  $F_{(5,68)} = 13.3566$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 0.0144$  vs control). In agreement with the behavior of *dBtk*<sup>M1/+</sup> (Fig. 1C, *dBtk*<sup>M1/+</sup>; ANOVA:  $F_{(2,58)} = 16.7450$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 0.4960$  vs control), heterozygotes for a lethal insertion (37042-*btk*<sup>L</sup>) in intron 4 (Fig. 1A) presented normal habituation (Fig. 1D, *btk*<sup>L/+</sup>; ANOVA:  $F_{(5,68)} = 13.3566$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 0.4495$  vs control). It should be noted that although the shock AF is intrinsic to calculation of the HI, AFs for all genotypes used herein are collectively presented on Table 1.

To ascertain that the habituation defect is indeed consequent of reduction in dBtk levels and not unrelated mutations on the chromosome, we examined hetero-allelic combinations of these independently isolated mutations. Significantly, hetero-allelics of the lethal *btk*<sup>L</sup> with both viable *btk*<sup>M1</sup> and *btk*<sup>M2</sup> insertions failed to habituate to 15 footshocks (Fig. 1D; ANOVA:  $F_{(5,68)} = 13.3566$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 1.2 \times 10^{-7}$  *btk*<sup>M1/L</sup>



**Figure 1.** *dBtk* is essential for footshock habituation. **A**, A schematic of the *Btk29A* (*dBtk*) genomic area where the transposon insertions used in this study reside. The open arrow demonstrates the direction of transcription. Filled boxes correspond to exons, whereas lines correspond to the indicated introns. The triangles show the MiMIC insertions. M<sup>1</sup>, The homozygous viable MI01270; M<sup>2</sup>, the homozygous viable MI02160; M<sup>L</sup>, the lethal insertion MI02966. **B**, Western blot analysis of head lysates from five 3- to 5-d-old female *btk* mutants and pan-neuronally expressing RNAis as indicated. The arrowhead points to a band which migrates as predicted for the long (type 2) dBtk protein, whereas the arrow points to the apparent short (type 1) isoform (Hamada-Kawaguchi et al., 2014). The following strains were used: C, Control (*y*<sup>1</sup>*w*<sup>\*</sup>); *btk*<sup>Ri-1</sup>, pan-neuronal expression under Elav-Gal4 of the BDSC 35159 RNAi-encoding transgene; *btk*<sup>Ri-2</sup>, pan-neuronal expression under Elav-Gal4 of the BDSC 25791 RNAi-encoding transgene; *btk*<sup>L/+</sup>, heterozygotes for the lethal (MI02966) insertion; *btk*<sup>M1/M1</sup>, homozygotes for the M<sup>1</sup> (MI01270) insertion; *btk*<sup>M2/M2</sup>, homozygotes for the M<sup>2</sup> (MI02160) insertion. The anti-dBtk antibody (Btk) recognizes two bands, the upper one of which is reduced the most in all mutants and RNAi-expressing animals, with the lower one also reduced, albeit to a lesser degree. Syntaxin 1 (*Syx*) is used as a loading control. **C–F**, Habituation indices quantify the difference in footshock avoidance following exposure to 15 stimuli from that of same genotype naive flies and are shown as the mean  $\pm$  SEM for the indicated number of repetitions (*n*). Stars indicate significant differences from controls as indicated in the text. **C**, Homozygous *btk*<sup>M1/M1</sup> mutants perform significantly different from mutant heterozygotes and *y*<sup>1</sup>*w*<sup>\*</sup> controls (**C**).  $n \geq 16$  for all groups. **D**, Complementation of the habituation failure among *dBtk* insertion mutants. Although the performance of *btk*<sup>L</sup> heterozygotes was not significantly different from that of the *y*<sup>1</sup>*w*<sup>\*</sup> controls (**C**), hetero-allelics of this lethal insertion over both viable M1 and M2 insertions presented significant habituation deficits.  $n \geq 9$  for all groups. **E**, The Btk inhibitor Ibrutinib induces habituation deficits in *y*<sup>1</sup>*w*<sup>\*</sup> control flies in a dose-specific manner. 0 represents *y*<sup>1</sup>*w*<sup>\*</sup> animals fed the DMSO vehicle and compared with their performance; 0.1  $\mu$ M Ibrutinib induced a significant deficit, but higher doses did not.  $n \geq 9$  for all groups. **F**, *dBtk* mutants do not present habituation deficits to 4 min exposure of the aversive odorant OCT.  $n \geq 12$  for all groups.

vs control and  $p = 1.1 \times 10^{-6}$  *btk*<sup>M2/L</sup> vs control), indicating that indeed reduced dBtk in the CNS does not support normal footshock habituation.

Because the human ortholog is implicated in cancers (Tillman et al., 2018) and the kinase domain is highly conserved (Gregory et al., 1987), covalent Btk inhibitors have been developed (Ysebaert and Michallet, 2014) and are available. Hence, dBtk activity was inhibited by feeding the commonly used inhibitor ibrutinib to control flies. Compared with vehicle treated animals, ibrutinib treatment resulted in strong abrogation of footshock habituation at low (Fig. 1E; 0.1  $\mu$ M, ANOVA:  $F_{(3,60)} = 5.4217$ ,  $p = 0.0024$ ; subsequent LSM:  $p = 0.0012$  vs 0  $\mu$ M), but not high concentrations (Fig. 1E; ANOVA:  $F_{(3,60)} = 5.4217$ ,  $p = 0.0024$ ; subsequent LSM:  $p = 0.5368$ , 1 vs 0  $\mu$ M, and  $p = 0.7965$  10 vs 0  $\mu$ M). The reason for this sharp concentration optimum is unclear at the moment, but at higher concentrations it may affect activities of potentially related proteins with opposing effects on habituation to footshock. Nevertheless, the results are consistent with the genetic evidence and strengthen the conclusion that dBtk kinase activity is required for habituation to repeated footshocks. Because, *dBtk* mutants were found to habituate prematurely in an olfactory jump reflex assay (Asztalos et al., 2007), we tested whether dBtk reduction resulted in deficient olfactory avoidance habituation (Semelidou et al., 2018). However, dBtk abrogation did not affect habituation to aversive odor (OCT) exposure (Semelidou et al., 2018), suggesting that the protein is specifically required for habituation to footshocks (Fig. 1F; ANOVA:  $F_{(3,62)} = 0.8339$ ,  $p = 0.4806$ ).

To independently validate these results, we used the TARGET system (McGuire et al., 2003) to abrogate the protein in adult animals by transgene-mediated RNA-interference (RNAi). In fact, adult-specific pan-neuronal (Fig. 2A; ANOVA:  $F_{(2,44)} = 6.1651$ ,  $p = 0.0045$ ; subsequent LSM:  $p = 0.0016$  and  $p = 0.018$  vs controls, respectively), dBtk attenuation recapitulated the footshock habituation failure of the mutants and also did not compromise habituation to the aversive odorant octanol (Fig. 2C; ANOVA:  $F_{(2,30)} = 1.3588$ ,  $p = 0.2734$ ). The latter supports the notion that dBtk functions specifically in footshock habituation. Identical results were obtained with the second RNAi-mediating transgene (Fig. 2B; ANOVA:  $F_{(2,35)} = 11.8202$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 0.00068$  and  $p = 0.00014$  vs controls, respectively) and a different pan-neuronal driver (Gouzi et al., 2011), Ras2Gal4 (Fig. 2D; ANOVA:  $F_{(2,27)} = 11.2630$ ,  $p = 0.0003$ ; subsequent LSM:  $p = 0.0002$  and  $p = 0.0008$  vs controls, respectively). Therefore, the failed habituation phenotype of *dBtk* mutants is not developmental in origin, but reflects an acute requirement for dBtk activity to facilitate habituation. Significantly, its abrogation in glia (Fig. 2E; ANOVA:  $F_{(2,46)} = 0.2643$ ,  $p = 0.7690$ ), or constitutively sparing the MBs from dBtk attenuation under MBGal80 (Fig. 2F; ANOVA:  $F_{(2,47)} = 0.0326$ ,  $p = 0.9679$ ; Krashes et al., 2007), did not affect habituation, suggesting that activity of the kinase is required within these neurons. Therefore, the habituation failure is not a consequence merely of the presence of the RNAi encoding transgenes, or drivers, but their induction not in glia, but within MB neurons, as suggested by Figure 2F and reported (Fig. 2G) expression of dBtk in these neurons.

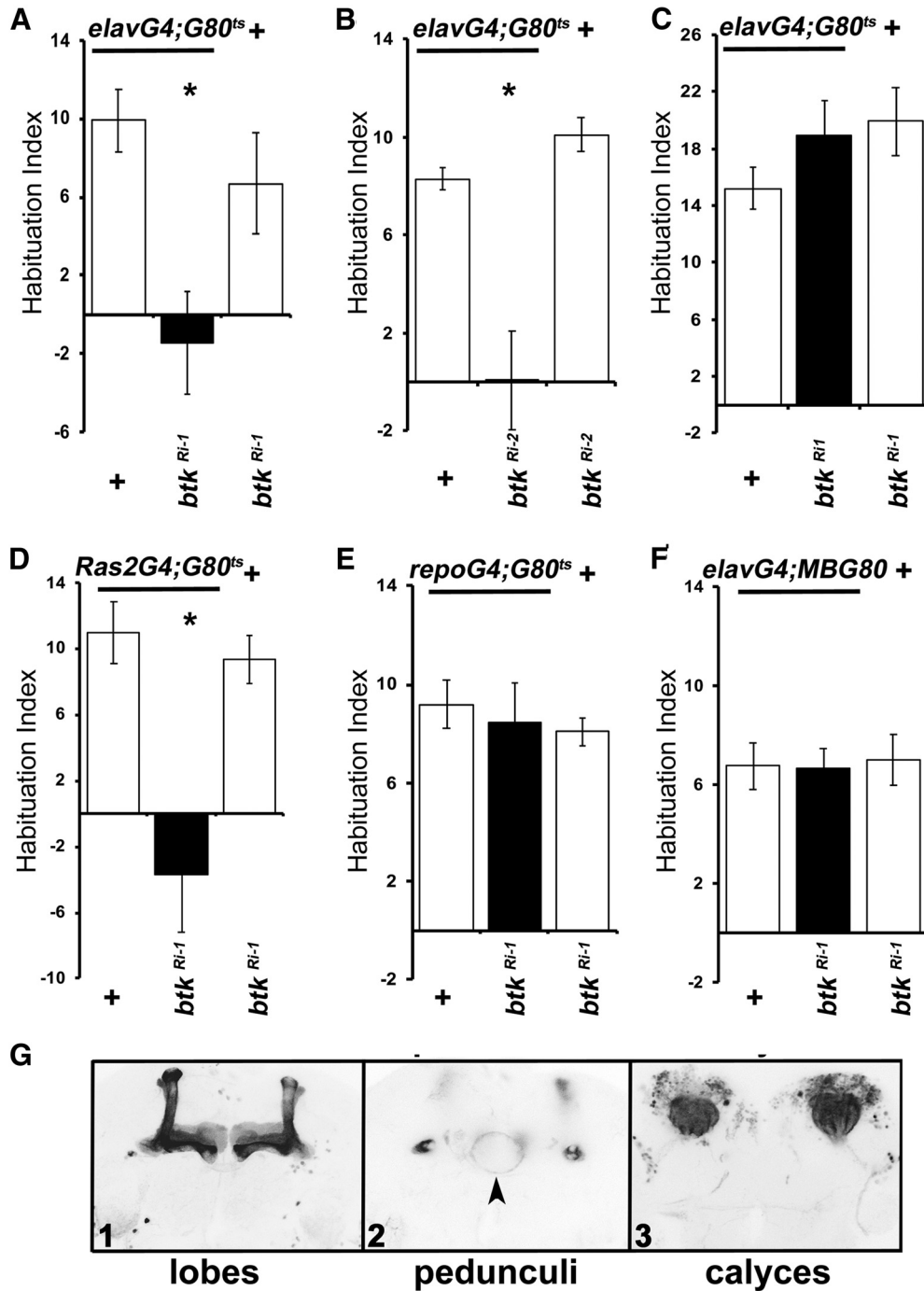
### dBtk functions within $\alpha'/\beta'$ mushroom body neurons to facilitate footshock habituation

Neurotransmission from the  $\alpha/\beta$  MB neurons is essential to prevent premature footshock habituation (Acevedo et al., 2007), but apparently not to facilitate its onset. So, we wondered where

within the adult CNS is dBtk activity required for facilitation of footshock habituation. Initially, to unequivocally establish that dBtk activity solely within the MBs is essential for facilitation of footshock habituation, we attenuated it therein with the strong pan-MB driver Leo-Gal4 (Messaritou et al., 2009). This abolished footshock habituation (Fig. 3A; ANOVA:  $F_{(2,34)} = 12.6248$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 0.0001$  and  $p = 0.0002$  vs controls, respectively) and the result was independently validated with another RNAi transgene (Fig. 3C; ANOVA:  $F_{(2,39)} = 60,9160$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 1.76 \times 10^{-12}$  and  $p = 6.55 \times 10^{-10}$  vs controls, respectively), establishing the necessity of dBtk within the MBs for the process. This conclusion was strengthened by reinstating dBtk specifically within the MBs of adult *btk*<sup>M1</sup> homozygotes, which fully reversed their deficient habituation (Fig. 3B; ANOVA:  $F_{(5,66)} = 18.3727$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 6 \times 10^{-6}$  leoMB-GAL4, *btk*<sup>M1</sup>/*btk*<sup>M1</sup>; GAL80ts/+ induced vs leoMB-GAL4, *btk*<sup>M1</sup>/+; GAL80ts/+ induced and  $p = 0.1918$  leoMB-GAL4, *btk*<sup>M1</sup>/*btk*<sup>M1</sup>; GAL80ts, UAS-*btk*<sup>short isoform</sup> induced vs leoMB-GAL4, *btk*<sup>M1</sup>/+; GAL80ts/+ induced). The transgene encodes the predominately neuronal type 1 short dBtk isoform (Tsikalas et al., 2014), containing all conserved domains except the N-terminal PH. However, if the UAS-*btk*<sup>short isoform</sup> was not induced in adult animals the mutant phenotype persisted (Fig. 3B; uninduced LSM: uninduced leoG4, *btk*<sup>M1</sup>/*btk*<sup>M1</sup>; G80<sup>ts</sup>/UAS*btk*  $p = 0.0367$  vs leoG4, *btk*<sup>M1</sup>/*btk*<sup>M1</sup>; G80<sup>ts</sup>/+ and  $p < 0.0001$  with leoG4, *btk*<sup>M1</sup>/+; G80<sup>ts</sup>/+ controls, respectively). The acute requirement for dBtk within adult MBs for footshock habituation was further supported by attenuating its levels under the pan-MB drivers 247-Gal4 (Fig. 3D; ANOVA:  $F_{(2,42)} = 11.0752$ ,  $p = 0.0001$ ; subsequent LSM:  $p = 0.0032$  and  $p = 4 \times 10^{-5}$  vs controls, respectively) and dnc-Gal4 (Fig. 3E; ANOVA:  $F_{(2,30)} = 12.4333$ ,  $p = 0.0001$ ; subsequent LSM:  $p = 0.0012$  and  $p = 0.0001$  vs controls, respectively), both of which precipitated pronounced habituation defects.

Given the role of neurotransmission from MB  $\alpha/\beta$  neurons in preventing premature habituation (Acevedo et al., 2007), we wondered whether dBtk acts within these neurons to inhibit this process and consequently facilitate habituation. Thus, dBtk was attenuated within  $\alpha/\beta$  neurons under the c772 (Fig. 3F; ANOVA:  $F_{(2,44)} = 0.6731$ ,  $p = 0.5156$ ) and c739 (Fig. 3G; ANOVA:  $F_{(2,43)} = 1.1146$ ,  $p = 0.3378$ ) drivers in  $\alpha\beta$  cortex or core neurons, respectively (Pavlopoulos et al., 2008), but habituation was not altered, eliminating this hypothesis. In addition, dBtk attenuation within  $\gamma$  neurons also did not affect footshock habituation (Fig. 3H; ANOVA:  $F_{(2,47)} = 0.0266$ ,  $p = 0.9737$ ). Significantly however, dBtk abrogation within  $\alpha'/\beta'$  MB neurons [Fig. 3I; ANOVA:  $F_{(3,44)} = 22.5186$ ,  $p < 0.0001$ ; subsequent LSM: induced (IN)  $p < 0.0001$  vs both controls] abolished footshock habituation. In contrast, if the RNAi-mediating transgene remained uninduced, habituation was indistinguishable from controls (Fig. 3I; LSM: uninduced (UN)  $p = 0.0891$  and  $p = 0.3397$  vs the controls, respectively). Therefore, dBtk is not developmentally, but rather acutely required within  $\alpha'/\beta'$  MB neurons to facilitate footshock habituation.

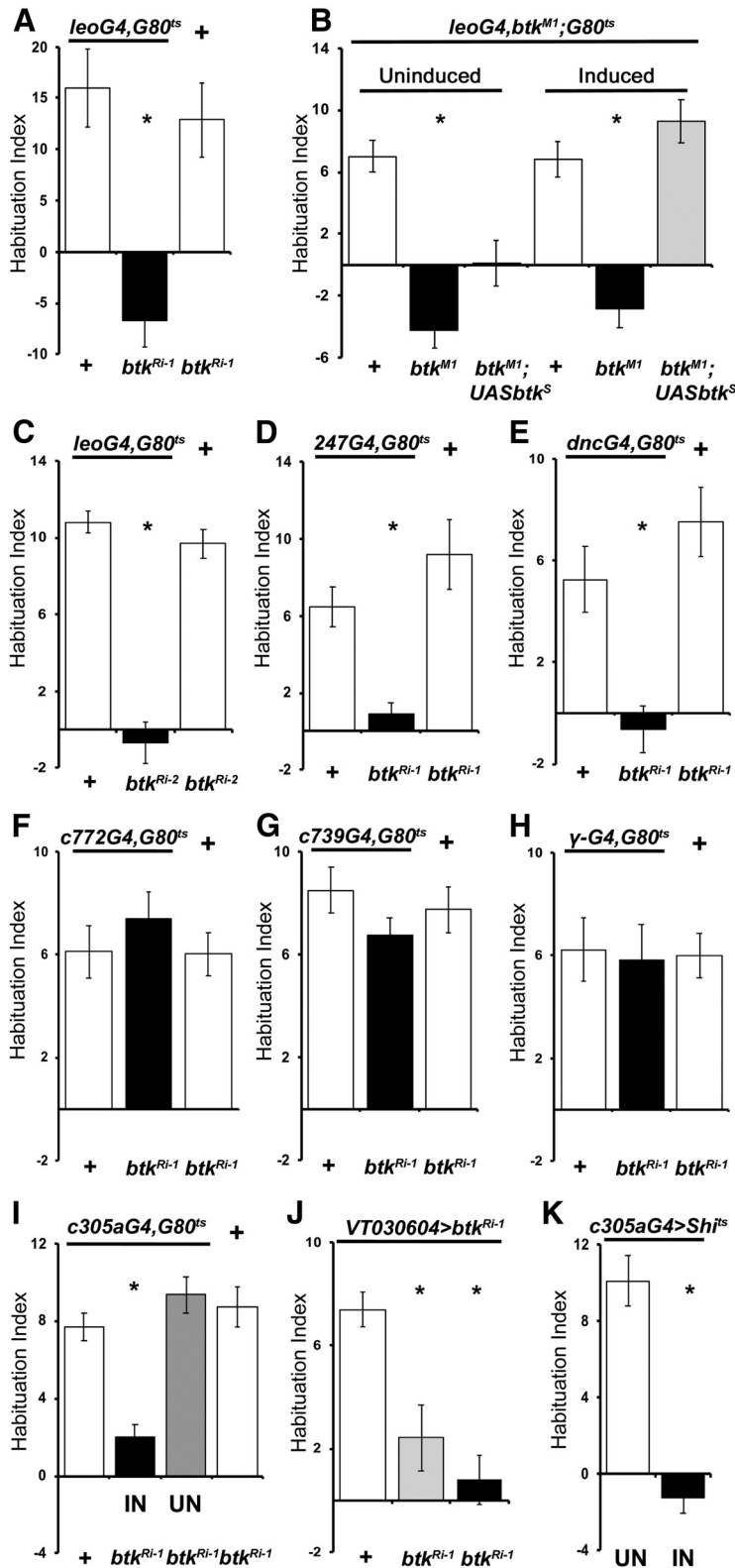
To verify this conclusion independently, we used a “split-Gal4” driver expressed specifically in  $\alpha'/\beta'$  MB neurons and varied the expression of the RNAi-mediating transgene (*btk*<sup>Ri-1</sup>) based on the optimum temperature for Gal4-mediated transcription (McGuire et al., 2004). Raising and maintaining the flies at 18°C, should not yield appreciable *btk*<sup>Ri-1</sup> expression and presented normal habituation within the range of controls (Fig. 1). In contrast, animals raised at 18°C, but kept at 30°C for 48 h before training to achieve maximal *btk*<sup>Ri-1</sup> expression failed to



**Figure 2.** Adult-specific abrogation of dBtk results in deficient footshock habituation. HIs quantifying the difference in footshock avoidance following exposure to 15 stimuli from that of same genotype naive flies are shown as the mean  $\pm$  SEM for the indicated number of repetitions ( $n$ ). All panels show the performance of animals expressing a *dbtk* RNAi-encoding transgene (*btk<sup>Ri-1</sup>/+*) under the indicated driver (black bar), the driver heterozygotes (left open bars), and the RNAi-mediating transgene heterozygotes (+, right open bars). driver/+, Progeny from the cross of the *w<sup>1118</sup>* background driver with 36303 *y<sup>1</sup>v<sup>1</sup>*, whereas for the *btk<sup>Ri-1</sup>/+*, the *y<sup>1</sup>v<sup>1</sup>* background of *btk<sup>Ri-1</sup>* was crossed to *w<sup>1118</sup>* so that the two controls have equivalent backgrounds as the experimentals. Asterisks indicate significant differences from controls as detailed in the text. **A**, Adult limited pan-neuronal expression of *btk<sup>Ri-1</sup>* eliminates habituation to 15 footshocks.  $n \geq 11$  for all groups. **B**, Adult-limited pan-neuronal expression of an independent RNAi-mediating transgene (*btk<sup>Ri-2</sup>*), also results in habituation defects.  $n \geq 9$  for all groups. **C**, Adult-limited pan-neuronal expression of *btk<sup>Ri-1</sup>* does not yield defects in habituation to 4 min of OCT exposure.  $n \geq 9$  for all groups. **D**, Flies expressing *btk<sup>Ri-1</sup>* under the pan-neuronal Ras2Gal4, Gal80<sup>ts</sup> driver present adult-specific habituation defects compared with controls.  $n \geq 9$  for all groups. **E**, Adult limited *btk<sup>Ri-1</sup>* expression in glia did not precipitate deficits.  $n \geq 14$  for all groups. **F**, Pan-neuronal expression of *btk<sup>Ri-1</sup>*, but at the exclusion of the MBs did not yield deficits in shock habituation.  $n \geq 12$  for all groups. **G**, dBtk is preferentially distributed within the MBs evidenced by mCD8-GFP expression under the control of the *btkG4* driver (BDSC, 49182). The arrowhead indicates faint expression in the ring neurons of the ellipsoid body at the level of the pedunculi.

habituate (Fig. 3J; ANOVA:  $F_{(2,55)} = 4.8324$ ,  $p = 0.0118$ ; subsequent LSM:  $p = 0.0032$  vs 18°C). Footshock habituation in animals raised and kept at 25°C was not abolished, but was highly suppressed compared with the performance of animals kept at

18°C (Fig. 3J; ANOVA:  $F_{(2,55)} = 4.8324$ ,  $p = 0.0118$ ; subsequent LSM:  $p = 0.0546$  vs 18°C). These results demonstrate that dBtk function in  $\alpha'/\beta'$  neurons is indeed essential to facilitate footshock habituation. This is not specific to dBtk, as conditionally



**Figure 3.** dBtk is acutely required within the  $\alpha'/\beta'$  mushroom body neurons to facilitate habituation to footshock. Habituation indices quantifying the difference in footshock avoidance following exposure to 15 stimuli from that of same genotype naive flies are shown as the mean  $\pm$  SEM for the indicated number of repetitions ( $n$ ). All panels show the performance of animals expressing a *dbtk* RNAi-encoding transgene (*btk<sup>RI-1</sup>/+*) under the indicated driver (black bar), the driver heterozygotes (left open bars), and the RNAi-mediating transgene heterozygotes (+, right open bars). **A**, Adult limited pan-MB expression of the independent *btk<sup>RI-1</sup>* transgene precipitates defective shock habituation compared with both controls.  $n \geq 11$  for all groups. **B**, Adult specific expression of *btk<sup>RI-1</sup>* under the independent pan-MB 247Gal4 driver results in abrogated shock habituation compared with controls.  $n \geq 8$  for all groups. **C**, Adult limited pan-MB expression of the independent *btk<sup>RI-2</sup>* transgene precipitates defective shock habituation compared with both controls.  $n \geq 10$  for all groups. **D**, Adult specific expression of *btk<sup>RI-1</sup>* under the independent pan-MB 247Gal4 driver results in abrogated shock habituation compared with controls.  $n \geq 14$  for all groups. **E**, A third MB-restricted driver (*dncGal4*) yields adult-specific shock habituation deficits when driving *UAS-btk<sup>RI-1</sup>*.  $n \geq 9$  for all groups. **F**, Adult-limited *btk<sup>RI-1</sup>* expression in  $\alpha'/\beta'$  MB neurons does not precipitate habituation deficits.  $n \geq 11$  for all groups. **G**, Adult-specific *btk<sup>RI-1</sup>* expression in  $\alpha'/\beta'$  MB neurons under the independent *c739Gal4* driver does not compromise shock habituation.  $n \geq 13$  for all groups. **H**, Adult-specific *btk<sup>RI-1</sup>* expression in  $\gamma$  MB neurons under *VT030604* also does not compromise shock habituation.  $n \geq 14$  for all groups. **I**, Adult-specific IN of *btk<sup>RI-1</sup>* expression in  $\alpha'/\beta'$  MB neurons abrogates shock habituation, whereas the uninduced transgene (UN) does not.  $n \geq 9$  for all groups. **J**, Expression of *btk<sup>RI-1</sup>* driven by the  $\alpha'/\beta'$ -specific *VT030604* driver present aberrant habituation (black bar) when induced for 2 d compared to their uninduced siblings. However, when flies were raised at 25°C (gray bar), they presented an ameliorated deficit compared with those raised at 18°C and uninduced.  $n \geq 14$  for all groups. **K**, Expression of *shi<sup>ts</sup>* in  $\alpha'/\beta'$  MB neurons compromises habituation under the restrictive conditions (black bar) compared with animals kept under permissive temperature ( $p = 9 \times 10^{-8}$ ).  $n \geq 11$  for all groups.

blocking neurotransmission from  $\alpha'/\beta'$  neurons with the thermosensitive transgenic dynamin *Shi<sup>ts</sup>* (Kitamoto, 2001), under *c305a-Gal4* completely blocked habituation (Fig. 3K; ANOVA:  $F_{(1,25)} = 56.7690$ ,  $p < 0.0001$ ; subsequent LSM:  $p = 9 \times 10^{-8}$ ). This is despite the small but significant shock avoidance reduction in the “induced” (IN) versus the “uninduced” (UN) state (Table 1), which does not contribute to the phenotype because of the normalization afforded by the calculation of the HI.

The collective results demonstrate two important points. Neurotransmission from  $\alpha'/\beta'$  MB neurons facilitates footshock habituation, whereas activation of their  $\alpha/\beta$  counterparts suppresses it (Acevedo et al., 2007).

**dBtk is also required within  $\alpha/\beta$  neurons to maintain stimulus responsiveness**

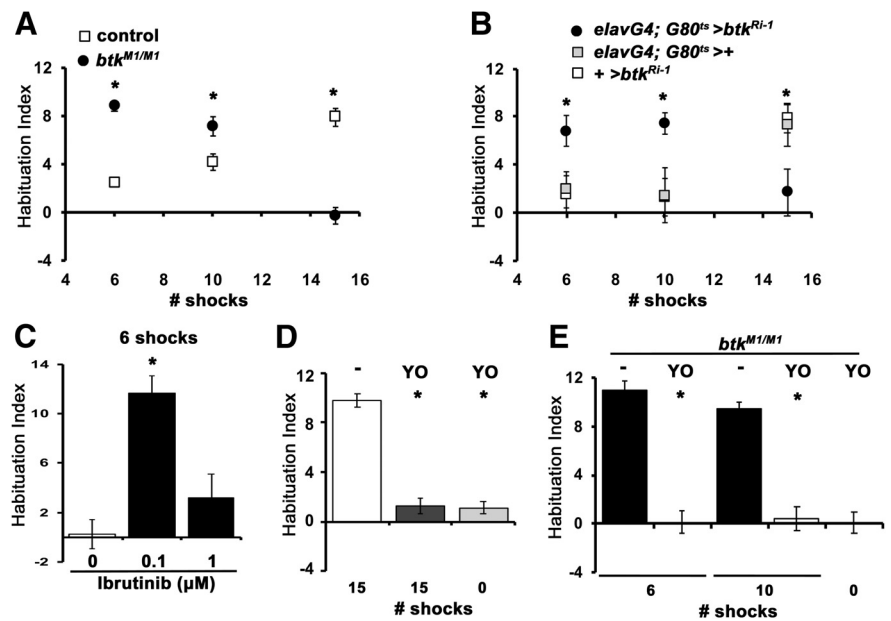
Three main reasons prompted us to investigate dBtk function in  $\alpha/\beta$  neurons. The protein appears present in these neurons (Fig. 2G), *dbtk* mutants habituate prematurely their olfactory jump reflex (Asztalos et al., 2007) and neurotransmission

transgene-bearing flies (gray bar) behave significantly different from controls, much like non-transgene bearing mutants homozygotes ( $p = 0.0014$  and  $p = 2.2 \times 10^{-6}$ , respectively). In contrast, a 2 d induction of the transgene (gray bar right side) completely reversed the deficient habituation ( $p = 0.1918$  vs  $p = 6 \times 10^{-6}$  for mutant homozygotes compare to controls; open bar).  $n \geq 8$  for all groups. **C**, Adult limited pan-MB expression of the independent *btk<sup>RI-2</sup>* transgene precipitates defective shock habituation compared with both controls.  $n \geq 10$  for all groups. **D**, Adult specific expression of *btk<sup>RI-1</sup>* under the independent pan-MB 247Gal4 driver results in abrogated shock habituation compared with controls.  $n \geq 14$  for all groups. **E**, A third MB-restricted driver (*dncGal4*) yields adult-specific shock habituation deficits when driving *UAS-btk<sup>RI-1</sup>*.  $n \geq 9$  for all groups. **F**, Adult-limited *btk<sup>RI-1</sup>* expression in  $\alpha/\beta$  MB neurons does not precipitate habituation deficits.  $n \geq 11$  for all groups. **G**, Adult-specific *btk<sup>RI-1</sup>* expression in  $\alpha/\beta$  MB neurons under the independent *c739Gal4* driver does not compromise shock habituation.  $n \geq 13$  for all groups. **H**, Adult-specific *btk<sup>RI-1</sup>* expression in  $\gamma$  MB neurons under *VT030604* also does not compromise shock habituation.  $n \geq 14$  for all groups. **I**, Adult-specific IN of *btk<sup>RI-1</sup>* expression in  $\alpha'/\beta'$  MB neurons abrogates shock habituation, whereas the uninduced transgene (UN) does not.  $n \geq 9$  for all groups. **J**, Expression of *btk<sup>RI-1</sup>* driven by the  $\alpha'/\beta'$ -specific *VT030604* driver present aberrant habituation (black bar) when induced for 2 d compared to their uninduced siblings. However, when flies were raised at 25°C (gray bar), they presented an ameliorated deficit compared with those raised at 18°C and uninduced.  $n \geq 14$  for all groups. **K**, Expression of *shi<sup>ts</sup>* in  $\alpha'/\beta'$  MB neurons compromises habituation under the restrictive conditions (black bar) compared with animals kept under permissive temperature ( $p = 9 \times 10^{-8}$ ).  $n \geq 11$  for all groups.



from  $\alpha/\beta$  neurons is essential for mediating responsiveness to footshocks and prevent premature habituation (Acevedo et al., 2007). Therefore, we wondered whether dBtk is also involved in maintenance of stimulus responsiveness within  $\alpha/\beta$  neurons. Surprisingly, in contrast to controls subjected to the same number of stimuli, *bt<sup>k</sup><sup>M1/M1</sup>* mutants habituated prematurely after 6 and 10 footshocks (Fig. 4A; ANOVA:  $F_{(5,104)} = 34.9318$ ,  $p < 0.0001$ ; subsequent LSM: 6 shocks  $p = 4 \times 10^{-10}$ , 10 shocks  $p = 0.0027$  vs controls), but failed to habituate to 15 shocks (LSM:  $p = 9 \times 10^{-16}$  vs controls). The deficits were phenocopied by acute pan-neuronal dBtk abrogation (Fig. 4B; ANOVA:  $F_{(8,156)} = 3.8010$ ,  $p = 0.0004$ ; subsequent LSM: 6 shocks  $p = 0.0088$  and  $p = 0.0193$ , 10 shocks  $p = 0.0121$  and  $p = 0.0259$ , 15 shocks  $p = 0.0057$  and  $p = 0.0125$  vs controls, respectively) and induced upon inhibition of dBtk activity with Ibrutinib in control flies with a similar dose–response as for failed habituation to 15 stimuli (Fig. 4C; ANOVA:  $F_{(2,28)} = 10.1984$ ,  $p = 0.0005$ ; subsequent LSM:  $p = 0.0002$ , 0.1 vs 0  $\mu\text{M}$  and  $p = 0.2240$ , 1 vs 0  $\mu\text{M}$ ). Therefore, as for habituation failure to 15 shocks, the premature avoidance attenuation of the mutants is also not developmental in origin. If the attenuated avoidance is indeed habituation, it should be dishabituated (Rankin et al., 2009). We used an 8 s puff of YO, which has been shown effective for habituation to aversive odors (Paranjpe et al., 2012; Semelidou et al., 2018). In fact, the premature footshock avoidance attenuation was reversed by post-training exposure to YO (Fig. 4E; ANOVA:  $F_{(4,43)} = 45.5267$ ,  $p < 0.0001$ ; subsequent LSM: 6 shocks  $p = 1.1 \times 10^{-11}$ , 10 shocks  $p = 1.5 \times 10^{-9}$ , AVD after YO  $p = 1.9 \times 10^{-9}$ ), which also suffices to dishabituate control flies after typical habituation to 15 shock stimuli (Fig. 4D; ANOVA:  $F_{(2,30)} = 84.8182$ ,  $p < 0.0001$ ; subsequent LSM: YO dishabituation,  $p = 9.28 \times 10^{-12}$ ; AVD,  $p = 6.27 \times 10^{-12}$ ). Because control flies do not habituate to six shocks (Fig. 4A,B), dishabituation after six shocks was not performed as unnecessary.

Further confirmation that both aberrant habituation phenotypes result from dBtk attenuation and not another mutation on the *bt<sup>k</sup><sup>M1</sup>* chromosome was obtained by adult-specific pan-MB abrogation of the protein under Leo-Gal4. This also yielded both premature habituation to 6 and 10 footshocks and failure to habituate to 15 stimuli and significantly both deficits were absent if the attenuating transgene was uninduced (Fig. 5A; ANOVA:  $F_{(5,119)} = 8.6993$ ,  $p < 0.0001$ ; subsequent LSM, 6 shocks,  $p = 0.0002$ ; 10 shocks,  $p = 0.0304$ ; 15 shocks,  $p = 1 \times 10^{-5}$  vs uninduced). Importantly, expression of the dBtk-encoding transgene (UAS-*bt<sup>k</sup><sup>S</sup>*) under Leo-Gal4 in *bt<sup>k</sup><sup>M1</sup>* homozygotes fully restored both, the ability to prevent premature habituation to 6 and 10 footshocks and to facilitate habituation after 15 stimuli, but not if the transgene remained uninduced (Fig. 5B;

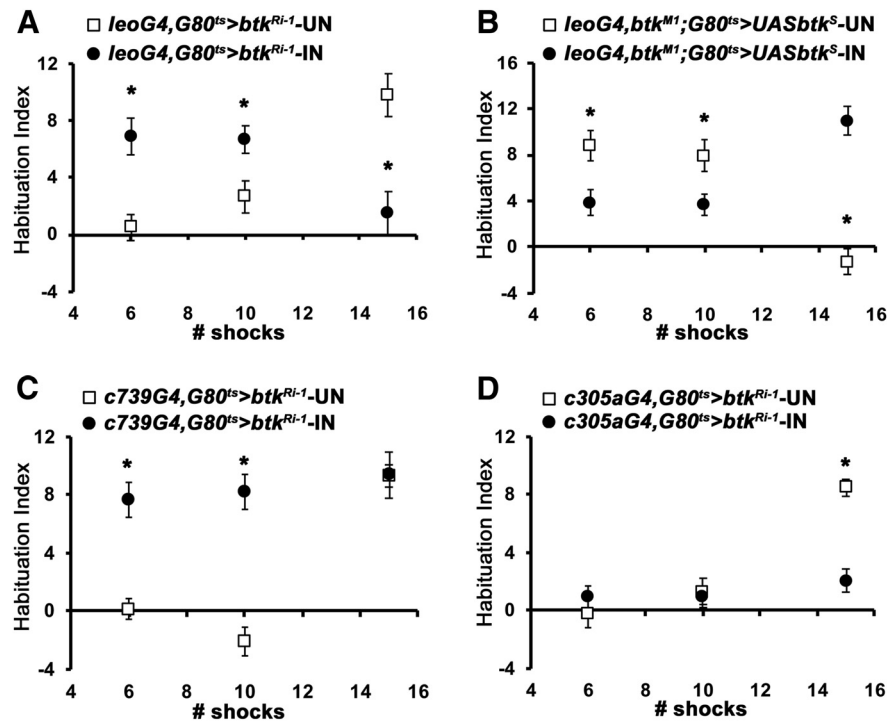


**Figure 4.** dBtk is acutely required within  $\alpha/\beta$  MB neurons to inhibit premature habituation to footshock. Habituation indices quantifying the difference in footshock avoidance following exposure to the indicated number of stimuli from that of same genotype naive flies are shown as the mean  $\pm$  SEM for the indicated number of repetitions ( $n$ ). Filled circles represent the mean performance of animals with abrogated dBtk, whole open squares the respective controls. Stars indicate significant differences from controls as detailed in the text. **A**, Footshock habituation of controls ( $y^1w^*$ ) and *bt<sup>k</sup><sup>M1/M1</sup>* mutants after prior experience of 6, 10, and 15 shocks. The performance of controls is significantly different from that of the mutants after 6, 10, and 15 footshocks.  $n > 13$  for all groups. **B**, Adult-limited pan-neuronal expression of *bt<sup>k</sup><sup>Ri-1</sup>* recapitulates both the premature habituation after 6 and 10 shocks and the habituation failure after 15 stimuli. So that the controls have similar genetic backgrounds, progeny from the cross of the  $w^{1118}$  background *ElavGal4; Gal80<sup>ts</sup>* with *36303 y<sup>1</sup>v<sup>1</sup>*, whereas for the *bt<sup>k</sup><sup>Ri-1</sup>/+*, the  $y^1v^1$  background *bt<sup>k</sup><sup>Ri-1</sup>* was crossed to  $w^{1118}$ .  $n \geq 11$  for all groups. **C**, Ibrutinib promotes premature habituation after 6 shocks at 0.1  $\mu\text{M}$ , but not at 1  $\mu\text{M}$  relative to the performance of vehicle-treated  $y^1w^*$  animals.  $n \geq 8$  for all groups. **D**, Dishabituation in  $y^1w^*$  flies after 8 s of YO puff. White bar, Habituation after 15 shocks; dark gray bar, dishabituation with YO after experiencing 15 shocks; light gray bar, the effect on avoidance of YO on naive flies before testing their shock avoidance. YO exposure has significant effects on reversing the habituated response, but YO exposure of naive flies does not affect their shock avoidance (0).  $n \geq 10$  for all groups. **E**, The reduced avoidance of *bt<sup>k</sup><sup>M1/M1</sup>* mutants after 6 and 10 footshocks is *bona fide* premature habituation because it is reversible (dishabituated) by a single puff of YO after the respective footshocks as indicated. Black bars denote habituation without YO and the open bars after odor presentation. 0 denotes shock avoidance of naive flies after YO exposure. YO exposure results in significant dishabituation after 6 and 10 shocks.  $n \geq 8$  for all groups.

ANOVA:  $F_{(5,77)} = 13.3452$ ,  $p < 0.0001$ ; subsequent LSM, 6 shocks,  $p = 0.0036$ ; 10 shocks,  $p = 0.0161$ ; 15 shocks,  $p = 6 \times 10^{-10}$  vs uninduced). This is consistent with the notion that dBtk functions to maintain habituation latency within  $\alpha/\beta$  MB neurons and to facilitate footshock habituation within their  $\alpha'/\beta'$  counterparts.

To confirm the independent function of dBtk in both preventing premature habituation in  $\alpha/\beta$  and facilitating it in  $\alpha'/\beta'$  neurons, the protein was selectively abrogated therein. Significantly, animals with abrogated dBtk in adult  $\alpha/\beta$  neurons under *c739Gal4* habituated prematurely to 6 and 10 stimuli (Fig. 5C; ANOVA:  $F_{(5,77)} = 17.1806$ ,  $p < 0.0001$ ; subsequent LSM: 6 shocks  $p = 8.9 \times 10^{-6}$ , 10 shocks  $p = 2.9 \times 10^{-8}$ , 15 shocks  $p = 0.9504$  vs control), whereas attenuation in  $\alpha'/\beta'$  neurons under *c305aGal4* did not affect habituation latency, but blocked habituation to 15 shocks (Fig. 5D; ANOVA:  $F_{(5,82)} = 12.2529$ ,  $p < 0.0001$ ; subsequent LSM: 6 shocks  $p = 0.3433$ , 10 shocks  $p = 0.777$ , 15 shocks  $p = 3.7 \times 10^{-7}$  vs control).

Therefore, dBtk has two functionally distinct and apparently independent neuronal type-specific roles. It engages mechanisms that maintain latency, thus preventing premature habituation within  $\alpha/\beta$  neurons and potentially distinct mechanisms that facilitate habituation in their  $\alpha'/\beta'$  counterparts. Alternatively,



**Figure 5.** Differential roles for dBtk in latency and habituation in  $\alpha/\beta$  and  $\alpha'/\beta'$  MB neurons. Habituation indices quantifying the difference in footshock avoidance following exposure to the indicated number of stimuli from that of same genotype naive flies are shown as the mean  $\pm$  SEM for the indicated number of repetitions ( $n$ ). Filled circles represent the mean performance of animals with abrogated dBtk; whole open squares, the respective controls. Asterisks indicate significant differences from controls as detailed in the text. **A**, Adult-limited pan-MB expression of *bt<sup>kRI-1</sup>* (IN) yields premature habituation after 6 and 10 stimuli and failed habituation after 15 shocks compared to in-genotype controls of siblings with the transgene silent (UN).  $n > 18$  for all groups. **B**, The complementary experiment of adult-limited pan-MB expression of UAS-*bt<sup>kS</sup>* rescued (IN) the premature habituation after 6 and 10 stimuli, as well as the habituation deficit after 15 footshocks compared to siblings not expressing the transgene (UN).  $n \geq 12$  for all groups. **C**, Adult-limited expression of *bt<sup>kRI-1</sup>* in  $\alpha/\beta$  MB neurons (IN) recapitulates the premature habituation after 6 and 10 shocks but not the failed habituation after 15 stimuli, compared to siblings not expressing the transgene (UN).  $n \geq 10$  for all groups. **D**, Adult-limited expression of *bt<sup>kRI-1</sup>* in  $\alpha'/\beta'$  MB neurons (IN) recapitulates the failed habituation to 15 footshocks, but not the premature habituation to 6 or 10 stimuli compared to siblings not expressing the transgene (UN).  $n \geq 11$  for all groups.

upon activation of these neurons by the footshock stimuli, dBtk may mediate neurotransmission from both types, which in the case of  $\alpha/\beta$  neurons is required to maintain the value of the stimulus and prevent premature habituation (Acevedo et al., 2007) and  $\alpha'/\beta'$  neurons to facilitate it (Fig. 3I,K). Moreover, since *bt<sup>k</sup>* mutants present habituation deficits specifically to footshock, but not odor stimuli, it is possible that the kinase is involved in second messenger pathways transducing dopaminergic signals that apparently communicate footshock information to the MBs and result in their activation (Cervantes-Sandoval et al., 2017; Horiuchi, 2019).

**Antipsychotics rescue the deficient habituation of dBtk mutants**

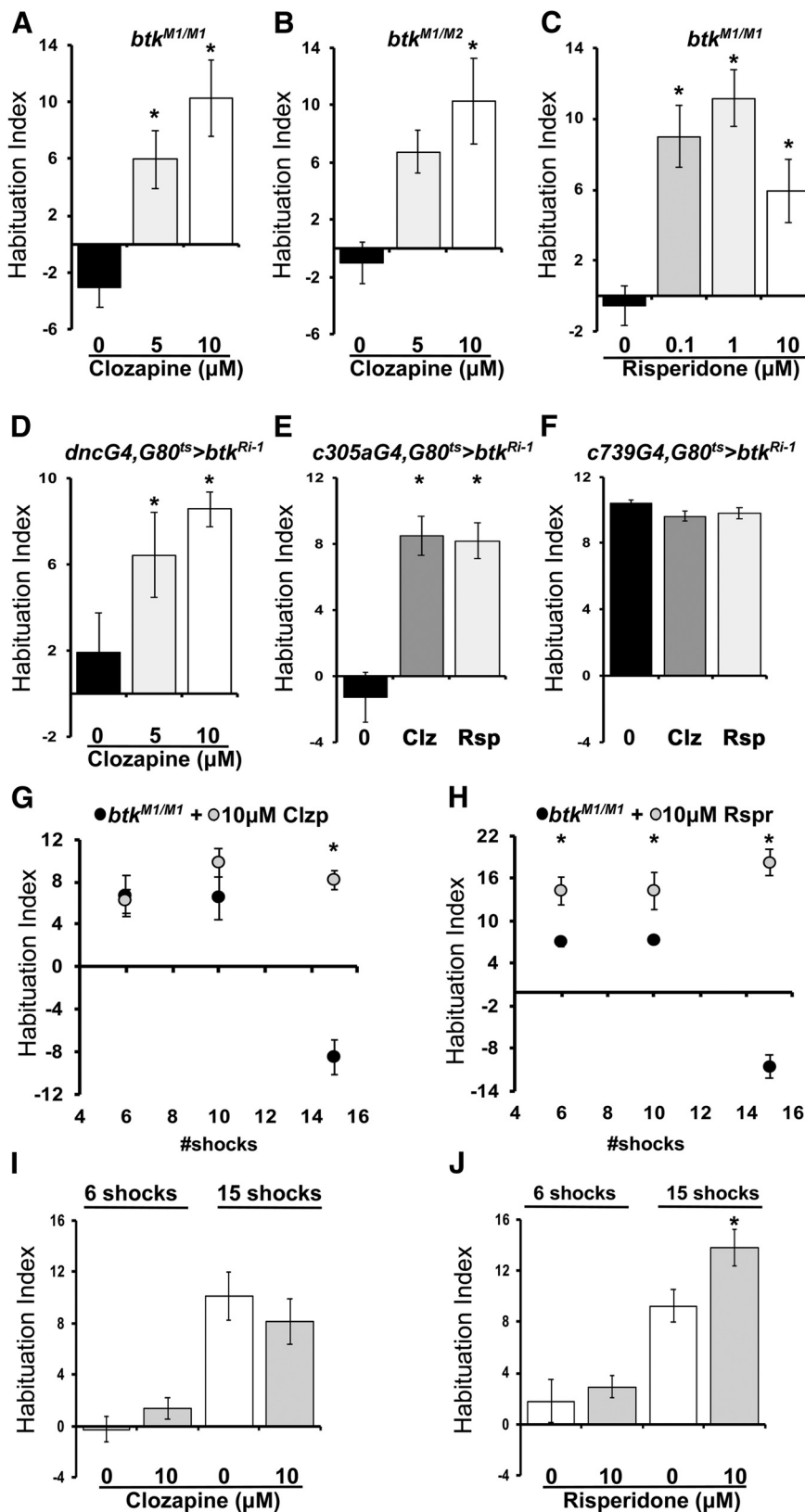
We assumed that the inability to habituate upon dBtk attenuation may result from persistent neurotransmission from  $\alpha'/\beta'$ . This could be due to excess dopaminergic signaling or inability of the mutants to downregulate the response to dopaminergic input. To differentiate between these possibilities, we initially sought to inhibit dopaminergic inputs pharmacologically. Interestingly, excess dopaminergic signaling has been linked with schizophrenia in humans (Brisch et al., 2014; Kesby et al., 2018), a condition also associated with habituation failures and defects (Gillberg, 2003; Ludewig et al., 2003; Barkus et al., 2014; McDi-

armid et al., 2017). In fact, pharmaceuticals used to treat this condition are thought to act mostly as monoamine receptor antagonists including dopaminergic ones (Robinson, 2007; Brunton et al., 2010).

Therefore, to investigate whether monoamine receptors are involved, we sought to antagonize them in the *bt<sup>k</sup>* mutants, with the typical tricyclic antipsychotic clozapine, thought to primarily address dopamine, but also serotonin receptors and the atypical benzisothiazole risperidone, which is thought to be a more pronounced serotonergic antagonist, but also efficaciously antagonizes additional monoamine receptors (Brunton et al., 2010). Interestingly, administration of clozapine for 16–18 h before habituation training reversed in a dose-dependent manner the inability of *bt<sup>kM1</sup>* homozygotes (Fig. 6A; ANOVA:  $F_{(2,28)} = 11.6054$ ,  $p = 0.0002$ ; subsequent LSM:  $5 \mu\text{M } p = 0.0041$ ,  $10 \mu\text{M } p = 0.0001$  vs  $0 \mu\text{M}$ ) and *bt<sup>kM1/M2</sup>* hetero-allelics (Fig. 6B; ANOVA:  $F_{(2,30)} = 4.7128$ ,  $p = 0.0172$ ; subsequent LSM:  $5 \mu\text{M } p = 0.0648$ ,  $10 \mu\text{M } p = 0.0048$  vs  $0 \mu\text{M}$ ) to habituate to 15 footshocks. A similar treatment with risperidone also reversed the defective habituation of *bt<sup>kM1</sup>* homozygotes, although at the high concentration of  $10 \mu\text{M}$ , the effect of the drug appeared reduced (Fig. 6C; ANOVA:  $F_{(3,44)} = 9.0647$ ,  $p = 0.0001$ ; subsequent LSM:  $0.1 \mu\text{M } p = 0.0005$ ,  $1 \mu\text{M } p = 1.2 \times 10^{-5}$ ,  $10 \mu\text{M } p = 0.0066$  vs  $0 \mu\text{M}$ ). Assuming that clozapine antagonizes dopamine receptors, dBtk loss appears to affect their levels or activity. Since phenotypic reversal

was elicited by risperidone, at least at the lower concentrations and serotonergic neurotransmission has not been reported to mediate footshock information to the MBs, it is likely that this drug also addresses dopaminergic receptors.

Importantly, clozapine treatment also reversed the defective footshock habituation of animals with pan-MB RNAi-mediated dBtk abrogation (Fig. 6D; ANOVA:  $F_{(2,26)} = 5.2147$ ,  $p = 0.0132$ ; subsequent LSM:  $5 \mu\text{M } p = 0.0531$ ,  $10 \mu\text{M } p = 0.0038$  vs  $0 \mu\text{M}$ ) suggesting that the drug reaches these neurons and mediates the observed phenotypic reversal upon acute dBtk attenuation therein. Significantly, both pharmaceuticals reversed the habituation defect to 15 shocks in animals with dBtk abrogation specifically in  $\alpha'/\beta'$  MB neurons (Fig. 6E; ANOVA:  $F_{(2,27)} = 19.1763$ ,  $p < 0.0001$ ; subsequent LSM:  $10 \mu\text{M}$  clozapine  $p < 0.0001$ ,  $10 \mu\text{M}$  risperidone  $p < 0.0001$  vs untreated-0). As expected, dBtk attenuation in  $\alpha/\beta$  neurons did not yield deficient habituation, and the drugs did not affect normal habituation to 15 footshocks (Fig. 6F; ANOVA:  $F_{(2,26)} = 1.5268$ ,  $p = 0.2376$ ). These results confirm that the habituation failure is specifically driven by dBtk attenuation within the  $\alpha'/\beta'$  neurons and likely results from consequent excess or elevated signaling from one or more monoamine receptors within these neurons. In fact, Dop1R2 (Crittenden et al., 1998) and 5HT1B have been reported to be expressed in these neurons (Ries et al., 2017). Moreover, dBtk abrogation



**Figure 6.** Antipsychotics selectively rescue the defective habituation of *btk* mutants. Habituation indices quantifying the difference in footshock avoidance following exposure to the indicated number of stimuli from that of same genotype naive flies are shown as the mean  $\pm$  SEM for the indicated number of repetitions (*n*). Asterisks indicate significant differences from controls as detailed in the text. **A**, Clozapine restores the defective habituation of *btk<sup>M1/M1</sup>* homozygotes. Compared with vehicle-treated mutants that do not habituate to 15 footshocks (0), habituation was significantly improved after treatment with 5 and 10  $\mu$ M clozapine. *n*  $\geq$  9. **B**, Clozapine restores the defective habituation of *btk<sup>M1/M2</sup>* hetero-allelics, which present a strong habituation defect if treated only with vehicle (0). Clozapine at 5 and 10  $\mu$ M restored habituation to 15 footshocks. *n*  $\geq$  8 for all groups.

in  $\alpha'/\beta'$  neurons did not affect habituation to 15 shocks, further suggesting that the drugs specifically address consequences of dBtk abrogation within their  $\alpha'/\beta'$  counterparts.

As *btk* mutants present both impaired latency and failed habituation, we wondered whether the antipsychotics can reverse both phenotypes. Interestingly, 10  $\mu$ M clozapine did not alter the premature habituation of *btk<sup>M1</sup>* homozygotes, but potentially reversed their failure to habituate (Fig. 6G; ANOVA:  $F_{(5,95)} = 7.8245$ ,  $p < 0.0001$ ; subsequent LSM: 6 shocks  $p = 0.8085$ , 10 shocks  $p = 0.1382$ , 15 shocks  $p = 7.3 \times 10^{-7}$  vs control). Risperidone also potentially suppressed the habituation failure in the mutants (Fig. 6H; ANOVA:  $F_{(5,73)} = 39.2195$ ,  $p < 0.0001$ ; subsequent LSM: 6 shocks  $p = 0.00014$ , 10 shocks  $p = 0.0017$ , 15 shocks  $p = 1 \times 10^{-19}$  vs control), but interestingly it increased significantly the shock avoidance of the mutants (Table 1), suggesting that it may actually induce hyperactivity, increased locomotion or enhance the aversion of the 45 V shock. In agreement, whereas 10  $\mu$ M clozapine did not affect avoidance (Table 1), or habituation to 6 or 15 stimuli in control flies (Fig. 6I; ANOVA:  $F_{(3,52)} = 14.7186$ ,  $p < 0.0001$ ; subsequent LSM: 6 shocks  $p = 0.3096$ , 15 shocks  $p = 0.3398$  vs control), risperidone (10  $\mu$ M) consistently elevated 45 V avoidance in control

**C**, Risperidone restores the defective habituation of *btk<sup>M1</sup>* homozygotes. The strong habituation defect to 15 shocks of vehicle-treated mutant homozygotes was reversed after treatment with 0.1, 1, and 10  $\mu$ M risperidone. *n*  $\geq$  9 for all groups. **D**, The habituation defect precipitated by adult-limited pan-MB abrogation of dBtk is barely reversible by 5  $\mu$ M, but restored with 10  $\mu$ M clozapine compared with vehicle-treated animals (0). *n*  $\geq$  8 for all groups. **E**, Clozapine (10  $\mu$ M) and risperidone (10  $\mu$ M) restore the defective habituation phenotype precipitated by adult-limited abrogation of dBtk in  $\alpha'/\beta'$  neurons. *n*  $\geq$  9 for all groups. **F**, Clozapine (10  $\mu$ M) and risperidone (10  $\mu$ M) do not affect the performance of flies where adult-limited abrogation of dBtk in  $\alpha'/\beta'$  neurons. *n*  $\geq$  9 for all groups. **I**, Clozapine (10  $\mu$ M) in control flies does not induce premature habituation after 6 shocks, or a change in habituation after 15 such stimuli relative to vehicle (0)-treated controls. *n*  $\geq$  9 for all groups. **J**, Risperidone (10  $\mu$ M) in control flies does not induce premature habituation after 6 shocks, but yields elevated habituation after 15 such stimuli relative to vehicle (0)-treated controls. *n*  $\geq$  9 for all groups. **G**, Clozapine selectively restores the defective, but not the premature habituation of *btk<sup>M1</sup>* homozygotes. Clozapine did not change the premature habituation to 6 or 10 stimuli, but reversed the habituation failure after 15 footshocks. *n*  $\geq$  10 for all groups. **H**, Risperidone restores the defective habituation, but also enhances the premature habituation of *btk<sup>M1</sup>* homozygotes. Risperidone rescued the habituation defect after 15 stimuli, but also resulted in significantly higher habituation scores after 6 and 10 footshocks. *n*  $\geq$  10 for all groups.



flies (Table 1). However, this did not affect habituation to 6 shocks in control animals, but appeared to mildly enhance habituation after 15 stimuli (Fig. 6F; ANOVA:  $F_{(3,53)} = 16.9637$ ,  $p < 0.0001$ ; subsequent LSM: 6 shocks  $p = 0.5796$ , 15 shocks  $p = 0.0205$  vs control). Collectively these results suggest distinct mechanisms of action for the two antipsychotics in *Drosophila*, in agreement with their proposed activities in vertebrates. In fact, risperidone is a potent serotonin and dopamine antagonist, but unlike clozapine presents measurable activities against adrenergic and histamine receptors (PDSB Ki database; Roth et al., 2000), which may underlie the increase in avoidance or locomotion upon its administration.

## Discussion

The response dynamics to repeated footshocks define two phases. Avoidance is initially maintained for 8–10 stimuli, followed by its rapid attenuation to an apparently asymptotic habituated response by the 15<sup>th</sup> stimulus (Acevedo et al., 2007; Figs. 4A,B, 5A,B, controls). Neurotransmission from  $\alpha/\beta$  MB neurons is required to maintain stimulus responsiveness in the initial latency phase (Acevedo et al., 2007) and significantly we now demonstrate that synaptic activity from their  $\alpha'/\beta'$  counterparts is essential to facilitate the subsequent habituation phase (Fig. 2K). Therefore, the two habituation phases are mediated by distinct MB neurons.

Because neurotransmission from different MB neurons is required both to prevent and to induce the habituated response, it is likely that it engages distinct  $\alpha/\beta$  and  $\alpha'/\beta'$  MB output neurons (MBONs; Takemura et al., 2017). We hypothesize that these potentially antagonistic signals are relayed to neurons driving the choice to avoid or ignore the footshocks in a manner akin to neurons toggling attractive and aversive odor responses in *Drosophila* (Yamazaki et al., 2018). It is unclear at the moment how the activities of the two types of MB neurons are coordinated upon repeated stimulation as indicated by the response dynamics (Acevedo et al., 2007; Fig. 3A,C–F). However, the current data predict that synaptic transmission from  $\alpha/\beta$  neurons requisite for maintenance of habituation latency precedes  $\alpha'/\beta'$  activation, which facilitates habituation onset. Nevertheless, how neurotransmission from  $\alpha/\beta$  neurons is attenuated after a relatively set number of stimuli and why it precedes activation of  $\alpha'/\beta'$  neurons is unclear at the moment. Significantly, abrogation of dBtk in adult  $\alpha/\beta$  neurons yielded strong latency attenuation (Fig. 3E), whereas loss from  $\alpha'/\beta'$  neurons did not affect latency, but eliminated habituation (Fig. 3F). Therefore, although  $\alpha/\beta$  and  $\alpha'/\beta'$  activities may be coordinated, they appear independent.

Importantly, we describe the first mutant with specific defects in footshock habituation and demonstrate that dBtk activity is acutely required in adult MBs for both phases (Figs. 1B,C, 3A), in accord with its apparent expression within  $\alpha/\beta$  and  $\alpha'/\beta'$  neurons. The premature habituation upon dBtk abrogation suggests reduced or dysregulated neurotransmission from  $\alpha/\beta$  neurons (Acevedo et al., 2007; Fig. 2K), whereas the habituation defect is consistent with attenuated neurotransmission from their  $\alpha'/\beta'$  counterparts. As Btk is involved in regulation of actin cytoskeleton dynamics (Corneth et al., 2016) and given the involvement of cortical actin in neurotransmitter release (Rust and Maritzen, 2015), the kinase may modulate directly or indirectly the responsiveness to afferent signaling and subsequent neurotransmission to MBONs from both types of MB neurons. In agreement with this notion, dBtk loss specifically from  $\alpha/\beta$  and  $\alpha'/\beta'$  neurons

phenotypically mimics *Shibire<sup>ts</sup>*-dependent silencing of their synaptic output (Figs. 3I–K, 5C,D).

Alternatively, dBtk may be involved in footshock signal reception at least by the  $\alpha'/\beta'$  neurons, which specifically respond to the dopamine and serotonin receptor antagonists clozapine and risperidone upon attenuation of the kinase. Given that footshock signals are relayed to the MBs by dopamine (Cognigni et al., 2018) and these neurons contain at least one dopamine (Crittenden et al., 1998) and serotonin (Ries et al., 2017) receptors, dBtk loss may alter the number or activity of these receptors within these neurons. Interestingly, mammalian Btk is implicated in regulation of G-protein-coupled receptor (GPCR) signaling (Corneth et al., 2016) and importantly, clozapine and risperidone address and antagonize primarily the typical serotonin and dopamine GPCRs (Naheed and Green, 2001; Brunton et al., 2010). This is in accord with the notion that in  $\alpha'/\beta'$  neurons dBtk negatively regulates dopamine and/or serotonin receptor signaling or levels. It follows that dBtk loss would increase the levels or activity of one or more of these receptors, altering MB activation threshold and reducing synaptic transmission to efferents mediating the habituated response. Clozapine and risperidone-mediated antagonism of this putative  $\alpha'/\beta'$  overactivation upon repetitive footshock may restore regulated neurotransmission mediating normal habituation. In contrast, in  $\alpha/\beta$  neurons, which do not appear to respond to dopamine or serotonin receptor antagonism, dBtk may regulate neurotransmitter release via its modulation of actin dynamics (Corneth et al., 2016). Impaired neurotransmission from  $\alpha/\beta$  neurons may also underlie the reduced latency to suppress the olfactory jump response of *btk* mutants (Asztalos et al., 2007).

Therefore, we propose that dBtk may differentially regulate neuronal activities in  $\alpha/\beta$  and  $\alpha'/\beta'$  neurons. In  $\alpha/\beta$  by positively regulating neurotransmitter release, which is impaired upon dBtk loss, leading to shortened latency and premature habituation. In contrast, in  $\alpha'/\beta'$  neurons dBtk could have primarily a postsynaptic role, by negatively regulating the number or downstream signaling of dopamine and/or serotonin receptors. Elevated intra- $\alpha'/\beta'$  monoamine receptor signaling upon dBtk loss may lead to dysregulation of downstream synaptic activity and functional silencing of neurotransmission required to facilitate footshock habituation onset. Ongoing experiments aim to test these hypotheses.

It is intriguing that schizophrenia patients also present habituation defects manifested as failures in prepulse inhibition (Braff et al., 2001; Akdag et al., 2003; Ludewig et al., 2003; Meincke et al., 2004; van Os and Kapur, 2009), where a weak prestimulus inhibits the reaction to a following strong startling stimulus. These defects are thought to reflect inability to devalue inconsequential stimuli (Braff et al., 2001; Swerdlow et al., 2008) and can be reversed with antipsychotics including clozapine and risperidone implicating excessive or dysregulated dopaminergic and/or serotonergic signaling similar to *btk* mutants. Recent genome-wide association studies (GWASs) suggested linkage of polymorphisms in a number of genes to schizophrenia and other neuropsychiatric disorders (Schizophrenia Working Group of the Psychiatric Genomics Consortium, 2014; Fromer et al., 2016; Roussos et al., 2016), but human Btk was not among them. Nevertheless, given that habituation deficits in flies and humans are reversible with antipsychotics, it is possible that mutations in *Drosophila* orthologs of genes linked to schizophrenia by GWAS may also present defective footshock habituation and provide expedient experimental validation of their effects on signaling within and between CNS neurons.



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