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Introduction

KCNQ (Kv7) channels mediate a range of important physiological functions and are a hotspot of genetic diseases and therefore target for existing and novel drug treatments. In human cardiac muscle, KCNQ1 loss of function mutations result in the most common form of cardiac arrhythmia, Long QT syndrome, while gain of functions mutations cause Short QT and atrial fibrillation [1,2]. KCNQ mutations also result in adult onset type II diabetes [3,4]. In the nervous system KCNQ3 can heteromultimerise with either KCNQ2 or KCNQ5 subunits to form a channel that mediates a M-current, a current that can be suppressed by muscarinic acetylcholine receptor activation. Because the M-current operates at resting membrane potential it is well poised to regulate membrane excitability so that when it is open it acts as a brake on neuronal excitability, with mutations causing a form of epilepsy. The M-current has been shown to be a key regulator of neuronal plasticity underlying associative memory and ethanol response in mammals. Previous work has shown that many of the molecules and plasticity mechanisms underlying changes in alcohol behaviour and addiction are shared with those of memory. We show that the single KCNQ channel in Drosophila (dKCNQ) when mutated show decrements in associative short- and long-term memory, with KCNQ function in the mushroom body z//ineurons being required for short-term memory. Ethanol disrupts memory in wildtype flies, but not in a KCNQ null mutant background suggesting KCNQ maybe a direct target of ethanol, the blockade of which interferes with the plasticity machinery required for memory formation. We show that as in humans, Drosophila display age-related memory impairment with the KCNQ mutant memory defect mimicking the effect of age on memory. Expression of KCNQ normally decreases in aging brains and KCNQ overexpression in the mushroom body neurons of KCNQ mutants restores age-related memory impairment. Therefore KCNQ is a central plasticity molecule that regulates age dependent memory impairment.

Abstract

In humans KCNQ2/3 heteromeric channels form an M-current that acts as a brake on neuronal excitability, with mutations causing a form of epilepsy. The M-current has been shown to be a key regulator of neuronal plasticity underlying associative memory and ethanol response in mammals. Previous work has shown that many of the molecules and plasticity mechanisms underlying changes in alcohol behaviour and addiction are shared with those of memory. We show that the single KCNQ channel in Drosophila (dKCNQ) when mutated show decrements in associative short- and long-term memory, with KCNQ function in the mushroom body z//ineurons being required for short-term memory. Ethanol disrupts memory in wildtype flies, but not in a KCNQ null mutant background suggesting KCNQ maybe a direct target of ethanol, the blockade of which interferes with the plasticity machinery required for memory formation. We show that as in humans, Drosophila display age-related memory impairment with the KCNQ mutant memory defect mimicking the effect of age on memory. Expression of KCNQ normally decreases in aging brains and KCNQ overexpression in the mushroom body neurons of KCNQ mutants restores age-related memory impairment. Therefore KCNQ is a central plasticity molecule that regulates age dependent memory impairment.

KCNQ Channels Regulate Age-Related Memory Impairment

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Figure 1. KCNQ signalling is required in the mushroom body \( \alpha \) and \( \beta \) neurons for short-term memory. A–C. Adult brains containing a Gal4 enhancer trap (KCNQNP3423) in the KCNQ gene locus revealed broad neuronal expression of KCNQ (labelled by membrane bound GFP in green (B)) especially in the fly memory structures of the mushroom body \( \alpha \) and \( \beta \) neurons and surrounding neurons known to be visualised by DLG-A (Ruiz-Canada et al., 2002) staining (in magenta (A), co-localised structures in white (C)). D. Initial (2 min) memory was reduced in the KCNQ mutant (black bar) and flies with reduced KCNQ levels (dark grey bars) in all neurons (Elav-Gal4, uas-KCNQ-RNAi) (\( p < 0.05 \)) compared with controls (CSw-, KCNQ control, and Gal4, +, white bars) but did not lead to any change in memory (\( p > 0.05 \)) between the remaining genotypes. E. KCNQ mutants and flies with reduced KCNQ in the mushroom body (OK107-Gal4 or MB247-Gal4, uas-KCNQ-RNAi), DPM (amn-Gal4, uas-KCNQ-RNAi) (\( p < 0.001 \)) or all (Elav-
Materials and Methods

Drosophila Stocks

The KCNQ deletion mutant contained an imprecise excision of the EP2074 element (KCNQ<sup>0</sup>) that removes all the 5′ and transmembrane regions of the channel and therefore is a null [13]. The KCNQ control was a precise excision of the element (KCNQ<sup>p</sup>) leaving the gene completely intact [13]. uas-KCNQ flies allowed Gal4 promoter driven overexpression of KCNQ [13] while uas-KCNQ-RNAi (Bloomington stock 27252) allowed Gal4 targeted knockdown of the channel. The KCNQ stocks were kind gifts of Dr Rolf Bodmer. Elav-Gal4, uas-mCD8-GFP and OK107-Gal4 [16] were gifts from Dr Leslie Griffith. OK107-Gal4, Gal80<sup>0</sup> [17] was a gift from Dr Yi Zhong. c305a-Gal4 [18], MB247-Gal4 [19] and Amn(c316)-Gal4 [20] stocks were gifts of Dr Scott Waddell. Wildtype flies were Canton S w- (CSw-) from a stock previously maintained in the Waddell lab. All KCNQ mutant, Gal4 and uas lines were out crossed with the relevant CSw- line prior to behavioural analysis. All genotypes and all other crosses were out crossed with the relevant genotypes to ensure that the strains were genetically fit. CSw- lines were out crossed with the relevant genotypes to ensure that the strains were genetically fit. CSw-0.05) memory at 30°C with these promoters.

Immunohistochemistry

Adult fly brains were dissected in HL3.1 (70 mM NaCl, 5 mM KCl, 10 mM NaHCO<sub>3</sub>, 115 mM sucrose, 4 mM MgCl<sub>2</sub>, 5 mM trehalose, 1.5 mM CaCl<sub>2</sub>, and 5 mM HEPES, pH 7.5) and isolated brains were fixed in 4% paraformaldehyde in HL3.1 for 30 min before being washed in HL3.1 [21]. The samples were permeabilised in HL3.1 with 0.1% triton X (HL3.1-Tx) for 1 hr, and then blocked for 1 hr in HL3.1-Tx with 0.1% BSA and 2% normal donkey serum (HL3.1-Tx-BSA-NDs). In order to visualise the mushroom body and antennal lobe, brains were incubated with (1:2000) rabbit anti-Drosophila DLG (PDZ1-2) a protein known to be highly expressed in these memory-related structures [22] overnight at 4°C in HL3.1-Tx-BSA-NDs. After washing three times in HL3.1-Tx for 20 min, the brains were incubated with anti-rabbit Alexa-648 conjugated secondary antibody (1:400 in HL3.1-Tx-BSA-NDs) for 2 hr at room temperature. Finally the brains were washed three times HL3.1-Tx before being mounted in Vectorshield (Vector Laboratories). Samples were stored at 4°C in the dark until examination using a Leica TCS SP5 confocal microscope. The endogenous KCNQ expression pattern was determined by visualising membrane targeted GFP expressed using KCNQ-Gal4 reporter lines (KCNQ<sup>NPS4235</sup>-Gal4, uas-mCD8-GFP).

Olfactory Aversive Conditioning

All experiments were performed at 25°C and 70% humidity under red light using the olfactory aversive conditioning protocol [23]. Groups of ~100 1–4 day old male and female flies received either 1 cycle of training during which they were exposed sequentially to one odour (conditioned stimulus, CS+; 3-octanol (1:74) or 4-methylcyclohexanol (1:57) diluted in mineral oil) for 1 min paired with electric 60 V DC shock (US) and then to a second odour (CS-; the reciprocal odour) for 1 min without electric shock separated by a 30 sec rest period when they were exposed to fresh air. Memory was measured after 1 (~2 min memory) training session at the choice point of the T-maze. To measure STM, flies were trained with 1 training cycle were stored for 1 hr and then allowed to distribute in the T-maze. LTM was assessed by giving flies either 5 cycles of spaced training cycles separated by 15 min rest intervals and then storing the flies for 24 hr before distribution in the T-maze. A performance index (PI) was calculated as the number of flies that distributed in the CS- arm minus the flies in the CS+ arm, divided by the total number of flies. Therefore a PI of 1.0 would be equivalent of 100% distribution where all the flies avoided the CS+ (perfect memory), while a 50:50 distribution would give a PI of zero (no memory). To test the effect of ethanol on Drosophila learning, flies were kept in bottles containing instant media (Formula 4–24 (R); Carolina biological supply company, Burlington, NC, USA) containing 10% ethanol in water containing a small amount of blue dye (0.05% Bromophenol blue) ~12 hr before testing. The controls were given the same water-blue dye solution but lacking ethanol. The blue dye was used to monitor whether the flies had actually drunk the ethanol solution; this was confirmed as all the flies had blue abdomens prior to the test. For OK107-Gal4, Gal80<sup>0</sup> experiments [17] flies were raised at 18°C and then shifted to 30°C allowing KCNQ transgene expression 1–2 days prior and during behavioural testing. Olfactory acuity was quantified by exposing naïve flies to the odour versus air in the T-maze during a 2 min test trial. The performance index was calculated by counting the number of flies avoiding odour divided by total number of flies.
number flies. Shock reactivity was quantified by placing grids in each arm of the T-maze, and applying shock via the grid in one arm of the maze during a 2 min test trial. The performance index was calculated by counting the number of flies avoiding shock divided by total number flies \[24\]. Ethanol avoidance was quantified by placing a solution of 40% ethanol in the odour cup of one arm of the T-maze during a 2 min test trial. The performance index was calculated by counting the number of flies avoiding 40% ethanol divided by total number flies. All statistical analysis for behavioural data were performed and plotted with Graphpad Prism software.

Figure 3. KCNQ mediates age-related memory impairment. A. Quantitative RT-PCR data show a dramatic age dependent reduction (p<0.05) in KCNQ expression in adult brains (20 flies per n). B. 1 hr memory after 1 cycle training was compared between young (1–5 days old, white bars) and aged (25–30 days, black bars) adults. 2-way ANOVA indicates significant differences in memory due to age (p = 0.0013) and genotype (p = 0.0008). Post-hoc analysis revealed that memory becomes significantly impaired in aged as opposed to young CSw- wildtype (p<0.01) and KCNQ control (p<0.05) flies. KCNQ mutant flies had equally low (p>0.05) memory whether young or old. C. Overexpression of KCNQ in the mushroom body rescues memory impairment of young and old KCNQ mutant flies. 2-way ANOVA indicates significant differences in memory due to age (p<0.01) and genotype (p<0.001). Post-hoc analysis revealed that memory becomes significantly impaired in aged as opposed to young CSw- wildtype (p<0.01), while the memory of KCNQ mutant; OK107-Gal4, uas-KCNQ rescue flies stays similarly high (p>0.05) in young and old flies as opposed to KCNQ mutant with Gal4 or uas alone (KCNQ mutant; OK107-Gal4orKCNQ mutant; uas-KCNQ) whose memory was similarly low in young and old flies (p>0.05). doi:10.1371/journal.pone.0062445.g003

Quantitative RT-PCR
Age-matched flies were frozen in liquid nitrogen and decapitated by vortexing. Heads were collected and an equal number of heads from each genotype were homogenised. Trizol was added directly in to the homogenised heads and RNA was extracted according to manufacturer’s instructions (Invitrogen). RNA was DNAase treated (Ambion Inc) and reverse-transcribed (Retroscript, Ambion). KCNQ mRNA was measured using a TaqMan Kit[KCNQ - Dm01846741_g1] and was normalised to Rpl23 (Rpl23-Dm02151827_g1) mRNA as a control allowing standardisation between the samples for aging experiments. The cDNA
concentration was measured using Roche’s Light Cycler system and using multiplexing on a Stratagene Mx3000P system (Stratagene). All statistical analysis of data were performed and plotted with Graphpad Prism software.

**Results**

**KCNQ Signalling Regulates Mushroom Body Dependent Associative Memory**

Adult flies were tested for associative memory using the olfactory aversive conditioning [23]. Anatomically, neurons critical for *Drosophila* memory include those of the mushroom body that are labelled by OK107-Gal4 [16]; these can be subdivided into α, β and γ neurons labelled by MB247-Gal4 (which expresses strongly in α/β neurons but only weakly in γ neurons, [19]) and α'/β’ neurons labelled by c305a-Gal4 [18]. The mushroom body-associated dorsal paired medial (DPM) neurons are also important for memory. They express the memory gene *annexia* (ann) and can be labelled by *ann(316)-Gal4* that expresses in the DPM [20]. KCNQ is broadly expressed in the brain [12,15]. Using a *Gal4* promoter enhancer trap within the KCNQ gene to drive GFP expression (Figure 1B), it appears KCNQ is expressed in the adult mushroom body α/β and surrounding neurons (Figure 1C), structures that are known to mediate memory formation and ethanol behaviour [24-26]. Compared with controls (Figure 1D), the *KCNQ* mutant had reduced initial memory 2 min after training, as did flies with pan-neural *KCNQ* knockdown. *KCNQ* mutants and flies with pan-neural, DPM or mushroom body neuron deficient *KCNQ* completely lacked the ability to form short-term memory (STM) assessed 1 hr after training (Figure 1E).

In order to map this STM phenotype further, we selectively knocked-down *KCNQ* in different parts of the mushroom body, finding that the α/β neurons that appear to express KCNQ (Figure 1C) were required for KCNQ’s role in STM (as opposed to α'/β’ neurons (Figure 1E) which do not seem to express KCNQ (Figure 1C)). *KCNQ* overexpression in any part of the mushroom body, DPM or all neurons did not change memory measured at 2 min or 1 hr (Figure 1D–E). We also found that expression of *KCNQ* in α/β neurons using MB247-Gal4 in a fly otherwise completely lacking *KCNQ* rescued the *KCNQ* mutant STM defect to normal (Figure 1F). Acute reduction of *KCNQ* levels in the mushroom body was sufficient to decrease 1 hr memory compared with controls (Figure 1G), showing that KCNQ is required post developmentally to mediate physiological changes underlying memory. We then wished to determine the role of KCNQ in long-term memory (LTM) which is formed after spaced training and lasts about 7 days and is protein synthesis and CREB dependent [17,27]. We found that the *KCNQ* mutant showed a drastic reduction in LTM (Figure 2A). No difference in ability of the flies to sense the odour (Figure S1A–B) or shock (Figure S1C) was observed between genotypes, showing that *KCNQ* mutants do not change peripheral sensory processing, but rather the memory defects are due to loss of KCNQ function in the mushroom body.

**Drosophila Display Age Dependent Memory Deficits that are Rescued by Mushroom Body KCNQ Expression**

Fly associative memory is known to decrease with age [28] and *KCNQ* expression decreases in old fly hearts leading to age related cardiac impairments [13]. Therefore we decided to test whether or not KCNQ was involved in age dependent memory decline. We first determined *KCNQ* expression in heads over the lifespan of *Drosophila* (~50 days, [28]) using quantitative RT-PCR. We found that by 25 days *KCNQ* expression had declined to about 10% of the level of 5 day olds (Figure 3A). Therefore, we tested STM of young (1–5 day old) as opposed to aged (25–30 day old) flies. Whereas control flies displayed an age dependent decrease in 1 hr STM (Figure 3B), *KCNQ* nulls were completely unable to form STM whether they were young or old. Previous experiments have implicated the ann DPM and mushroom body neurons in mediating the effect of age on 1 hr memory [28–30], with expression of a *PKA* transgene in mushroom body neurons restoring age-related memory impairment. We therefore over-expressed *KCNQ* in the mushroom body (Figure 3C) of *KCNQ* mutants and demonstrate rescue of age-dependent memory impairment. These experiments are consistent with decreases in KCNQ signalling being central forage dependent decrements in memory. This experiment also confirms that in young flies expression of *KCNQ* using mushroom body *Gal4* lines (Figure 1F and 3C) in a fly otherwise completely lacking *KCNQ* rescues the *KCNQ* mutant STM defect to normal.

![Figure 4. Ethanol disrupts memory in wildtype flies an effect removed by the KCNQ mutation. A. Flies received an overnight (~12 hr) exposure to 10% ethanol and were tested for 2 min memory. 2-way ANOVA showed a significant effect due to genotype and ethanol (p<0.05). Post-hoc analysis showed that in KCNQ control ethanol caused a reduction (p<0.05) in memory compared to water. The reduction in memory was removed by the KCNQ mutation that had similarly low memory with or without ethanol (p>0.05). B. Ethanol content of KCNQ mutant and control (CSw- wildtype) flies exposed to 10% ethanol solution for ~12 hr was similar (p>0.05, unpaired t-test, 20 flies per n). doi:10.1371/journal.pone.0062445.g004](image-url)
Ethanol Disrupts Memory an Effect Mimicked by KCNQ Mutation

Previous work has shown that many of the molecules and plasticity mechanisms underlying changes in ethanol behaviour and addiction are shared with those of associative memory with ethanol known to disrupt synaptic plasticity and memory in humans [29,30]. Furthermore ethanol has been demonstrated to directly inhibit the M-current in dopaminergic neurons of the ventral tegmental area (VTA) a region of the brain important for ethanol reinforcement [26,31]. Likewise it has been shown that dKCNQ shows conserved blockade by ethanol, with reduction in KCNQ function causing increased ethanol sensitivity and tolerance via changes in dopamine neurons [15]. Consequently we investigated whether or not ethanol disrupted fly memory. Wildtype flies were exposed to 10% ethanol solution for ~12 hr and then immediately tested for 2 min memory (Figure 4A), although the flies did not appear sedated or intoxicated after the exposure or during the memory test, ethanol was found to reduce their memory. As KCNQ may be a direct target of ethanol that is required for memory, we tested the KCNQ mutant and found that this resulted in a loss of the reduction in memory. This suggests that KCNQ is the plasticity molecule blocked by ethanol interfering with memory. No change in ethanol content was found between genotypes and was ~10 mM at the time of the memory test (Figure 4B), this would be sufficient to block a significant proportion of neural KCNQ channels (IC50 = 19.8 mM, [15]).

Discussion

Drosophila KCNQ displays conserved electrophysiological and pharmacological properties with mammalian neuronal KCNQ2/3 channels, with both mediating a slowly activating and non-inactivating Kv current called a M-current due to its suppression by muscarinic acetylcholine receptors [12,14]. In the hippocampus, expression of a dominant negative human KCNQ2 transgene was found to suppress the M-current [7]. This resulted in a decrease in the afterhyperpolarization (after a train of action potentials, the increase in Ca2+ activates K+ channels leading to a pronounced hyperpolarization) and caused deficits in associative memory. Kv channels have also been implicated in plasticity underlying fly memory [32–35] consistent with the conspicuous memory. Kv channels have also been implicated in plasticity and memory formation, thereby leading to alcohol induced amnesia or blackout. Consistent with this proposition we found that ethanol disrupts fly memory, an effect that was removed in the KCNQ mutant background (Figure 4A). This suggests that KCNQ is a key molecule that ethanol interacts with in the plasticity machinery involved in memory. Given the conserved role of mammalian KCNQ in ethanol response and memory [15], we suggest that it is likely this will also be the case for mammals.

Fly memory is known to reduce with age [28], as does KCNQ expression and function in the heart [13]. We found that KCNQ brain expression dramatically decreases with age and this is accompanied by an age-dependent decrease in memory (Figure 3A). Young KCNQ mutant flies that have low levels of KCNQ comparable to the low levels of KCNQ in aged wildtype flies have comparably reduced levels of memory. As mushroom body neurons are known to be important for mediating changes in memory performance with age [28,29] and mushroom body knockdown of KCNQ completely removed 1 hr memory (Figure 1E), we overexpressed a KCNQ transgene in the mushroom body (Figure 3C) of a fly otherwise completely lacking KCNQ and found that this restored age-related memory impairment with young and old flies having similarly high memory. In summary we show reduction in KCNQ function in mushroom body neurons mediates age-dependent cognitive decline.

In mammals, KCNQ specific modulators have been suggested to alleviate memory deficits associated with age related memory diseases such as Alzheimer’s disease [2]. It is not clear how KCNQ-mediated mechanisms may affect memory in aged animals. However, based on the contribution of KCNQ2/3 to hippocampal afterhyperpolarizations and memory [7], one candidate mechanism would involve reduced neuronal KCNQ as this modulates afterhyperpolarization duration that is known to change with memory and in aged animals [47–49]. Recently, KCNQ channels have been implicated in age-dependent decrements in the memory of primates [50], suggesting that KCNQ function in cognitive impairments accompanying aging are likely conserved from flies to humans. Furthermore, PKA signalling has been implicated in age-related memory impairment in flies and mammals [20,29,30,51] with the mammalian KCNQ channel open state being increased by PKA [30,32], suggesting that the mushroom body neuron KCNQ-mediated memory and age-dependent memory defects maybe due to an interaction with PKA.

The genetic and experimental tractability of Drosophila combined with its ~50 day lifespan and molecular conservation with human make it a convenient and powerful genetic model [26] to study further the age-dependent KCNQ cognitive deficits. We have shown that KCNQ neuronal function in memory and ethanol response are evolutionarily conserved with mammals, allowing further development of Drosophila models of KCNQ neuronal function and channelopathies to elucidate KCNQ...
signalling networks, mechanisms of aging and potential screening for new disease therapies.

Supporting Information

Figure S1 A. KCNQ channel mutants display normal olfactory acuity and shock reactivity. Experimental and control (CSw+ or CSw−) flies similarly (p>0.05) avoided OCT A and MCH B. Odour used in the memory assay. C. Experimental and control (CSw+ or CSw−) control flies similarly (p>0.05) avoided the arm of the T-maze delivering 60 V DC electric shock. Data in A-C were analysed by 1-way ANOVA with Bonferroni post-hoc test. (TIFF)

References


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Author Contributions

Conceived and designed the experiments: JMLH. Performed the experiments: SC BM JMLH. Analyzed the data: SC BM JMLH. Wrote the paper: SC BM JMLH.