

1 **Immune stimulation reduces sleep and memory ability**

2 **in *Drosophila melanogaster***

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26 **Abstract**

27 Psychoneuroimmunology studies the increasing number of connections between
28 neurobiology, immunology and behaviour. We establish *Drosophila melanogaster* as a
29 tractable model in this field by demonstrating the effects of the immune response on
30 two fundamental behaviours: sleep and memory ability.

31 We used the Geneswitch system to upregulate peptidoglycan receptor protein (PGRP)
32 expression, thereby stimulating the immune system in the absence of infection.
33 Geneswitch was activated by feeding the steroid RU486, to the flies. We used an
34 aversive classical conditioning paradigm to quantify memory and measures of activity
35 to infer sleep.

36 Immune stimulated flies exhibited reduced levels of sleep, which could not be
37 explained by a generalised increase in waking activity. The effects on sleep were more
38 pronounced for day compared to night sleep. Immune stimulated flies also showed a
39 reduction in memory abilities.

40 These are important results as they establish *Drosophila* as a model for immune-neural
41 interactions and provide a possible role for sleep in the interplay between the immune
42 response and memory.

43 **Keywords**

44 immune-neural interactions, imd, geneswitch, PGRP-LCa

45 **Introduction**

46 Psychoneuroimmunology, in vertebrates, studies the connections between
47 neurobiology, immunology and behaviour (Ader *et al.* 1991). These neural-immune
48 interactions have also been found in invertebrates (Demas *et al.* 2011). For example,
49 immune response negatively affects learning and memory in bees (Mallon *et al.* 2003;
50 Riddell & Mallon 2006; Gegear *et al.* 2006; Iqbal & Mueller 2007; Alghamdi *et al.*
51 2008). A tractable invertebrate model of these immune-neural links would provide a
52 stimulus to this field (Aubert 2007). The fruit fly, *Drosophila melanogaster*, has been
53 tremendously helpful to the analysis of associative learning (Kim *et al.* 2007) and
54 immunity (Lemaitre & Hoffmann 2007). In this paper we demonstrate immune-memory
55 links in *Drosophila* and further expand the paradigm by showing immune-sleep
56 interactions in flies.

57 Sleep is a resting state where the sleeper exhibits inattention to the environment and is
58 usually immobile (Siegel 2003). *Drosophila melanogaster* like vertebrates have been
59 shown to have a distinct sleep state. In flies, a sleep episode is defined as a period of
60 immobility lasting five minutes or longer (Hendricks *et al.* 2000; Shaw *et al.* 2000).
61 Such intervals are associated with reversible increases in arousal threshold, which can
62 be further augmented following sleep deprivation (Huber *et al.* 2004), are associated
63 with changes in brain electrical activity (Nitz *et al.* 2002; Alphen *et al.* 2013), and are
64 reduced by several drugs like caffeine and modafinil and are increased by
65 antihistamines (Hendricks *et al.* 2000; Shaw *et al.* 2000). As in mammals, sleep
66 deprivation leads to a rebound in quantity of sleep (Shaw *et al.* 2000).

67 Infections increase sleep in humans, most likely through induction of proinflammatory
68 cytokines (Bryant *et al.* 2004). Fruit flies infected with gram-negative bacteria also

69 show increased sleep (Kuo *et al.* 2010). On the contrary, Shirasu-Hiza infected flies
70 with gram-positive bacteria and observed that they slept less (Shirasu-Hiza *et al.* 2007).
71 The latter agrees with findings of increased immune gene transcription and resistance to
72 disease in sleep-deprived flies or in reduced sleep phenotype transgenic flies (Cirelli *et*
73 *al.* 2005; Williams *et al.* 2007).

74 Here, we activated the immune system non-pathogenically (Moret & Schmid-Hempel
75 2000; Mallon *et al.* 2003; Riddell & Mallon 2006; Alghamdi *et al.* 2008; Richard *et al.*
76 2008). This has the advantage that it separates the effect of the immune response from
77 any direct effect of the pathogen, for example, parasite manipulation of the host
78 (Adamo & Webster 2013). We used Geneswitch (Osterwalder *et al.* 2001) to up-
79 regulate peptidoglycan receptor protein LCa (PGRP-Lca) in adult flies. PGRP-Lca is a
80 pattern recognition protein that recognizes gram-negative bacteria, setting off the IMD
81 immune pathway and leading to the expression of antimicrobial peptides (Gottar *et al.*
82 2002). Geneswitch is activated in the presence of the steroid RU486. We used an
83 aversive classical conditioning paradigm to measure memory abilities of flies (Mery &
84 Kawecki 2005). Sleep was measured using the *Drosophila* Activity Monitoring System
85 2 (DAMS2, Trikinetics, Waltham, MA).

86

87 **Methods and Materials**

88 The Geneswitch line $w^{1118}; P\{w^{+mW.hs=Switch1}\}bun^{Switch1.32}$ (hereafter referred to as
89 *GS1.32*) drives expression of RU486-activated GAL4 in adult fat bodies (Gottar *et al.*
90 2002) (<http://flystocks.bio.indiana.edu>). The three genotypes used were *GS1.32>PGRP-*
91 *Lca*($w^{1118};GS1.32/+; UAS-PGRP-Lca/+$), and the control genotypes *GS1.32/+(w^{1118};*
92 *GS1.32/+; +/+)* and *+/PGRP-Lca (+/+;UAS-PGRP-Lca/+)*.

93 Flies were maintained in vials containing agar, sugar, and Brewer's yeast media in a 12
94 h: 12 h light: dark cycle at 25°C. Males and females were selected at eclosion and flies
95 were 1–3 days old at the beginning of the experiment. Both sexes were used for the
96 memory assay. As is common in fly research, only males were used for the sleep assay
97 as they sleep for over twice as long as females (Isaac *et al.* 2010).

98

99 Geneswitch

100 In the Geneswitch system, the DNA binding domain of the GAL4 protein is fused to the
101 activation moiety of p65 through a mutant progesterone receptor ligand binding
102 domain. Thus, Geneswitch is a chimeric ligand-stimulated activator of transcription. In
103 the absence of ligand, the Geneswitch is in the “off” state. In the presence of the
104 antiprogestin RU486 the Geneswitch molecule changes to an active conformation, in
105 which it binds, as a dimer, to UAS sequences and activates transcription of downstream
106 genes. In flies, Geneswitch mediated expression can be detectable 3–5 hr after feeding
107 on RU486, reaching maximal levels 21–48 hr later (Roman *et al.* 2001; Osterwalder *et*
108 *al.* 2001).

109 20 ml of RU486 (Sigma Aldrich) 10mM stock solution (0.13 g of RU486 in 32 ml of
110 80% ethanol) was mixed with 980 ml molten *Drosophila* food (200 μ M final
111 concentration). For the memory assay, flies were fed for two days with RU486 before
112 the start of the training and returned to the RU486 food after training. For the sleep
113 assay, flies were placed in vials containing RU486 food for two days to allow feeding.
114 After two days flies were immediately loaded into tubes containing more of the RU486
115 food. For all lines we have flies fed with RU486 and genetically identical animals
116 cultured on fly medium supplemented with an equal amount of vehicle (80% ethanol)
117 that lacked RU486.

118

119 Memory assay

120 Each sample was a single sex group of 50 adult flies. This memory assay was described
121 previously (Mery & Kawecki 2005). Conditioning consisted of 5 training sessions
122 separated by 20min intervals. In each training session flies were first exposed for 30s to
123 one odorant simultaneously with mechanical shock delivered every 5s. This period was
124 followed by a 60s rest period (no odour and no shock). Then, for 30s another odorant
125 was delivered, without shock. Flies were either conditioned against 3-octanol or 4-
126 methylcyclohexanol (both 0.6ml/l of paraffin).

127 24 hours after the conditioning period flies were transported to the choice point of a T-
128 maze, where they were allowed to choose between the two odors for 60s. The memory
129 score was the proportion of individuals choosing the correct odour, i.e. not the one they
130 were trained against. One hundred and fifteen replicates were carried out, distributed
131 between the genotype, sex, RU486 (presence/absence) and odour used. The data was
132 normalised using a box-cox transformation.

133

134 Sleep assay

135 Fly locomotor activity was monitored by the *Drosophila* Activity Monitoring System 2
136 (DAMS2, Trikinetics, Waltham, MA), at 25°C, continuously for seventy-two hours
137 under a 12:12 light:dark cycle. Output from DAMS2 was the number of times a fly
138 crossed an infrared beam in a given 1 min period (bin). A sleep episode (bout) was
139 defined as 5 or more consecutive bins of immobility. 384 flies were tested, divided
140 between genotype and RU486 (presence/absence).

141

142 Data analysis for sleep assay

143 The DAMS2 output was converted to three measures; 1) Sleepbins per hour: number of
144 minutes when a fly is asleep in an hour, 2) Mean waking activity: the mean activity
145 taking into account only those bins that are classified as ‘waking’ and 3) Bouts of sleep:
146 the number of sleep episodes.

147 Flies sleep differently during the day and night (Ishimoto *et al.* 2012). Therefore for
148 each dependent sleep variable, two ANOVAs one for day and one for night was run.
149 The independent variables were genotype and RU486 (presence/absence). The
150 important term here is an interaction term between genotype and RU486. If this was
151 significant, the genotypes responded differently to the treatments. To discover which
152 genotypes were significantly different two further ANOVAs were performed, one for
153 genotypes GS1.32>PGRP-Lca vs GS1.32/+ and one for genotypes GS1.32>PGRP-Lca
154 vs +/- PGRP-Lca. If the interaction terms in both these ANOVAs are significant
155 GS1.32>PGRP-Lca (the immune stimulated genotype) responses differently to the

156 control genotypes. Using a Bonferroni correction the significance level α was reduced
157 to 0.0083 (0.05/6). All analysis was carried out using STATA12.

158

159 Zone of inhibition assay

160 Our treatment line had previously been shown to upregulate the immune response
161 (Gottar *et al.* 2002). However we used the zone of inhibition assay to confirm increased
162 immune response in our treated flies. This assay measures antibacterial activity: it is
163 based on the ability of immune proteins to inhibit bacterial growth when placed onto an
164 agar plate seeded with bacteria (*Arthrobacter globiformis* 125 μ l of an overnight
165 culture per 50ml of agar). Thirty seven GS1.32>PGRP-Lca flies, 17 fed RU486 and 20
166 not fed RU486 were used. Each fly was homogenized in 30 μ l of ringer solution. Five
167 microlitres of the supernatant from the centrifuged solution (1300g for 10 min at 4°C)
168 were pipetted into a hole on the agar plate. This was incubated for 48hrs (30°C). The
169 resultant ZOI were measured as the mean of three diameters.

170 **Results**

171 Feeding RU486 to GS1.32>PGRP-Lca flies increased their antibacterial activity by
172 26% ($t = -2.3263$, $df = 29.202$, $p = 0.02715$).

173 Immune stimulation effects on memory

174 Genotype had a significant effect on memory score ($F_{2,109} = 22.46$, $p < 0.0001$). Neither
175 sex, whether RU486 was used, nor odour used had a significant effect on memory
176 score. GS1.32>PGRP-Lca flies, showed a 11.4% decrease in memory scores when fed
177 RU468 relative to those not fed RU468 of the same genotype (interaction between
178 genotype and RU486 was significant $F_{2,109} = 5.76$, $p = 0.0042$). See Figure 1. As
179 feeding RU486 to GS1.32>PGRP-Lca flies leads to an increased immune response,
180 immune stimulation decreases memory scores.

181 Immune stimulation effects on sleep

182 Immune stimulated males (GS1.32>PGRP-Lca fed with RU486) showed a 23%
183 decrease in sleep during the day relative to controls (GS1.32>PGRP-Lca vs GS1.32/+:
184 $F_{1,4607} = 136.29$, $p < 0.00001$, GS1.32>PGRP-Lca vs +/- PGRP-Lca: $F_{1,4535} = 26.87$, $p <$
185 0.00001) and a 9% decrease at night (GS1.32>PGRP-Lca vs GS1.32/+: $F_{1,4607} = 85.53$,
186 $p < 0.00001$, GS1.32>PGRP-Lca vs +/- PGRP-Lca: $F_{1,4535} = 8.49$, $p = 0.0036$). See
187 Figure 2. There was no corresponding change in mean waking activity during the day
188 ($F_{2,6839} = 0.5$, $p = 0.6044$), or during the night (GS1.32>PGRP-Lca vs GS1.32/+: $F_{1,4607}$
189 $= 63.34$, $p < 0.00001$, GS1.32>PGRP-Lca vs +/- PGRP-Lca: $F_{1,4535} = 1.96$, N.S.). There
190 was no change in the number of sleep bouts during the day (GS1.32>PGRP-Lca vs
191 GS1.32/+: $F_{1,4607} = 6.42$, $p = 0.0113$, GS1.32>PGRP-Lca vs +/- PGRP-Lca: $F_{1,4535} =$
192 10.43 , $p = 0.0012$) and a small but significant increase (0.5%) at night (GS1.32>PGRP-

193 Lca vs GS1.32/+ : $F_{1,4607} = 16.38$, $p < 0.00001$, GS1.32>PGRP-Lca vs +/- PGRP-Lca:

194 $F_{1,4535} = 7.56$, $p = 0.0060$).

195 **Discussion**

196 Immune stimulated adult flies exhibit reduced levels of sleep both during day and
197 night. Immune stimulated flies have slightly more fragmented sleep at night, as
198 evinced by an increase in the number of sleep bouts. Immune stimulation also leads to
199 a reduction in memory abilities.

200 The reduction in sleep cannot be explained simply in terms of a generalised increase
201 in activity. Stimulating the immune response had no effect on mean waking activity
202 during the day or night, but immune-stimulated flies slept less than the non-stimulated
203 controls.

204 Our sleep results agree with a previous study by Shirasu-Hiza showing a similar
205 outcome after gram-positive bacterial infections (Shirasu-Hiza *et al.* 2007). However
206 Kuo (Kuo *et al.* 2010) found that when they infected flies with gram-negative
207 bacteria, the flies slept more. The discrepancies in sleep were explained by Kuo *et al.*
208 as being due to different types of infection. Our work did not use an infection but
209 rather a direct stimulation of the immune response. By upregulating PGRP-Lca we
210 reproduced the immune response associated with gram-negative bacteria. This
211 suggests that if type of infection were the cause of the discrepancies, our results
212 would have just mirrored those of Kuo *et al.* The discrepancies between these two
213 previous results are difficult to explain as the experiments differed in numerous other
214 methodical aspects, e.g. strength of infection, lighting paradigm, when the phenotype
215 was measured.

216 Although Imd is one of the canonical immune pathways in insects, over-expression of
217 the Imd pathway can also lead to apoptosis (Georgel *et al.* 2001; Leulier *et al.* 2003).
218 It cannot be excluded that our results could be caused by a side effect: apoptosis of the

219 fat body by the Imd pathway rather than its main effect of immune response. This will
220 be examined in future work.

221 We have shown that immune response decreases sleep and memory in *Drosophila*
222 *melanogaster*. We propose a possible link between all three systems as an interesting
223 area for future research. One of the main hypotheses on sleep function is that sleep
224 periods are favourable for brain plasticity and in the adult brain for learning and
225 memory (Maquet 2001). Like humans, flies with a fragmented sleep show impaired
226 learning compared with flies with consolidated sleep (Seugnet *et al.* 2008). Flies also
227 exhibit decreases in learning after 6 or 12 hours of sleep deprivation (Seugnet *et al.*
228 2006 p. 200). We propose sleep as an intermediate between immunity and memory.
229 We hypothesise that it is not the activation of the immune system *per se* that affects
230 memory in flies, but rather that immune stimulation reduces the length and quality of
231 sleep that in turn, reduces memory ability. However, with our current data, we cannot
232 exclude that in flies the level of immune activation has a direct effect on memory.

233 Our results establishes *Drosophila* as a model for immune-neural interactions. As well
234 as the potential use as a model for mammalian neural-immune links, this work has
235 direct impact on recent concern for insect foragers and the role of multiple stressors in
236 their decline (Gill *et al.* 2012).

237

238

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241 setting up the memory assay. Thanks to E. Green (Genetics, University of Leicester)
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- 336

338 Figure Legends

339

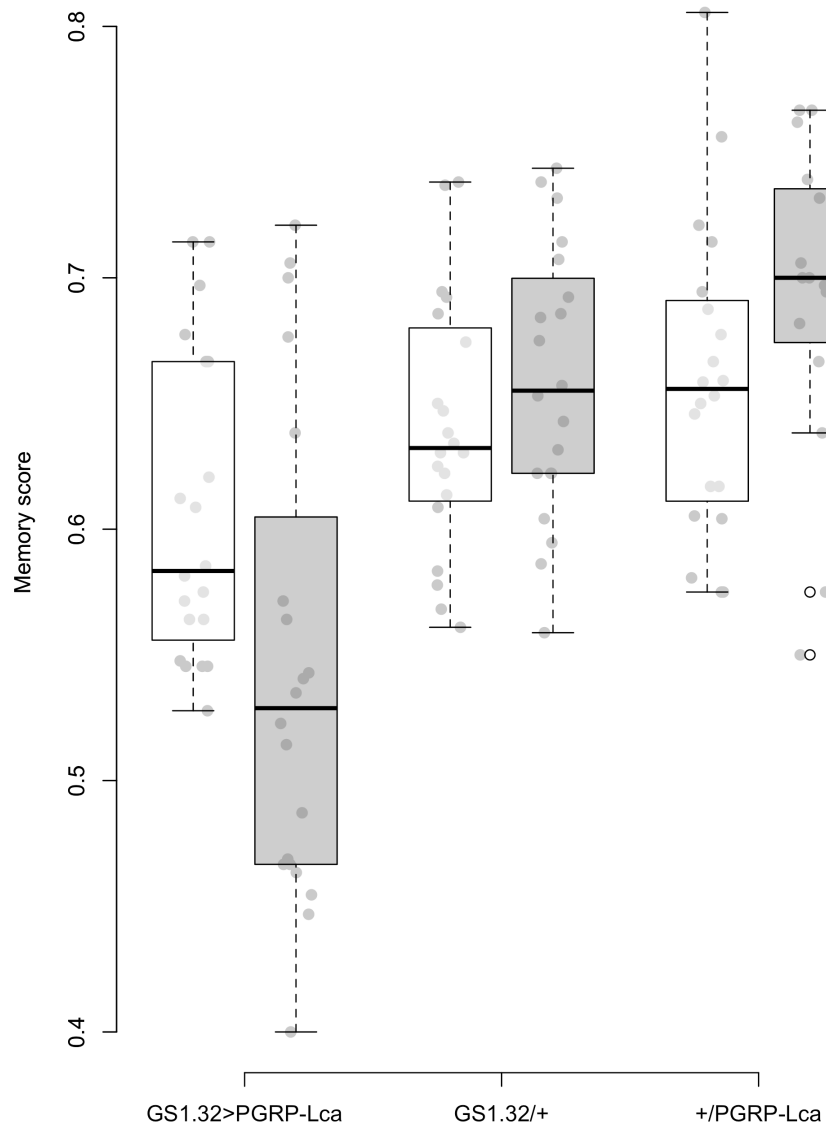
340 Figure 1. Memory score for each genotype. Memory score is the proportion of flies
341 that choose the odour they were not trained against. The white boxes represent the
342 mean memory score for the RU486⁻ flies. The grey boxes represent the RU486⁺ flies.
343 The grey dots are the individual data points.

344

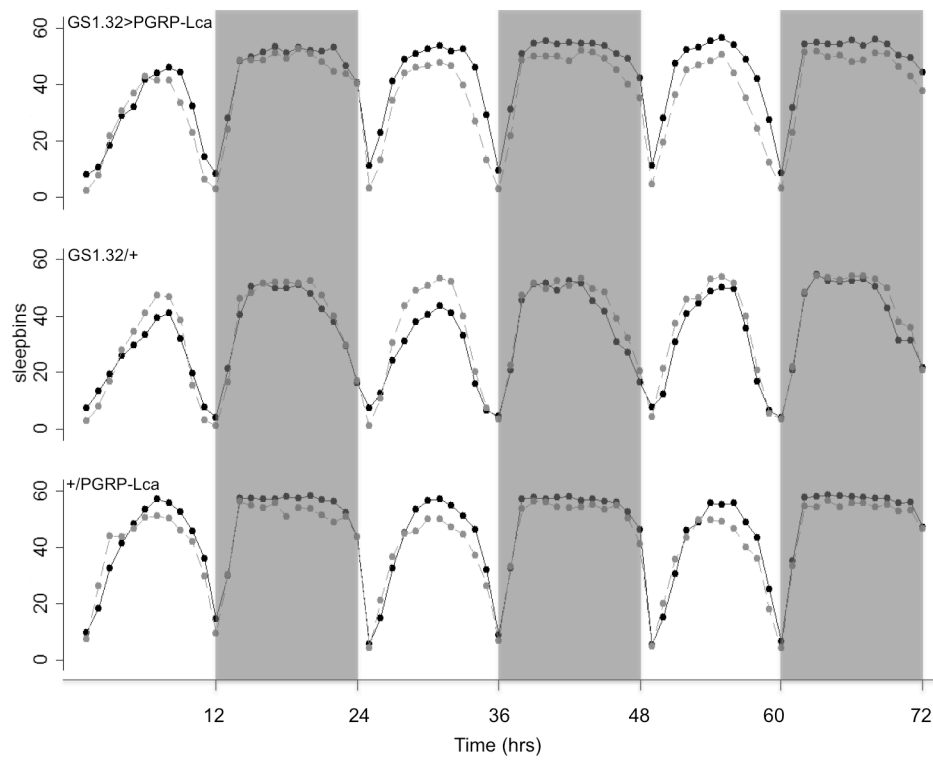
345

346 Figure 2. Sleepbins for each genotype. The black points represent the means of
347 sleepbins for the RU486⁻ flies. The grey points represent the RU486⁺ flies. The
348 shaded times are night (lights off).

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350



351
352 Figure 1



353
354 Figure 2
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356