

## RESEARCH ARTICLE

# Activity-dependent gene expression in honey bee mushroom bodies in response to orientation flight

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### SUMMARY

The natural history of adult worker honey bees (*Apis mellifera*) provides an opportunity to study the molecular basis of learning in an ecological context. Foragers must learn to navigate between the hive and floral locations that may be up to miles away. Young pre-foragers prepare for this task by performing orientation flights near the hive, during which they begin to learn navigational cues such as the appearance of the hive, the position of landmarks, and the movement of the sun. Despite well-described spatial learning and navigation behavior, there is currently limited information on the neural basis of insect spatial learning. We found that *Egr*, an insect homolog of *Egr-1*, is rapidly and transiently upregulated in the mushroom bodies in response to orientation. This result is the first example of an *Egr-1* homolog acting as a learning-related immediate-early gene in an insect and also demonstrates that honey bee orientation uses a molecular mechanism that is known to be involved in many other forms of learning. This transcriptional response occurred both in naïve bees and in foragers induced to re-orient. Further experiments suggest that visual environmental novelty, rather than exercise or memorization of specific visual cues, acts as the stimulus for *Egr* upregulation. Our results implicate the mushroom bodies in spatial learning and emphasize the deep conservation of *Egr*-related pathways in experience-dependent plasticity.

Key words: immediate-early gene, mushroom bodies, honey bee, orientation flight, learning.

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### INTRODUCTION

Honey bees provide an advantageous system for molecular studies of ecologically relevant learning. Honey bees are central place foragers that depend on ephemeral and scattered floral resources. Foraging workers must form robust long-term memories of floral and hive locations, as well as the odor and appearance of preferred floral sources (Robinson and Dyer, 1993; Menzel et al., 2006). Young bees that have not yet initiated foraging behavior (pre-foragers) prepare for the navigational challenges of foraging by performing short learning flights called orientation flights, during which bees acquire information about landmarks and learn to associate position of the sun, time of day and directionality (Winston, 1987; Capaldi and Dyer, 1999). Bees with prior foraging experience will also perform re-orientation flights after the relocation of a colony (Winston, 1987). This information is necessary for later foraging behavior (Becker, 1958), during which bees must accurately navigate between floral sources and the hive. Orientation flights thus offer an exceptional opportunity for molecular analysis because they are discrete, natural learning events driven by innate behavior.

*Early growth response protein 1* (*Egr-1*; also known as *zif268*, *NGFI-A*, *Krox-24* or *zenk*) is a canonical immediate-early gene (IEG), a transcription factor whose expression is activity-dependent and associated with learning and novelty detection in many vertebrate systems (Knapska and Kaczmarek, 2004). *Egr-1* expression is induced in the hippocampus of rodents engaged in spatial learning tasks (Bozon et al., 2002). Recent investigation of IEG activity in insects has yielded growing evidence for activity- and learning-related expression similar to that in vertebrates (Alaux

and Robinson, 2007; Ghosal et al., 2010; Kiya and Kubo, 2011; Kiya et al., 2008; Sen Sarma et al., 2010). However, the insect homolog of *Egr-1*, here named *Egr*, has mainly been studied for its role in muscle development (Volk, 1999), though it has been shown to be upregulated in the brains of flies after seizure, suggesting that it may also be induced by neuronal activity (Guan et al., 2005).

Brain regions involved in vertebrate, particularly mammalian, spatial learning have been identified and extensively studied (Mizumori et al., 2004; Moser et al., 2008). Less is known about the neural basis for spatial learning in insects, but there is evidence to suggest that the mushroom bodies, a region of the insect brain involved in sensory integration and memory, support spatial learning in some species (Mizunami et al., 1998; Farris, 2008).

We explored whether honey bee orientation involves a molecular mechanism that is known to be involved in many other forms of learning, characterized by *Egr* expression. We used this IEG to identify the mushroom bodies as a brain region that is active during orientation flights. To further explore the relationship between *Egr* expression and orientation flights, we conducted a series of behavioral manipulations designed to isolate various aspects of orientation flight experience. To do this, we tested a set of hypotheses related to the ability of exercise, motor learning, specific visual cues and visual environmental perception to induce upregulation of *Egr* (Table 1).

### MATERIALS AND METHODS

#### Phylogenetic analyses

We obtained the nucleotide and amino acid sequences for *Egr-1*, *stripe* (original *Drosophila* name for *Egr* ortholog), *GB50091* and

Table 1. Hypotheses tested to determine what aspects of the orientation flight experience are necessary for *Egr* induction in the honey bee mushroom bodies

Hypothesis	Prediction	Result	Supported?
I. <i>Egr</i> is upregulated by exercise alone	<i>Egr</i> will be upregulated by exercise in the absence of environmental novelty	<i>Egr</i> was not upregulated in the mushroom bodies of bees after performing vigorous wing fanning inside the hive (Fig. 4)	No
II. <i>Egr</i> is upregulated by motor learning associated with flight	<i>Egr</i> will not be upregulated by re-orientation flight in foragers with prior flight experience	<i>Egr</i> was upregulated in the mushroom bodies of foragers after a re-orientation flight (Fig. 5A)	No
III. <i>Egr</i> is upregulated by exposure to specific visual cues	<i>Egr</i> will not be upregulated by flight in environments devoid of visual cues	<i>Egr</i> was upregulated in the mushroom bodies of bees after an orientation flight in deprived environments, and after flight in a white enclosure (Fig. 5B,C)	No
IV. <i>Egr</i> is upregulated by exposure to environmental novelty	<i>Egr</i> will not be upregulated by exercise when environmental novelty cannot be detected	<i>Egr</i> was not upregulated in the mushroom bodies of bees after flight without visual information (Fig. 6)	Yes

AGAP005288 in *Mus musculus*, *Drosophila melanogaster*, *Apis mellifera* and *Anopheles gambiae*, respectively, from GenBank. Sequence orthology was examined using CLUSTALW2 alignments (Goujon et al., 2010) and displayed with Boxshade 3.21 ([http://www.ch.embnet.org/software/BOX\\_form.html](http://www.ch.embnet.org/software/BOX_form.html)).

### Animals

Single-cohort colonies were created as in Robinson et al. (Robinson et al., 1989). Frames of honeycomb containing pupae were collected from 25–30 colonies with naturally mated queens and stored in a dark, humid incubator at 34°C. Adults were removed from the comb within 24 h of emergence, marked on the thorax with paint (Testors, Rockford, IL, USA) and introduced to an experimental colony. Each colony was composed by placing a laying queen and ~3000 one-day-old adult workers bees into a small Styrofoam hive box with several frames with honeycomb containing nectar and pollen. To ensure that bees were naïve to flight, the entrance to each colony was blocked, and colonies were kept inside at 21–23°C and protected from light for 5–6 days after assembly. In single-cohort colonies composed of young bees, a subset will undergo precocious maturation to produce a foraging workforce (Huang and Robinson, 1996). A previous study found that when flight is restricted for several days, bees on the cusp of foraging become ‘primed’ to make an orientation flight, exiting the colony to do so as soon as the entrance is opened (Capaldi and Dyer, 1999). This process therefore creates a convenient population of young bees ready to perform orientation flights. All experiments, other than the test of Hypothesis II, were performed using young, pre-forager bees. On the day of experimentation, colonies were moved outside and allowed to acclimate for at least 30 min.

### Collections and gene expression analyses

Bees that were sampled for molecular analysis were first placed in individual cages and returned to the hive for 30 min (in one experiment, 60 min). Expression of *Egr-1* typically peaks 30 min after the initiating stimulus in vertebrate studies (Zangenehpour and Chaudhuri, 2002). Bees were then flash-frozen in liquid nitrogen (for qRT-PCR analysis) or chilled on ice until brain dissection (for *in situ* analysis).

*In situ* hybridization was performed as previously described (Velarde et al., 2006). Whole brains were dissected in cold physiological saline (Fahrbach et al., 1995), frozen in OCT medium, and cut in 12 µm frontal sections. Sections were mounted on slides, fixed in 4% paraformaldehyde and hybridized overnight with a digoxigenin (DIG)-labeled RNA probe. After incubation with anti-

DIG alkaline phosphatase, slides were developed in NBT/BCIP (nitro-blue tetrazolium chloride and 5-bromo-4-chloro-3'-indolyl phosphate *p*-toluidine salt) until a clear signal was visible.

qRT-PCR on mushroom body tissue was performed as described in Lutz et al. (Lutz et al., 2012). Mushroom bodies were isolated as described in that study after treatment of flash-frozen heads with RNAlater-ice (Life Technologies, Carlsbad, CA, USA). RNA was extracted from isolated mushroom bodies or other brain regions using Picopure extraction kits (Arcturus, Grand Island, NY, USA). Abundance of transcript was analyzed with a SYBR Green probe, quantified relative to a genomic DNA standard curve, and normalized to *elf-s8*, a constitutively expressed endogenous control gene used in a previous study (Alaux et al., 2009). In each experiment, expression levels are expressed as ratios, relative to the control or sham-treated group for that experiment. Results shown are an average of two trials (for the initial investigation of orientation flight, three trials;  $N=6-8$  for all groups). Two-factor ANOVAs were performed for multiple trials using PROC MIXED in SAS (SAS Institute, Cary, NC, USA).

Sequences of primers used to generate the probe for *in situ* hybridization, as well as for qRT-PCR, are listed in Table 2.

### Is *Egr* upregulated by orientation flight?

To observe and collect bees performing orientation flights, groups of six to 10 bees were allowed to exit a colony, and an observer confirmed that the flight pattern and duration were typical of orienting bees (Winston, 1987). Control bees were captured while exiting the hive and held for 6–7 min in individual cages next to the hive. The 6–7 min holding period allowed control bees to experience sunlight, outdoor odors and other environmental stimuli for an amount of time roughly equivalent to an orientation flight. Because control bees were collected while exiting the hive, it is presumed that these individuals would have performed an orientation flight had they not been captured.

The same conditions used in this experiment were adapted in order to test specific hypotheses for the upregulation of *Egr* (Table 1), as described in the following sections.

### Hypothesis I – is *Egr* upregulated by exercise in the absence of flight?

We tested this hypothesis by studying exercise in a familiar environment. Fanning bees (bees exhibiting vigorous, flight-like wing movement while remaining stationary) were collected by setting up a colony as described for naïve orientation flight collections, but in a small glass-walled hive housed indoors; only

Table 2. Primers used to produce RNA probe for *in situ* hybridization and qRT-PCR

Primer	Primer name	Sequence 5'–3'
ISH probe	<i>Egr-ISH-F</i>	TAA TAC GAC TCA CTA TAG GGC GCA AGT ACC CGA ATC GAC
	<i>Egr-ISH-R</i>	GAT TTA GGT GAC ACT ATA GGC TTC TTC TCG TCG CTC CTC
qRT-PCR	<i>Egr-F</i>	GCA AAC GGT GCA GCT CAG T
	<i>Egr-R</i>	CCG CAT ACG ATC GAA TTC G
	<i>eif3-s8-F</i>	TGA GTG TCT GCT ATG GAT TGC AA
	<i>eif3-s8-R</i>	TCG CGG CTC GTG GTA AA

1000 worker bees were added. After 7 days, the colony was heated to 40°C for 45 min. This induced some bees to beat their wings vigorously at the hive entrance to provide ventilation (Southwick and Moritz, 1987). Bees that engaged in this fanning behavior for at least 2 min were collected, along with several inactive bees to serve as controls.

#### Hypothesis II – is *Egr* upregulated by motor learning associated with flight?

We tested this hypothesis by inducing re-orientation flights by foragers. Foragers, bees with extensive flight experience, perform additional re-orientation flights after a hive is moved, or after moving to a new nest site as part of a swarm (Winston, 1987). These re-orientation flights allow bees to learn their new environment. To induce re-orientation, small colonies with natural age demographics were closed after sunset and transported to a new location at a distance outside the maximum foraging range (Capaldi and Dyer, 1999; Capaldi et al., 2000). This treatment induces bees with foraging experience to perform additional orientation flights in their new environment. The morning after the colony move, orienting foragers and controls were collected as described above, including observations of flight structure and duration. Returning bees were examined to ensure that they had not gathered a pollen or nectar load, indicating that they were performing an orientation flight and not foraging.

#### Hypothesis III – is *Egr* upregulated by exposure to specific visual cues?

We tested this hypothesis by studying orientation flights in deprived conditions. Bees will perform orientation flights even in indoor arenas of limited size, devoid of distant landmarks (Brandon and Coss, 1982); such environments are in extreme contrast to the sensorially enriched outdoors. The experimental conditions described above for testing response to a typical first orientation flight were repeated for tethered bees in a small indoor flight enclosure relatively devoid of visual patterning. Temperature in the enclosure was 29.5°C during the day and 24°C at night, and lights were on a 12 h:12 h light:dark cycle.

Bees used in tethered flight experiments came from colonies prepared as for the naïve orientation flight collections. Bees were captured as they exited the colony entrance in preparation for an orientation flight and then immediately placed on ice just until anesthetized. One end of a thread was then fixed to the thorax using superglue; the other end was fixed to the end of a rod. Tethered bees were placed on the floor of a white box and allowed to recover until they were able to walk in a coordinated manner. Flight was induced by lifting the bee's feet from the ground (Feller and Nachtigall, 1989); this action was repeated as often as necessary to sustain flight for 2 min, usually zero to two additional times. Controls were treated in the same manner as foragers, but restrained in an individual plastic cage rather than being induced to fly.

#### Hypothesis IV – is *Egr* upregulated by exposure to environmental novelty?

We tested this hypothesis by studying flight in the absence of visual input. The tethered flight experiment described above was repeated using red lighting, which is not visible to bees (Peitsch et al., 1992).

## RESULTS

#### Honey bee *Egr* is a homolog of the vertebrate *Egr* gene family

We investigated *GB50091*, a 2448 bp transcript encoding a putative 815 amino acid protein. This gene is an ortholog of the *Drosophila* gene *stripe*; the genes are reciprocal best BLAST hits and share 53.8% sequence identity (e-value: 1e-95). Like *stripe*, the honey bee gene is a homolog of genes in the *Egr* gene family in vertebrates, including *Egr-1*, with which it shares 44.5% sequence identity (e-value: 2e-46). The DNA-binding domain, which consists of three zinc-finger domains (Pavletich and Pabo, 1991), is highly conserved between bee, fly, mouse and mosquito (Fig. 1). The amino acid sequence identity between the honey bee and *Drosophila* zinc-finger domains is 94%; between the honey bee and mouse zinc-finger domains it is 89%. Because of the apparent functional similarity (see below) to the vertebrate gene *Egr-1*, we have used the name *Egr* to refer to the honey bee gene.

#### *Egr* expression is rapidly and transiently increased in the mushroom bodies of orienting bees

We first assessed the spatial distribution of *Egr* mRNA in the honey bee brain, using *in situ* hybridization and qPCR. *In situ* analysis revealed visible *Egr* mRNA expression only in scattered Kenyon cells in the mushroom bodies (Fig. 2). Similarly, qRT-PCR analysis showed that *Egr* expression was enriched in the mushroom bodies compared with the rest of the brain (Fig. 3).

We next assessed activity-dependent changes in *Egr* expression in the mushroom bodies of pre-forager bees after a single orientation flight. In many vertebrate species, activity-dependent *Egr-1* mRNA expression peaks ~30 min after the relevant stimulus and returns to baseline after 60 min (Zangenehpour and Chaudhuri, 2002). We investigated the temporal dynamics of *Egr* expression in orienting bees by analyzing transcript abundance in the mushroom bodies of bees collected 30 min and 1 h after flight.

Orienting bees showed an almost twofold upregulation of *Egr* in comparison to controls ( $P < 0.005$ ; Fig. 3) 30 min after flight. Because control bees were collected as they attempted to perform an orientation flight, and were exposed to sunlight and environmental odors, we hypothesized that increased *Egr* expression in the mushroom bodies was caused by an aspect of the orientation flight other than preparedness or exposure to stimuli.

The orientation flight, although most significant as an example of spatial learning, can be broken down into several component experiences: exercise, motor learning, exposure to visual cues and exposure to novelty. Any of these components could conceivably induce an IEG response. We performed a series of experiments to

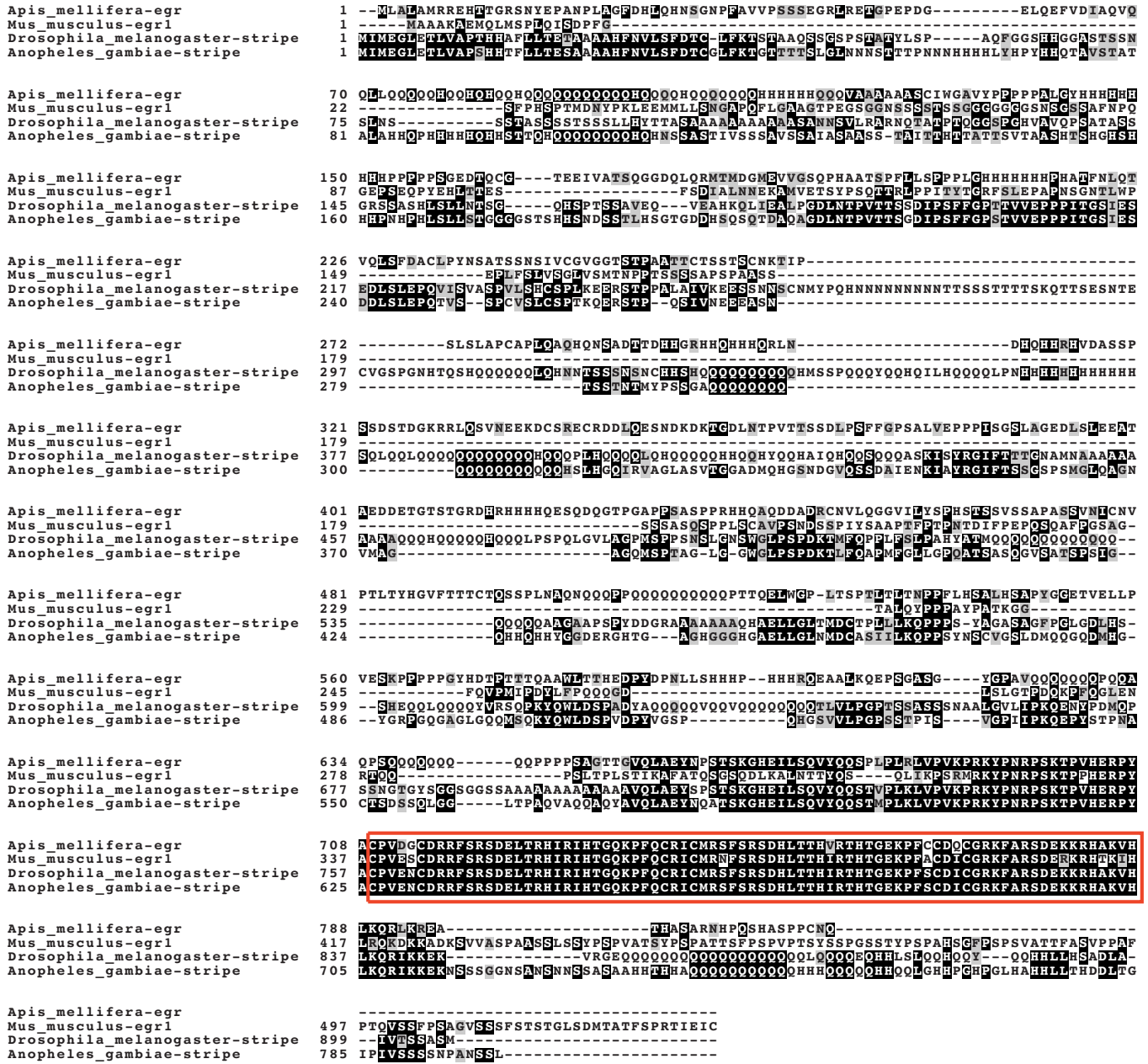


Fig. 1. Alignment of *Egr* with a vertebrate homolog in mouse (*Mus musculus*), and orthologs in fruit fly (*Drosophila melanogaster*) and mosquito (*Anopheles gambiae*). Conserved residues are highlighted in dark gray. The DNA-binding domain is outlined with a red box.

test predictions stemming from specific hypotheses on the relationship between the experience of an orientation flight and upregulation of *Egr* (Table 1).

***Egr* is not upregulated by exercise in the absence of flight**

Because flight in honey bees represents an abrupt increase in metabolic demand, we wondered whether flight mediates an exercise-dependent increase in *Egr* expression. In rodents, *c-fos*, *arc* and *Egr-1* expression in the hippocampus increase in response to exercise (Rhodes et al., 2003; Clark et al., 2011). We devised a behavioral manipulation that decoupled the exercise associated with rapid wing movement from real flight. The exercise examined was wing fanning inside the hive entrance to ventilate the hive after heat stress. Pre-forager bees that were collected while fanning

experienced a level of activity roughly comparable with that during flight, but in a familiar environment (Yang et al., 2010).

Fanning bees showed no *Egr* upregulation in response to this form of exercise compared with controls ( $P=0.859$ ; Fig. 4). As a positive control, bees from the same colony did show the expected *Egr* upregulation in response to an orientation flight ( $P<0.001$ ). Although expression in fanning bees appears to be lower than unmanipulated controls from the same colony, this difference is not significant. This experiment suggests that exercise alone cannot induce *Egr* upregulation in the mushroom bodies.

***Egr* is not upregulated by motor learning associated with flight**

Experienced foragers will perform additional orientation flights after participating in a swarm, or after a colony is moved by humans



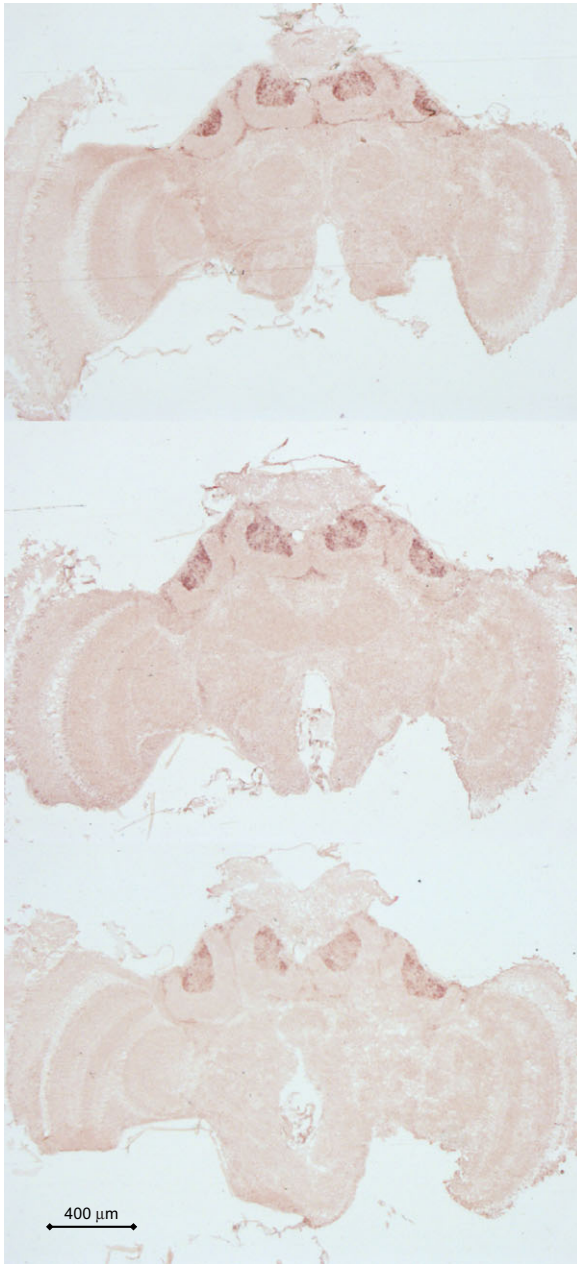


Fig. 2. Frontal sections of a brain from an orienting bee. Dark red staining indicates *Egr* expression in mushroom body neuron somata. Staining was not observed in any other brain region.

(Winston, 1987). We took advantage of this to examine *Egr* expression in the mushroom bodies of foragers that performed a re-orientation flight after a colony move. If *Egr* upregulation was a response to the motor learning aspect of orientation flights performed by pre-foragers, a similar upregulation would not be expected to occur in re-orienting foragers with several days of prior flight experience.

*Egr* upregulation is not caused by the motor learning aspect of orientation flights, because re-orienting foragers exhibited upregulation of mushroom body *Egr* expression in comparison to controls ( $P < 0.0005$ ; Fig. 5A). It is thus likely that *Egr* is upregulated in response to some other experience common to both orientation and re-orientation flights.

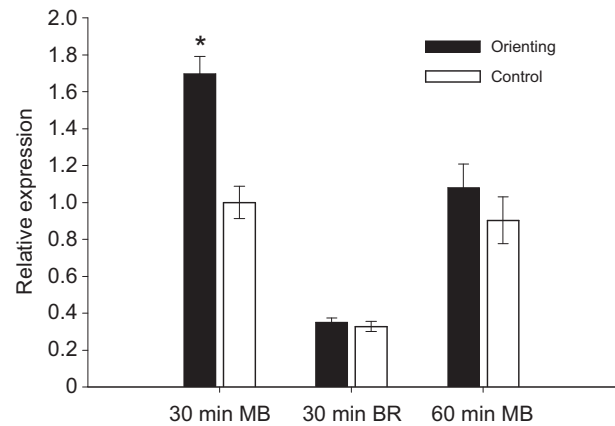


Fig. 3. Expression of *Egr* in the brains of orienting and control bees collected either 30 or 60 min post-collection, measured using qRT-PCR. MB, mushroom bodies; BR, all brain tissue other than mushroom bodies. *Egr* showed nearly twofold upregulation in mushroom bodies 30 min after orientation flight ( $P < 0.0001$ ). *Egr* was not differentially expressed between orienting bees and controls in BR samples at 30 min, or in MB 60 min after orientation flight.

#### ***Egr* upregulation does not require exposure to specific visual cues**

To examine the relationship between feature memorization and *Egr* upregulation, we investigated the transcriptional response to orientation flight in visually deprived environments. We collected pre-forager bees after they performed an orientation flight in a small indoor flight arena with white floors and bare screen walls and ceiling. Bees that performed an orientation flight in this deprived environment still showed upregulation of mushroom body *Egr* expression compared with controls ( $P < 0.0001$ ; Fig. 5B).

To examine the effect of even more extreme visual deprivation during flight, pre-forager bees preparing for an orientation flight were tethered to a restraint that allowed freedom of movement and induced to fly in a blank, white arena. Bees that experienced 2 min of tethered flight in a white box devoid of visual cues also showed increased *Egr* expression compared with tethered and untreated controls ( $P < 0.0001$ ; Fig. 5C). Although tethered bees were exposed to a novel environment during flight, exposure to specific visual cues was not necessary to induce *Egr* upregulation.

#### **Exposure to environmental novelty is necessary for *Egr* upregulation**

IEG expression in many vertebrate species is upregulated in response to the novelty of a stimulus (Tischmeyer and Grimm, 1999; Clayton, 2000), and orientation flights expose bees to novel stimuli. To test the necessity of visual novelty to induce *Egr* upregulation, we designed an experiment to give bees the experience of flight without visual novelty. Pre-forager bees experienced tethered flight as described above, but in perceived darkness (Peitsch et al., 1992). These tethered bees showed no upregulation of *Egr* in comparison with controls ( $P = 0.303$ ; Fig. 6). Orienting bees from the same colony showed upregulation, as in experiments reported above ( $P < 0.005$ ). These results suggest that visual environmental novelty during the orientation flight is necessary to induce *Egr* upregulation in the mushroom bodies.

#### **DISCUSSION**

We investigated the transcriptional response of *Egr*, an insect homolog of *Egr-1*, in the brains of honey bees after the performance

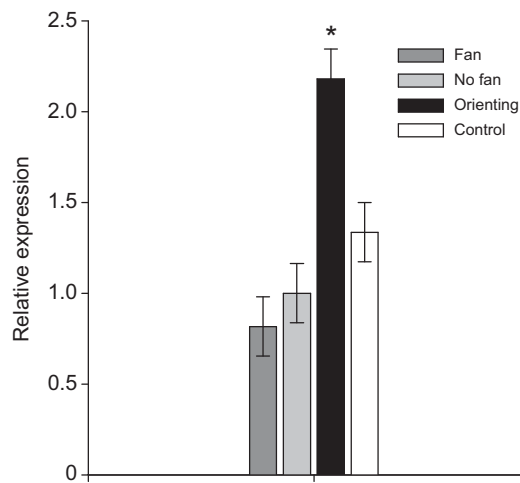


Fig. 4. *Egr* is not upregulated by exercise alone. Expression of *Egr* in the mushroom bodies of bees vigorously fanning their wings for ~2 min to ventilate the hive, compared with orienting bees and controls. *Egr* was upregulated in response to flight ( $P < 0.001$ ), but not fanning ( $P = 0.859$ ).

of orientation flights and related behaviors. We found that a single orientation flight upregulates *Egr* expression exclusively in the mushroom bodies. By contrast, this increased expression was not seen in bees exiting the hive for a flight. *Egr* expression in the mushroom bodies is thus associated with the experience of, rather than the anticipation of, a naturally occurring learning event.

To our knowledge, we are the first to report that *Egr* displays activity-dependent expression in an insect, providing evidence for a highly conserved role as an IEG. Despite extensive literature linking the *Egr* gene family to neuronal activation and learning in vertebrates, there has been little work on the role of this gene in the *Drosophila* nervous system, and no study linking this gene to novelty or learning.

In vertebrates it has been difficult to distinguish between *Egr* expression induction resulting from the behavior or stimulus of interest and that resulting from exercise, stress or other confounding factors (Knapska and Kaczmarek, 2004). We found that flight in

several contexts and environments was sufficient to cause this upregulation, as long as the environment was novel, but exercise was insufficient to cause *Egr* upregulation without the visual perception of environmental novelty. These results demonstrate that it is possible for the perception of an experience to trigger *Egr* expression. Similarly, there are some genes in the mushroom bodies and optic lobes of honey bees that are responsive to changes in the perception of flight distance, rather than the actual distance flown (Sen Sarma et al., 2010).

*Egr-1* is hypothesized to play a role in memory consolidation by promoting structural neuroplasticity in the vertebrate brain following exposure to novel or salient stimuli (Knapska and Kaczmarek, 2004; Tischmeyer and Grimm, 1999; Moorman et al., 2011). Many regulatory targets of *Egr-1* have been identified, and several are involved in synaptic formation and remodeling. Given this new demonstration of the deep evolutionary conservation of *Egr* as a mediator of experience-dependent plasticity, it is plausible that *Egr* may function similarly in the bee brain. A limitation of the present study is that it did not examine potential downstream targets of honey bee *Egr*. However, because of its homology with vertebrate *Egr-1*, *Egr* is a good candidate for a functional link between orientation flights and the resulting dendritic spine remodeling on mushroom body neurons (Brandon and Coss, 1982). Future investigations of downstream targets of neuronal *Egr* in the honey bees or other insects may yield a more complete picture of how environmental stimuli act through molecular signaling to promote neuroanatomical change. Future studies should also involve determination of the behavioral and downstream molecular consequences of manipulations of *Egr* signaling through pharmacological manipulations or RNAi knockdown. Such work could provide a more causal link between *Egr*, downstream signaling and the resulting consolidation of learning that is believed to occur in vertebrates (Bozon et al., 2003; Davis et al., 2003; Pérez-Cadahía et al., 2011).

Our results suggest that the interaction between flight activity and visual novelty is sufficient to stimulate *Egr* upregulation. To clarify whether flight activity, in combination with visual novelty, is necessary for *Egr* upregulation, future experiments could examine gene expression or homing behavior after bees are transported mechanically in a manner that imitates flight.

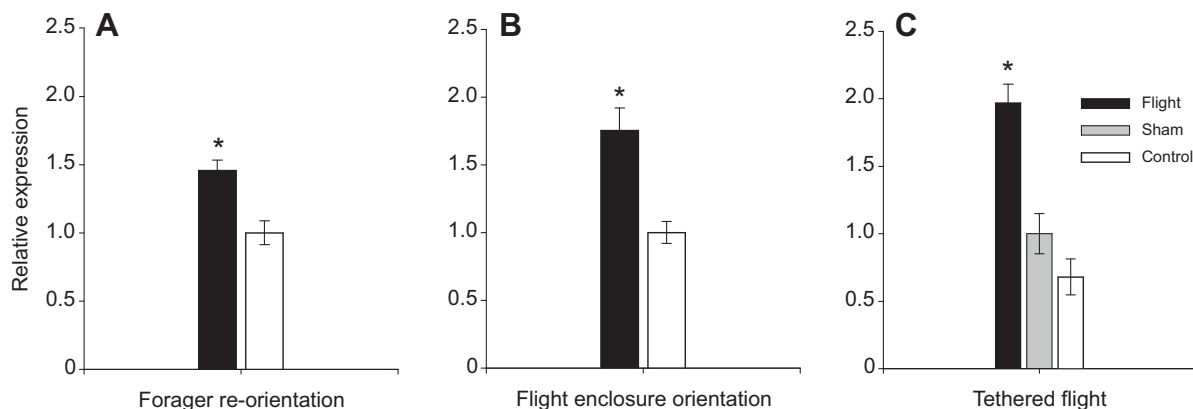


Fig. 5. (A) *Egr* is not upregulated by motor learning associated with flight. Expression of *Egr* in the mushroom bodies of re-orienting and control foragers. *Egr* showed upregulation in response to re-orientation by experienced foragers in a novel environment ( $P < 0.0005$ ). (B,C) *Egr* is not upregulated by exposure to specific visual cues. (B) Expression of *Egr* in the mushroom bodies of bees orienting in an indoor flight enclosure, compared with controls. *Egr* was upregulated in response to an orientation flight in this more deprived environment ( $P < 0.0001$ ). (C) Expression of *Egr* in the mushroom bodies of tethered bees flying for 2 min in a white enclosure. Flight indicates bees that experienced flight while tethered. Sham bees were anesthetized and tethered, but not allowed to fly. Controls were simultaneously collected, unmanipulated bees. *Egr* was upregulated in response to flight ( $P < 0.0001$ ).

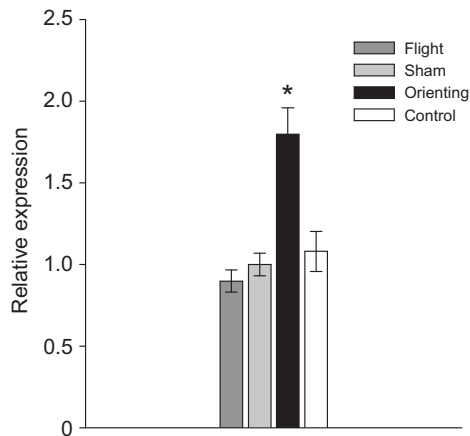


Fig. 6. *Egr* is upregulated by exposure to environmental novelty, but not by exercise alone. Expression of *Egr* in the mushroom bodies of tethered bees flying for 2 min in a red light, compared with orienting bees and controls. *Egr* was upregulated in response to flight ( $P < 0.005$ ), but not to tethered flight in red light ( $P = 0.303$ ).

Our molecular and neuroanatomical results support the hypothesis that the mushroom bodies are involved in spatial learning in honey bees, which contrasts with previous research on *Drosophila* (Neuser et al., 2008; Wolf et al., 1998; Putz and Heisenberg, 2002; Sitaraman et al., 2008; Zars et al., 2000; Ofstad et al., 2011). However, prior work in other insect species besides *Drosophila* has suggested a possible link between the mushroom bodies and spatial learning. Cockroaches with lesioned mushroom bodies perform poorly on a spatial learning task similar to the Morris water maze (Mizunami et al., 1998). In honey bees, mushroom body neuropil undergoes age-related expansion roughly coincident with orientation flights in both workers and reproductives (Fahrbach et al., 1995; Brandon and Coss, 1982; Withers et al., 1993; Fahrbach et al., 1997), and as mentioned above, one study found rapid remodeling of dendritic spines in the mushroom bodies after an orientation flight (Brandon and Coss, 1982). In addition, Kiya et al. (Kiya et al., 2007) found evidence of neuronal activation in the mushroom bodies of re-orienting foragers. *Drosophila* have minimal mushroom bodies that receive olfactory input almost exclusively (Heisenberg, 2003), while those of honey bees comprise a substantial portion of total brain volume and receive significant visual and olfactory input. These differences may relate to species differences in foraging ecology, as argued in the following paragraph.

It has been recently suggested that *Drosophila* spatial learning may not involve the mushroom bodies because this species, in sharp contrast to honey bees, has no need for long-term spatial memories (Zeil, 2012). Hymenopterans (which include bees) and other insects that must navigate a fairly constant environment for days, weeks or months would be predicted to exhibit mushroom body involvement in spatial learning. In addition, insect species whose feeding ecologies depend more heavily on navigation have relatively larger, more elaborate mushroom bodies than *Drosophila* (Farris, 2008; Mizunami et al., 1998). Given the similarity in mushroom body structure in hymenopterans and the shared ecological pressures that may have shaped them (Farris, 2008), a role in spatial learning is more likely to be shared in bees, ants and wasps, and perhaps other species with elaborate mushroom bodies as well. Our findings suggest that *Egr* can be a useful tool to understanding the ability of large and small brains to acquire the spatial information needed to navigate successfully.

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## AUTHOR CONTRIBUTIONS

C.C.L. was involved in all aspects of the study. G.E.R. was involved in conception and design of the study, interpretation of the findings, and revising the article.

## COMPETING INTERESTS

No competing interests declared.

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