Adaptations during Maturation in an Identified Honeybee Interneuron Responsive to Waggle Dance Vibration Signals

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Abstract

Honeybees are social insects, and individual bees take on different social roles as they mature, performing a multitude of tasks that involve multi-modal sensory integration. Several activities vital for foraging, like flight and waggle dance communication, involve sensing air vibrations through their antennae. We investigated changes in the identified vibration-sensitive interneuron DL-Int-1 in the honeybee Apis mellifera during maturation by comparing properties of neurons from newly emerged adult and forager honeybees. Although comparison of morphological reconstructions of the neurons revealed no significant changes in gross dendritic features, consistent and region-dependent changes were found in dendritic density. Comparison of electrophysiological properties showed an increase in the firing rate differences between stimulus and nonstimulus periods in foragers compared with newly emerged adult bees. The observed differences in neurons of foragers compared with newly emerged adult honeybees suggest refined connectivity, improved signal propagation, and enhancement of response features possibly important for the network processing of air vibration signals relevant for the waggle dance communication of honeybees.

Key words: adaptation; honeybee; maturation; neuron morphology

Significance Statement

In the darkness of the hive, honeybees inform each other about profitable food sources using stereotypic movements accompanied by specific sound patterns produced by wing-beats. Here we present a study of an identified vibration-sensitive neuron, named DL-Int-1, in the honeybee brain focusing on structural and functional adaptations by comparing data from young, newly emerged adult and mature forager honeybees. We found region-dependent changes in the morphological structure of DL-Int-1 as well as specific changes in its response properties, which suggest an adaptation process during maturation leading to a refinement in network connectivity and improved processing of waggle dance signals in the honeybee brain.

Introduction

Perception of vibrations and sounds is very important for social insects (Hunt and Richard, 2013) and among them, honeybees are unique in that they use air-borne vibrations for communication (Kirchner, 1997). Among several intra-hive communication behaviors linked to airborne vibration sensing (Barth et al., 2005; Hunt and Richard, 2013; Nieh, 2010), the waggle dance behavior, which is used to communicate the distance, direction and profitability of food sources, has been extensively studied.
in the honeybee *Apis mellifera* (von Frisch 1965, 1967; Kirchner and Towne 1994; Brockmann and Robinson, 2007; Hrmcic et al., 2011; Couvillon, 2012). Waggle dance behavior consists of alternative repetitions of two movements: a straight onward movement called the “waggle phase” during which honeybees produce air vibrations by oscillating their abdomen from side to side and beating their wings; and a curved movement called the “return phase” during which they return to the starting point of the onward phase. The neural mechanisms underlying the processing and decoding of the waggle dance vibration signals have so far not been uncovered.

Air vibrations behind dancing honeybees are detectable only up to 15–20 mm (Michelsen et al., 1987) and consist of low (12–25 Hz) and high (200–300 Hz) frequencies from two sources (Wenner, 1962). Vibrations produced by the wagging abdomen only contain low frequencies, whereas jets of air vibration produced by wing beats have a pulse train pattern and contain both low and high frequencies (Michelsen et al., 1987; Michelsen, 2003). Both kind of vibrations have been shown to be relevant for waggle dance communication (Michelsen et al., 1989, 1992). Honeybees can detect air vibrations using various mechanosensory organs on their bodies. Among them, the Johnston’s Organ (JO) located in the pedicel of the antennae (Fig. 1a) has been shown to be the primary sensory organ for detecting near-field vibrations of the waggle dance (Dreiler and Kirchner, 1993). Sensory afferents of the JO project into the honeybee brain, specifically in medial posterior protocerebral lobe (mPPL) and the antennal mechanosensory and motor center (AMMC), which consists of the dorsal lobe (DL) and dorsal subesophageal ganglion (dSEG; Fig. 1a; Ai et al., 2007).

More than 10 groups of interneurons belonging to three categories have been identified and characterized in these regions (Ai et al., 2009, 2017; Ai, 2010, 2013) and have been shown to respond to antennal vibrations similar to those produced by air vibration jets of the waggle dance (Ai et al., 2017). In particular, a group of GABAergic interneurons called DL-Int-1 has been studied intensively and has been characterized in detail (Ai et al., 2009, 2017; Ai, 2013). However, neural responses to low-frequency vibrations produced by abdomen wagging has not yet been characterized.

DL-Int-1 somata are located in the rind of the protocerebrum and have single neurites branching and projecting to the DL, the dSEG and the mPPL, where they further branch into dense arborizations that run close to afferents (Fig. 1b; Ai et al., 2009). DL-Int-1 are spontaneously active and respond to vibration stimuli applied to the ipsilateral antenna. Their responses to vibration stimuli are characterized by on-phasic excitation to stimulus onset, tonic inhibition during continuous stimulation, and rebound spiking after vibration offset (Ai et al., 2009). DL-Int-1 neurons are thought to play a role in encoding the duration of the waggle phase (Ai et al., 2017), which correlates with the distance of the advertised food source from the hive (von Frisch, 1967).

As they mature, adult honeybees engage in four primary social roles—cleaners, nurses, food storers, and foragers—and perform different tasks in different roles (Seeley, 1996). Several studies have investigated the neural basis of such behavioral versatility by studying structural changes in the honeybee brain with age and social role, mainly focusing on the mushroom body (Groh, et al., 2006, 2012; Groh and Meierzhagen, 2010). Although most developmental changes in the honeybee brain occur during pupal and larval stages (Devaud and Masson, 1999; Ganeshina et al., 2000), considerable age-dependent and experience-dependent anatomic changes have been described at the level of subregions in the adult honeybee antennal lobe (Winnington et al., 1996; Sigg et al., 1997; Morgan et al., 1998; Brown et al., 2004; Andrione et al., 2017; Arenas et al., 2013) and the mushroom body (Withers et al., 1993, 1995; Durst et al., 1994; Fahrbach et al., 1998; Wolschin et al., 2009), as well as at the level of single mushroom body neurons (Farris et al., 2001). In addition, electrophysiological properties of honeybee neurons also mature with age and experience in the antennal lobe (Wang et al., 2005) and in the mushroom body (Kiya et al., 2007). Adaptations in neural processing could be especially crucial during the transition to foraging, because, compared with in-hive activities, foraging entails several new and complex behaviors such as attending to waggle dancers, sensing the waggle dance vibration signals and decoding target location, using such information on foraging trips, and advertising newly found locations to hivemates through the waggle dance. Honeybees start following waggle dances only after 1 week after emergence (Ai et al., 2018b). Mechanosensory neurons in the JO of the antennae become more responsive to high-frequency waggle dance vibrations as honeybees mature from newly emerged adults to foragers (Tsujii et al., 2007). It is unclear, however, to what extent neurons in central circuits processing waggle dance vibration signals show such adaptation. We therefore investigated morphological and electrophysiological changes of neurons in the primary mechanosensory center of the honeybee, focusing on DL-Int-1 neurons. To identify maturation-related adaptations in DL-Int-1, we analyzed and compared reconstructed morphologies and electrophysiological properties of neurons from newly emerged adult bees and foragers.
Materials and Methods

Honeybees (Apis mellifera) reared at Fukuoka University between 2012 and 2014 were used in this study. Experiments were conducted on more than 300 bees for investigating neurons in the primary mechanosensory center of the honeybee brain. Collected data included electrophysiological recordings and laser scanning microscopy images, which were stored in a database and classified into multiple neuron groups based on electrophysiological and morphological characteristics (Ai et al., 2017, their Table 1). In the current study, we used DL-Int-1 data from the database belonging to honeybees of two stages of maturation:
- **Newly emerged adults (age 1-3 d):** female honeybees shortly after emerging from their cell in the hive. Before the experiments, these bees were kept in isolated cages containing sugar solution and pollen.
- **Foragers (older than 10 d):** female honeybees returning from foraging with pollen on their hindlegs.

Experimental procedure

The experimental procedure for generating image stacks and electrophysiological response traces from
honeybee vibration-sensitive interneurons has been presented in detail by Ai et al. (2017) and we describe it here briefly. After immobilization and head fixing using bee's wax, the frontal surface of the honeybee brain was exposed by cutting away a small rectangular window between the compound eyes. Borosilicate glass electrodes filled at the tip with a dye were inserted into the primary mechanosensory center to record from individual neurons. Three dyes were used: Lucifer yellow CH dilithium salt (catalog number L0259, Sigma-Aldrich), Dextran tetramethylrhodamine solution (3000 molecular weight, anionic, lysine fixable; catalog number D3308, ThermoFisher Scientific), and AlexaFluor 647 hydrazide (catalog number A20502, ThermoFisher Scientific), and AlexaFluor 647 hydrazide (catalog number A20502, ThermoFisher Scientific). With the electrode stump inserted into a vibration-sensitive interneuron, sinusoidal vibration stimuli of frequency 265 Hz and duration 1 s were applied to the right antenna and responses were recorded intracellularly. Electrical signals were amplified using an amplifier (MEZ8301, Nihon Kohden), filtered to remove frequencies higher than 20 kHz and recorded using Spike2 (Cambridge Electronic Design; RRID: SCR_000903) at a sampling rate of 20.833 kHz. After recording electrical activity, a hyperpolarizing current (2–5 nA for 2–10 min) was applied to inject the dye into the neuron. Thereafter, the brains were dissected out, fixed in 4% paraformaldehyde for 4 h at room temperature, and then rinsed in phosphate buffer solution, dehydrated, and cleared in methyl salicylate for subsequent observation and imaging.

The cleared specimen containing intracellularly stained neurons were viewed from the posterior side of the brain under a confocal laser-scanning microscope (LSM 510, Carl Zeiss) with a Zeiss Plan-Apochromat 25×/numerical aperture 0.8 oil lens objective (working distance, 0.57 mm). Image stacks of the AMMC and the mPPL were taken at a resolution of 0.36 μm on the imaging plane using 1-μm-thick optical sections and stitched together digitally to obtain image stacks of complete neurons.

**Morphological subregions of DL-Int-1**

To refer to specific subregions of the DL-Int-1 morphology, we adopt the following definitions (Fig. 1c; Ai et al., 2017):

- **Soma and primary neurite (SPN):** consists of the soma and its primary neurite until bifurcation.
- **Main branch (MB):** consists of the two daughter branches of the primary neurite until they bifurcate.
- **Dorsal branch (DB):** consists of the remaining dendritic arborization originating from the dorsal end of the MB.
- **Ventral branch (VB):** consists of the remaining dendritic arborization originating from the ventral end of the MB.
- **Whole arborization (WA):** consists of the MB, the DB, and the VB.

**Reconstruction of morphologies**

The reconstruction procedure has been described in detail by Ikeno et al. (2018). Briefly, image stacks with single dye-filled neurons were de-convolved to reduce image blurring and noise. Regions of each image stack containing the dendritic subtrees emerging from the dorsal and ventral daughter branches of SPNs (Ikeno et al., 2018, their Fig. 6E) were identified based on continuity of branching structure and dendritic thickness and were converted into custom image masks. Applying these masks, two image stacks were created that separately contained the identified dorsal and ventral subtrees of SPN. Morphologies of these subtrees were reconstructed from their image stacks by segmentation, pruning and smoothing using SIGEN software (Minemoto et al., 2009; RRID:SCR_016284) and combined to form the reconstructed morphologies of the neurons.

### Table 1. Scalar morphometric measures showing significant differences

| Measure                                      | Morphological subregion | Newly emerged (NE, n = 6) | Forager (F, n = 6) | Change in median value ([F-NE]/NE) | p     |
|----------------------------------------------|-------------------------|---------------------------|-------------------|------------------------------------|-------|
| Width (along x), μm                         | Main branch             | 10.4, 22.1, 26.5          | 25.2, 34.1, 61.4  | +54.3%                             | 0.004 |
| Height (along y), μm                        | Dorsal branch           | 152, 236, 263            | 212, 268, 294     | +13.6%                             | 0.041 |
| Total dendritic volume, \( \times 10^3 (\mu m)^3 \) | Main branch             | 0.183, 0.409, 0.803, 0.597 | 0.301, 0.829, 0.967 | +103%                             | 0.041 |
| Average partition asymmetry                  | Whole arborization       | 0.661, 0.653             | 0.561, 0.576, 0.636 | -12.86%                           | 0.041 |
| Maximum centrifugal order                   | Ventral branch          | 13, 30, 43               | 12, 16, 20        | -46.7%                             | 0.043 |
| Hausdorff fractal dimension                  | Ventral branch          | 1.14, 1.24, 1.35         | 1.11, 1.14, 1.2    | -8.07%                             | 0.041 |

Summary statistics of six scalar morphometric and topological parameters that show significant differences for at least one morphological subregion. The triplets in columns three and four represent minimum, median, and maximum values. P values were calculated using Mann–Whitney U test and a cutoff of 5% was used. Summary statistics for all 19 scalar measures and for all 4 subregions are provided in Extended data Table 1-1, Table 1-2, Table 1-3, and Table 1-4. Morphologies showed significant differences for a few scalar measures, with width, total dendritic volume, and maximum centrifugal order showing large changes.
Morphological comparison using scalar measures

Morphologies of DL-Int-1 neurons from newly emerged adult and forager honeybees were compared using 19 widely used metric and topological measures (Uylings and van Pelt, 2002; Scorcioni et al., 2008; Peng et al., 2014). These measures were calculated using a modified version of the BTMORPH software v2.2.1 (Torben-Nielsen 2014; RRID: SCR_003566; code: https://github.com/wachtler-lab/btmorph.v2), Vaa3d v.3.447 (Peng et al., 2014; RRID: SCR_002609), and pyVaa3d (code: https://github.com/ajkswamy/pyVaa3d, v0.4). Mann–Whitney U test was used for calculating the significance of differences between the maturation levels with a cutoff of 5%.

Spatial registration

Preliminary visual comparisons indicated that DL-Int-1 morphologies had differences in translation, rotation and scaling that could have resulted from structural differences between honeybee brains as well as from fixation and dehydration artifacts caused during experimentation. Therefore, we coregistered all DL-Int-1 morphologies to a common frame of reference using the Reg-MaxS-N software (Kumaraswamy et al., 2018; code https://doi.org/10.12751/g-node.fee4e7; RRID:SCR_016257). Reg-MaxS-N estimates and removes differences between morphologies by translation, rotation, and scaling, successively refining estimates of differences at multiple spatial resolutions. In this study we used spatial resolutions of 160, 80, 40, and 20 μm. Morphologies from newly emerged adult and forager bees were coregistered in two steps. First, newly emerged adult and forager morphologies were coregistered separately using Reg-MaxS-N (Kumaraswamy et al., 2018). Then, the two resulting groups of morphologies were brought to the same frame of reference by coregistering the unions of the points of all morphologies in a maturation group using Reg-MaxS (Kumaraswamy et al., 2018).

To control for parameter choice during spatial registration, the procedure above was repeated using multiple parameter sets. Newly emerged adult and forager morphologies were each coregistered separately using three initial references to generate three sets of registered morphologies for each maturation level. Taking all possible combinations of these sets, nine sets of all 12 morphologies were created, which were in turn registered together. All other parameters remained the same for the nine sets (for all parameters, see Extended data Fig. 2–1).

Morphological comparison using spherical shells

The radial distribution of dendritic length was compared between the two maturation levels by dividing the space containing the morphologies into spherical shells of thickness 20 μm, similar to Sholl analysis (Sholl 1953; Uylings and van Pelt, 2002; Langhammer et al., 2010; Garcia-Segura and Perez-Marquez, 2014), which has been shown to be effective in analyzing morphologies (Cuntz et al., 2008; Luebbe et al., 2015; O’Neill et al., 2015). As a natural extension of Sholl analysis, we used the measure dendritic length to quantify changes in dendritic arborization during maturation. For every shell, we calculated PDLshell, which is the percentage of dendritic length contained in the shell. Using two-way ANOVA (Wobbrock et al., 2011), we tested whether, in each shell, PDLshell (1) was significantly different between newly emerged adults and foragers and (2) showed no significant dependence between the effects caused by maturation and registration parameters. The tests used a cutoff level of significance of 5% after Bonferroni correction (Bland and Altman, 1995; McDonald, 2014). This analysis was not applied to MB morphologies because most of them had no branching points and comprised of single stretches of dendrites spanning less than 50 μm.

Morphological comparison using 3D voxels

To compare the morphologies with an even finer spatial granularity, we analyzed non-overlapping 3D voxels of size 20 μm. For each voxel, we calculated PDLvoxel, which is the percentage dendritic length of a morphology contained in the voxel. Note that, because all the voxels had the same volume, changes in PDLvoxel are proportional to changes in average dendritic density. The same criteria as in the previous analysis were used for identifying voxels for which PDLvoxel changed significantly during maturation, independent of registration parameters. To visualize the changes in dendritic density, we calculated for each voxel the normalized change in PDLvoxel as follows:

\[
\Delta PDL^{norm}_{voxel} = PDL_{voxel}^{n} - PDL_{voxel}^{f}\div PDL_{voxel}^{m},
\]

where, for a given voxel, PDL_{voxel}^{n} is the average PDL_{voxel} for forager morphologies across registration parameters and honeybee samples, PDL_{voxel}^{f} is the average PDL_{voxel} for newly emerged adult morphology across registration parameters and honeybee samples; and PDL_{voxel}^{m} is the average PDL_{voxel} across all maturation levels, registration parameters, and honeybee samples.

Morphological comparison using proximal and distal partitions

We divided the space containing the morphologies into proximal and distal partitions based on the remoteness of morphological nodes from their roots, which was quantified using the measure %PL:

\[
%PL = 100 \times \frac{PL_{root}}{PL_{root} + PL_{term}^{max}},
\]

where PL_{root} is the distance along the dendritic tree, also called path-length, between the node and the root; and \( PL_{term}^{max} \) is the maximum of the path-lengths between the node and all terminals in the sub-tree emanating from the node. A voxel was classified to be distal if the median value of %PL, calculated across maturation levels, registration parameters and honeybee samples...
was more than 90. The significance of differences in $PDL_{\text{voxel}}$ between newly emerged adults and foragers were calculated separately for proximal and distal partitions using aligned rank transform (ART) two-way ANOVA (Wobbrock et al., 2011) and the same criteria as in previous analysis.

Analysis of electrophysiology

The physiological response of DL-Int-1 to continuous vibration stimuli applied to the antenna consisted of on-phasic excitation followed by a tonic inhibition and offset rebound (Ai et al., 2009). We defined four time periods for analyzing the electrophysiological activity of DL-Int-1 (Fig. 4b):

- **Spontaneous activity**: 3 s period preceding stimulus onset.
- **On-phasic response**: first 75 ms after stimulus onset.
- **Inhibitory response**: from the end of on-phasic response until stimulus offset.
- **Rebound response**: a 75 ms period after a delay of 25 ms from stimulus offset.

Raw data of electrophysiological recordings were read from Spike2 files using NEO v0.5 (Garcia et al., 2014; RRID:SCR_000634), stored using the NIX format v1.4.5 (Stoewer et al., 2014; RRID:SCR_016196) and analyzed using custom Python scripts. Trials were time-aligned to stimulus onset and time-resolved estimates of average firing rates were generated using adaptive kernel density estimation (Shimazaki and Shinomoto, 2010; Implementation: https://github.com/cooperlab/AdaptiveKDE). The distribution of spike train features such as spike rates and spike times were visualized using the “violinplot” function of the Python package seaborn (Waskom et al., 2018). This function uses kernel density estimation to estimate continuous distributions using Gaussian kernels and Scott’s formula for bandwidth calculation (Härdle et al., 2004, p 73). Mann–Whitney $U$ test was used for calculating the significance of differences in response features with a cutoff of 5%.

We quantified the strength of inhibition relative to spontaneous activity by calculating Relative Inhibition, defined as follows:

$$\text{Relative Inhibition} = \frac{\text{Firing Rate during Inhibitory Response period}}{\text{Firing Rate during Spontaneous Activity period}}$$

Computational environment, code, and data availability

Data preprocessing and analysis were conducted on a desktop computer with an 8-core Intel i7 Processor, 16 GB of RAM running Ubuntu 16.04. The data used for this study are available online on the repository GIN (https://doi.org/10.12751/g-node.e70cb4). Analysis of morphologies and electrophysiological activities was done using custom Python (RRID:SCR_008394) scripts, which are available online (https://github.com/wachtlerlab/GJMorph and https://github.com/wachtlerlab/GJEphys, respectively; Extended Data 1).

Results

Data collection

Sharp electrodes were inserted into DL-Int-1 neurons in the honeybee brain to record electrophysiological activity as well as to inject dye for imaging neuron morphology. Only about 10% of electrode insertions yielded useful data because honeybee brains were not transparent enough for visually targeted electrode insertion and DL-Int-1 neurons were encountered in about one-third of such insertions (Ai et al., 2017). Furthermore, maintaining the electrode within the neuron long enough to obtain sufficient electrophysiological data were difficult especially for newly emerged adult honeybees, as their brains were soft and infirm. Our data of DL-Int-1 neurons from newly emerged adults were therefore limited to six samples with sufficient data for analysis. For the comparative analysis, we chose six forager samples from our database matching the response pattern of the neurons from newly emerged adults.

Morphological adaptations

The four subregions of DL-Int-1 morphology—the WA, MB, DB, and VB (see Materials and Methods, Reconstruction of morphologies)—were compared separately to investigate changes during maturation.

Analysis 1: scalar morphometrics

We first compared the morphologies using whole-cell scalar measures, which detect net overall changes in morphological subregions as they combine data from all dendrites. Table 1 lists the measures that showed significant differences between the morphologies of newly emerged adult and forager DL-Int-1 neurons for at least one subregion (for summary statistics of WA, MB, DB, and VB, see Extended data Table 1-1, Table 1-2, Table 1-3, and Table 1-4, respectively). MB and VB showed significant differences between newly emerged and forager DL-Int-1 neurons for two measures each, whereas DB and WA had one measure each with a significant difference. The changes were neither consistent across morphological subregions nor highly significant ($p$-values between 1 and 5%) and the number of measures showing significant differences were consistent with the number false-positives expected. Hence, at the level of whole-neuron morphological measures, significant changes could not be detected. However, these results did not exclude the possibility of localized changes in dendritic arborization. Therefore we investigated the morphologies at finer spatial scales.

Analysis 2: radial distribution of dendritic length

Before detailed spatial analysis, DL-Int-1 morphologies of newly emerged adults and foragers were coregistered to a common frame of reference (see Materials and Methods, Spatial registration) to establish spatial correspondence. Figure 2 compares the radial distributions of $PDL_{\text{shell}}$ between newly emerged adults and foragers for WA, DB, and VB and highlights those spherical shells for which $PDL_{\text{shell}}$ changed significantly during maturation, independent of registration parameters (see Materials
and Methods, Morphological comparison using spherical shells). WA showed reductions in $PDL_{shell}$ at 110 μm, whereas DB showed an increase at 170 μm during maturation. VB showed reductions up to 130 μm and increases for between 190 and 270 μm during maturation. These comparisons support a consistent redistribution of $PDL_{shell}$ over shells, with reductions in proximal regions and increases in distal regions of the morphology.

To exclude that the observed pattern of proximal reduction and distal increase in dendritic density was a result of a residual scaling difference because of incomplete convergence of the iterative registration process, we repeated the coregistration of the morphologies with different starting conditions, using versions of the newly emerged adult morphologies that were artificially scaled up by 10 or 15%. In both cases, the results were the same as without the scaling (data not shown), confirming that the observed differences in the spatial distributions of dendritic length were not caused by scaling differences.

**Analysis 3: local dendritic length**

To investigate the observed changes in the radial distribution of dendritic length at finer spatial detail, we compared the morphologies at the scale of voxels of size 20 μm using $PDL_{voxel}$ (see Materials and Methods, Morphological comparison using 3D voxels). Figure 3a–c visualizes the magnitude and spatial distribution of normalized change in $PDL_{voxel}$ (see Materials and Methods, Morphological comparison using 3D voxels; Extended data Fig. 3-1 for distributions). Consistent with indications from the previous analysis, some proximal voxels showed reductions in $PDL_{voxel}$, whereas some distal voxels showed increases. To quantify the observed changes more concretely, we divided the space containing the morphologies into proximal and distal partitions (Fig. 3d, Extended data Figure 3-2; see Materials and Methods, Morphological comparison using proximal and distal partitions). Pooling values across voxels in each partition, we used two-way ANOVA to test for significant changes in $PDL_{voxel}$ during maturation independent of registration parameters (Fig. 3e, Extended data Figure 3-3, see Materials and Methods, Morphological comparison using proximal and distal partitions). WA, DB, and VB showed significant reductions of 8.5, 11.3, and 11.9%, respectively, in median $PDL_{voxel}$ for the proximal partition. Whereas WA and DB did not show a significant change in median $PDL_{voxel}$ for the distal partition, VB showed a significant reduction of 18.3%. Thus, there was a region-dependent reduction in the dendritic density of DL-Int-1 with more subregions showing a reduction for proximal parts of the arborization than for distal parts.

**Electrophysiological adaptations**

Comparison of time-resolved firing rate estimates of the responses of DL-Int-1 neurons (see Fig. 5a) indicated increased spontaneous activity and a remarkable increase in firing rate just after stimulus offset in foragers compared with newly emerged adults. These observations were quantified by comparing the firing rates during the four activity periods (Fig. 4b) as well as the spike timing during on-phasic response (Fig. 4c). Figure 5b summarizes the comparison of firing rates for the four periods. Average spontaneous firing rate showed a sig-
significant increase of 39.6%. Average firing rates during on-phasic and inhibitory response periods did not show significant changes, but average firing rate during rebound response nearly doubled, increasing by 94.75%. Thus, DL-Int-1 responses showed stronger spontaneous activity and rebound response.

DL-Int-1 is GABAergic and likely part of a disinhibitory network (Ai et al., 2017; Kumaraswamy et al., 2017). Therefore, increased spontaneous rate and poststimulus rebound in foragers compared with newly emerged adults is expected to result in enhanced strength of the inhibitory signal indicating antennal vibration. To quantify the signal strength of inhibition in DL-Int-1 relative to the level of spontaneous activity, we calculated Relative Inhibition (see Materials and Methods, Analysis of electrophysiology) by plotting the firing rates during the two response periods against each other (Fig. 5c). In general, higher spontaneous spiking was associated with higher spike rates during inhibition in DL-Int-1 neurons, but the difference in firing rates between spontaneous activity and inhibitory response was larger in foragers than in newly emerged adults. Quantitatively, Relative Inhibition was 0.42 ± 0.24 in newly emerged adults, but 0.66 ± 0.18 in foragers (mean ± SE; p-value: 1.34%, Welch’s unequal variance t test). Additionally, least-squares regression in Figure 5c indicated that the difference became larger with firing rate level in foragers (slope of 1.42 for foragers vs 1.06 for newly emerged adult), indicative of faster response dynamics in foragers compared with newly emerged adults.

In addition to changes in activity levels during different periods, comparison of firing rate profiles during on-phasic response (Fig. 5a, inset) indicated a change in the timing of the excitation peak. We investigated this by comparing the first spike latency, first interspike interval (ISI), second ISI, and third ISI during on-phasic response (Fig. 5d). All four spike-timing features showed a systematic reduction during maturation, with average values of first spike latency and first ISI showing significant reductions of 1.76 and 4.01 ms, respectively. Thus, spike timing of DL-Int-1 neurons during on-phasic excitation showed a systematic reduction indicative of response speed-up.

Discussion
In this study, we have compared morphological and physiological properties of an identified vibration-sensitive interneuron, DL-Int-1, between newly emerged adult and forager honeybees. Although comparisons of whole-cell scalar morphometric measures showed no major differences in broad dendritic structure and gross morphological features, detailed spatial analyses revealed region-dependent reduction in dendritic density.
with stronger reductions in proximal parts than in distal parts. This is consistent with findings from previous studies that investigated changes during maturation in the honeybee antennal lobe (Devaud and Masson, 1999) and mushroom body (Farris et al., 2004), which concluded that most of the process of dendritic maturation is completed before emergence, although minor age-dependent and age-independent changes continue for the first few weeks. Such region-dependent changes have also been shown in dendritic arborizations of Kenyon cells in the adult honeybee (Farris et al., 2001) as well as in the paper wasp (Jones et al., 2009). Comparison of the electrophysiological responses of DL-Int-1 to vibration stimuli between newly emerged adult and forager honeybees showed increased spontaneous activity and stronger poststimulus rebound during maturation, whereas the qualitative pattern of response remained unchanged. Similar response enhancements have been reported for odor representation in honeybees (Wang et al., 2005), where odor-dependent activity patterns in antennal lobe glomeruli were similar in newly emerged adult and forager honeybees, with older neurons showing higher spiking rates and more active glomeruli.

**Effect on network connectivity**

Although the DB and VB subregions of DL-Int-1 showed reductions in dendritic density during maturation in the proximal parts, only VB showed reduction in dendritic density in the distal parts. The observed changes thus indicate region-dependent pruning in distal parts. DB and VB arborize in different sets of brain neuropils; the DB arborizes in the mPPL and the DL, whereas the VB arborizes in the DL and the dSEG. DB and VB could therefore be the areas where DL-Int-1 connects to different networks and the observed changes in morphology during maturation might reflect a refinement of the network connectivity.

The AMMC region of the honeybee brain is a center for multisensory integration, especially for waggle dance signals produced by wing beats (Ai and Hagio, 2013; Brockmann and Robinson, 2007). DL-Int-1 arborizes with fine terminals and boutons in the AMMC (Ai et al., 2009), indicating the presence of synaptic inputs and outputs in the region (Petralia et al., 2016). The observed decreases in dendritic density could be associated with changes in the synaptic ultra-structure, similar to that shown in the honeybee mushroom body (Groh et al., 2012; Muenz et al., 2015). However, more studies with synaptic labeling and higher resolution imaging are required to clarify such changes.

**Enhanced inhibition**

The observed changes in electrophysiological activity of DL-Int-1 could reflect an enhancement of features relevant for network processing of high-frequency components of waggle dance vibration signals in the honeybee primary mechanosensory center. Spontaneous firing rates were significantly higher in foragers than in newly emerged adults, while firing rates during inhibitory responses were similar. Thus, the inhibitory response to vibration stimuli was relatively stronger in foragers than in newly emerged adults. Because DL-Int-1 is itself inhibitory and possibly part of a disinhibitory network processing waggle dance air vibration signals (Ai et al., 2017, 2018a; Kumaraswamy et al., 2017), the observed strengthening of relative inhibition could result in more effective disinhibition. Furthermore, the strength of post-inhibitory rebound doubled during maturation. Because inhibition coupled with post-inhibitory rebound has been suggested to play an important role in processing temporal signals in insects (Ai et al., 2018a) and specifically in detecting temporal features (Hedwig 2016; Alluri et al., 2016; Naud et al., 2015; Yamada et al., 2018), our results suggest improved detection of information encoded in the temporal features of waggle dance air vibration signals in forager honeybees.
Effects of morphological changes on physiology

The observed changes in neuron morphology of DL-Int-1 neurons are consistent with a refinement process during maturation that may lead to improved propagation and processing of vibration signals in foragers compared with newly emerged adults. The broad structure of the DL-Int-1 did not show major changes during maturation, with the dendritic branches of the DB and VB extending to similar regions in the honeybee brain. Under this condition and assuming unchanged membrane parameters, reduced dendritic length in proximal regions could indicate lower electrical resistance, and thus, for passive propagation, lower signal attenuation through the DB and VB (Rall and Rinzel, 1973; Ferrante et al., 2013). However, clarification of these effects using multi-compartmental neuron simulations is currently limited by the lack of data about membrane parameters. The potential for such simulations is nonetheless high because morphological reconstructions for several newly emerged adult and forager DL-Int-1 neurons are available.

Changes in response properties: neuron or network?

The response of DL-Int-1 neurons to vibration stimuli applied to the antennae is the combined effect of its inputs and its own intrinsic electrophysiological properties (Ai et al., 2017). In this study, significant increases were seen in spontaneous activity and the strength of inhibition relative to spontaneous activity, as well as in the strength of post-inhibitory rebound. These changes are likely because of maturation of the electrophysiological properties of DL-Int-1 as well as its connected neuronal network. Specifically, the adaptations in the strength of inhibition relative to spontaneous activity could have a stronger dependence on network factors as DL-Int-1 is inhibitory.

Figure 5. Analysis of electrophysiological properties. a, Comparison of the average firing rate profiles of newly emerged adult and forager DL-Int-1 neurons. Smoothed estimates of time-resolved average firing rates were calculated from responses aligned to stimulus onset using adaptive kernel density estimation (Shimazaki and Shinomoto, 2010). Solid lines indicate average firing rates, whereas shaded regions indicate 95% confidence intervals. Inset, Average firing rate during on-phasic response with expanded time scale. b, Comparison of firing rates during four activity periods. The filled areas represent firing rate distributions estimated using kernel density estimation (see Methods, Analysis of electrophysiology). The distributions were normalized to have equal areas. Horizontal white markers indicate mean values of the distributions. The numbers below the distributions are p values calculated using Mann–Whitney U test. P-values <5% are highlighted in green. Firing rates during spontaneous activity and rebound response showed significant increases. c, Comparison of the strength of inhibition relative to spontaneous activity by plotting the firing rates during the two periods against each other. Lines were fit using linear least-squares regression. The dashed line indicates the line of slope 1. d, Comparison of the timing of the first four spikes of the response using first spike latency, first ISI, second ISI, and third ISI. The distributions were estimated using Kernel Density estimation and normalized to have the same area. The numbers under the distributions are p values calculated using Mann–Whitney U test. Horizontal white markers indicate mean values, which are also shown above the distributions with sample numbers in parentheses. Spiking response was faster in foragers compared with newly emerged adults with significant reductions in first spike latency and first ISI.
and is believed to be inhibited in turn as part of a disinhibitory network in the honeybee primary mechanosensory center (Ai et al., 2017; Kumaraswamy et al., 2017). Further, the JO sensory neurons, which transduce anten

al vibrations of the waggle dance and project close to the dendrites of DL-Int-1, show stronger responses to anten

nal deflection in foragers than in newly emerged adults (Tsujuchi et al., 2007). This adaptation in the responses of JO neurons can also play a role in shaping the response properties of DL-Int-1 during maturation. Clarification of the contribution of these sources would be beneficial for further understanding the role of DL-Int-1 in networks that process air vibration jets of the waggle dance in the primary mechanosensory center of the honeybee.

Genetically programmed aging or foraging experience?

In this study we have quantified morphological and physiological changes in DL-Int-1 as honeybees mature from newly emerged adult bees (1–3 d old) to forager bees (>10 d old). There are two major factors that could cause such changes during maturation: genetically programmed aging and foraging experience. Further studies with age

controlled older honeybees with no foraging experience are required to elucidate the effect of these factors on DL-Int-1 maturation.

Linking observed changes to behavior

After successful foraging, honeybees return to their hive and perform the waggle dance, during which they pro

duce patterns of air vibration pulses. Follower bees detect these pulses and gain information about the distance and direction of the advertised food sources (Michelsen et al., 1992; Landgraf, 2013). It has been argued that DL-Int-1 plays a role in the networks encoding information from air vibration jets of the waggle dance into neural signals (Ai et al., 2017; Kumaraswamy et al., 2017). The observed changes in DL-Int-1 suggest that neurons and networks processing waggle dance communication signals under

gro functional and structural refinement as the honey

bee matures that could prepare the bees to better process those important signals as foragers.

References

Ai H (2010) Vibration-processing interneurons in the honeybee brain. Front Syst Neurosci 3:19

Ai H (2013) Sensors and sensory processing for airborne vibrations in silk moths and honeybees. Sensors 13:9344–9363.

Ai H, Hagio H (2013) Morphological analysis of the primary center receiving spatial information transferred by the waggle dance of honeybees. J Comp Neurol 521:2570–2584.

Ai H, Nishino H, Itoh T (2007) Topographic organization of sensory afferents of Johnston’s organ in the honeybee brain. J Comp Neurol 502:1030–1046.

Ai H, Rybak J, Menzel R, Itoh T (2009) Response characteristics of vibration-sensitive interneurons related to Johnston’s organ in the honeybee, Apis Mellifera. J Comp Neurol 515:145–160.

Ai H, Kae K, Kumaraswamy A, Ikeno H, Wachtler T (2017) Interneurons in the honeybee primary auditory center responding to waggle dance-like vibration pulses. J Neurosci 37:10624–10635.

Ai H, Kumaraswamy A, Kohashi T, Ikeno H, Wachtler T (2018a) Inhibitory pathways for processing the temporal structure of sensory signals in the insect brain. Front Psychol 9:1517.

Ai H, Shinya T, Ayaya O, Koji H, Sakashi M, Naoyuki T (2018b) How do the honeybees learn waggle dance? International Union for the Study of Social Insects (IUSSI2018), Guarujá, Brazil, August.

Aliuri RK, Rose GJ, Hanson JL, Leary CJ, Vasquez-Opazo GA, Jakob JA, Wilkerson J (2016) Phasic, suprathreshold excitation and sustained inhibition underlie neuronal selectivity for short-duration sounds. Proc Natl Acad Sci U S A 113:E1927–E1933.

Andrieone M, Timberlake BF, Vallortigara G, Antolini R, Haase A (2017) Morphofunctional experience-dependent plasticity in the honeybee brain. Learn Mem 24:622–629.

Arenas A, Ramirez GP, Balbuena MS, Farina WM (2013) Behavioral and neural plasticity caused by early social experiences: the case of the honeybee. Front Physiol 4:41.

Barth F, Hmrc M, Tautz J (2005) Vibratory and airborne-sound signals in bee communication (Hymenoptera). In: Insect Sounds and Communication, pp 421–436. Boca Raton, FL: CRC.

Bland JM, Altman DG (1995) Multiple significance tests: the Bonferroni method. BMJ 310:170

Brockmann A, Robinson GE (2007) Central projections of sensory systems involved in honey bee dance language communication. Brain, Behav Evol 70:125–136.

Brown SM, Napper RM, Mercer AR (2004) Foraging experience, glomerulus volume, and synapse number: a stereological study of the honey bee antennal lobe. Journal Neurobiol 60:40–50.

Cannon RC, Turner DA, Pyapali GK, Wheel HV (1998) An on-line archive of reconstructed hippocampal neurons. J Neurosci Methods 84:49–54.

Couvillon MJ (2012) The dance legacy of Karl von Frisch. Insectes Sociaux 59:297–306.

Guntz H, Forstner F, Haag J, Borst A (2008) The morphological identity of insect dendrites. PLoS Comput Biol 4:1–7.

Devaud JM, Masson C (1999) Dendritic pattern development of the honeybee antennal lobe neurons: a laser scanning confocal mi

croscopic study. J Neurobiol 39:461–474.

Dreller C, Kirchner WH (1993) Hearing in honeybees: localization of the auditory sense organ. J Comp Physiol 173:275–279.

Durst C, Eichmüller S, Menzel R (1994) Development and experience lead to increased volume of subcompartments of the honeybee mushroom body. Behav Neural Biol 62:259–263.

Fahrbach S, Darrell M, Capaldi EA, Farris SM, Robinson GE (1998) Experience-expectant plasticity in the mushroom bodies of the honeybee. Learn Mem 5:115–123.

Farris SM, Abrams AI, Strausfeld NJ (2004) Development and morphol

ogy of class II Kenyon cells in the mushroom bodies of the honeybee, Apis Mellifera. J Comp Neurol 474:325–339.

Farris SM, Robinson GE, Fahrbach SE (2001) Experience- and age

related outgrowth of intrinsic neurons in the mushroom bodies of the adult worker honeybee. J Neurosci 21:6395–6404.

Ferrante M, Migliore M, Ascoli GA (2013) Functional impact of den

ritic branch-point morphology. J Neurosci 33:2156–65.

Ganeshina O, Schäfer S, Malun D (2000) Proliferation and pro

liferation of synaptic complexes in the mushroom bodies of the honeybee. J Comp Neurol 417:349–365.

Garcia S, Guarino D, Jallet F, Jennings T, Pröpper R, Rautenberg PL, Rodricks CC, Sobolev A, Wachtler T, Yger P, Davison AP (2014) Neo: an object model for handling electrophysiology data in mul
tiple formats. Front Neuroinform 8:10.

Garcia-Segura LM, Perez-Marquez J (2014) A new mathematical function to evaluate neuronal morphology using the Sholl analysis. J Neurosci Methods 226:103–109.

Groh C, Ahrens D, Rössler W (2006) Environment- and age

dependent plasticity of synaptic complexes in the mushroom bodies of honeybee queens. Brain Behav Evol 68:1–14.

Groh C, Lu Z, Meinertzhagen IA, Rössler W (2012) Age-related plasticity in the synaptic ultrastructure of neurons in the mushroom body calyx of the adult honeybee Apis Mellifera. J Comp Neurol 520:3509–3527.

Groh C, Meinertzhagen IA (2010) Brain plasticity in diptera and hymenoptera. Front Biosci 2:268–288.
Härde W, Wervatz A, Müller S, Sperlich S (2004) Nonparametric and semiparametric models. Berlin; Heidelberg: Springer.

Hedwig BG (2016) Sequential filtering processes shape feature detection in crickets: a framework for song pattern recognition. Front. Physiol 7:46.

Hmcir M, Maia-Silva C, McCabe SI, Farina WM (2011) The recruiters of thoracic vibrations during the honey bee waggle dance related to food source profitability. J Exp Biol 214:4055–4064.

Hunt JH, Richard FJ (2013) Intracellular vibroacoustic communication in social insects. Insectes Sociaux 60:403–417.

Ikeno H, Kumaraswamy A, Kai K, Ikeno H, Wachtler T, Ai H (2018) A segmentation scheme for complex neuronal arbors and application to vibration sensitive neurons in the honeybee brain. Front Neuroinform 12:21.

Jones TA, Donlan NA, O'Donnell S (2009) Growth and pruning of mushroom body Kenyon cell dendrites during worker behavioral development in the paper wasp, Polyscia Equestris (Hymenoptera: Vespidae). Neurobiol Learn Mem 92:485–495.

Kirchner WH (1997) Acoustical communication in social insects. In: Orientation and Communication in Arthropods (Lehrer M, ed), pp. 273–300. Basel, Switzerland: Birkhäuser; Springer.

Kirchner WH, Towne WF (1994) The sensory basis of the honeybee’s dance language. Sci Am 270:74–80.

Kiya T, Kunieda T, Kubo T (2007) Increased neural activity of a mushroom body neuron subtype in the brains of forager honeybees. PLoS One 2:e3711.

Kumaraswamy A, Kai K, Ai H, Ikeno H, Wachtler T (2018) Spatial registration of neuron morphologies based on maximization of volume overlap. BMC Bioinformatics 19:143.

Kumaraswamy A, Maksutov A, Kai K, Ai H, Ikeno H, Wachtler T (2017) Network simulations of interneuron circuits in the honeybee primary auditory center. bioRxiv 159533.

Landgraf T, Kellner CJ, Benda J, Wachtler T, Jan G (2014) A negative feedback signal that is triggered by peril curbs honey bee recruitment. Curr Biol 20:310–315.

O’Neill KM, Akum BF, Dhawan ST, Kwon M, Langhammer CG, Firestein BL (2015) Assessing effects on dendritic arborization using novel Sholl analyses. Front Cell Neurosci 9:285.

Peng H, Bria A, Zhou Z, Iannello G, Long F (2014) Extensible visualization and analysis for multidimensional images using Vaa3D. Nat Protoc 9:193–208.

Petralia RS, Wang YX, Mattson MP, Yao PJ (2016) The diversity of spine synapses in animals. NeuroMolecular Med 18:497–539.

Rall W, Rinzel J (1973) Branch input resistance and steady attenuation for input to one branch of a dendritic neuron model. Biophys J 13:648–687.

Scorcioli R, Polavaram S, Ascoli GA (2008) L-Measure: a web-accessible tool for the analysis, comparison and search of digital reconstructions of neuronal morphologies. Nat Protoc 3:866–876.

Seeley TD (1996) The wisdom of the hive: the social physiology of honey bee colonies. Cambridge, MA: Harvard UP.

Shimazaki H, Shimomoto S (2010) Kernel bandwidth optimization in spike rate estimation. J Comp Neurosci 29:171–182.

Sholl DA (1953) Dendritic organization in the neurons of the visual and motor cortices of the cat. J Anat 97:387–406.

Stoewer A, Kellner CJ, Benda J, Wachtler T, Jan G (2014) File format and library for neuroscience data and metadata. Front Neuroinform Abstr 27:00027.

Tobaben-Nielsen B (2014) An efficient and extendable python library to analyze neuronal morphologies. Neuroinformatics 12:619–622.

Tsujii T, Sivan-Loukianova E, Eberl DF, Kitagawa Y, Kadowaki T (2007) Dynamic range compression in the honey bee auditory system toward waggle dance sounds. PLoS One 2:e2324.

Uylings HBM, van Pelt J (2002) Measures for quantifying dendritic arborizations. Network 13:397–414.

von Frisch K (1965) Tanzsprache und ориентierung der bienen. Berlin; Heidelberg: Springer.

van Frisch K (1967) The dance language and orientation of bees. Cambridge, MA: Harvard UP.

Winnington AP, Napper RM, Mercer AR (1996) Structural plasticity of identified glomeruli in the antennal lobes of the adult worker honey bee, Apis mellifera (L.). J Comp Physiol 51:1244–1254.

Waskom M, Botvinnik O, O’Kane D, Hobson P, Ostblom J, Lukauskas S, Gerperline DC, Augspurger T, Halchenko Y, Cole JB, Warnenov Jen, de Ruiter J, Pye C, Hoyer S, Vanderplas J, Villalba S, Kunter G, Quintero E, Bachant P, Martin M, et al., (2018) Mwaskom/seaborn: V0.9.0 (July, 2018). Zenodo.

Winmer AM (1962) Sound production during the waggle dance of the honey bee. Animal Behav10:79–95.

Wittington AP, Napper RM, Mercer AR (1996) Structural plasticity of identified glomeruli in the antennal lobes of the adult worker honey bee, Apis mellifera (L.). J Comp Physiol 51:1244–1254.

Withers GS, Pearlman SE, Robinson GE (1995) Effects of experience and juvenile hormone on the organization of the mushroom bodies of honey bees. J Neurobiol 26:130–144.

Withers GS, Pearlman SE, Robinson GE (1993) Selective neuroanatomical plasticity and division of labour in the honeybee. Nature 364:238–240.

Wobbrock JO, Findlater L, Gergle D, Higgins JJ (2011) The aligned rank transform for nonparametric factorial analyses using only ANOVA procedures. Proceedings of the 2011 Annual Conference on Human Factors in Computing Systems—CHI 11, Vancouver, BC, Canada, May.

Wolsch F, Munch D, Amdam GV (2009) Structural and proteomic effects of queenlessness on the maturation of the honey bee, Apis mellifera (L.). J Comp Physiol 189:165–174.

Wu TY, Zhang J, Sato K, Srinivasan MV (2005) Maturation of odor representation in the honeybee antennal lobe. J Insect Physiol 51:1244–1254.

Xu Y, Zhai Q, Liu S, Yang H (2011) Evaluating accuracy of soma localization and library for neuroscience data and metadata. Front Neuroinform Abstr 27:00027.

Yamada D, Ishimoto H, Li X, Kohashi T, Ishikawa Y, Kamikouchi A (2018) GABAergic local interneurons shape female fruit fly response to mating songs. J Neurosci 38:4329–4347.