The organization of the gravity-sensing system in zebrafish

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Abstract

Motor circuits develop in sequence from those governing fast movements to those governing slow. Here we examine whether upstream sensory circuits are organized by similar principles. Using serial-section electron microscopy in larval zebrafish, we generated a complete map of the gravity-sensing (utricular) system spanning from the inner ear to the brainstem. We find that both sensory tuning and developmental sequence are organizing principles of vestibular topography. Patterned rostrocaudal innervation from hair cells to afferents creates an anatomically inferred directional tuning map in the utricular ganglion, forming segregated pathways for rostral and caudal tilt. Furthermore, the mediolateral axis of the ganglion is linked to both developmental sequence and neuronal temporal dynamics. Early-born pathways carrying phasic information preferentially excite fast escape circuits, whereas later-born pathways carrying tonic signals excite slower postural and oculomotor circuits. These results demonstrate that vestibular circuits are organized by tuning direction and dynamics, aligning them with downstream motor circuits and behaviors.

Introduction

Central neuronal circuits mediate the transformation from sensory inputs to motor outputs. Motor output can be described by both the direction of movement and its temporal dynamics—fast and rapidly fatiguing versus slow and sustained contractions. Fast and slow motor behaviors emerge sequentially, set up by early-born and late-born motor circuits, respectively[1]. In zebrafish, spinal motor neurons differentiate in sequence from those controlling fast movements to those controlling slow[2]. Similarly, premotor neurons in spinal cord and brainstem governing fast escape movements develop earlier than premotor neurons driving slower locomotion[3-5]. Developmental sequence also influences motor circuit connectivity in flies, linking the timeframe of motor circuit formation to temporal dynamics of movement[6,7]. Both fast and slow behaviors can be elicited by a wide range of sensory inputs. It remains unclear whether sensory systems are also organized around principles set up by developmental sequence that align with the speed-dependent motor circuit architecture.

The vestibular system encodes information about head movement in space. Both translational acceleration and orientation with respect to gravity are encoded by hair cells arrayed underneath otoliths in the inner ear. Striolar hair cells carry high-pass signals which are predominantly relayed by irregular-firing vestibular afferents with phasic kinetics, whereas extrastriolar hair cells excite predominantly regular-firing vestibular afferents with tonic or phasic-tonic dynamics[8]. Both types of vestibular afferents relay head movement information into the brainstem, where they excite a variety of central targets. Central vestibular nucleus neurons innervated by vestibular afferents project directly to oculomotor and spinal neurons to drive behaviors[9]. This tight connection from sensory input to motor output suggests that vestibular circuits, like motor and premotor circuits, might be organized around speed-dependent principles. However, it has been difficult to link the patterning of central premotor circuits to hair cell organization because vestibular afferents are not known to be spatially organized by tuning direction or their temporal dynamics.

We examined the architecture of the gravity-sensing system in larval zebrafish. Zebrafish begin to exhibit simple vestibular functions, such as postural control and vestibulo-ocular reflexes, as early as 3 days post fertilization[10-12] (dpf). Over the following days and weeks,

vestibular behaviors improve and refine[13,14]. The small size of the larval zebrafish brain makes it
tractable for reconstruction with serial-section electron microscopy (ssEM) at synaptic
resolution. In these animals, only the utricular otolith is responsible for gravity sensation[10,15], and
accordingly we reconstructed all utricular hair cells, afferents, and four classes of central neurons
receiving utricular inputs.
Directional tuning of utricular hair cells was topographically mapped onto the
rostrocaudal axis of the utricular afferent ganglion. Afferents carrying inferred rostral or caudal
tilt information excited compensatory central vestibulo-ocular reflex (VOR) circuits to stabilize
eye position. Similarly, afferents carrying information about head movement in inferred
ipsilateral or contralateral directions excited distinct elements of Mauthner cell escape circuits.
Along with these directional maps, we identified cellular signatures of vestibular hair cells and
afferents, including cilia length, synapse counts, and myelination, that indicated their
developmental sequence. Inferred developmental sequence was mapped onto the mediolateral
axis of the utricular ganglion and also correlated with central vestibular targets controlling
distinct behaviors. Early-born sensory pathways are preferentially connected to drive early-born
fast motor circuits, whereas later-born pathways govern movements mediated by later-born
slower motor circuits. Collectively, these data revealed a sensorimotor transformation organized
around movement speed, where phasic and tonic vestibular signals are preferentially used to
regulate fast and slow movements respectively. Together, the directional and temporal tuning of
vestibular circuits pattern the entire vestibulomotor transformation.
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Results
Imaging the utricular system at synaptic resolution
Gravity and inertial forces are sensed by hair cells in the inner ear. The otolith, or in mammals
the otoconial matrix, slides relative to the macula during head tilt or translation to deflect hair
cell cilia. Utricular hair cells, which in larval zebrafish serve as the sole gravitational sensors[10,15],
synapse onto the peripheral process of utricular afferents (schematic, Fig. 1a). These afferents,
whose cell bodies reside within the vestibular ganglion, project axons that bifurcate and synapse
in several brainstem nuclei that mediate behaviors like escape, posture, and the vestibulo-ocular
reflex (VOR). We obtained an ultrathin section library of the larval zebrafish at 5.5 days post
fertilization (dpf), which had originally been imaged at $18.8 \times 18.8 \times 60.0$ nm^3 per voxel or 56.4
$\times 56.4 \times 60.0$ nm^3 per voxel depending on the region[16]. We re-imaged the peripheral and central
areas of the right utricular circuit at $4.0 \times 4.0 \times 60.0$ nm^3 / voxel (Fig. 1b), sufficient resolution to
visualize hair cell cilia (Fig. 1c) and vestibular afferent processes (Fig. 1d). The new images
were aligned to the lower-resolution data and used to produce a reconstruction of the gravity-
sensing system, including 91 utricular hair cells, 105 ganglion afferents, and ~45 target neurons
in the ipsilateral vestibular brainstem (Figs. 1e, f, g; Supplemental Video 1).
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Characterization of direction tuning in utricular afferent circuits
We first established the tuning direction of every hair cell in the utricular macula. Hair cells are
maximally depolarized by head tilts in the direction of their kinocilium relative to the cluster of
stereocilia[8]. We measured directional tuning for each hair cell by drawing a vector from the
center of mass of the stereocilia bases to the kinocilium base (schematic, Fig. 2a). Hair cell
direction tuning vectors were displayed relative to soma position to yield a sensory map of the
entire utricular macula (Fig. 2b), with vectors colorized by their inferred directional tuning.

Consistent with prior observations[17,18], the line of polarity reversal (LPR) was found towards the
lateral edge. On either side of the LPR, directional tuning varied smoothly from rostral tilt
sensitive to caudal tilt sensitive. As a result, hair cells located near each other tended to have
more similar tuning than those far apart (Fig. 2c).
97
Organization of the utricular ganglion by directional tuning
We next asked whether this organization of hair cells by directional tuning was reflected in the
organization of the utricular afferent ganglion. Despite a rich literature of physiological
responses in utricular afferents recorded in their axonal processes[19-21], the bone surrounding this
ganglion in adult animals has historically prevented somatic recordings, rendering its sensory
topography unknown. To create a map of utricular ganglion topography, we reconstructed all
processes postsynaptic to the hair cell ribbon synapses (Fig. 2a, bottom), which are by definition
utricular afferents, back to their soma locations in the utricular ganglion. Out of 944 ribbon
synapses, 929 (98.4%) were apposed to afferent processes that could be followed out of the
macula. A total of 105 utricular afferents were reconstructed, with an average of 3.0 ± 1.5
discrete hair cells contacting each afferent, and 3.4 ± 1.4 afferents contacting each hair cell
(mean $\pm$ SD). In a 3D reconstruction of all afferents, there was a visible rostrocaudal gradient to
their organization (Fig. 2d; note that this reconstruction is tilted slightly relative to a true
horizontal view so that the afferent ganglion is visible beneath the central axon projections).
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Afferents whose somata were rostrally located within the ganglion were innervated by hair cells
in the rostral utricular macula, whereas afferents in the caudal portion of the ganglion were
innervated by hair cells in the caudal macula. Consistent with previous reports[22], each afferent
received input from hair cells exclusively on one side or the other of the LPR. To infer
directional tuning of each utricular afferent, we weighted its convergent hair cell vectors by their
number of ribbon synapses. Here and throughout the manuscript, this anatomically-inferred
directional tuning will be referred to more simply as directional tuning, but we note that actual
directional tuning might differ based on variations in synaptic weight, process morphology, and
other biological variables that cannot be assessed with EM.
122
The resulting map revealed a sensory tuning topography in the utricular ganglion, with rostrally
located afferents encoding rostral head tilts and caudally located afferents encoding caudal head
tilts (Fig. 2d, e; Supplemental Video 2). Quantifying hair cell to afferent connectivity revealed a
strong correlation between the rostrocaudal position of each afferent soma and the hair cells that
innervate it (Fig. 2f). However, unlike in the hair cell macula (Fig. 2b), afferents innervating hair
cells lateral to the LPR are intermingled with those innervating hair cells medial to the LPR (Fig.
2g, note red arrows), both in this horizontal view and in sagittal and coronal views
(Supplemental Fig. 2a, b). Nonetheless, these afferents with inferred contralateral tilt tuning
formed a segregated axon bundle in the brainstem (Fig. 2d). Afferent somata located near each
other typically had more similar tuning than afferents that were further apart (Fig. 2h), although
the strength of the relationship is less pronounced than in the hair cells themselves (cf. Fig. 2c).
134
Central organization of directional tuning
We next asked how directional tuning was organized centrally in three utricular afferent target
populations: the Mauthner escape circuit, the VOR circuit, and the vestibulospinal (VS) postural
circuit. Our high-resolution reimaged territory in the brainstem allowed us to identify the

postsynaptic targets of utricular afferents. These central brainstem targets were then
reconstructed and characterized based on soma location and axon projection. Although most
central vestibular neurons project axons outside the vestibular brainstem, into lower resolution
regions that were not reimaged, they could be reconstructed over long distances if their axons
were myelinated[16]. Within these criteria, we characterized utricular input to the ipsilateral
Mauthner cell, 4 commissural escape neurons projecting to the contralateral Mauthner cell, 23
VOR neurons, and 19 VS neurons. We note that additional commissural, VOR, and VS neurons
presumably exist with insufficiently myelinated axons for adequate reconstruction, and therefore
the set sampled here is likely biased towards earlier-born neurons. Additional neurons of less
certain identity will be described in a future publication.
149
The Mauthner cells are a specialized pair of reticulospinal neurons that trigger fast escape
movements in response to multiple types of sensory input. Mauthner cells develop by 8 hours
after fertilization, excite primary (fast, early-born) spinal motor neurons, and drive fast escape
movements by 1 dpf[23-25]. We identified and reconstructed utricular inputs both onto the
ipsilateral Mauthner cell and onto brainstem neurons whose axons cross the midline and appear
to synapse onto the contralateral Mauthner cell. In total, 18 utricular afferents contacted the
Mauthner cell lateral dendrite (example, Fig. 3a) with a total of 52 synaptic contacts. All
utricular afferent synapses were clustered together tightly on ventral sub-branches of the
Mauthner cell lateral dendrite (Fig. 3b), as reported[26]. In addition, seven utricular afferents
synapsed on a commissurally projecting neuron population that contacted the contralateral
Mauthner (Fig. 3c), which we term "commissural escape" neurons.
161
Interestingly, no utricular afferents diverged to contact both the ipsilateral Mauthner cell and this
commissural escape population. Utricular afferents presynaptic to the Mauthner cell were
innervated by hair cells in the medial portion of the macula, whereas afferents that excited the
commissural-projecting neurons were innervated by hair cells in the lateral portion of the
macula, on the far side of the LPR. The inferred tuning vectors of all afferents contacting the
Mauthner cell were averaged to yield an inferred tuning to tilt in the ipsilateral and rostral
direction (Fig. 3c, d). Tuning vectors of commissural escape neurons pointed in the opposite
direction, contralaterally and caudally (Fig. 3c, d; average difference between Mauthner and
commissural neuron inferred tuning, 196°). Therefore, these commissural escape neurons are
presumably similarly tuned to inertial movements as the contralateral Mauthner cell that they
contact.
173
Vestibular stimuli can elicit locomotion[27,28], but escape behaviors have primarily been evoked
with auditory or mixed auditory-vestibular stimuli that do not allow isolation of the vestibular
component[29,30]. We reasoned that the utricular afferents exciting the Mauthner cell are tuned to
report ipsilateral head tilts and therefore contralateral translational movements, with the
predicted consequence that a translational movement to the left should trigger escape bends to
the left (Fig. 3e). This type of circuit would be useful to detect predator movement towards the
zebrafish. While the water flow of a predator bow wave can be detected by lateral line circuits[31],
there is little known about whether the head deflection itself can also elicit escapes. To test this
prediction, we delivered a large amplitude translational stimulus optimized for speed (~10 ms, >
1 g) to freely moving zebrafish larvae while recording behavior at 508 frames/s. This stimulus
evoked a classic short-latency escape response in 35% of trials (Fig. 3f; 28 escapes in 81 trials

from 27 animals; escapes defined as C-bends within 10 ms of peak acceleration[29]). Animals
escaped in the direction predicted by the circuit in Fig. 3e on 64% of trials (18/28). However, due
to sled limitations, this stimulus involved bidirectional movement to maximize the accelerative
force. We therefore repeated this experiment with a second stimulus that was slightly slower but
optimized to be unidirectional (~20 ms, 0.8 *g*). This stimulus elicited a similar frequency of
escapes (28%: 56 escapes in 202 trials from 105 animals). Notably, successful escapes were
strongly directionally biased: animals turned in the direction of peak acceleration on 95% (53/56)
of escapes (Fig. 3g, shaded regions). Turns in the "wrong" direction occurred on trials where
animals were accelerated rostrally or caudally (Fig. 3g). To test for utricular dependence, both
stimuli were presented to the utricle-deficient *rock solo* fish line, an otogelin mutant[10,32]. Utricle-
deficient animals escaped about half as often as their sibling controls (Fig. 3h). Because
experiments were carried out under infrared light, the remaining escapes were likely triggered by
the lateral line system[31]. Based on these anatomical and behavioral results, we conclude that this
utricular-activated escape circuit allows for computation of the direction of head deflections,
such as occur during predator approach.
200
Next, we evaluated directional tuning in three vestibular nuclei. VOR neurons of the superior
vestibular nuclei (SVN) and tangential nuclei collectively govern vertical and torsional eye
movements. We reconstructed 12 neurons in the SVN, which inhibits the ipsilateral trochlear and
oculomotor nuclei[9], and 11 neurons of the tangential nucleus[12], which excites contralateral
trochlear and oculomotor neurons (Fig. 4a, b). Of these VOR neurons, 22/23 had inferred tuning
in the ipsilateral rostral or caudal directions (insets, Fig. 4a, b), consistent with their well-
described roles in the VOR[9]. This inferred tuning also aligns with the angles of the anterior and
posterior semicircular canals, which contribute to rotational VOR behaviors later in
development[9]. Notably, the SVN and tangential neurons tuned for rostral vs caudal tilt also
projected axons with largely distinct trajectories. SVN axons with inferred rostral tilt encoding
traveled more laterally than those encoding caudal tilt (boxed inset, Fig. 4a), and presumably
inhibit the motor neurons that drive downward eye rotation via the inferior rectus and superior
oblique muscles[33]. Tangential axons with inferred rostral tilt encoding traveled ventrally and are
presumed to activate the activate the eyes-up pathway through motor neurons that control the
superior rectus and inferior oblique, whereas those with inferred caudal tilt encoding traveled
dorsally where they likely activate the eyes-down pathway through motor neurons of the inferior
rectus and superior oblique[34-36] (Fig. 4b).
218
Similarly, we characterized the tuning of afferent inputs to the VS population, which is involved
in postural control (Fig. 4c)[37]. We extended our previously reported connectivity from
myelinated utricular afferents[38] to include the unmyelinated afferents. The 19 VS neurons
received input from a total of 61 afferents. In contrast to VOR nuclei, VS neurons primarily
received input from more rostral tilt sensitive afferents (Fig. 4c). VS neurons were typically
contacted by a greater number of distinct utricular afferents than VOR pathway neurons were
(medians: 8 afferents per VS neuron, 25th-75th %ile, 3.5 – 15.5; 4 afferents per VOR neuron, 25th
-75th %ile, 3-7; Wilcoxon-Mann-Whitney between the number of distinct utricular afferents
contacting VS and VOR neurons, $p = 0.012$; U, U' = 85, 245). Thus VS neurons are contacted
by a large number of predominantly rostral tilt sensitive utricular afferents.
229

Little is known about subcellular organization of vestibular afferent input onto central neurons.
The concentration of utricular inputs onto a portion of dendrite in the Mauthner cell (Fig. 3b) led
us to use the high resolution of EM to examine whether utricular inputs were similarly
concentrated in VOR and VS neurons. Dendrograms of utricular afferent input to VOR and VS
neurons revealed that most utricular input is not evenly distributed across the dendritic arbor but
instead arrives on a small subset of branches (examples, Fig. 4d). To quantify the concentration
of synapses, we carried out Monte Carlo simulations of synaptic distribution with either
randomly distributed synapses or locations weighted by distance from afferents (see Methods).
We found that utricular afferent synapses were located more closely to each other than expected
by random chance, likely related to the limited spatial range over which utricular afferents
interact with vestibular dendrites (Fig. 4e, Supplemental Fig. 5). This result indicates that
branch-specific computation may occur in vestibular nucleus neurons, perhaps in conjunction
with localized cerebellar input[39].
243
244 These analyses revealed a rostrocaudal map of direction tuning in the utricular afferent ganglion,
245 as well as patterns of direction tuning in central brainstem targets of the utricular afferents. We
246 next considered the organization of temporal kinetics arising from distinct types of hair cells.
247
Temporal dynamics and developmental sequence of utricular hair cells
In addition to their directional tuning, hair cells are characterized as striolar or extrastriolar,
based on morphological differences in soma shape and ciliary lengths[40-43]. Striolar hair cells
typically drive phasic, adapting, irregular-firing afferents with high-pass sensitivity, whereas
extrastriolar hair cells drive tonic or tonic-phasic regular-firing afferents that are less dependent
on stimulus frequency[8,19,42,44-46]. Fish and frogs do not express the classical Type I striolar cell
shape seen in amniotes, but striolar hair cells can still be recognized based on the length of the
kinocilium relative to the tallest stereocilium[43,47]. In the adult zebrafish utricle, striolar hair cells
are estimated to have kinocilia and tallest stereocilium lengths around 5 µm, whereas
extrastriolar hair cells have a kinocilium estimated at 6-8 µm and tallest stereocilium 2-3 µm[17].
We reconstructed the kinocilium and tallest stereocilium of each hair cell (Fig. 5a) and plotted
the relationship between their lengths. In one group of hair cells, both kinocilium and
stereocilium were > 4.8 µm and the kinocilium to stereocilium length (K/S) ratio ranged from 1.1
– 1.7; these were identified as striolar (Fig. 5b). In a second group, the kinocilium was > 5 µm
but the tallest stereocilium was < 5 µm, yielding K/S ratios from 1.75 – 3.3, which we identified
as extrastriolar. These ciliary lengths are close to those in the adult zebrafish, suggesting that
these are mostly mature hair cells. Finally, at the tail end of the ciliary length distributions, we
identified hair cells in which the kinocilium was < 5 µm; we classified these as immature, though
they presumably exist on a continuum with established hair cells (Fig. 5b). Some of these
immature hair cells were also characterized by less cytoplasm, with fewer mitochondria and
vesicles, consistent with this identification.
269
We marked the position of each cell by its kinocilium and found that striolar hair cells straddled
the LPR while extrastriolar hair cells populated the rest of the macula, as expected[43,48] (Fig. 5c).
Utricular hair cells differentiate in a temporal sequence, with roughly radial development from
the center or striola to the periphery[49-53]. We wondered whether hair cell ribbon counts, which
generally increase during development[54], could establish additional signatures of this ongoing
sequence of differentiation. Counts of the total number of ribbon synapses (example, Fig. 2c)

showed that hair cells located centrally in the macula had the highest number of ribbons per cell,
with fewer ribbons found in peripheral hair cells (Fig. 5c). Accordingly, striolar hair cells overall
had the largest number of ribbons, followed by extrastriolar and immature hair cells (Fig. 5d).
Even within the extrastriolar population, hair cells with more ribbons were located more
centrally, and hair cells with fewer more peripherally, consistent with a radial growth pattern.
Thus, both ciliary lengths and ribbon synapse number are consistent with a radial pattern of
growth in the macula, providing proxies for developmental sequence.
283
Organization of the utricular ganglion by developmental sequence
We next asked whether similar proxies for developmental sequence could be identified in the
utricular afferent ganglion neurons. Myelination is responsive to neuronal activity[55,56], and
evidence suggests that it initiates first in early-born neurons[57,58]. Of utricular afferents, 16 of 105
(15.2%) were myelinated throughout most of their central and peripheral processes; the
remaining afferents were mostly or entirely unmyelinated (examples, Fig. 1d), similar to early
developmental stages in other animals[59-61]. Because the entire vestibular nerve is myelinated by
adulthood[22,59], the myelinated afferents we observed are likely to be early-born, instead of a
specialized category of afferents. Consistent with this idea, we found that afferent somata that are
myelinated (Fig. 5e) and that receive the highest numbers of ribbon contacts from hair cells (Fig.
5f) occupy the most lateral edge of the nascent ganglion, the site of the earliest-born vestibular
afferents[62-64]. Later-born afferents are added more medially, forming a half-shell around the
earliest born somata[62-64]. These observations are supported by quantification of mediolateral
position (Fig. 5g) and the ribbon synapse count (Fig. 5h) in myelinated versus unmyelinated
afferents. Thus, afferent myelination and soma position are proxies for developmental sequence.
299
Do early born hair cells preferentially connect with early born afferents? We found that
myelinated afferents received over half of their input from striolar hair cells, and little input from
immature hair cells (Fig. 5i). This was true whether quantified by the number of distinct hair
cells providing input or the total number of synaptic ribbon connections (Supp. Fig. 3d, e). In
contrast, unmyelinated afferents predominantly received input from extrastriolar hair cells, and
had a higher proportion of input from immature hair cells (Fig. 5i). At the level of individual
neurons, afferents with laterally located somata tended to receive input from striolar hair cells,
whereas medially located somata were the most likely to receive immature hair cell input (Fig.
5j).
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We conclude from these results that early-born afferents, marked by early myelination,
preferentially receive input from early-born, predominantly striolar hair cells, and occupy a
lateral position in the ganglion. Later-born afferents contact later-born, predominantly
extrastriolar hair cells, and occupy progressively more medial positions, forming a half-shell
around the early-born afferents. At the same time, directional tuning is preserved in the
rostrocaudal axis, with rostrally located afferents contacting rostrally located hair cells and vice
versa (Fig. 2). Thus, the utricular ganglion is organized by directional tuning in the rostrocaudal
axis and developmental sequence in the mediolateral axis. Moreover, because inferred afferent
temporal dynamics are aligned with developmental order, the ganglion mediolateral axis is also
organized from phasic, striolar-dominated afferents at the lateral edge to tonic, extrastriolar-
dominated afferents more medially. Therefore, both the spatial and temporal tuning of vestibular

signals are topographically organized in the utricular ganglion, though more loosely than at the
macula.
323
Organization of central circuits by developmental sequence
Building on our observation that developmental sequence is an organizing principle of the
utricular afferent ganglion and its connectivity with peripheral hair cells, we asked whether
developmental sequence might also be linked to central connectivity. The downstream targets of
vestibular afferents become functional at different times in development: Mauthner cells are the
first to form connections and drive escape behavior, whereas VOR and VS neurons driving eye
and postural movements are born later[37,65]. Does the developmental sequence of utricular
afferents predict their patterns of connectivity to brainstem targets? We mapped the hair cells
contributing input to afferents that drove escape, VOR, or VS neurons (Fig. 6a). Central escape
neurons (the Mauthner cell and the commissural escape neurons) received most of their input
from afferents contacting striolar hair cells. In contrast, afferents exciting VOR and VS neurons
received input from a much broader territory of the hair cell macula (Fig. 6a). We quantified hair
cell input to these pathways by weighting each central synaptic contact by its afferent's fraction
of input from striolar, extrastriolar, or immature hair cells. Around two-thirds of utricular inputs
to the escape circuits arose from striolar pathways, whereas VOR and VS neurons received a
much larger portion of their inputs from extrastriolar regions (Fig. 6b). Similarly, the Mauthner
cell and commissural escape neurons also received a higher proportion of their utricular inputs
from myelinated afferents, in comparison to VS and VOR populations (Fig. 6a, c). Consistent
with the interpretation that myelinated afferents are early-born and more mature, they also
diverge to more postsynaptic target nuclei (Myelinated afferents diverging to Mauthner, SVN,
tangential, and VS neurons: 9/16; to three targets, 3/16; to two targets, 1/16. Unmyelinated
afferents diverging to all four targets, 0/90; to three targets, 9/90; to two targets, 25/90.
Supplemental Table 2.) Therefore, early-born hair cells preferentially signal via early-born
afferents to early-born brainstem populations, which in turn drive early-born spinal motor
circuits for escape. Similarly, later-born hair cells excite later-born circuit elements for posture
and oculomotor control.
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Developmental sequence may play a significant role not just across but also within neuronal
populations. We found that three VS neurons whose axons travel more medially in the brainstem
before joining the rest (VS_{med}) receive more input from afferents carrying immature hair cell
information, whereas four VS neurons whose axons travel more ventrally (VS_{vent}) receive more
input from afferents carrying striolar information, as compared to other VS neurons (Fig. 6d).
Based on these inputs, VS subpopulations with distinct axonal trajectories are predicted to
exhibit different temporal dynamics, with more phasic information carried by the VS_{vent} and
more tonic by VS_{med}. Though the postsynaptic targets of these subpopulations of VS neurons are
not known, this result may help identify circuits underlying later refinement of postural control[14].
360
Collectively, our anatomical analyses demonstrate that directional tuning and developmental
sequence pattern the entire vestibulomotor transformation (Fig. 7). The rostrocaudal axis of hair
cell organization is largely preserved in the afferent ganglion, leading to distinct pathways that
drive responses to pitch. Afferents encoding contralateral tilt are intermingled at the level of the
ganglion but form spatially segregated pathways that underlie head direction computation in the
escape circuit (Fig. 7a). At the same time, early-born hair cells contact early-born utricular

afferents, which in turn preferentially drive escape behaviors mediated by fast, early-developing
motor circuits. Later-born sensory pathways support postural and oculomotor behaviors
mediated by a mixture of fast and slow muscles (Fig. 7b). These results also demonstrate that
striolar signals, which are carried mostly by irregular-firing afferents with predicted high-pass,
phasic encoding properties, establish connections with circuits driving rapid-onset reflexes via
fast motor pools.
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375 **Discussion**
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Here we show that the utricular afferent ganglion, which carries gravity sensation into the brain,
is organized along two axes: a rostrocaudal axis for directional tuning, and a mediolateral axis
for development. Afferents with different directional tuning excite distinct brainstem
populations. Further, we demonstrate that early-born afferents preferentially receive information
from early-born and striolar hair cells, yielding a gradient from phasic to tonic signals in the
ganglion. This developmental organization aligns vestibular signals with downstream motor
circuits. Brainstem neurons governing the fastest motor circuits, which underlie early-developing
escape behaviors, are excited mostly by utricular afferents with inferred phasic firing and early
development. Oculomotor and postural circuits, which drive slower motor neurons, are excited
preferentially by afferents with inferred tonic firing and late development. Collectively, these
results demonstrate that the vestibular circuit is organized by both directional tuning and
temporal dynamics to mediate transformation into motor outputs.
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390 *Developmental sequence as an organizing principle in sensorimotor transformations*
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We demonstrate here that the utricular afferent ganglion is patterned both by directional
sensitivity in the rostrocaudal axis, and by developmental sequence, correlating to temporal
dynamics, in the mediolateral axis. As far as we are aware, this is the first demonstration of any
topography in the vestibular afferent ganglia. A similar mediolateral gradient has been described
for the nearby lateral line afferent ganglion, where early-born afferents are positioned more
laterally than later-developing afferents[66]. Early-born lateral line afferents also exhibit larger
soma size, lower excitability, and more dorsal central projections than later-born afferents[67].
Thus, they are well placed to mediate coarser, large-amplitude stimuli, similar to our observation
that early-born utricular afferents carry striolar information with inferred phasic responses. The
mechanisms shaping topographic patterning in these two sensory systems may have been
templates for tonotopic organization in the auditory cochlear afferents that evolved later[68].
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The link between developmental sequence and motor control has been shown most robustly in
zebrafish[3-5,69] and fruit flies[6,7], suggesting that sequential developmental from fast to slow is an
ancient principle of motor control. Building on these results, several classes of spinal neurons in
mice have also been shown to differentiate by subtype according to their birth order[70,71].
However, whether these subtypes are tied to different speeds of movement remains to be
explored.
410
Our work demonstrates that developmental sequence also links sensory inputs to motor outputs.
The vestibular system is tightly coupled to motor control, and therefore it seems plausible that its

organization relies on related principles. Our data show that the Mauthner cell preferentially
receives utricular input from striolar-driven afferents (65% of utricular input). Notably, these
afferents are the earliest-born vestibular pathways; similarly, the Mauthner cell drives the escape
reflex as early as 2 dpf via the earliest-born spinal motor and premotor neurons[25]. Thus, the
entire escape reflex arc runs from from early-developing hair cells, to early-developing afferents,
to early-developing hindbrain reticulospinal neurons, to early-developing spinal elements and
muscles. Both oculomotor and postural behaviors appear around 3-4 dpf in zebrafish[12,15,72-74],
with continued maturation at later stages[14]. Correspondingly, the VS and VOR neurons develop
after the Mauthner cell in zebrafish[37] (D. Schoppik, personal communication), and these
populations appear to develop at roughly the same time in amniotes[75]. In mouse, VS neurons
preferentially synapse with slow motor neurons[76], consistent with our observation that they
receive more extrastriolar and later-developing input than escape circuits do. It is plausible that
developmental sequence may be significant in circuit assembly beyond the vestibulomotor
pathways examined here.
427
We find that VS neurons can be subdivided based on axon trajectory, with differing proportions
of striolar, extrastriolar, and immature input across groups. The fact that the VS_{med} population,
whose axons plunge directly towards the midline, get a higher proportion of utricular input from
immature hair cells suggests that they might be later-born and participate in later-onset postural
functions. Zebrafish refine their postural control over the pitch axis from 4 to 15 dpf by
increasing use of fins and improved bout timing mediated in part by VS neurons[13,14,37]. We
speculate that these VS_{med} neurons may contribute to this refinement by preferentially connecting
with fin motor neurons, which are located near the midline[77]. In mammals, the slower-onset
portion of postural reflexes is mediated in part by a subset of lateral vestibular nucleus neurons
(LVN_C) which collateralize to contact pontine reticulospinal neurons[78]. These similarities
suggest that a VS population encoding tonic head movements could target multiple downstream
targets for fine motor control. In contrast, the higher proportion of striolar inputs to VS_{vent}
suggests a function in rapid postural control. Together, our findings suggest a relationship within
the VS population between the developmental time course of sensory innervation, axon
trajectory, and the speed of behavioral responses.
443
444　Although our reimaging did not extend to the oculomotor and trochlear nuclei, the dorsal
445　trajectory of caudal-tilt sensitive (nose-up) tangential neurons corresponds well with the dorsal
446　position of inferior rectus and superior oblique (eyes-down) motor neurons in these areas[36].
447　Based on the earlier development of eyes-down motor neurons[35], we hypothesize that tangential
448　and SVN pathways contacting these motor neurons may be earlier born as well.
449
Mauthner cell computations
The directional tuning of afferent inputs to the Mauthner cell and the commissural escape
neurons was opposite in direction. Thus each Mauthner cell receives two forms of utricular
inputs tuned to ipsilateral tilt: monosynaptic innervation from ipsilateral vestibular afferents and
disynaptic innervation from the contralateral side. The identity of these commissural escape
neurons receiving utricular input is not known. Based on their anatomy, they are not similar to
excitatory spiral fiber neurons, whose axons target the Mauthner axon cap[79], nor to inhibitory
commissural neurons underlying left-right choice in the auditory system, whose axons target the
Mauthner cell body[80]. Instead, these commissural utricular neurons appear to target the Mauthner

ventral dendrite (Fig. 3c). Vestibular commissural neurons in goldfish have previously been
interpreted as inhibitory, because their activation diminishes the amplitude of the antidromically
triggered spike[81,82]. We speculate that these zebrafish commissural escape neurons are instead
excitatory, because they carry signal that is similar to the predicted monosynaptic vestibular
afferent inputs to that Mauthner cell. Alternatively, these commissural neurons might provide an
inhibitory signal that sharpens lateralization, as described in the auditory and lateral line
systems[80]. Further high-resolution imaging or physiology could resolve this question.
466
Additionally, because utricular inputs to the Mauthner cell and these commissural neurons are
predominantly striolar, they are expected to be phasic and high-pass, with exquisite spike
timing[44,83]. Irregular-firing afferents carrying striolar signals are thought to be kinetically well
suited to signal translational movements, which are typically brief, in contrast to the lower-
frequency movements generated by tilt[84]. In keeping with this prediction, we identified a
vestibular-dependent escape behavior triggered by rapid translation. This behavior may be
important for survival: a predator approaching from the right would cause a fast translational
head displacement to the left. This would activate both the ipsilateral tilt tuned afferents on the
right side and the contralateral tilt tuned afferents on the left side of the animal (Fig. 3e). Via the
direct and indirect pathways described here, these would summate to activate the right side
Mauthner cell, which triggers contraction of the left side of the body, leading to a correctly
directed escape away from the predator.
479
Central vestibular tuning
We find that VOR circuit neurons in the tangential and SVN all receive inputs tuned to ipsilateral
tilt, with subpopulations aligned to either rostral or caudal tilt directions for control of vertical
and torsional eye movements. The shared ipsilateral sensitivity explains why these
subpopulations were not differentiated in zebrafish imaging studies of vestibular tuning in the
roll axis[85,86]. Our anatomical data predict that VS neurons exhibit greater sensitivity to rostral tilt,
consistent with our physiological results, in which VS neurons are preferentially sensitive to
caudal translation[38]. Some VS neurons are contacted by divergently tuned afferents in the
rostrocaudal axis and are predicted to have correspondingly more complex tuning responses, as
seen in mammals[87,88]. These results align well with our observation that complex central tuning
in VS neurons is constructed by feedforward excitation from afferents with differing tuning[38].
We also discovered that vestibular afferents innervating the most lateral portion of the macula
(contralateral tilt sensitive) form a separate axon bundle in the hindbrain (Fig. 2d and
Supplemental Video 3). This result implies differential targeting of this pathway, consistent with
the commissural utricular neurons in the escape circuit. The contralateral tilt-sensitive pathway is
also thought to drive local feedforward inhibition to amplify target tuning by a push-pull
mechanism[89].
497
Our observation that VOR and VS neurons receive similar mixtures of striolar and extrastriolar
input are consistent with physiological analyses suggesting that both populations receive similar
amounts of synaptic input from irregular and regular firing vestibular afferents[83,90,91]. In those
analyses, irregular-firing (phasic) afferent inputs appeared to dominate, whereas in our
anatomical analyses, striolar-driven afferents make up less than half the overall utricular input.
We suggest that synaptic contacts onto VOR and VS neurons from irregular afferents are
stronger in amplitude than those from regular-firing afferents, consistent with their

morphologically larger synaptic contacts[92]. However, irregular-firing inputs preferentially
encode high-frequency head movements, which are less common[93]. Therefore, their dominance
in nerve stimulation experiments may reflect synaptic weights but not their overall influence on
typical activity. Indeed, loss of striolar hair cells has little effect on oculomotor and basic
postural behaviors[94], consistent with our observation that those pathways draw mostly on
extrastriolar inputs. There are also likely to be significant variations across vertebrates in the
balance between phasic and tonic vestibular signals, based on characteristics of movement[95].
512
Our analyses could not test whether developmental sequence *governs* connectivity, as opposed to
the possibility that a genetically distinct class of early-born afferents preferentially connects to
escape circuits. Future experiments manipulating circuit formation may shed light on this
question. Our data also necessarily represent a single snapshot in time. Is the observed circuit,
aligned with developmental sequence, likely to be maintained in adulthood? We suggest that it
likely is, on the basis of the functional correspondence between early-born hair cells and early-
born motor neurons which both display high-pass dynamics, while late-born hair cells and motor
neurons display slower dynamics[96-98]. However, the spatial organization of the ganglion may
become distorted by continued development; further experiments will be needed to address this
question. It will also be important to evaluate how later-born hair cells in the striolar region
integrate into the circuit. If they drive activity in later-born central circuits, their afferents may
change central temporal dynamics. Alternatively, if they integrate with early-born circuits as
other striolar hair cells do, they may serve to augment existing signals.
526
527
Materials and Methods
Ultrathin (60 nm thick) serial sections from a 5.5 dpf larval zebrafish were a generous loan from
J. Lichtman and F. Engert. Using the published $18.8 \times 18.8 \times 60$ nm^3 per voxel and $56.4 \times 56.4 \times$
60 nm^3 per voxel reference map and reconstructions (Hildebrand et al., 2017), we re-imaged the
right side of the fish, covering the utricular hair cells, utricular afferents (identified by their
peripheral processes reaching the utricular macula), and a rostrocaudal extent of the brainstem
that covered several major vestibular nuclei at $4.0 \times 4.0 \times 60$ nm^3 per voxel. The volume covered
~100 µm in the rostrocaudal axis, 150 µm in the mediolateral axis, and 100 µm in the
dorsoventral axis, in an irregular shape designed to capture the afferent peripheral and central
processes (Fig. 1b). Imaging was carried out at the Washington University Center for Cellular
Imaging on a Zeiss Merlin 540 FE-SEM with a solid-state backscatter detector. WaferMapper
software[99] was used to control an ATLAS scan engine for automated focus and acquisition[100].
The resulting images were aligned onto the original $56.4 \times 56.4 \times 60$ nm^3 per voxel dataset using
affine and elastic transformations in FIJI's TrakEM2 plugin[101,102], with custom support from
UniDesign Solutions.
543
The entire image volume was hosted in a CATMAID instance[103,104]. Vestibular circuit neurons
and hair cells were reconstructed as skeletons, *i.e.* node points without volume fills. All utricular
afferents were identified by stepping section by section through the entire anterior macula twice
and marking every hair cell ribbon synapse. Ribbon synapses were identified by the
characteristic large dark ribbon structure surrounded by vesicles (Fig. 2a). Every utricular
afferent was followed as far as possible, in most cases to the corresponding soma in the

vestibular ganglion. Only 1.6% (15/944) of processes adjacent to ribbons could not be followed
to a soma due to the quality or ambiguity of the images.
552
Hair cell kinocilia and the tallest stereocilia were traced from the apical surface of each hair cell
to their distal tips. The kinocilium was recognizable based on its distinctive structure (see Fig.
2a). Ciliary length was calculated as the sum of the Euclidean point-to-point distances. Positions
of all cilia were plotted at the epithelial plane and a three-dimensional tuning vector for each hair
cell was derived from the center of mass of all stereocilia to the kinocilium. Hair cell vector
lengths were typically short in the dorsal-ventral axis relative to their extent in the other two axes
(around one-third the normalized vector magnitude of the other two axes; Supp. Fig. 1b),
consistent with the mostly horizontal orientation of the utricular macula. Therefore, for the
purposes of analysis we focused exclusively on their projection in the horizontal plane. Ganglion
soma position was quantified in three dimensions for all analyses. During fixation, differential
shrinkage caused a small tissue separation that is visible as a gap in the horizontal projection of
the utricular ganglion reconstruction (Fig. 2g, upper right), but there was no loss of tissue.
Tuning similarity was calculated as the cosine of the difference between the tuning directions of
each pair of hair cells or afferents. Hair cell distance was determined by the 3D Euclidean
distance between their kinocilia. Afferent distance was determined by the 3D Euclidean distance
between their soma centers. Only hair cells on the medial side of LPR or afferents innervating
the medial side of LPR were included for analysis of tuning similarity.
570
From the afferent somata in the utricular ganglion, afferent axons were then followed into the
brain. A total of 105 afferents were successfully reconstructed by two experienced annotators
(N.S. and M.W.B.) and all tracing was reviewed (M.W.B.). Central synapses were identified by
close appositions, thickening of the presynaptic membrane, and clustered vesicles (e.g. Fig. 3a,
4a). The Mauthner cell was previously reconstructed, as were most of the VS neurons[16,38].
Additional VS neurons were identified by reconstruction of utricular target neurons to the point
that they joined up with previously traced myelinated axons. VOR neurons were identified based
on their utricular input and their characteristic axonal projection patterns in the medial
longitudinal fasciculus. Because much of the axonal projections lay outside the reimaged
territory, only myelinated portions of axons could be reconstructed with confidence. Therefore,
we were not able to follow some VOR axons all the way to the trochlear and oculomotor nuclei.
Utricular commissural neurons were identified by their axons which crossed the midline and
traveled to the contralateral Mauthner cell. We note that we have identified a large number of
additional utricular target neurons in the brainstem that either do not fit into these categories
(e.g., commissural neurons) or cannot be confidently identified due to the difficulty in extending
their axons into lower-resolution territory. These neurons will be described in a future
publication. Therefore, the set of brainstem neurons analyzed here is likely to be strongly biased
to early-born, or at least early-myelinated, and is not a complete description of all VS or VOR
neurons. Nonetheless, the 19 VS neurons identified here is a large proportion of the 27 identified
by retrograde labeling[37].
591
Behavioral data were acquired in 5-6 dpf larvae from wild-types or the line *rock solo^AN66*, an
otogelin mutation[32]. Animals were visually verified as having normal (+/-) or absent (-/-)
utricular otoliths. Animals were free-swimming in a small dish with infrared transillumination
and imaged at 508 frames/s with a HiSpec-1 2g monochrome camera mounted on a Scientifica

SliceScope with a 2X objective. The translational stimulus was delivered with an air-floated sled
(Aerotech ABL 1500WB) and designed for large amplitude acceleration and jerk, to maximize
responses of irregular otolith afferents[105]. Due to the exploratory nature of this experiment, there
was no attempt to separate larvae into responders and non-responders as has been done for
acoustic stimuli[29], but animals were selected for behavioral tests based on whether they exhibited
some response to dish tapping. Images were analyzed with ZebraZoom[106] (zebrazoom.org) to
extract the smoothed tail angle.
603
604 Synaptic clustering was measured as the average distances between synapses on the same
605 dendritic arbor. Vestibular neuron arbor skeletons were first simplified for analysis by
606 regularizing the distance between nodes to ~1 um. Each synapse was associated with the nearest
607 skeleton node, and distances between synapses were calculated along the Euclidean length of the
608 skeleton.
609
To understand how the observed mean distance between synapses compared to random
distributions of synapses on the vestibular neuron arbors, we used several Monte Carlo models.
For each model, the synapses of each vestibular neuron were redistributed across the arbor
100,000 times. The mean distance between synapses was measured the same way as the non-
randomized mean distances was measured. The measure used for each of the 100,000 iterations
was the average of the mean synaptic distances of 43 vestibular neurons observed for that
iteration. Confidence intervals were calculated as the bounds containing 95% of the results of the
100,000 experimental randomizations.
618
619 The first model (unweighted distribution) compares the observed clustering of synapses to what
620 would be expected if synapses of each vestibular neuron were randomly distributed across the
621 entire arbor of that neuron. In this model, each vestibular neuron skeleton node had an equal
622 probability of being randomly assigned to one of the synapses of that vestibular neuron.
623
In the second model (50 µm-from-afferent weighted distribution) we restricted nodes likely to be
assigned to a synapse using the proximity of the nodes to the synaptic terminals of afferent
axons. 105 afferent axons were skeletonized as above. We then counted the number of afferent
synapse nodes within 50 µm of each vestibular neuron skeleton node (proximity score),
subtracting 1 to remove the influence of synaptic afferents presynaptic to a given vestibular
neuron node. We next shuffled synaptic locations while maintaining the likelihood of synaptic
connectivity given this proximity score. For example, if a skeleton node with a score of 95 has a
7% chance of being a synaptic locus, whereas a skeleton node with a score of 60 has a 1%
chance, then nodes with scores of 95 will be given a 7-fold greater chance of being assigned a
synapse in the redistribution than nodes with scores of 60.
634
In the final, most restrictive model (< 5 µm-from-afferent weighted distribution), we attempt to
use proximity to afferent terminals to recapitulate the clustering we observe in the real data.
Proximity scores were calculated as +1 for each vestibular neuron node 0 µm from a presynaptic
node. The added score decreased linearly to 0 at 5 µm distance. The rest of the 5 µm model was
the same as the 50 µm model. Code for these analyses is available at
https://github.com/bagnall-lab under "Liu_Connectome".
641

642   Analyses and statistics were carried out in Igor Pro 6 (Wavemetrics) or Matlab R2017a
643   (Mathworks). Statistical tests were carried out as reported in text, two-tailed where relevant, and
644   typically with nonparametric analyses due to the non-normal distribution of parameters.
645
646   *Data availability*
647   Quantification of hair cell to afferent and afferent to central target connectivity are provided
648   (Supp. Table 1, 2). All images and reconstructions are hosted and publicly available at
649   http://zebrafish.link/hildebrand16/data/vestibular_right
650

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Z.L., Y.J., and M.W.B. conceived the project, performed the experiments and analyzed the data. J.L.M. and D.G.C.H. carried out the alignment of the serial-section EM image data. J.L.M. designed and carried out the Monte Carlo simulations. Z.L., M.W.B., Y.J., and N.S. completed the reconstructions. Z.L. and M.W.B. wrote the manuscript with input from all other authors.

Declaration of Interests:

The authors declare no competing interests

930 **Figure legends**
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Figure 1. High-resolution serial-section electron microscopy of the gravity-sensing system.
a, Schematic of the gravity-sensing system in fish. Hair cells in the utricular macula (HC, gray)
are inertial sensors of head tilt and translation, exciting the peripheral process of utricular
afferents (green). These afferents, whose cell bodies are located in the utricular ganglion, project
to brainstem neurons involved in escape (Mauthner cell, black), posture (vestibulospinal [VS]
cell, blue), and oculomotor (VOR) reflexes (superior vestibular nucleus [SVN] and tangential
nucleus [Tan], brown). Dashed line indicates midline.
b, Coronal section through the head of a 5.5 dpf zebrafish. The region reimaged at high
resolution is visible as an L-shaped territory (dashed outline) covering the right utricle and hair
cells, utricular ganglion, and ipsilateral brainstem. The reimaged territory extended across 1757
coronal sections (105 µm in the rostrocaudal axis). Scale bar, 100 µm.
c, Electron micrograph of two hair cells in the utricular macula, with portions of their cilia. Scale
bar, 3 µm.
d, Section of the vestibular nerve, peripheral processes. At this developmental stage, some axons
are myelinated (pseudocolored dark green) while others are not yet (light green). Scale bar, 1
µm.
e, Horizontal projection of reconstructed brainstem targets (Mauthner, VS, SVN, Tangential)
colorized as in **a**.
f, Sagittal projection of utricular hair cells, afferents, and brainstem targets, as in **a**.
g, Coronal projection as in **f**.
952

Figure 2. The utricular afferent ganglion is organized in the rostrocaudal axis by directional
tuning
a, Top, Electron micrograph of utricular hair cell with stereocilia (black) and kinocilium (red)
marked. Right, schematic of the tuning vector derived from cilia positions, viewed from above.
Bottom, EM image of hair cell synaptic ribbons (arrowheads) apposed to a utricular afferent.
Scale bars, 1 µm.
b, Horizontal projection of the utricular macula, showing tuning direction vectors for all 91 hair
cells. Dashed line represents the line of polarity reversal (LPR). Vectors are colored by their
directional tuning to facilitate visualization. Note a slight asymmetry in the colorization; this was
chosen to ensure hair cells from the medial and lateral sides of the LPR are represented in
different colors. R, C = rostral, caudal = nose-down and nose-up pitch, respectively. M, L =
medial, lateral = contralateral and ipsilateral roll, respectively. The same conventions are used
throughout the paper.
c, Hair cells located near each other have more similar directional tuning, which falls off sharply
and within ~30 µm, indicating orderly array of vectors. Data from hair cells medial to the LPR
only.
d, Horizontal view of reconstructions of all 105 utricular ganglion afferents including somata
(larger spheres, left) and their postsynaptic contacts in the utricular macula (smaller spheres,
right). Afferents are colorized by inferred direction tuning as in **b**. View is slightly tilted to aid in
visualization of ganglion, which is otherwise obscured by centrally projecting axons. Afferents
with inferred contralateral head tilt tuning (red) form a segregated axon bundle in the brainstem
(left).

e, Sagittal view of reconstructions shown in **d**.
f, Correlation of rostrocaudal soma position between synaptically connected utricular hair cells
and afferents. Circle size reflects the number of synaptic ribbon connections (range: 1-19).
Significance of linear correlation, t-test, $p = 1.3 \times 10^{-102}$.
g, Horizontal projection of inferred afferent tuning vectors, relative to soma position in the
afferent ganglion. Each vector indicates an afferent's tuning direction, calculated by weighting
by the number of ribbon inputs it receives from each hair cell. Colors as in **b**.
h, Afferents located close to each other have similar directional tuning, but the relationship is
looser than in hair cells (**c**). Data are from afferents innervating the macula medial to the LPR
only.
985
Figure 3. Escape circuits compute head movement direction from utricular input
a, Example electron micrograph of two utricular afferents (pseudocolored green) synapsing onto
the lateral dendrite of the ipsilateral Mauthner cell (yellow). Chemical synapses are recognizable
by clustered presynaptic vesicles and synaptic density. The very tight apposition between the
upper afferent and the Mauthner is likely an electrical synapse (gap junction). Scale bar, 0.5 µm.
b, Coronal projection of Mauthner cell skeleton reconstruction (black) with utricular afferent
input synapses (gray). The Mauthner cell nucleus is represented by a sphere. Inset, expanded
view of utricular inputs onto the lateral dendrite.
c, Horizontal projection of reconstructions of both Mauthner cells (gray, blue) and the four
commissural utricular neurons (red). All four commissural utricular neurons make synaptic
contacts on the contralateral Mauthner (small red circles) and are therefore termed commissural
escape neurons. Colors indicate inferred directional tuning.
d, Polar plot of inferred direction tuning of utricular input to Mauthner cell and four commissural
escape neurons. As in Fig. 2, directional tuning is indicated in the context of head tilt. However,
otoliths are inertial sensors and therefore equally sensitive to head translation, which deflects the
hair cells in the opposite direction (*e.g.,* rightward head tilt and leftward translation are
identically processed by otoliths).
e, Schematic of predicted Mauthner cell computation of head translation. A predator approaching
from the right will cause a head deflection to the left. Deflection of the utricular otolith by inertia
to the right relative to hair cells (blue arrow) would depolarize the ipsilateral tilt / contralateral
translation pathway (blue; medial to the LPR). These utricular afferents excite the ipsilateral
Mauthner cell, promoting an escape movement to the left, away from the predator. Commissural
escape neurons, in contrast, will respond to rightward head movements (red hair cells and dashed
lines) and are predicted to activate the contralateral Mauthner cell, promoting escapes to the
right.
f, Example of behavioral response in a free-swimming larva subjected to rapid translation. High-
speed videography captures the onset of escape and characteristic C-bend. Bottom, quantification
of tail angle (black) and translational stimulus (gray). Scale bars, 1 rad, 100 ms, and 1 *g*.
g, Escape responses to a unidirectional translational stimulus are plotted relative to the larval
heading angle at the start of the translational stimulus. In accordance with our circuit predictions,
larvae accelerated to the right (heading direction 0-180˚) escape to the right, whereas larvae
accelerated to the left escape left. Escapes only occurred in the "incorrect" direction when
animals were accelerated in predominantly rostral or caudal directions.

h, With both types of stimulus, utricle-deficient *rock solo* larvae escaped at approximately half
the rate as their heterozygous siblings. N = 52 sibling and 55 -/- fish. Chi-squared test, p =
0.0006 (1 degree of freedom [df], chi-squared value = 11.65, 266 total observations).
1022
Figure 4. Structure and tuning of central VOR and VS utricular targets
a, Horizontal projection of 12 SVN reconstructed neurons. Neurons are colorized based on their
inferred directional tuning. Polar plot inset shows the directional tuning vectors of each neuron.
Gray bar indicates location of coronal plane where axon trajectories are shown in boxed inset.
Scale for reconstructions as in **c**. Electron micrograph (bottom) shows a utricular afferent
(pseudocolored green) contacting two dendrites of SVN neurons (yellow). Scale bar, 500 nm.
b, As in **a**, for 11 tangential nucleus reconstructed neurons. Sagittal projection inset (bottom)
shows the divergence of tangential neuron axons in the dorsoventral axis.
c, As in **a** but for 19 VS neurons.
d, Dendrograms of three example neurons from the SVN, tangential, and VS populations. The
soma is represented by a black circle and each dendrite is represented by lines. Gray circles
indicate synaptic inputs from utricular afferents, which appear disproportionately concentrated
on a small number of dendrites. Axons are truncated for purposes of scale.
e, Quantification of synaptic clustering, measured as distances between synapses, using three
Monte Carlo models compared to actual data. Synaptic locations were modeled as randomly
distributed across the arbor (unweighted model), preferentially weighted within 50 µm of
afferent axons, or preferentially weighted within 5 µm of afferent axons. The observed level of
synaptic clustering is shown in red. See Methods for detailed description.
1041
Figure 5. The utricular afferent ganglion is organized in the mediolateral axis by developmental
sequence and temporal dynamics.
a, Representation of kinocilium reconstruction across successive images to obtain the total
length. The neighboring tallest stereocilium was also reconstructed (not shown).
b, Plot of length of the kinocilium vs. the tallest stereocilium for all 91 hair cells (dots). Striolar
hair cells are identified by their long stereocilia and lower K/S ratios (dark brown) whereas
extrastriolar hair cells have much higher K/S ratios (tan). Hair cells characterized as immature
have very short kinocilia and stereocilia (blue-green).
c, Horizontal projection of the utricular macula, showing number of synaptic ribbons in each hair
cell. Circle diameter reflects synaptic ribbon count; hair cells with larger numbers of ribbons
tend to be located more centrally.
d, Quantification of synaptic ribbon counts across hair cell categories. Box plot represents
medians ± 25%ile; whiskers indicate 10-90%iles. Immature hair cells form fewer ribbon
synapses than striolar or extrastriolar hair cells. Significance was tested with ANOVA (p = 4.8 x
10^{-5}; df = 90, F statistic = 11.14) and then pairwise with Wilcoxon-Mann-Whitney (striola –
extrastriolar, p = 0.60 [U, U' = 552, 644]; striolar – immature, p = 4.3 x 10^{-4} [U, U' = 302, 66];
extrastriolar – immature, p = 1.3 x 10^{-4} [U, U' = 668.5, 163.5]).
e, Horizontal projection of all utricular afferents, colorized by whether they are myelinated
(black) or not (gray).
f, Horizontal projection map of all utricular ganglion somata by position; circle diameter reflects
the number of hair cell ribbon synaptic inputs that each one receives (cf. Fig. 2d).
g, Myelinated afferent somata are located more laterally in the utricular ganglion. Each dot
represents one afferent soma. Wilcoxon-Mann-Whitney, p = 2.9 x 10^{-4} [U, U' = 1105, 319].

h, Myelinated afferents are contacted by significantly more hair cell ribbons than unmyelinated
afferents. Wilcoxon-Mann-Whitney, $p = 1.8$ x 10^{-10} [U, U' = 1422, 2]. See also Supp. Fig. 3c.
i, Myelinated afferents receive the majority of their input from striolar hair cells, whereas
unmyelinated afferents receive most of their input from extrastriolar and developing hair cells.
The distributions are significantly different (chi-squared test, $p < 1$ x 10^{-10}, df = 3, chi squared
value = 133.4). See also Supp. Fig 3d, e..
j, The weighted fraction of inputs each utricular afferent receives from the different hair cell
classes. Afferents are ordered based on soma position from most medial to most lateral. Red dots
identify afferents receiving input from hair cells lateral to the LPR (contralateral tilt sensitive).
There is a gradient from laterally positioned afferents with predominantly striolar (phasic) inputs,
to medially positioned afferents with predominantly extrastriolar (tonic) or immature inputs.
1076
Figure 6. Early developing afferent pathways with fast kinetics preferentially drive early
developing central neurons for fast escapes.
a, Hair cells in the utricular macula that excite afferents connected to Mauthner escape, VOR, or
VS circuits (top, middle, and bottom). Hair cells are colorized if they contribute input to a
pathway or gray if they do not. Right, afferents connected to escape, VOR, or VS circuits,
colorized by their myelination status (black: myelinated; gray, unmyelinated).
b, Quantification of the contribution of striolar, extrastriolar, and immature hair cells to these
central pathways. Striolar hair cells preferentially drive rapid escape circuits, both via the direct
afferent input to the ipsilateral Mauthner and the afferent input to the commissural escape
pathway, whereas VOR and VS circuits receive input from a mixture of pathways, with
extrastriolar inputs dominating. Numbers in parentheses indicate the number of central neurons
in each category. Wilcoxon-Mann-Whitney test, striolar contribution to escape vs non-escape
neurons, $p = 0.041$ (U, U' = 141, 39).
c, Quantification of the contribution of synapses from myelinated and unmyelinated afferents to
each central pathway. Afferents driving escape pathways are largely myelinated at this age,
whereas afferents driving VOR and postural pathways are more mixed. Chi-squared test for all
groups, $p = 6.0$ x 10^{-8} (df = 3, chi squared = 36.45). Follow-up chi-squared: VOR vs escape, 1.4
x 10^{-4} (df = df = 1, chi squared = 14.47); VS vs escape, 1.7 x 10^{-8} (df = 1, chi squared = 31.75).
d, Subsets of VS neurons, identified by axon trajectories, are predicted to exhibit different
temporal kinetics. VS neurons with axons that approach the midline before descending (VS$_{med}$,
greens) receive mostly extrastriolar (tonic) input with a large contribution from immature hair
cells, whereas VS neurons with ventral axon trajectories (VS$_{vent}$, dark reds) receive mostly
striolar (phasic) input and none from immature pathways. The skeleton reconstruction is
projected at a mixed horizontal/sagittal angle to facilitate visualization of these groups. See also
representation of axon trajectories in Fig. 4d.
1102
Figure 7. Directional tuning and developmental sequence are organizing principles of
vestibulomotor connectivity.
a, Summary schematic of organization by directional tuning. Hair cells in the utricular macula,
left, project via afferents that maintain rostrocaudal organization but not mediolateral
organization. The utricular afferent ganglion is organized rostrocaudally, but contralateral tilt
sensitive afferents are intermingled. These afferents project with different patterns to distinct
brainstem targets, conferring directional sensitivity in the mediolateral (escape) or rostrocaudal
(VOR, posture) pathways. Colors indicate directional tuning as previously.

b, Summary schematic of organization by temporal kinetics. Early-born, striolar hair cells make
synaptic connections to early-born afferents, whose cell bodies are positioned laterally in the
utricular ganglion, and typically myelinated by the larval stage examined here. These early-born
afferents, carrying phasic information about head movement, preferentially excite escape
pathways, which consist of early-born, fast reticulospinal and spinal motor neurons and muscles.
Postural and VOR reflex pathways rely more on the tonic and phasic-tonic signals arising from
extrastriolar, slightly later born pathways. We speculate that circuits carrying immature input,
like VS_{med}, may project to motor circuits governing slower and more refined control of
movement.
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Supplemental Videos
Supplemental Video 1. The 3D reconstructed utricular circuits in the larval zebrafish, related to Figure 1. Rendering of the utricular hair cells (gray), afferents (blues), Mauthner cell (black), and central vestibular neurons (oranges) to show the entire gravity-sensing system. Utricles and eyes (gray volumes) are included for reference.

Supplemental Video 2. The 3D reconstruction of utricular afferents and the utricular macula, related to Figure 2. Utricular afferent neurons are colorized according to rostrocaudal position as in Fig. 2d. The positioning of afferent somata reflects a systematic innervation pattern from the afferents to the hair cells.

Supplemental Video 3. The 3D reconstruction of Mauthner and utricular commissural circuits, related to Figure 3. The right Mauthner (blue) and vestibular commissural (red) neurons receive projections from afferents that innervate the medial side of LPR (blue and yellow, tuned to ipsilateral tilt) and the lateral side of LPR (red, tuned to contralateral tilt). Note that the contralateral tilt (red) afferent axons exhibit a different trajectory in the brainstem than the ipsilateral tilt afferents.

Supplemental Video 4. The 3D reconstruction of the superior vestibular and tangential circuits, related to Figure 4.

Supplemental Tables
Supplemental Table 1. A grid of complete connectivity from utricular hair cells to utricular afferents. Numbers in each cell represent the number of ribbon synapses from a hair cell onto an afferent. Table also includes the computed tuning of each hair cell in radians.

Supplemental Table 2. A grid of connectivity from utricular afferents to identified central neurons in the brainstem. Numbers indicate the number of synaptic contacts. Where relevant, distinct release sites are counted as distinct contacts, but these are interpreted and less straightforward to quantify than ribbon synapses. Note that there are many afferent contacts onto central neurons not included in this table because they have not been reconstructed or identified. For example, some commissural neurons of unclear identity are not included. Table also includes the computed tuning of each afferent in radians.
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Figure 1

a
VOR
Utricle
SVN
M
HC
VS
Utric. ganglion
Tan
Escape Posture

b
D
M — L
V
Utric. gang.
utricle

c

d

e
R
M — L
C
50 μm

f
D
C — R
V
50 μm

g
D
M — L
V
utricular ganglion
utricle
hair cells
50 μm

Figure 2

a

b

LPR

Direction tuning
color code

R
M — L
C

R
M — L
C

10 μm

c

Tuning similarity

Hair cell dist. (μm)

d

R
M — L
C

central
projections

afferent
ganglion

Afferent
reconstructions

synapses with
hair cells

25 μm

e

D
C — R
V

25 μm

f

R   Hair cell position   C

R

Afferent position

C

r² = 0.40

○ 5 ribbons
◯ 10 ribbons

20 μm

g

R
M — L
C

10 μm

h

Tuning similarity

Afferent dist. (μm)

Figure 3

a

b
D
M ┼ L
V

c
25 μm
R
M ┼ L
C
R
M ┼ L
C
Mauthner cell
Commissural escape

d
Rostral
Contra ← → Ipsi
Caudal

e
Predator from right
Head translates left
Ipsi tilt path, escape left
inertial movement
Predator from left
Head translates right
Contra tilt path, escape right
Escape left   Escape right

f
10 ms   14 ms

g
Escape left
Escape right
0   90   180   270   360
Initial heading direction
accel

h
Escapes (%)
***
50/131
26/135
+/- siblings   rock solo

a SVN

mid

axons

5 μm

R
M—L
C

b Tangential

mid

5 μm

C—D—R
V

25 μm

c VS

mid

50 μm

D
M—L
V

20 μm

d

axon

SVN

axon

Tan.

axon

VS

25 μm

e

<5 μm-from-afferent
weighted model

<50 μm-from-afferent
weighted model

observed
distance

unweighted model

Fraction of of simulations

Average mean distance between synapses (μm)

Figure 5

a Measuring ciliary lengths in 3D

b
Kinocilium (μm) vs Stereocilium (μm)
- Striola
- Extrastriola
- Immature

c
LPR
ribbons
- striola: 5
- extrastriola: 15
- immature: 24

d
Number of ribbons
*** ***
striola, extrastriola, immature

e Utricular ganglion / Myelinated / Unmyelinated / Utricular macula — 25 μm

f Utricular ganglion — 5, 10, 20 — 10 μm

g Lateral / Medial — 10 μm — *** — Myelin. Unmyelin.

h # of hair cell synapses — *** — Myelin. Unmyelin.

i % of hair cells — Myelin. Unmyelin.

j
Weighted input fraction — Lateral to LPR
striola / extrastriola / immature
Medial (inferred late-born) — Individual afferents — Lateral (inferred early-born)

Figure 6

a

Inputs to escape circuit

Inputs to VOR circuit

Inputs to VS circuit

b

c

d

LPR

striola
extrastriola
immature

Escape | VOR | Posture
(1) (4) (23) (19)

% of hair cells (weighted)

Myel.
Unmyel.

% afferent synapses

Mauthner
Commiss.
SVN, Tan.
VS

VS_med
VS
VS_vent

50 μm

% of hair cells (weighted)

VS_med
VS
VS_vent

Figure 7

a

Rostral ipsi tilt

Contra tilt

Caudal ipsi tilt

Utricular macula

Utricular ganglion

R
M — L
C

Directional escape

Directional VOR

Directional posture

b

Striola / early developing / irregular firing

● Striola
● Extrastriola
● Immature

Extrastriola / later developing / regular firing

Motor control

phasic → Preferential to fastest motor circuits

tonic → Preferential to medium speed circuits

tonic → Preferential to slow/late circuits?

Escape

VOR
Posture

Fins?
Slow posture?