

RESEARCH

Open Access



Reduced olfactory acuity in recently flightless insects suggests rapid regressive evolution

Stefanie Neupert, Graham A. McCulloch, Brodie J. Foster, Jonathan M. Waters and Paul Szyszka*

Abstract

Background: Insects have exceptionally fast smelling capabilities, and some can track the temporal structure of odour plumes at rates above 100 Hz. It has been hypothesized that this fast smelling capability is an adaptation for flying. We test this hypothesis by comparing the olfactory acuity of sympatric flighted versus flightless lineages within a wing-polymorphic stonefly species.

Results: Our analyses of olfactory receptor neuron responses reveal that recently-evolved flightless lineages have reduced olfactory acuity. By comparing flighted versus flightless ecotypes with similar genetic backgrounds, we eliminate other confounding factors that might have affected the evolution of their olfactory reception mechanisms. Our detection of different patterns of reduced olfactory response strength and speed in independently wing-reduced lineages suggests parallel evolution of reduced olfactory acuity.

Conclusions: These reductions in olfactory acuity echo the rapid reduction of wings themselves, and represent an olfactory parallel to the convergent phenotypic shifts seen under selective gradients in other sensory systems (e.g. parallel loss of vision in cave fauna). Our study provides evidence for the hypothesis that flight poses a selective pressure on the speed and strength of olfactory receptor neuron responses and emphasizes the energetic costs of rapid olfaction.

Keywords: Regressive evolution, 'Use it or lose it', Evolutionary lability, Energetic cost, Selective pressure, Flight, Temporal resolution, Olfactory acuity, Plecoptera, Electroantennogram

Background

The origin of flight posed novel challenges for animals' sensory systems, including the need for rapid processing of environmental information, because flying animals move faster and therefore experience more rapid changes in sensory stimuli. Accordingly, insects that fly faster, have evolved faster responding photoreceptor cells [1, 2]. This need for rapid sensing seems to be particularly pronounced for olfaction [3, 4], because both the speed of wind-borne odour plumes and the rate of odour

concentration fluctuations increase with increasing distance from the ground [5, 6]. Indeed, olfactory receptor neurons of flighted insects can respond to odorants rapidly (within 3 ms and with sub-millisecond precision) [7], they can resolve fast odorant fluctuations (above 100 Hz) [8–10], corollary discharge from flight motor circuits enhances the temporal resolution of the insect olfactory system [11], and flying fruit flies can identify and respond to odorants within just 85 ms [12]. While it seems obvious that flight must have generated selective pressure for rapid olfactory transduction (the transformation of olfactory stimuli into action potentials), there is still only little direct evidence with which to assess this hypothesis [4, 13–16]. This lack of research on the evolutionary lability

*Correspondence: paul.szyszka@otago.ac.nz
Department of Zoology, University of Otago, Dunedin 9054, New Zealand



© The Author(s) 2022, corrected publication 2022. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

of temporal acuity of olfaction stands in stark contrast to the many studies on the evolution of the specificity of olfaction, which demonstrate rapid evolutionary adaptation of the olfactory system to the animals' chemical environments [13, 17–24].

While broad scale evolutionary trends in sensory systems are well documented, the pace at which such systems can evolve is poorly understood. Darwin [25] observed that loss of nonfunctional phenotypes (reductive evolution) is a repeated phenomenon in nature [26, 27]. Dramatic examples of such reductive evolutionary processes include rapid deterioration of inactivated genetic material (e.g. [28]), and parallel losses of pigmentation and eyes in diverse cave fauna (e.g. [29, 30]). If maintenance of rapid olfactory transduction is key to the success of flighted insects, this begs the question of what happens to this sensory capacity when such lineages become secondarily wing-reduced. Indeed, flight loss has evolved independently in nearly every order of winged

insects, and in some clades it has occurred repeatedly [31–33].

Under the 'use it or lose it' hypothesis, we propose that olfactory acuity becomes rapidly reduced in insect lineages that no longer use flight, in the same way that wings themselves become reduced. Here, we use a wing-dimorphic member of the early-diverging winged insect order Plecoptera as a model to test the hypothesis that rapid olfaction is a requirement specifically for flighted insects, but not for flightless lineages. The New Zealand *Zelandoperla fenestrata* Tillyard stonefly complex comprises both full-winged (flighted) and wing-reduced (non-flying) lineages that co-occur widely [34, 35] (Fig. 1A, B). Flight loss in *Z. fenestrata* is believed to be an evolutionary adaptation to high winds [36], with wing reduced populations typically found above the alpine treeline, or in lowland deforested regions [37–39]. Recent genomic analyses of this species complex indicate that wing reduction has evolved recently (likely within the last

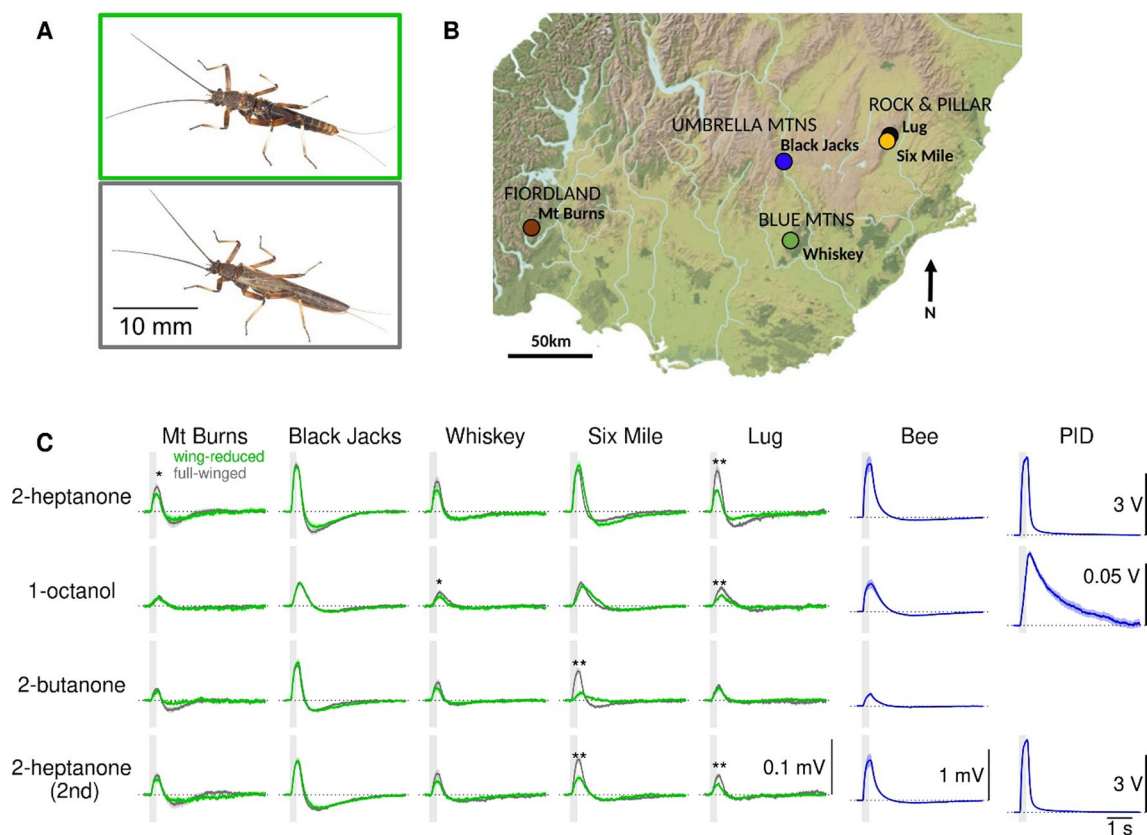


Fig. 1 Odorant-evoked antennal responses are weaker in wing-reduced than in full-winged stoneflies. **A** Wing-reduced (green box) and full-winged (grey box) *Zelandoperla fenestrata* ecotypes. **B** Topographic map of the five sampling sites in New Zealand. **C** Antennal signal traces (mean \pm SEM) for the five stonefly populations (grey: full-winged; green: wing-reduced). Number of antennae for wing-reduced/full winged stoneflies of the five populations (from left to right): 10/10, 24/23, 11/13, 11/14, 6/8. Signal traces of honey bee antennae (12 antennae) for scale and a photoionisation detector (PID) for visualizing the stimulus dynamics (10 recordings; PID signals for 2-butanone saturated and are not shown). Grey vertical bars indicate odorant valve opening time (300 ms). Horizontal dotted lines show 0 V

15,000 years) and independently in different regions [37], and revealed differential expression of olfaction-related genes between full-winged versus wing-reduced stonefly ecotypes [39, 40]. Together, this makes *Z. fenestrata* a replicated model system for testing the hypothesis that flightless lineages exhibit secondarily reduced temporal olfactory acuity relative to that of flighted lineages.

Results

To test the hypothesis that flight loss leads to a reduction of temporal olfactory acuity we compared odorant-evoked antennal responses between co-occurring full-winged and wing-reduced stonefly lineages (measured with electroantennogram recordings, EAG) (Fig. 1A). We sampled stoneflies from five genetically distinct stream populations [39] (Fig. 1B).

Antennal recordings in full-winged versus wing-reduced stonefly ecotypes revealed differences in the strength of antennal responses to odorants in four out of five stonefly populations (there was no difference in the Black Jacks population, Fig. 1C, Additional file 1: Fig. S1A). When differences in response strength were detected, wing-reduced stoneflies always showed weaker responses than full-winged stoneflies. Those weaker responses occurred for some but not all odorant stimuli, and odorants that evoked weaker responses in wing-reduced stoneflies, differed across stonefly populations. In the Six Mile population, wing-reduced stoneflies showed weaker responses to the second but not to the first set of 2-heptanone stimuli. This could indicate a faster deterioration or adaptation in antennae of wing-reduced stoneflies, however, we did not investigate the cause of this effect.

In addition to weaker antennal responses, wing-reduced stoneflies showed slower response onset and offset times relative to full-winged stoneflies (Fig. 2A, B). Slower responses occurred in wing-reduced stoneflies from three out of five populations (Mt Burns, Black Jacks, Six Mile), they occurred for some but not all odorant stimuli, and odorants that evoked slower responses, differed across the different stonefly populations. In the Lug population, there were no differences in the response onset or offset times, and in the Whiskey population one odorant evoked a faster response offset time in wing-reduced stoneflies.

We probed the capability of stonefly antennae to resolve rapidly fluctuating odorant stimuli (Additional file 1: Fig. S1B) and recorded responses to 3-s long 10-Hz pulse series. However, antennae from neither full-winged nor wing-reduced stoneflies could track 10-Hz odorant pulses, indicating that antennal responses in stoneflies have a lower temporal resolution than antennal responses

in more derived insect groups [9, 41, 42], including honey bees [8] (Additional file 1: Fig. S1B).

Discussion

Weaker and slower antennal responses in wing-reduced stoneflies

Wing-reduced stonefly ecotypes consistently displayed weaker antennal (EAG) responses and slower response onset and offset times relative to their full-winged counterparts. A weaker antennal response is associated with a lower action potential rate (within individual olfactory receptor neurons [43] or across a population of olfactory receptor neurons [44]), and a lower action potential rate reduces the temporal acuity of encoding the onset of an odorant stimulus [7, 45]. Therefore, the weaker antennal responses in wing-reduced individuals suggest these lineages have reduced temporal olfactory acuity.

The response dynamics of olfactory receptor neurons are shaped by multiple transduction (odorant-receptor (un)binding, receptor (de)activation), and adaptation processes, and the degree to which those processes contribute to a receptor neuron's response dynamic depends on receptor type, odorant, and odorant dynamic [46–48]. Therefore, the finding that wing-reduced stoneflies from different streams show distinct, odorant-specific patterns of reduced response speed (Figs. 1C, 2) may indicate that those different transduction and adaptation processes were affected differently in the different lineages and suggests that these shifts evolved independently in each population. This inference is reinforced by recent genomic analyses, which indicate flight loss has independently evolved in each population [37, 39]. The current study thus provides additional evidence for the repeated evolution of flight loss, with different forms of reduced olfactory acuity detected in these independently flightless lineages.

Flightless *Z. fenestrata* stonefly ecotypes have evolved within the last 15,000 years [39], indicating the reduction of olfactory acuity has likewise occurred rapidly. Recent genomic studies indicate that clusters of the adaptive loci often underpin rapid adaptation [49–51]. If olfactory genes are closely genetically linked to the gene underpinning wing loss in *Z. fenestrata* this may facilitate parallel shifts in flight ability and olfaction. The exact genomic mechanism of wing loss in *Z. fenestrata* is not known, and likely varies across populations [39, 52]. However, a recent study suggests that the developmental supergene *doublesex* may play an important role, and this gene is closely physically linked to at least one odorant binding protein [39]. Future genomic studies promise to further unravel the potential role that genetic linkage may play in the parallel reductions of flight and olfactory ability in this species.

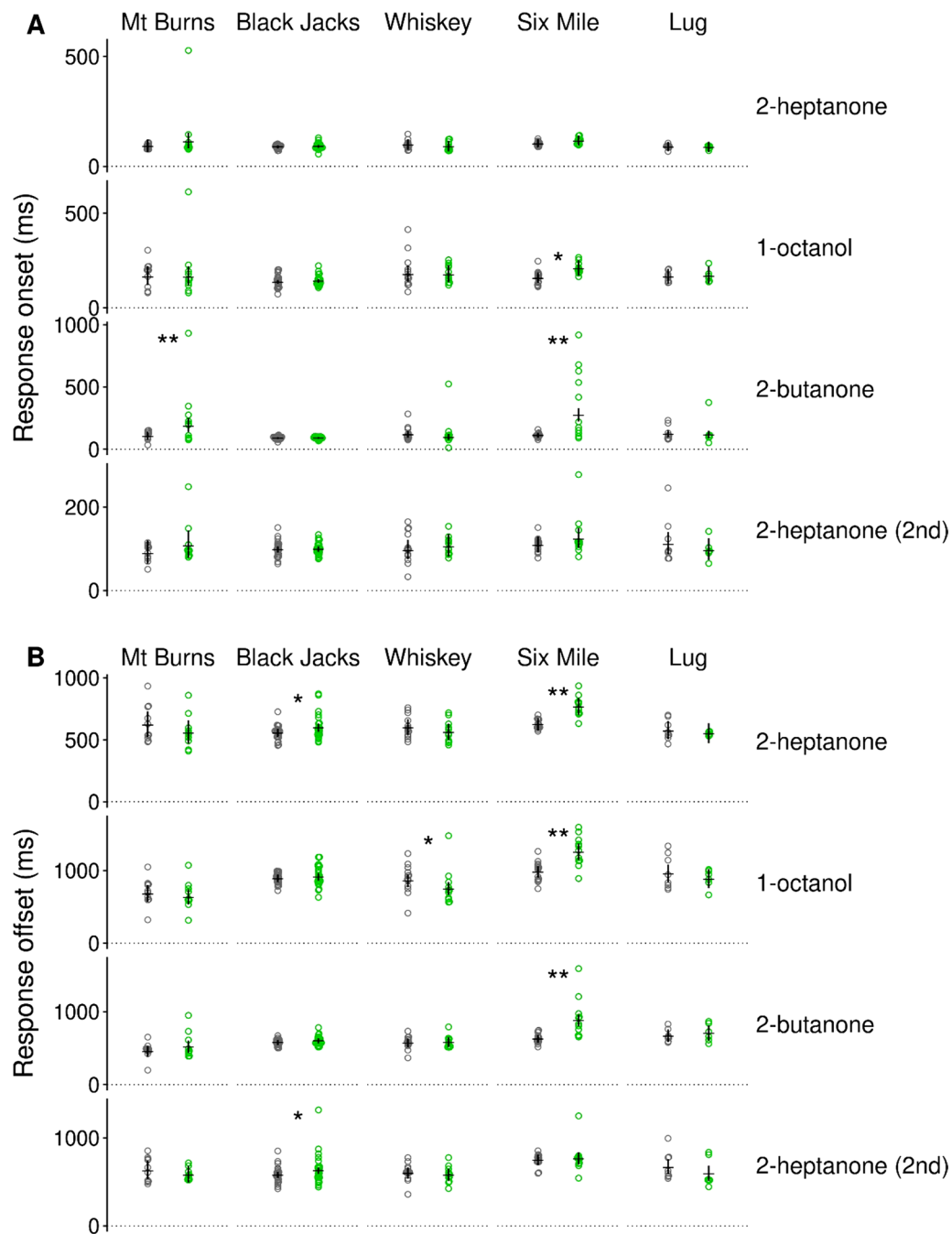


Fig. 2 Odorant-evoked antennal responses are slower in wing-reduced than in full-winged stoneflies. **A** Antennal response onset time (time to 10% of signal maximum after valve opening, includes the time odorant need to reach the antenna), and **B** response offset time (time to 10% of signal maximum after the maximum). Grey: full-winged; green: wing-reduced. Circles show individual antennae. Horizontal black lines show means and vertical black lines show 95% credible intervals. * or ** greater than 95% or 99% certainty for differences between antennae of full-winged and wing-reduced stoneflies

Selective pressure for rapid olfactory transduction in flying insects

The faster onsets and offsets of odour-evoked antennal responses detected in full-winged stoneflies are likely

to facilitate long-range search behaviour. When searching for an odorous target (e.g. a mating partner), flying insects encounter rapidly fluctuating odour plumes [53–55]. Because the randomness of air flow destroys most of

the directional information present in an odour plume (e.g. concentration gradients), flying animals need to use reactive or infotactic strategies for odour plume tracking, both of which require the animal to rapidly detect and react to the target odour [56–60]. A fast onset of the antennal response would shorten the time to detect and react to the target odour, while fast offset of the antennal response would facilitate the detection of subsequent odour stimuli. In addition to facilitating odour plume tracking, rapid olfaction enables the perceptual segregation of mixed odorants from different sources based on short differences in the arrival of odorants [10, 61–66].

In line with the ecological requirement for rapid olfaction, insect olfactory receptor neurons can respond to odorants rapidly (within 3 ms) [7, 8]. Olfactory receptor neurons can respond rapidly because olfactory receptors are ion channels which are composed of odorant-specific olfactory receptors (OR) and co-receptors (Orco) [67–69]. Missbach et al. [14] suggested that the OR/Orco system represents a specific adaptation associated with insect flight (but see [13] for an opposing view). While OR genes likely evolved in the common ancestor of modern insects, perhaps as an adaptation to terrestrial conditions [13, 24], the OR/Orco system may have arisen more recently and may have facilitated rapid olfaction in winged insects [15]. The finding that even full-winged stoneflies had a relatively slow odorant pulse tracking capability (below 10 Hz; Additional file 1: Fig. S1B) compared to that of strong-flying insects (above 100 Hz [8]) may reflect the generally weaker flying ability of this early-diverging insect clade [70].

Energetic costs of rapid olfactory transduction and mechanisms of its slackening

The reduced strength and speed of antennal responses in wing-reduced stoneflies suggests that olfactory transduction is energetically costly, which in turn generates a selective pressure for slackening olfactory transduction in flightless insects. The strength of an antennal (EAG) response correlates with the receptor current amplitude and action potential number [43, 44]. Because both ion currents and action potentials are energetically costly [71], the energetic cost of antennal responses increases with response strength. Likewise, a rapidly responding receptor neuron is likely to be energetically costly, because it requires a low membrane time constant which in turn, requires an energetically costly high membrane conductance [72].

Besides fast activation, high temporal resolution requires fast deactivation of olfactory receptor neurons, so that they can respond to following odorant pulses. Deactivation is thought to be mediated by removal of odorants from the OR/Orco complex through

odorant-binding proteins and odorant-removing proteins [73]. A putative odorant-removing protein, *Pinocchio* [74], is significantly more expressed in the notum of wing-reduced than in full-winged stoneflies from Lug Creek [40]. If *Pinocchio* is similarly overexpressed in the antennae of wing-reduced stoneflies, this may explain the weaker odour-evoked antennal responses detected in wing-reduced stoneflies. Future comparisons of antenna morphologies, antennal expression of olfaction-related genes [75], and the number, membrane conductance, and action potential threshold of olfactory receptor neurons will further elucidate the mechanistic basis of this reduction in olfactory acuity.

Conclusions

The findings of the current study highlight not only the need for rapid olfactory processing in flying insects [3], but also that olfactory acuity can be rapidly reduced when no longer required (when flight ability is lost). The locally and independently wing-reduced lineages analysed here have diverged from their winged counterparts only very recently in evolutionary terms (during the current interglacial, less than 15,000 years ago) [37, 39]. This rapid reductive evolution of sensory ability echoes the rapid reduction of wings themselves, and also represents a neurobiological parallel to the rapid phenotypic shifts seen under sharp selective gradients in other systems (e.g. loss of vision in cave fauna) [29, 30]. Broadly, these findings emphasize the energetic costs of sensory acuity, and the key role of natural selection in shaping neurobiological shifts. Additionally, this multidisciplinary analysis highlights the potential for future studies to further elucidate evolutionary changes in sensory systems.

Materials and methods

Stonefly sampling

We sampled stoneflies from zones of ecotypic overlap from five genetically independent stream populations in New Zealand [37, 39]: Mount Burns, Black Jacks Creek, Whiskey Creek, Six Mile Creek and Lug Creek (Fig. 1B). Final instar nymphs were collected by hand from under stones or wood in stream cascades and rapids. Nymphs were subsequently reared in the laboratory in Styrofoam cups at 11 °C under a natural day:night cycle, in water from their natal stream. We collected nymphs of full-winged and wing-reduced ecotypes from the same locations, and reared them under the same conditions, so that any differences between them are unlikely to reflect environmental variation. After emerging as adults, stoneflies were sexed based on genitalia, and morphologically characterized as either full-winged or wing-reduced.

Antennal responses to olfactory stimuli

We used electroantennograms (EAGs) of detached antennae as a proxy for the amplitude and dynamics of olfactory receptor neuron responses. The amplitude of EAG signals increases with increasing receptor current amplitudes and with increasing number of activated receptor neurons [43, 44, 76]. Note that EAG signals do not accurately reflect spike rates, but there is a positive correlation between receptor current amplitude and spike rate [43, 46, 77]. We chose EAG recordings over single neuron recordings, because we could not collect the high number of stoneflies that would have been needed for single neuron recordings. For the experiments shown in Figs. 1, 2 and Additional file 1: S1, we used 1- to 5-day old adult male stoneflies and female honey bees. For the experiments shown in Additional file 1: Fig. S2 we used male and female stoneflies. We used a razor blade to cut off a 5 mm long section of the distal antennae. Antennae were mounted with conductive gel (GEL+, Ritex, Germany) on a four-channel silver electrode [66]. Five minutes later, the antennae were placed at a distance of 2 mm in front of the outlet of the olfactory stimulator. To eliminate between-session variability (e.g., due to humidity or circadian rhythms in antennal responsiveness [78]), the left and right antennae of one full-winged and one wing-reduced stonefly were recorded simultaneously. EAG signals were differentially amplified against the reference electrode using 1000× gain, recorded in AC-coupled mode and low-pass-filtered at 1 kHz (MA 103 preamplifier and MA 102 four-channel amplifier, Universität zu Köln). The distal tips of the four antennae were mounted on a common central electrode that was connected to the inverted inputs of the preamplifiers. Under this approach, positive EAG signals reflect excitatory responses (activation of olfactory receptor neurons) and negative EAG signals reflect inhibitory responses.

Olfactory stimulation

We used a custom-made 6-channel olfactory stimulator (same approach as in [79] but built with materials as described in [80]). This stimulator provided a constant airflow (volume flow rate = 4.8 L/min, flow speed = 100 cm/s). The antennae were exposed to this constant airflow throughout the whole duration of the recordings before and between the odorant stimuli. To apply odorants, odorant-laden air (300 mL/min) was injected into the constant carrier air stream and, simultaneously, the same amount of clean air was withdrawn to keep the total airflow constant. The odorant air stream was produced by bottled compressed air and the carrier air stream was produced by an aquarium pump. Both air streams were filtered with active carbon filters (HN4S-AUN, Parker). The stimulation was controlled with the

data acquisition system Micro 3 1401 and Spike2 software version 8.03 (CED).

All odorants were kept in glass vials with PTFE septum screw caps (20 mL EPA vial, JG Finneran). We presented the following olfactory stimuli in the following sequence: odourless blank (empty vial), 2-heptanone, 1-octanol, and 2-butanone (these three odorants purchased from Sigma-Aldrich). We chose the odorants for the following reasons: 2-heptanone because it induces robust antennal responses in different insect species [8]; 1-octanol because it was used in a previous EAG study of stoneflies [81], 2-butanone because of its fast stimulus dynamics [61]. We also trialed propionic acid (Sigma-Aldrich), which had evoked particularly strong antennal responses in a previous study [81], and we tested water, but we ultimately excluded these two odorants following the detection of non-biological artifacts (Additional file 1: Fig. S2).

We used a photoionisation detector (miniPID B, Aurora) to measure the dynamics of odorant concentration change (Fig. 1C, Additional file 1: Fig. S1B). To vary the strength of olfactory stimuli we presented each odorant at up to four different pulse durations (15, 30, 150 and 300 ms valve open time, Additional file 1: Fig. S1A). Because the rise time of the odorant concentration was larger than 150 ms, these different stimulus durations resulted in different maximum concentrations. Note that odorant-specific differences in stimulus dynamics are a consequence of interactions between odorants and the surfaces of both the olfactory stimulator and the photoionisation detector [47, 80]. We also recorded antennal responses in honey bees (Fig. 1C, Additional file 1: Fig. S1) as a scale for comparing stoneflies' antennal responses to other insects (see temporal resolution of antennal responses of honey bees, locusts, moths, and cockroaches in a previous study [8]).

Each olfactory stimulus (odorant/pulse duration combination) was presented ten times at an inter-stimulus interval of 5 s. Ten seconds after the last 300 ms long stimulus we presented fluctuating 10-Hz stimuli by repetitively opening the valve for 50 ms at a frequency of 10 Hz over a period of 3 s (Additional file 1: Fig. S1B). This 10-Hz stimulus served to quantify the temporal resolution. Five seconds after the 10-Hz stimulus the next odorant started. At the end of each experiment we presented another 10 stimuli of 2-heptanone (300 ms) to test whether the antennae were still responding. Stimulus protocols varied between stream populations in the number of stimuli, and these differences could explain differences in antennal responses between stream populations. We excluded 22 antennae that showed sudden baseline shifts.

Data analysis

EAG signal data were exported from Spike2 and then further analyzed using R (version 3.6.3) [82]. Each recording was cut into 5 s traces, starting 0.2 s before and ending 4.8 s after each stimulus onset. Each trace was baseline-corrected by subtracting the median voltage during the 0.2 s time window before valve opening from the entire trace. To increase the signal-to-noise ratio, we calculated the median trace over each set of 10 stimulations with the same odorant/pulse duration combination. To reduce the noise, we applied a running median filter with a window size of 11 ms on the traces.

To assess whether an antenna shows stimulus-induced responses to the last set of 2-heptanone stimulations, we defined a response threshold as two times the standard deviation of the last 3 s of the trace. If the stimulus induced response of an antenna to these 2-heptanone stimulations did not exceed the response threshold for at least 50 consecutive ms between 50 and 1000 ms after valve opening, we excluded all recordings from this antenna from further analysis (68 of 76 full-winged antennae and 62 of 77 wing-reduced antennae were analyzed).

We quantified different response parameters evoked by 2-heptanone, 1-octanol, and 2-butanone between stonefly ecotypes for a given odorant and collection site: (1) response strength as the mean response in the time window of 25 ms before and 25 ms after the maximum signal between 50 ms and 1 s after valve opening; (2) response onset: time between valve opening and 10% of maximum signal (before signal maximum); (3) response offset time: time between valve opening and 10% of maximum signal (after signal maximum); and (4) temporal resolution for 10-Hz stimulations: power spectral densities on a 3 s time window starting 0.4 s after the first valve opening using the *multitaper* R package [83] with the *sine taper* method.

To quantify if there are differences in response strength between ecotypes, we ran linear mixed models with log₂ transformed response strength as the response variable. To avoid negative values, we added an offset of 0.01 to each mean response prior to the log₂ transformation. We included ecotype (full-winged or wing-reduced) and pulse duration as explanatory variables and added an interaction between both variables. To account for repeated measurements of the same antennae, we included antenna identity as a random factor.

To test if there are differences in response timing, i.e. in response onset time or offset time between stonefly ecotypes for a given odorant and collection site, we ran linear models. We included the log₂ transformed response timing of interest (either response onset, or response offset) for 300 ms long pulses as the response

variable and ecotype (full-winged or wing-reduced) as explanatory variable.

Inferences for all types of models were drawn using Bayesian statistics. We used an improper prior distribution (flat prior) and simulated 10,000 random draws from the posterior distribution using the function *sim* from the R package *arm* [84].

We used model estimates as the mean and the 2.5% and 97.5% quantiles as the lower and upper limit of the 95% credible interval. We calculated the proportion of simulated values from the posterior distribution that are bigger for one ecotype (e.g. A) over the other (e.g. B). A resulting proportion of 0.99 would mean that we are 99% certain that ecotype A has a larger parameter value than ecotype B. For plotting, we back-transformed the quantiles in the original scale. For the mean response strength, we subtracted the offset of 0.01 again. We marked differences between full-winged and wing-reduced ecotypes in a given odorant/pulse duration configuration of >95% and >99% certainty with one and two asterisks, respectively.

Abbreviations

EAG: Electroantennogram; OR: Olfactory receptor; Orco: Olfactory receptor co-receptors; PID: Photoionisation detector.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12862-022-02005-w>.

Additional file 1: Figure S1. Antennal responses to different stimulus durations and pulse series. **Figure S2.** Biological and non-biological antennal signals are independent from each other.

Acknowledgements

We thank Michael Thoma for comments on the manuscript and for pointing us to the possibility that propionic acid evokes non-biological EAG responses, and we thank the participants of the “Behavioural Ecology and Evolution” seminar (Dept. Zoology, University of Otago) and of the “Molecular Basis of Animal Evolution” course (California Institute of Technology) for feedback on the pre-print of this study [85].

Author contributions

Conceptualization, PS and JMW; Methodology, PS and SN; Software, SN; Validation, SN and PS; Formal Analysis, SN; Investigation, PS; Resources, SN, GAM, BJF, JMW, and PS; Data Curation, SN and PS; Writing—Original Draft, PS, JMW; Writing—Review & Editing, PS, SN, JMW, GAM, and BJF; Supervision, PS; Funding Acquisition, PS. All authors read and approved the final manuscript.

Funding

P.S. has been partly funded by the Marsden contract UOO2114. The photoionisation detector was funded by a University of Otago Research Grant (#3435, #18817) to P.S. Locations with ecotype mixtures were identified and sampled based on ongoing ecological genomic research funded by Marsden contracts UOO1412 and UOO2016 (Royal Society of New Zealand) to J.M.W. and G.A.M.

Availability of data and materials

The datasets generated and analysed during the current study are available at <https://doi.org/10.12751/g-node.yq9qml>.

Declarations

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

Received: 4 July 2021 Accepted: 8 April 2022

Published: 16 April 2022

References

- O'Carroll DC, Bidwell NJ, Laughlin SB, Warrant EJ. Insect motion detectors matched to visual ecology. *Nature*. 1996;382(6586):63–6.
- Howard J, Dubs A, Payne R. The dynamics of phototransduction in insects—a comparative study. *J Comp Physiol A*. 1984;154(5):707–18.
- De Bruyne M, Baker TC. Odor detection in insects: volatile codes. *J Chem Ecol*. 2008;34:882–97.
- Getahun MN, Wicher D, Hansson BS, Olsson SB. Temporal response dynamics of *Drosophila* olfactory sensory neurons depends on receptor type and response polarity. *Front Cell Neurosci*. 2012;6(November):54.
- Allen LH. Turbulence and wind speed spectra within a Japanese larch plantation. *J Appl Meteorol*. 1968;7(1):73–8.
- Yee E, Chan R, Kosteniuk PR, Chandler GM, Biloft CA, Bowers JF. The vertical structure of concentration fluctuation statistics in plumes dispersing in the atmospheric surface layer. *Bound Layer Meteorol*. 1995;76(1–2):41–67.
- Egea-Weiss A, Renner A, Kleineidam CJ, Szyszka P. High precision of spike timing across olfactory receptor neurons allows rapid odor coding in *Drosophila*. *iScience*. 2018;4:76–83.
- Szyszka P, Gerkin RC, Galizia CG, Smith BH. High-speed odor transduction and pulse tracking by insect olfactory receptor neurons. *Proc Natl Acad Sci USA*. 2014;111(47):16925–30.
- Schuckel J, Meisner S, Torkkeli PH, French AS. Dynamic properties of *Drosophila* olfactory electroantennograms. *J Comp Physiol A*. 2008;194(5):483–9.
- Szyszka P, Stierle JS, Biergans S, Galizia CG. The speed of smell: odor-object segregation within milliseconds. Louis M, editor. *PLoS ONE*. 2012;7(4):e36096.
- Chapman PD, Burkland R, Bradley SP, Houot B, Bullman V, Dacks AM, et al. Flight motor networks modulate primary olfactory processing in the moth *Manduca sexta*. *Proc Natl Acad Sci USA*. 2018;115(21):5588–93.
- Bhandawat V, Maimon G, Dickinson MH, Wilson RI. Olfactory modulation of flight in *Drosophila* is sensitive, selective and rapid. *J Exp Biol*. 2010;213(21):3625–35.
- Brand P, Robertson HM, Lin W, Pothula R, Klingeman WE, Jurat-Fuentes JL, et al. The origin of the odorant receptor gene family in insects. *eLife*. 2018. <https://doi.org/10.7554/eLife.38340>.
- Missbach C, Dweck HKM, Vogel H, Vilcinskis A, Stensmyr MC, Hansson BS, et al. Evolution of insect olfactory receptors. *eLife*. 2014. <https://doi.org/10.7554/eLife.02115>.
- Thoma M, Missbach C, Jordan MD, Grosse-Wilde E, Newcomb RD, Hansson BS. Transcriptome surveys in silverfish suggest a multistep origin of the insect odorant receptor gene family. *Front Ecol Evol*. 2019;7:281.
- Getahun MN, Thoma M, Lavista-Llanos S, Keesey I, Fandino RA, Knaden M, et al. Intracellular regulation of the insect chemoreceptor complex impacts odor localization in flying insects. *J Exp Biol*. 2016;219(September):jeb.143396.
- Conchou L, Lucas P, Meslin C, Proffit M, Staudt M, Renou M. Insect odorscapes: from plant volatiles to natural olfactory scenes. *Front Physiol*. 2019;10:972.
- Andersson MN, Löfstedt C, Newcomb RD. Insect olfaction and the evolution of receptor tuning. *Front Ecol Evol*. 2015;3:53.
- Hansson BS, Stensmyr MC. Evolution of insect olfaction. *Neuron*. 2011;72:698–711.
- Auer TO, Khallaf MA, Silbering AF, Zappia G, Ellis K, Álvarez-Ocaña R, et al. Olfactory receptor and circuit evolution promote host specialization. *Nature*. 2020;579(7799):402–8.
- Prieto-Godino LL, Rytz R, Bargeton B, Abuin L, Arguello JR, Peraro MD, et al. Olfactory receptor pseudo-pseudogenes. *Nature*. 2016;539(7627):93–7.
- Silbering AF, Rytz R, Grosjean Y, Abuin L, Ramdya P, Jefferis GSXE, et al. Complementary function and integrated wiring of the evolutionarily distinct *Drosophila* olfactory subsystems. *J Neurosci*. 2011;31(38):13357–75.
- Hayden S, Bekaert M, Crider TA, Mariani S, Murphy WJ, Teeling EC. Ecological adaptation determines functional mammalian olfactory subgenomes. *Genome Res*. 2010;20(1):1–9.
- Robertson HM, Warr CG, Carlson JR. Molecular evolution of the insect chemoreceptor gene superfamily in *Drosophila melanogaster*. *Proc Natl Acad Sci USA*. 2003;100(24):14537–42.
- Darwin C. On the origin of species. 1st ed. London: Murray; 1859. p. 502.
- Fong DW, Kane TC, Culver DC. Vestigialization and loss of nonfunctional characters. *Annu Rev Ecol Syst*. 1995;26:249–68.
- Porter ML, Crandall KA. Lost along the way: the significance of evolution in reverse. *Trends Ecol Evol*. 2003;18:541–7.
- Kuo CH, Ochman H. The extinction dynamics of bacterial pseudogenes. *PLoS Genet*. 2010. <https://doi.org/10.1371/journal.pgen.1001050>.
- Jeffery WR. Adaptive evolution of eye degeneration in the Mexican blind cavefish. *J Hered*. 2005;96:185–96.
- Protas ME, Trontelj P, Patel NH. Genetic basis of eye and pigment loss in the cave crustacean, *Asellus aquaticus*. *Proc Natl Acad Sci USA*. 2011;108(14):5702–7.
- Roff DA. The evolution of flightlessness in insects. *Ecol Monogr*. 1990;60(4):389–421.
- Roff DA. The evolution of flightlessness: is history important? *Evol Ecol*. 1994;8(6):639–57.
- Waters JM, Emerson BC, Arribas P, McCulloch GA. Dispersal reduction: causes, genomic mechanisms, and evolutionary consequences. *Trends Ecol Evol*. 2020. <https://doi.org/10.1016/j.tree.2020.01.012>.
- McCulloch GA, Wallis GP, Waters JM. Do insects lose flight before they lose their wings? Population genetic structure in subalpine stoneflies. *Mol Ecol*. 2009;18(19):4073–87.
- McLellan ID. A revision of *Zelandoperla* Tillyard (Plecoptera: Gripopterygidae: Zelandoperlinae). *N Z J Zool*. 1999;26(3):199–219.
- McCulloch GA, Foster BJ, Ingram T, Waters JM. Insect wing loss is tightly linked to the treeline: evidence from a diverse stonefly assemblage. *Ecography (Cop)*. 2019;42(4):811–3.
- McCulloch GA, Foster BJ, Dutoit L, Ingram T, Hay E, Veale AJ, et al. Ecological gradients drive insect wing loss and speciation: The role of the alpine treeline. *Mol Ecol*. 2019;28(13):mec.15114.
- Foster BJ, McCulloch GA, Vogel MFS, Ingram T, Waters JM. Anthropogenic evolution in an insect wing polymorphism following widespread deforestation. *Biol Lett*. 2021. <https://doi.org/10.1098/rsbl.2021.0069>.
- McCulloch GA, Foster BJ, Dutoit L, Harrop TWR, Guhlin J, Dearden PK, et al. Genomics reveals widespread ecological speciation in flightless insects. *Syst Biol*. 2020. <https://doi.org/10.1093/sysbio/syaa094>.
- McCulloch GA, Oliphant A, Dearden PK, Veale AJ, Ellen CW, Waters JM. Comparative transcriptomic analysis of a wing-dimorphic stonefly reveals candidate wing loss genes. *EvoDevo*. 2019;10(1):21.
- Lemon W, Getz W. Temporal resolution of general odor pulses by olfactory sensory neurons in American cockroaches. *J Exp Biol*. 1997;200(Pt 12):1809–19.
- Bau J, Justus KA, Cardé RT. Antennal resolution of pulsed pheromone plumes in three moth species. *J Insect Physiol*. 2002;48(4):433–42.
- Mayer MS, Mankin RW, Lemire GF. Quantitation of the insect electroantennogram: measurement of sensillar contributions, elimination of background potentials, and relationship to olfactory sensation. *J Insect Physiol*. 1984;30(9):757–63.
- Kapitskii SV, Gribakin FG. Electroantennogram of the American cockroach: effect of oxygen and an electrical model. *J Comp Physiol A*. 1992;170(5):651–63.
- Jeanne JM, Wilson RI. Convergence, divergence, and reconvergence in a feedforward network improves neural speed and accuracy. *Neuron*. 2015;88(5):1014–26.
- Nagel KI, Wilson RI. Biophysical mechanisms underlying olfactory receptor neuron dynamics. *Nat Neurosci*. 2011;14(2):208–16.

47. Martelli C, Carlson JR, Emonet T. Intensity invariant dynamics and odor-specific latencies in olfactory receptor neuron response. *J Neurosci*. 2013;33(15):6285–97.
48. Kim AJ, Lazar AA, Slutskiy YB. System identification of *Drosophila* olfactory sensory neurons. *J Comput Neurosci*. 2011;30(1):143–61.
49. Lotterhos KE, Yeaman S, Degner J, Aitken S, Hodgins KA. Modularity of genes involved in local adaptation to climate despite physical linkage. *Genome Biol*. 2018;19(1):1–24.
50. Yeaman S. Genomic rearrangements and the evolution of clusters of locally adaptive loci. *Proc Natl Acad Sci USA*. 2013;110(19):E1743–51.
51. Zong SB, Li YL, Liu JX. Genomic architecture of rapid parallel adaptation to fresh water in a wild fish. *Mol Biol Evol*. 2021;38(4):1317–29.
52. Veale AJ, Foster BJ, Dearden PK, Waters JM. Genotyping-by-sequencing supports a genetic basis for wing reduction in an alpine New Zealand stonefly. *Sci Rep*. 2018;8(1):1–12.
53. Celani A, Villermaux E, Vergassola M. Odor landscapes in turbulent environments. *Phys Rev X*. 2014;4(4):1–17.
54. Kree M, Duplat J, Villermaux E. The mixing of distant sources. *Phys Fluids*. 2013;25(9):091103.
55. Riffell JA, Shlizerman E, Sanders E, Abrell L, Medina B, Hinterwirth AJ, et al. Sensory biology. Flower discrimination by pollinators in a dynamic chemical environment. *Science*. 2014;344(6191):1515–8.
56. Vergassola M, Villermaux E, Shraiman BL. 'Infotaxis' as a strategy for searching without gradients. *Nature*. 2007;445(7126):406–9.
57. Vickers N. Mechanisms of animal navigation in odor plumes. *Biol Bull*. 2000;198(2):203–12.
58. Van BF, Dickinson MH. Article plume-tracking behavior of flying *Drosophila* emerges from a set of distinct sensory-motor reflexes. *Curr Biol*. 2014;24:274–86.
59. Mafra-Neto A, Cardé RT. Fine-scale structure of pheromone plumes modulates upwind orientation of flying moths. *Nature*. 1994;369(6476):142–4.
60. Murlis J, Elkinton J, Carde R. Odor plumes and how insects use them. *Annu Rev Entomol*. 1992;37(1):505–32.
61. Sehdev A, Mohammed YG, Triphan T, Szyszka P. Olfactory object recognition based on fine-scale stimulus timing in *Drosophila*. *iScience*. 2019;13:113–24.
62. Stierle JS, Galizia CG, Szyszka P. Millisecond stimulus onset-asynchrony enhances information about components in an odor mixture. *J Neurosci*. 2013;33(14):6060–9.
63. Saha D, Leong K, Li C, Peterson S, Siegel G, Raman B. A spatiotemporal coding mechanism for background-invariant odor recognition. *Nat Neurosci*. 2013;16(12):1830–9.
64. Baker TCT, Fadamiro HY, Cosse AAA. Moth uses fine tuning for odour resolution. *Nature*. 1998;393(6685):530–530.
65. Andersson MN, Binyameen M, Sadek MM, Schlyter F. Attraction modulated by spacing of pheromone components and anti-attractants in a bark beetle and a moth. *J Chem Ecol*. 2011;37(8):899–911.
66. Nikonov AA, Leal WS. Peripheral coding of sex pheromone and a behavioral antagonist in the Japanese beetle, *Popillia japonica*. *J Chem Ecol*. 2002;28(5):1075–89.
67. Sato K, Pellegrino M, Nakagawa T, Nakagawa T, Vosshall LB, Touhara K. Insect olfactory receptors are heteromeric ligand-gated ion channels. *Nature*. 2008;452(7190):1002–6.
68. Butterwick JA, del Marmol J, Kim KH, Kahlson MA, Rogow JA, Walz T, et al. Cryo-EM structure of the insect olfactory receptor Orco. *Nature*. 2018;560(7719):447–52.
69. Wicher D, Schäfer R, Bauernfeind R, Stensmyr MC, Heller R, Heinemann SH, et al. *Drosophila* odorant receptors are both ligand-gated and cyclic-nucleotide-activated cation channels. *Nature*. 2008;452(7190):1007–11.
70. Marden JH, Kramer MG. Surface-skimming stoneflies: a possible intermediate stage in insect flight evolution. *Science*. 1994;266(5184):427–30.
71. Niven JE, Laughlin SB. Energy limitation as a selective pressure on the evolution of sensory systems. *J Exp Biol*. 2008;211:1792–804.
72. Niven JE. Neuronal energy consumption: biophysics, efficiency and evolution. *Curr Opin Neurobiol*. 2016;41:129–35.
73. Leal WS. Odorant reception in insects: roles of receptors, binding proteins, and degrading enzymes. *Annu Rev Entomol*. 2013;58(1):373–91.
74. Rollmann SM, Mackay TFC, Anholt RRH. Pinocchio, a novel protein expressed in the antenna, contributes to olfactory behavior in *Drosophila melanogaster*. *J Neurobiol*. 2005;63(2):146–58.
75. McCormick AC, Grosse-Wilde E, Wheeler D, Mescher MC, Hansson BS, De Moraes CM. Comparing the expression of olfaction-related genes in gypsy moth (*Lymantria dispar*) adult females and larvae from one flightless and two flight-capable populations. *Front Ecol Evol*. 2017;5(SEP):115.
76. Nagai T. Summation and gradient characteristics of local electroantennogram response of the European corn borer *Ostrinia nubilalis*. *Pestic Biochem Physiol*. 1985;24(1):32–9.
77. Cao L-H, Jing B-Y, Yang D, Zeng X, Shen Y, Tu Y, et al. Distinct signaling of *Drosophila* chemoreceptors in olfactory sensory neurons. *Proc Natl Acad Sci USA*. 2016;113(7):E902–11.
78. Nagari M, Szyszka P, Galizia G, Bloch G. Task-related phasing of circadian rhythms in antennal responsiveness to odorants and pheromones in honeybees. *J Biol Rhythms*. 2017;32(6):593–608.
79. Szyszka P, Demmler C, Oemisch M, Sommer L, Biergens S, Birnbach B, et al. Mind the gap: olfactory trace conditioning in honeybees. *J Neurosci*. 2011;31(20):7229–39.
80. Raiser G, Galizia CG, Szyszka P. A high-bandwidth dual-channel olfactory stimulator for studying temporal sensitivity of olfactory processing. *Chem Senses*. 2016;42(2):bjw114.
81. Rebora M, Piersanti S, Frati F, Salerno G. Antennal responses to volatile organic compounds in a stonefly. *J Insect Physiol*. 2017;98:231–7.
82. R Core Team. R: a language and environment for statistical computing [Internet]. Vienna: R Foundation for Statistical Computing; 2020. Available from: <https://www.R-project.org>.
83. Rahim KJ. Applications of multitaper spectral analysis to nonstationary data. 2014.
84. Gelman A, Su Y-S. arm data analysis using regression and Multilevel Hierarchical Models [Internet]. 2018. Available from: <https://cran.r-project.org/package=arm>.
85. Neupert S, McCulloch GA, Foster BJ, Waters JM, Szyszka P. Adaptive evolution of olfactory degeneration in recently flightless insects. *bioRxiv*. 2020. <https://doi.org/10.1101/2020.04.10.035311>.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

