Emerging technologies revolutionise

- ² insect ecology and monitoring
- 3
- 4
- 5 Authors: Roel van Klink^{1,2}, Tom August³, Yves Bas⁴, Paul Bodesheim⁵, Aletta Bonn^{1,6,7},
- 6 Frode Fossøy⁸, Toke T. Høye⁹, Eelke Jongejans¹⁰, Myles H. M. Menz¹¹, Andreia
- 7 Miraldo¹², Tomas Roslin¹³, Helen E. Roy³, Ireneusz Ruczyński¹⁴, Dmitry Schigel¹⁵, Livia
- 8 Schäffler¹⁶, Julie K. Sheard¹⁷, Cecilie Svenningsen¹⁸, Georg F. Tschan¹⁶, Jana
- 9 Wäldchen¹⁹, Vera M. A. Zizka¹⁶, Jens Åström⁸, Diana E. Bowler^{1,6,7}
- 10
- 11
- 12 ¹German Centre for Integrative Biodiversity Research (iDiv) Halle Jena Leipzig, Puschstrasse 4,
- 13 04103, Leipzig, Germany
- 14 ²Martin Luther University-Halle Wittenberg, Department of Computer Science, , 06099, Halle
- 15 (Saale), Germany
- 16 ³UK Centre for Ecology & Hydrology, , Benson Lane, OX10 8BB, Wallingford, UK
- ⁴Muséum national d'Histoire naturelle, Centre d'Écologie et des Sciences de la Conservation, 57
- 18 rue Cuvier, 75005, Paris, France
- 19 ⁵Friedrich Schiller University Jena, Institute for Computer Science, Ernst-Abbe-Platz 2, 07743,
- 20 Jena, Germany
- ⁶Helmholtz Centre for Environmental Research UFZ, , Permoserstr. 15, 04318, Leipzig,
- 22 Germany

23 ⁷Friedrich Schiller University Jena, Institute of Biodiversity, Dornburger Str. 159, 07743, Jena,

24 Germany

- ⁸Norwegian Institute for Nature Research, Department for aquatic biodiversity, P.O. Box 5685
- 26 Torgarden, 7485, Trondheim, Norway
- ⁹Aarhus University, Department of Ecoscience and Arctic Research Centre, C.F. Møllers Allé 4-
- 28 8, 8000, Aarhus, Denmark
- ¹⁰Radboud University, Animal Ecology and Ecophysiology, Heyendaalseweg 135, 6525 AJ,
- 30 Nijmegen, The Netherlands
- ¹¹Max Planck Institute for Animal Behaviour, Department of Migration, Am Obstberg 1, 78315,
- 32 Radolfzell, Germany
- 33 ¹²Swedish Museum of Natural Sciences, Department of Bioinformatics and Genetics,
- 34 Frescativägen 40, 114 18, Stockholm, Sweden
- ¹³Swedish University of Agricultural Sciences (SLU), Department of Ecology, Ulls väg 18B,
- 36 75651, Uppsala, Sweden
- ¹⁴Mammal Research Institute, Polish Academy of Sciences, , Stoczek 1, 17-230, Białowieża,
- 38 Poland
- ¹⁵Global Biodiversity Information Facility (GBIF), Secretariat, Universitetsparken 15, 2100,
- 40 Copenhagen Ø, Denmark
- 41 ¹⁶Leibniz Institute for the Analysis of Biodiversity Change , Museum Koenig Bonn,
- 42 Adenauerallee 127, 53113, Bonn, Germany
- 43 ¹⁷University of Copenhagen, Center for Macroecology, Evolution and Climate, GLOBE Institute,
- 44 Universitetsparken 15, bld. 3, 2100, Copenhagen Ø, Denmark
- ¹⁸University of Copenhagen, Natural History Museum of Denmark, Oester Voldgade 5-7, 1350,
- 46 Copenhagen, Denmark
- 47 ¹⁹Max Planck Institute for Biogeochemistry, Department of Biogeochemical Integration, Hans-
- 48 Knoell-Str. 10, 07745, jena, Germany

49	Corres	sponding author: Van Klink, R. (roel.klink@idiv.de)	
50			
51	Key w	ords	
52	Entom	ology; barcoding; automated monitoring; radar; computer vision; eDNA	
53			
54			
55	Highlights		
56			
57	-	We appraise four emerging tools and technologies (computer vision, acoustic	
58		monitoring, radar and molecular methods) that provide unprecedented	
59		opportunities for insect monitoring and the study of insect ecology.	
60	-	These technologies have various benefits over traditional insect monitoring	
61		methods, including increased resolution of data collection across space and time,	
62		and a broader taxonomic coverage.	
63	-	At the same time, each technology has its limitations, some of which can be	
64		overcome with further methodological developments.	
65	-	Key issues regarding open science and international standards need to be	
66		addressed.	
67	-	While technology can never replace the knowledge of entomological specialists,	
68		we expect that integration of data across technologies, along with expert	
69		knowledge, will become commonplace in the future.	
70			
71			
72			

73 Glossary

Term	Definition
Acoustic sensor	A device that detects and records sounds.
AI (Artificial intelligence)	Scientific field of computer science interested in (partially) reproducing human skills, such as thinking, acting, or interpreting data, with computational algorithms. Often used as a synonym for machine learning (see below).
Citizen science	The participation of the general public in scientific processes. Participation can occur at different levels of involvement and expertise, and at different stages of the process (study design, data collection and/or interpretation).
Computer vision	Scientific field of computer science that develops algorithms for analysing image or video data to produce descriptions of the depicted content, e.g., a categorization via numerical representations.
(Convolutional) neural networks Deep learning	Group of machine learning methods that require large datasets for training, often used for image analysis and pattern recognition, where each network consists of connected nodes and layers that process input data to obtain desired outputs. General term that denotes the training and application of deep

	(convolutional) neural networks for a specific task, often used as a
	synonym for machine learning with deep neural networks.
Edge computing	Data processing done near the site of data collection, instead of transferring the data to a remote location for processing and analysis.
Environmental DNA (eDNA)	DNA obtained from environmental samples such as water from lakes or rivers, soil, faeces, air etc.
FAIR data	Data that are findable, accessible, interoperable and reusable.
Machine learning	Scientific field of computer science for developing predictive algorithms that learn patterns in data to make predictions. The algorithms learn from example training data rather than being programmed explicitly.
Metabarcoding	The recognition of species from their genetic structure, applied to multiple taxa contained in bulk or mixed samples. A common genetic region used for barcoding is mitochondrial cytochrome c oxidase subunit 1 (CO1).
Multi-sensor station	Observation device equipped with multiple sensors and recording units.
Radar	Device emitting radio waves in a certain direction to record the time, intensity and other features of the electromagnetic pulses that return from objects

	Traditional monitoring	Observations, usually by sight or trap, and species identification by
		human eye in the field or in the lab.
75		
76		
77		
78		
79		

80 Abstract

81 Insects are the most diverse group of animals on Earth, but their small size and high 82 diversity have always made them challenging to study. Recent technological advances 83 have the potential to revolutionise insect ecology and monitoring. We describe the state-84 of-the-art of four technologies (computer vision, acoustic monitoring, radar, and 85 molecular methods), and assess their advantages, current limitations and future 86 potential. We discuss how these technologies can adhere to modern standards of data 87 curation and transparency, their implications for citizen science, and their potential for 88 integration among different monitoring programs and technologies. We argue that they 89 provide unprecedented possibilities for insect ecology and monitoring but it will be 90 important to foster international standards via collaboration.

- 91
- 92

Technological advancement for insect monitoring 93

94 Insects are the most diverse group of eukaryotic organisms on Earth, comprising an estimated 95 80% of all animal life [1]. This staggering diversity, of which at least 80% remains undescribed, 96 forms a major challenge to studying insects and monitoring their responses to environmental 97 changes [2]. Recent reports of long-term declines in insect biomass and abundances [3,4], in 98 combination with the emergence of new technologies [5–7], have led to calls for [8], and the 99 establishment of, new research projects for monitoring populations and assemblages of insects 100 and other invertebrates [9,10]. 101 Traditionally, the monitoring of insects involves the killing of insects, followed by time-consuming 102 sorting and species identification by specialists [11]. Often the number of individuals and the 103 taxonomic diversity within a sample are so large that only a subgroup of taxa are identified, or 104 taxa are only identified to a coarse taxonomic level. Hence, there is a heavy bias towards 105 research on well-resolved groups, such as butterflies and ground beetles, whereas the diversity 106 of other taxa, e.g. most Diptera is often ignored [e.g. 12]. Additionally, the required human 107 labour for both data collection and processing limits the number of locations or the frequency of 108 sampling in traditional insect monitoring programs. 109 Recent development of technologies that employ novel detection and identification methods, 110 often in combination with citizen science, has opened up exciting new avenues for tracking 111 insect populations and assemblages [5-7,13]. These technologies, which include automated 112 image- and sound recognition, radar, and molecular methods, have the potential to radically 113 increase the spatial, temporal and taxonomic coverage of monitoring programs. They also allow 114 new questions to be asked about insect population dynamics, phenology and biotic interactions 115 (Box 1). At the same time, these technologies come with their own set of limitations, and are in 116 parallel development in different projects and countries. To ensure efficient progress, there is a 117

need for large-scale collaboration to develop international databases, metadata standards, and

open communication on hardware and software development, to ensure adherence to FAIRdata principles.

- 120 This paper is a collaborative effort of researchers from different European countries, aiming to
- 121 review emerging tools and technologies for insect monitoring and outlining a research agenda
- 122 that harnesses their potential. We provide an overview of (1) the state-of-the-art of these
- technologies, their advantages, current limitations, and future potential, (2) how the data
- 124 collected using these technologies can adhere to modern standards of data curation and
- transparency, (3) how citizens can participate in projects using these new technologies, and (4)
- 126 the potential for integration and synergies among technologies.
- 127
- 128

Box 1: Seeing the unseen using new technologies

Species interaction networks:

Interactions between species are often hard to detect due to the time, place or scale at which interactions take place, but modern technologies can help make the unseen visible. Molecular methods are being used to identify both the predators and food of insects directly, i.e. by analysing faeces [14], gut contents [15] (also from blood meals [16]), or parasite presence [17], and indirectly, using DNA left after resource visitation, such as on flowers [18]. Also computer vision has been used to quantify insect resource use and foraging behaviour [19].

Quantifying ecosystem services: Technologies are already being used to quantify insect pollination. Computer vision is applied to images taken by cameras fixed above plants [19–21] and metabarcoding can be used on pollen or flowers to identify flower visitors [19,22,23]. Computer vision or acoustic monitoring may also prove useful in studying the decomposition

of dung, carrion or dead plant matter, but has to our knowledge not yet been applied.

Tracking species movements and occurrences from local to continental scales:

For many ecological questions, as well as for biodiversity conservation, public health and crop protection, it is essential to track the whereabouts of specific insect species. Several of the technologies we discussed can help doing this. At the smallest scales, computer vision can track insects, such as pollinators, as they forage for resources [19,20], and eDNA can detect traces of past insect visits [18]. At regional to continental scales, different technologies can be used to detect the occurrence and movement of beneficial species [24], pest species [25–27], disease vectors [28], invasive species [29–31], and protected species [32].

Energy and biomass fluxes within and across habitats:

The movement of insects creates fluxes of nutrients and energy across large distances and across ecosystem boundaries (linking e.g. aquatic and terrestrial systems). Tracking these fluxes is now possible in 4 dimensions in a non-invasive and unbiased way [33,34]. Vertically-looking radar has been used to quantify high altitude insect migrations [35], and vertical photography and lidar can show insect biomass fluxes at closer ranges [33,36].

130

131

132

¹³⁴ Four technologies that are revolutionising

135 entomology

136 Computer vision

137 Computer vision is a field of computer science that develops algorithms to extract information 138 from digital images and video (Fig. 1A). In ecology, computer vision is being used in diverse 139 ways to collect observations and automize species identification. For instance, it is being used 140 for automated and standardised sampling of biodiversity, using cameras aimed at an 141 environmental feature [19] or at a screen placed in the field (see Box 2), often in combination 142 with traps (e.g., light traps [37], sticky traps [38] or pheromone traps [39]) to increase detection 143 rates. Computer vision is also helping to digitise the vast museum collections of specimens to 144 mobilise historic occurrence records [40,41]. Images are also being collected by citizen 145 scientists and uploaded to portals for opportunistic observations [42]. Several of these 146 applications (e.g. www.iNaturalist.org, www.observation.org/apps/obsidentify/, and 147 www.pictureinsect.com) support automated identification. While the technology has yet to be 148 applied at a large scale for insect monitoring, the first applications show promising results (Box 149 2). 150 151 There are a number of potential advantages to using computer vision for sampling and 152 identifying insects over traditional techniques. First, computer vision methods are often non-

153 destructive, so individuals don't need to be interfered with or killed. Second, computer vision

technology can count and classify insects with less human labour and observer bias [6]. Third,

155 by reducing the necessity for taxonomic expertise, computer vision is creating opportunities to

156 expand the engagement of citizen scientists (Box 3). Last, computer vision can be used to

157 collect information on insect behaviour and interactions. For instance, fixed cameras have been

158 mounted over resources such as flowers to record the activity of insects, including plant-

159 pollinator interactions [19,20].

160

161 Computer vision uses machine learning algorithms, such as convolutional neural networks 162 (CNNs), which are trained to identify insects using a library of pre-classified images. Accuracy 163 rates can be over 90% at the species-level for some taxa, but strongly depend on taxon group 164 size and morphological similarity [43–48]. In addition to taxonomic identification, algorithms are 165 being used to count individuals in an image and estimate individual size, biomass and 166 movement [49,50].

167

168 Several technical challenges are currently hindering the widespread application of computer 169 vision in entomological research. A main challenge is the large amounts of training data 170 (reference libraries) needed, which may be taxon, region, and project specific. Computers have 171 difficulties identifying species with limited training data (typically rare species), and tend to 172 overpredict species with a disproportionately large amount of training data (typically common 173 species). One solution to expand reference libraries is the development of apps for local experts 174 and citizens to submit training image data of species at different angles [51]. Another challenge 175 is the power consumption for the cameras and subsequent data transfer. This difficulty may be 176 reduced by the use of solar panels (Box 2), but this increases the risk of theft. For on-site 177 classification, internet connectivity is important, however, edge computing (local data 178 processing) enables classification directly on the device (e.g. the Seek app by iNaturalist 179 (https://www.inaturalist.org/pages/seek app) with the potential for real-time monitoring [19]. 180 Hence, while there still are challenges [6], already the opportunities of computer vision are 181 numerous and will likely transform insect monitoring in the coming decade (Fig. 2A).

182 Acoustic monitoring

183	A diverse range of insect taxa emit sounds that can be used for efficient monitoring. Like
184	computer vision, acoustic monitoring uses a field recorder to collect information (i.e., sounds), in
185	combination with machine learning algorithms for species identification (Fig. 1B). Insect sounds
186	may be sampled using stationary recorders or by mobile transects from cars or trains [52,53].
187	So far, these methods have mostly been applied to detect the chirping of insects such as
188	orthopterans and cicadas (Box 2), and have been tested on freshwater invertebrates [54,55],
189	and bees, hornets and mosquitoes based on their flight sounds [56,57], but they have a much
190	broader range of possibilities (Fig. 2B).
191	
192	One of the main advantages of acoustic monitoring over other sampling approaches, is that
193	insects can be detected over much longer ranges - sometimes more than 100 m [53].
194	Additionally, like computer vision, acoustic monitoring is nondestructive, rapid, and inexpensive
195	[58], and machine learning algorithms applied to the recorded sound circumvent observer
196	biases [53,59]. In addition to species presence, acoustic signals contain information on insect
197	behaviour, such as phenology, activity and courtship behaviour [52,53,60], and can provide
198	direct measures of ecological functions, such as pollination or wood-boring [57,61]. Recordings
199	of composite environmental sounds [62] - soundscapes - also contain rich information about the
200	state of biological assemblages related to species diversity [63].
201	

202 Identification of species from their sounds is still limited by the size of the reference libraries,

which are poorly developed compared to those for vertebrates [59]. Currently, these libraries are

- 204 only sufficiently large in temperate regions for some terrestrial vocalising insect groups, whereas
- 205 for the use of other insect sounds (especially flight sounds), reference libraries are largely
- 206 lacking (but see [56]). Citizen science schemes could, however, help build these acoustic

207 reference libraries [64]. There is also a strong need for research into the factors that influence

the detectability of insect sounds, including microphone type, weather, and vegetation

209 attenuation, to understand the sampling ranges. Nevertheless, acoustic monitoring has

210 underexplored potential for low-cost but large-scale monitoring (Fig 2B).

211

212 Radar

213 The application of remote sensing technologies for biodiversity monitoring has rapidly expanded 214 over the last decade. In entomology, radar monitoring uses terrestrial radar systems, including 215 weather surveillance radar, to detect insects in the airspace (Fig. 1C). It has been long known 216 that radar can detect large swarms of insects, but modern radar can provide detailed 217 information on flying insects, including size, shape, speed, trajectory and wing beat frequency 218 [65]. Specialised entomological radars can detect insects far above the ground, from 150 m 219 above ground level, with the potential to detect larger insects (i.e., >15 mg) up to 1.2 km above 220 ground level [65].

221

Advantages of monitoring insects by radar are that it's non-invasive, has large detection radius, and can operate day and night. Hence, radar observations are especially useful to study biomass fluxes [35], migratory behaviour [65], and population dynamics [24] (Box 1). Radar can also be used to reveal insect presence indirectly, by detecting signs of vegetation damage [25] or nest structures [66]. Data from weather surveillance radars have already been combined with local monitoring programs to document population declines in mayflies [67] and the movement of locust swarms [26].

229

230 Radar technologies have significant potential for large-scale monitoring of insects, even at the

continental-scale, using the existing networks of weather surveillance radars [34]. However,

they would benefit from improved algorithms for filtering biological targets from other airborne

233 particles, as well as increased knowledge of the reflective properties of insect taxa [68,69].

234

LiDAR (laser radar) has only recently been applied in entomology, but can be used to detect insects much closer to the ground than most radar systems, over sampling ranges of 10-600 m. Moreover, it has the potential to use the spectral reflectance to identify insects to much lower taxonomic levels [28,70,71]. As the technology develops, better taxonomic classification can be

achieved when libraries on spectral scatter become available for more taxa [13].

240

241

242 Molecular methods

Out of the modern technologies, molecular methods using genetic information are the most developed and most widely used so far. These methods can be used for many goals including the quick discovery of new species [72], the detection of endangered [32], invasive or pest species [73], the characterization of species interaction networks [18,74], and assessment of

taxonomic [22,75] and genetic diversity of whole assemblages [76,77].

248

249 The most common use of genetic information is based on DNA barcoding, i.e. amplification of a

short section of DNA from a specific gene or genes, providing adequate separation between

focal taxa. Barcoding was originally proposed for the identification of individual specimens [78].

- 252 Yet, advancements in laboratory protocols and high-throughput sequencing technologies now
- 253 enable DNA isolation, amplification and taxon identification from complex mixture samples (DNA

254 metabarcoding; Fig. 1D) [29]. Compared to traditional monitoring, metabarcoding can be time-255 and cost-efficient [72]) and is highly scalable, enabling simultaneous processing of many 256 samples and species. Metabarcoding methods can be applied directly to organismal samples, 257 using the storage medium [79] or homogenised bulk samples of collected insects [80]. 258 Alternatively, it is possible to detect the presence of species from DNA fragments in 259 environmental samples (eDNA), such as water [32], soil [81] or air [82]. Interactions between 260 insects and other taxa can be identified by using samples derived from animals guts, blood or 261 faeces [74] (Box 1). One of the most recent advances is the use of eRNA [83] to distinguish the 262 presence of living from dead individuals, since RNA is only present in metabolically active cells, 263 whereas DNA may derive from the remains of dead individuals. 264 265 Molecular methods overcome many of the observation biases associated with traditional 266 monitoring, however, they bring a new set of biases. Differences in DNA amounts and 267 extractability among insect taxa [79], or taxa-specific variation in PCR amplification [84,85], may 268 result in some species not being detected even when present in the sample. Size sorting within 269 a sample can help DNA amplification of small and rare species [86]. However, commonly used 270 markers, such as the CO1 gene, sometimes still fail to detect some insect taxa such as 271 Hymenoptera [87]. Amplification biases may also be circumvented by bypassing the PCR step 272 and directly sequencing the complete extracted DNA [75,88] or RNA (metatranscriptomics). 273 RNA sequencing also brings the potential to detect metabolic capacities and gene expression of 274 individuals or assemblages at the moment of sampling [89].

275

The primary outputs of DNA-technologies are gene-based operational units (operational taxonomic units [OTUs] and amplicon sequence variants [ASVs]; e.g. [90]), which represent clusters of organisms with similar DNA sequences. To link with existing knowledge of species and their ecology, these units must be mapped to reference databases, such as BOLD or GenBank.

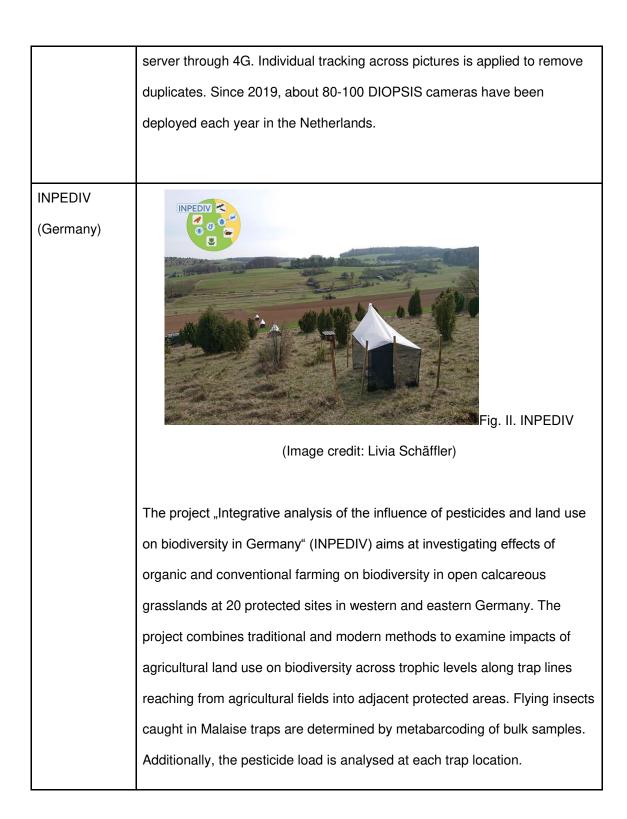
These reference databases are rapidly growing, BOLD now containing genetic data on 213,344 publicly available insect species. However, errors, synonymy, misidentifications and missing species can cause misclassifications. Nevertheless, international, national and taxon-specific initiatives are currently making strong progress on improving the taxonomic coverage of such reference libraries [80,91].

285

286

Box 2: Case in point: Pioneering monitoring projects

Case study	Brief description
DIOPSIS (Netherlands)	Fig. I. DIOPSIS. DIOPSIS (Digital Identification Of Photographically Sampled Insect Species) takes regular photos of a yellow screen that attracts insects and uses machine learning to recognize and count the photographed insects [92]. Photos are taken every 10 seconds or when movement is detected. If a photo is different from the previous one, it is stored locally and/or sent to a



Suivi des

Orthoptères Nocturnes

(France)



nocturnes

Fig. II. Suivi des Orthoptères Nocturnes.

In France, nocturnally vocalising bush crickets have been monitored by citizen scientists since 2006, as an add-on to the acoustic bat monitoring scheme Vigie-Chiro. For this purpose, Tadarida software was developed to detect both bat and insect calls and classify them into 79 classes including all common bat and bush-cricket species, using a random forest algorithm [93]. This nationwide monitoring scheme, with so far 16 349 individual sampling locations, has detected significant declines of four bush-cricket species, and an increase of *Phaneroptera nana*.

Insektmobilen

(Denmark /

Germany)

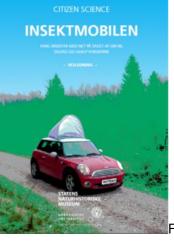
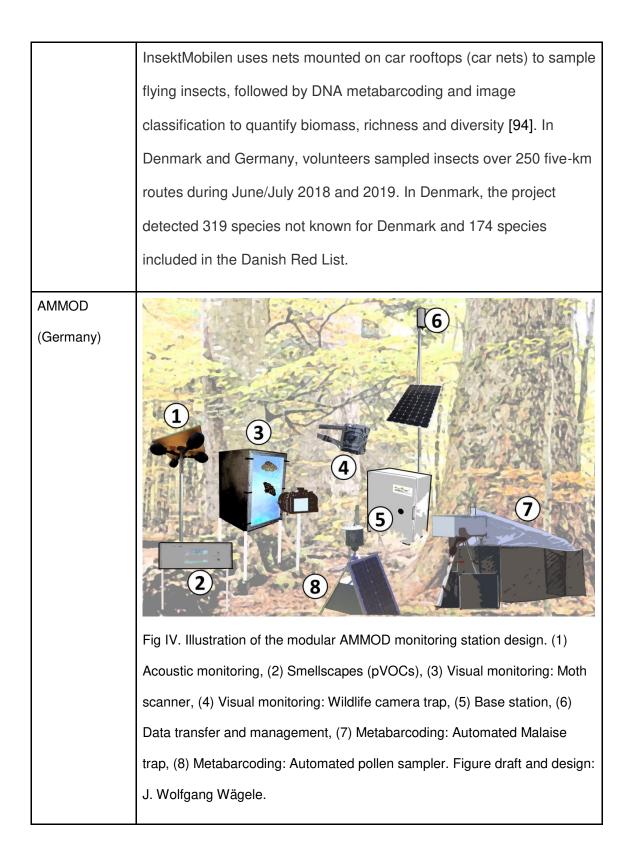


Fig. III. Insektmobilen (Image credit: Anders

Drud, Natural History Museum of Denmark.)



	AMMODs are Automated Multisensor stations for Monitoring of species
	Diversity [9]. Analogous to weather stations, these are autonomous
	samplers that monitor plants, birds, mammals and insects. The technology
	consists of six modules: (i) automatized visual monitoring and image
	analyses (mammals and moths), (ii) detection of smellscapes using volatile
	organic compounds, (iii) Malaise and pollen traps for metabarcoding, (iv)
	automated bioacoustic monitoring (birds and bats), (v) development of a
	base station, and (vi) data management and cross-platform analysis. Since
	2020, AMMOD is being tested at three sites in Germany.
288	
289	
290	

²⁹¹ Box 3: New technologies as opportunities to advance

292 citizen science

293

About 70-80 % of species records in Europe are collected by volunteers [95]and these data underpin many national and regional assessments of biodiversity change [96,97]. Historically, most insect monitoring was organised outside academia, especially by taxonomic specialists and natural history societies [98], and there is a long tradition of including lay people in the scientific data collection process for various insect taxa including butterflies and <u>mosquitoes</u> [95]. Recent technological developments have increased the opportunities for people, including non-specialists, to get involved, for example helping with digitization of museum collections (e.g. <u>Denmark</u>).

Out of the new technologies, computer vision has been most often integrated into citizen science; for example, a range of smartphone applications use computer vision to help users identify species (e.g. <u>iNaturalist</u>, <u>ObsIdentify</u>, and <u>Picture Insect</u>, <u>lepsnap</u>). Many of these applications use a so-called 'human-in-the-loop' approach - the technology helps users narrow down the likely species by suggesting the most visually similar species. Another way citizens have contributed to computer vision-based science is by helping to compile the training data needed for machine learning, for example in the <u>PollinatorWatch</u> project. In projects using DNA-technology, some rely on citizen scientists for the collection of the insect samples [94], which are subsequently processed by scientists. A few citizen science projects are starting to include citizens in the analysis steps e.g., the DNA&life project in Denmark [99].

Ecologists often debate the reliability of species observations from citizen science. However, the development of AI based apps [100] and DNA based methods [31] may help increase identification accuracy. For instance, AI tools may allow instant feedback on the likelihood of an observation. AI methods can also be used to develop intuitive field guides that may deviate from classic dichotomous identification keys. Some citizen science platforms already use crowd-sourced expert identification for validation of observation (such as expert crowd verification of iNaturalist or iRecord observations); however, the rate of manual validation is unable to keep pace with the rapidly growing number of submissions. In these cases, technologies could use active learning AI algorithms, which select only a subset of images for human validation for i) groundtruthing or training of the AI classifier, and ii) where the AI

classifier was most uncertain in its decision. Citizens with taxonomic expertise may also help compile the training datasets by identifying species on images or sounds.

New technologies have the potential to increase the accessibility and diversity of entomological citizen science. For instance, citizen science activities could be extended to volunteers with expertise in joint software development and data visualisation. Care, however, needs to be taken to avoid access barriers and unintended exclusion due to possible technology barriers or disconnect of data, people and wildlife. Overall, there could be considerable benefits from involving citizen scientists in the development and application of the tools through co-created projects and community partnerships [98].



Fig. V. Using automated identification technology to monitor insects can be a win-win situation for citizens and scientists. Using such tools, citizens can learn about species identity and ecology, and scientists can use the data collected to study, for example, species interactions, such as this lady beetle feeding on aphids on their host plant.

294

295 The road forward

296 The development of new technologies for insect ecology and monitoring is no goal in itself, but 297 must be guided by the needs of society, policy makers, as well as the scientific questions 298 scientists address (Box 1). Furthermore, they must meet the demands of modern science in 299 terms of data curation and transparency [101], and consider the possibility of involvement of 300 other stakeholders, such as citizens (Box 3), who have contributed 70-80 % of species 301 occurrence records in Europe [95]. There are also un(der)explored possibilities for integration 302 among technologies. In the next sections, we will outline the opportunities for how these 303 technologies can revolutionise insect ecology and monitoring. 304

305 Open Science

- 306 Insect data collected by traditional monitoring schemes or derived from museum specimens are
- 307 becoming increasingly accessible via data discovery platforms such as <u>www.GBIF.org</u>.
- 308 However, for data collected using the technologies discussed here, the norms and practices of

open science, as well as standards for data publishing have yet to evolve and to be agreedupon. To make these new technologies open and reproducible, both the underlying data and

- 311 processing steps must be FAIR: findable, accessible, interoperable, and reusable [101].
- 312

While generalist data repositories may function sufficiently well for basic data access and retrieval, specialist repositories are often needed for the most efficient data re-use. Data openness has been fostered for DNA-based technologies through International Nucleotide Sequence Database Collaboration (www.<u>INSDC</u>.org) data portals such as the Sequence Reed Archive (<u>SRA</u>). GBIF has also led the development of protocols to deal with sequence data to improve discoverability of DNA derived data [102]. For sharing species images, there are various citizen scientist platforms, but fewer for audio recordings (but see www.iNaturalist.org).

320

New technologies face practical problems about which form of data should be stored due to the typically large file sizes or novel data attributes. To ensure comparability over time, data should be stored in their original form, so the data can be reprocessed when reference libraries or technologies improve and enable better species detection and/or classification.

325

326 Standardisation and quality of data and metadata are key for interoperability and reusability.

327 Among the most widespread are Ecological Metadata Language (EML [103]) and Darwin Core

328 [104]. Yet, it is still unclear what metadata would be sufficient for reproducibility of data collected

329 by different technologies or different protocols [105]. Technological reproducibility also needs to

involve openness of hardware (type, model, as well as mechanical, electrical and optical

331 settings), and software (version, documentation), and the availability of analytical code as a

332 community norm. For DNA technologies, specific steps of the laboratory protocols, such as

333 preservation buffer, DNA polymerase, PCR enhancer, are essential for reproducibility [106] and

automated workflows are being proposed for standardisation [102,107].

335 The potential of technological integration

336	Each of the reviewed technologies has its own strengths and weaknesses, and new studies
337	should seek to combine the strengths of the different technologies, as well as integrate the
338	continued value of traditional monitoring methods. Combining different technologies could bring
339	a range of benefits: increased spatial, temporal or taxonomic coverage, a broader range of
340	biodiversity metrics, or simply more confident taxonomic assignment. Integration is also likely to
341	be the optimal solution for effective large-scale and long-term insect monitoring. Some
342	examples of complementary use of methods already exist. Below, we outline some possibilities:
343	
344	(1) Quantification of different biodiversity metrics in insect bulk samples
345	A combination of technologies applied to the same sample can increase the range of produced
346	biodiversity metrics. While molecular methods provide reliable biodiversity metrics such as
347	richness and diversity, traditional methods [108] and computer vision [50] still provide more
348	robust quantitative metrics such as biomass and abundance. Although we note that methods
349	are being tested to quantify species abundances and biomass from DNA samples [109].
350	
351	The onset of robotic techniques for the processing of individual insects from bulk samples [110]
352	may to a large extent replace the laborious work of manual species identification. Together,
353	computer vision, robotic sorting, and DNA-based identification may add both images and DNA-
354	sequences of previously unencountered taxa to reference libraries, provide all desired
355	biodiversity metrics, and discover new and rare species for further processing by taxonomic
356	specialists. So far, only a prototype for this approach exists [110], but this combination of
357	technologies can upscale biodiversity monitoring to unprecedented levels.
358	

359 (2) Increasing confidence of species identification

360 The integration of different technologies may improve identification accuracy and coverage of 361 the insects in a sample. Integration could occur either during the taxonomic classification step, 362 as a multi-sensor input for the neural networks (so-called cross-modal perception), which may 363 work especially well for combined visual and acoustic monitoring. Alternatively, integration may 364 occur as a final step to check for overall concordance of the classifications between the 365 technologies; for instance, following analysis of insect bulk samples by computer vision and 366 DNA analysis. Integrating optical and acoustic sensors may be especially useful for developing 367 pollinator indicators, which are especially urgent given their key role in ecosystems, but difficult 368 to accurately identify based on each technology alone.

369

370 (3) Filling the gaps: increased spatial, temporal and taxonomic coverage

371 Due to the decreased human labour needed, new technologies can increase the spatial,

372 temporal and taxonomic coverage of monitoring programs. To align with existing schemes, new

technologies could be initially set up to target current spatial and temporal gaps e.g., when and

374 where fewer people are active, such as in remote areas. Another way of upscaling monitoring to

375 large spatial scales with great potential is the use of (weather) radar. Although radar currently

376 largely lacks certainty about species identity, it could also be combined with vertical

377 photography [33] and aerial eDNA [82] to sample the same aerospace.

378

379 For assessment of whole ecological assemblages within a region, multi-sensor biodiversity

380 'weather' stations [9] may become particularly useful. These stations simultaneously use

381 multiple technologies and trap types to monitor a broad range of organisms, including insects,

382 plants and vertebrates (see AMMOD project in Box 2). Such monitoring is especially useful to

383 understand trophic links and for monitoring overall ecosystem health.

384

385

386 On-going role of traditional monitoring

387 Regardless of technological developments, new technologies cannot replace specialist 388 taxonomic knowledge and traditional methods [111]. Instead, new technologies should seek to 389 complement traditional monitoring, to alleviate workload and tedious tasks, and to increase the 390 spatial, temporal and taxonomic coverage of existing monitoring schemes. Furthermore, 391 combining metabarcoding and microscopy has been shown to increase the level of species 392 identification [108]. 393 Entomological expertise is also needed for building and improving reference libraries and the 394 validation of the results from automated monitoring. Moreover, there are still some groups that 395 can be poorly distinguished by modern technology, e.g. morphologically similar taxa (such as 396 ants) or taxa that are poorly distinguishable by commonly used barcoding genes [112]. As novel 397 methods continue to emerge and may eventually dominate the records, expert checks will 398 become crucial to ensure data quality. 399 Another area where human labour will remain essential is the detection of protected species, 400 which are rare and not allowed to be trapped, such as those under the European Commission 401 Habitats Directive for Annex I. For aquatic species, eDNA may be a viable option, but for 402 monitoring rare terrestrial habitat specialists, such as the hermit beetle Osmoderma eremita or 403 the Great Capricorn Beetle Cerambyx cerdo, human observations will remain essential.

404

406 Outstanding questions

407	٠	What standards are needed to ensure interoperability of data and reproducibility
408		of methods using each technology? How should these standards be fostered by
409		the scientific and biodiversity informatics community?
410		
411	•	How do these new technologies compare to traditional methods in accuracy and
412		informativeness? Are new technologies just quicker and cheaper? When should
413		we still use traditional taxonomy and sampling, and how can new technologies
414		and traditional sampling best complement each other?
415		
416	•	What are the sampling biases of new technologies, in terms of both taxonomic
417		accuracy (which groups can be monitored by each technology) as well as species
418		traits (which kinds of species might be missed by each technology)?
419		
420	•	How should the technologies be optimised to maximize both data quantity and
421		data quality, bearing in mind that many of the new technologies may rapidly
422		increase the rate of data collection, outpacing storage capacities and/or the rate at
423		which the data can be validated, processed and analysed?
424		
425	٠	How should data aggregation services (such as GBIF and national records) define
426		and label data from novel identification techniques, including the uncertainty in
427		species identity and non-validated insect records, so that these uncertainties are
428		transparent to the end-user?
429		

430	٠	How can new technologies overcome common biases associated with the effects
431		of insect activity on monitoring indices and provide unbiased density estimates?
432		
433	٠	What is the best sampling design to upscale and integrate different technologies
434		for large-scale and long-term insect monitoring?
435		
436	•	How can new technologies facilitate and enhance engagement with society,
437		promote experience of and learning about insects, and foster meaningful and
438		innovative citizen science?
439		
440	•	How can we ensure that the outputs of these technologies align with policy-
441		relevant indicators for ecological states and trends at relevant spatial and
442		temporal scales?
443		

444 Concluding remarks

445 The technological developments described in this paper provide unprecedented possibilities for 446 entomological research and monitoring. However, most of them are still in a proof-of-concept 447 stage and are not ready for large scale deployment, and none of them is free of biases (see 448 Outstanding Questions). While these technologies cannot replace specialist taxonomic 449 knowledge, they can help save time on species identification, and some can enable non-lethal 450 monitoring. Existing monitoring programs using traditional methods have proven invaluable for 451 understanding the extent of recent insect declines [3], and should be maintained to extend 452 historic time-series. Before new technologies can be deployed for large-scale insect monitoring, 453 international standards need to be developed via collaboration across borders, projects and

454 technologies. It will also be crucial to involve different stakeholders to develop policy-relevant
455 indicators, so that the data collected can be truly and broadly useful. The future of entomology
456 will be a collaboration between human and machine.

457

458 Acknowledgements

459 This paper is the result of a workshop funded by the Volkswagen Stiftung to DEB and RvK. 460 RvK, AB and DEB also acknowledge support by iDiv funded by the German Research 461 Foundation (DFG-FZT 118, 202548816). The LIB and consortium partners acknowledge funding 462 of INPEDIV by the Leibniz Association (project K120/2018) and of DINA by the BMBF (FKZ 463 16LC1901G). The members of the AMMOD Team would like to thank the BMBF for financing 464 the project (FKZ 16LC1903A). TTH was funded by EU Horizon 2020 Research and Innovation 465 programme (Grant Agreement no. 773554 (EcoStack)). FF, JC and JÅ were funded by 466 Norwegian environmental agency (ref: 18087129 - 2018/5765). IR was funded by the Polish 467 National Science Centre (DEC-2013/10/E/ NZ8/00725). AM was funded by the Knut and Alice 468 Wallenberg Foundation (KAW 2017.0088). HER was funded by the UK Natural Environment 469 Research Council award number NE/R016429/1 as part of the UK-SCAPE programme 470 Delivering National Capability. TR was funded by the European Research Council Synergy 471 Grant (856506 - LIFEPLAN). JKS was funded by the Danish National Research Foundation 472 (DNRF96). CS was funded by the Aage V. Jensen Nature Foundation. Gabriele Rada created 473 the figures. 474

475

476

477 References

- 478 1 Chapman, A.D. (2009) Numbers of Living Species in Australia and the World. 2nd edition,
- 479 Australian Biological Resources Services.
- 480 2 Cardoso, P. et al. (2011) The seven impediments in invertebrate conservation and how to
- 481 overcome them. *Biol. Conserv.* 144, 2647–2655
- 482 3 Van Klink, R. *et al.* (2020) Meta-analysis reveals declines in terrestrial but increases in
- 483 freshwater insect abundances. *Science* 368, 417–420
- 484 4 Hallmann, C.A. et al. (2017) More than 75 percent decline over 27 years in total flying insect
- 485 biomass in protected areas. *PLoS ONE* 12, e0185809
- 486 5 Bálint, M. et al. (2018) Environmental DNA Time Series in Ecology. Trends Ecol. Evol. 33,
- 487 945–957
- 488 6 Høye, T.T. *et al.* (2021) Deep learning and computer vision will transform entomology. *Proc.*489 *Natl. Acad. Sci.* 118, 2002545117
- 490 7 Tosa, M.I. et al. (2021) The Rapid Rise of Next-Generation Natural History. Front. Ecol. Evol.
- 491 0, 480
- 492 8 Saunders, M.E. et al. (2020) Moving On from the Insect Apocalypse Narrative: Engaging with
- 493 Evidence-Based Insect Conservation. *BioScience* 70, 80–89
- 494 9 Wägele, J.W. et al. (2022) Towards a multisensor station for automated biodiversity
- 495 monitoring. *Basic Appl. Ecol.* DOI: 10.1016/j.baae.2022.01.003
- 496 10 Lehmann, G.U.C. et al. (2021) Diversity of Insects in Nature protected Areas (DINA): an
- 497 interdisciplinary German research project. *Biodivers. Conserv. 2021 308* 30, 2605–2614
- 498 11 Montgomery, G.A. et al. (2021) Standards and Best Practices for Monitoring and
- 499 Benchmarking Insects. Front. Ecol. Evol. 0, 513
- 500 12 Van Klink, R. *et al.* (2015) Effects of large herbivores on grassland arthropod diversity.
- 501 *Biol. Rev.* 90, 347–366

502 13 Brydegaard, M. and Jansson, S. (2019) Advances in entomological laser radar. *J. Eng.*503 2019, 7542–7545

- 504 14 Mata, V.A. et al. (2021) Combining DNA metabarcoding and ecological networks to
- 505 inform conservation biocontrol by small vertebrate predators. *Ecol. Appl.* 31, e02457
- 506 15 Masonick, P. *et al.* (2019) No guts, no glory: Gut content metabarcoding unveils the diet
- 507 of a flower-associated coastal sage scrub predator. *Ecosphere* 10, e02712
- Massey, A.L. *et al.* Invertebrates for vertebrate biodiversity monitoring: Comparisons
 using three insect taxa as iDNA samplers. *Mol. Ecol. Resour.* n/a,
- 510 17 Hrcek, J. et al. (2011) Molecular detection of trophic links in a complex insect host-
- 511 parasitoid food web. Mol. Ecol. Resour. 11, 786–794
- 512 18 Tiusanen, M. et al. (2019) Flower-visitor communities of an arcto-alpine plant—Global
- 513 patterns in species richness, phylogenetic diversity and ecological functioning. *Mol. Ecol.* 28,
- 514 318–335

515 19 Bjerge, K. *et al.* Real-time insect tracking and monitoring with computer vision and deep

516 learning. Remote Sens. Ecol. Conserv. n/a,

517 20 Ratnayake, M.N. et al. (2021) Tracking individual honeybees among wildflower clusters

- 518 with computer vision-facilitated pollinator monitoring. *PLOS ONE* 16, e0239504
- 519 21 Pegoraro, L. *et al.* (2020) Automated video monitoring of insect pollinators in the field.
- 520 Emerg. Top. Life Sci. 4, 87–97
- 521 22 Thomsen, P.F. and Sigsgaard, E.E. (2019) Environmental DNA metabarcoding of wild
- flowers reveals diverse communities of terrestrial arthropods. Ecol. Evol. 9, 1665–1679
- 523 23 Lucas, A. et al. (2018) Generalisation and specialisation in hoverfly (Syrphidae)
- 524 grassland pollen transport networks revealed by DNA metabarcoding. J. Anim. Ecol. 87,

525 1008-1021

- 526 24 Wotton, K.R. *et al.* (2019) Mass Seasonal Migrations of Hoverflies Provide Extensive
- 527 Pollination and Crop Protection Services. *Curr. Biol.* 29, 2167-2173.e5

- 528 25 Ghulam Rasool, K. et al. (2020) Evaluation of some non-invasive approaches for the
- 529 detection of red palm weevil infestation. Saudi J. Biol. Sci. 27, 401–406
- 530 26 Amarjyothi, K. et al. (2022) Identification and Tracking of Locust Swarms by Indian
- 531 Doppler Weather Radar. *IEEE Geosci. Remote Sens. Lett.* 19, 1–4
- 532 27 Wang, R. et al. (2022) An automatic system for pest recognition and forecasting. Pest
- 533 Manag. Sci. 78, 711–721
- 534 28 Brydegaard, M. et al. (2020) Lidar reveals activity anomaly of malaria vectors during
- 535 pan-African eclipse. *Sci. Adv.* DOI: 10.1126/sciadv.aay5487
- 536 29 Piper, A.M. et al. (2019) Prospects and challenges of implementing DNA metabarcoding
- 537 for high-throughput insect surveillance. *GigaScience* 8, giz092
- 538 30 Comtet, T. et al. (2015) DNA (meta)barcoding of biological invasions: a powerful tool to
- 539 elucidate invasion processes and help managing aliens. *Biol. Invasions* 17, 905–922
- 540 31 Larson, E.R. et al. (2020) From eDNA to citizen science: emerging tools for the early

541 detection of invasive species. *Front. Ecol. Environ.* 18, 194–202

- 542 32 Doi, H. et al. Detection of an endangered aquatic heteropteran using environmental DNA
- 543 in a wetland ecosystem. R. Soc. Open Sci. 4, 170568
- 544 33 Ruczyński, I. et al. (2020) Camera transects as a method to monitor high temporal and
- 545 spatial ephemerality of flying nocturnal insects. *Methods Ecol. Evol.* 11, 294–302
- 546 34 Bauer, S. et al. (2017) From Agricultural Benefits to Aviation Safety: Realizing the
- 547 Potential of Continent-Wide Radar Networks. *BioScience* 67, 912–918
- 548 35 Hu, G. *et al.* (2016) Mass seasonal bioflows of high-flying insect migrants. *Science* 354,
 549 1584–1587
- 550 36 Brydegaard, M. et al. (2015), Daily evolution of the insect biomass spectrum in an
- agricultural landscape accessed with lidar., presented at the ILRC27, New York
- 552 37 Bjerge, K. et al. (2021) An Automated Light Trap to Monitor Moths (Lepidoptera) Using
- 553 Computer Vision-Based Tracking and Deep Learning. Sensors 21, 343

- 554 38 Gerovichev, A. *et al.* (2021) High Throughput Data Acquisition and Deep Learning for 555 Insect Ecoinformatics. *Front. Ecol. Evol.* 9,
- 556 39 Yalcin, H. (2015), Vision based automatic inspection of insects in pheromone traps., in
- 557 2015 Fourth International Conference on Agro-Geoinformatics (Agro-geoinformatics), pp.
- 558 333–338
- 559 40 Wilson, R.J. *et al.* Applying computer vision to digitised natural history collections for
- 560 climate change research: temperature-size responses in British butterflies. . 22-Dec-(2021) ,
- 561 bioRxiv, 2021.12.21.473511
- 562 41 Carranza-Rojas, J. et al. (2017) Going deeper in the automated identification of
- 563 Herbarium specimens. BMC Evol. Biol. 17, 181
- 564 42 Wäldchen, J. and Mäder, P. (2018) Machine learning for image based species
- identification. *Methods Ecol. Evol.* 9, 2216–2225
- Martineau, M. *et al.* (2017) A survey on image-based insect classification. *Pattern Recognit.* 65, 273–284
- 568 44 Knyshov, A. et al. (2021) Pretrained Convolutional Neural Networks Perform Well in a
- 569 Challenging Test Case: Identification of Plant Bugs (Hemiptera: Miridae) Using a Small
- 570 Number of Training Images. Insect Syst. Divers. 5, 3
- 571 45 Spiesman, B.J. et al. (2021) Assessing the potential for deep learning and computer
- 572 vision to identify bumble bee species from images. Sci. Rep. 2021 111 11, 1–10
- 573 46 Milošević, D. et al. (2020) Application of deep learning in aquatic bioassessment:
- 574 Towards automated identification of non-biting midges. *Sci. Total Environ.* 711, 135160
- 575 47 Valan, M. *et al.* (2019) Automated Taxonomic Identification of Insects with Expert-Level
- 576 Accuracy Using Effective Feature Transfer from Convolutional Networks. Syst. Biol. 68, 876–
- 577 895
- 578 48 Korsch, D. et al. (2021) Deep Learning Pipeline for Automated Visual Moth Monitoring:
- 579 Insect Localization and Species Classification, Gesellschaft für Informatik, Bonn.

- 580 49 Bruijning, M. et al. (2018) trackdem: Automated particle tracking to obtain population
- 581 counts and size distributions from videos in r. *Methods Ecol. Evol.* 9, 965–973
- 582 50 Schneider, S. *et al.* Bulk arthropod abundance, biomass and diversity estimation using 583 deep learning for computer vision. *Methods Ecol. Evol.* n/a,
- 584 51 Boho, D. et al. (2020) Flora Capture: a citizen science application for collecting
- 585 structured plant observations. *BMC Bioinformatics* 21, 576
- 586 52 Newson, S.E. et al. (2017) Potential for coupling the monitoring of bush-crickets with
- 587 established large-scale acoustic monitoring of bats. *Methods Ecol. Evol.* 8, 1051–1062
- 588 53 Jeliazkov, A. et al. (2016) Large-scale semi-automated acoustic monitoring allows to
- 589 detect temporal decline of bush-crickets. *Glob. Ecol. Conserv.* 6, 208–218
- 590 54 van der Lee, G.H. et al. (2020) Freshwater ecoacoustics: Listening to the ecological
- 591 status of multi-stressed lowland waters. *Ecol. Indic.* 113, 106252
- 592 55 Linke, S. et al. (2018) Freshwater ecoacoustics as a tool for continuous ecosystem
- 593 monitoring. Front. Ecol. Environ. 16, 231–238
- 594 56 Kiskin, I. et al. (2021) HumBugDB: A Large-scale Acoustic Mosquito Dataset.
- 595 ArXiv211007607 Cs Eess at http://arxiv.org/abs/2110.07607>
- 596 57 Kawakita, S. and Ichikawa, K. (2019) Automated classification of bees and hornet using
- 597 acoustic analysis of their flight sounds. *Apidologie* 50, 71–79
- 598 58 Hill, A.P. et al. (2018) AudioMoth: Evaluation of a smart open acoustic device for
- 599 monitoring biodiversity and the environment. *Methods Ecol. Evol.* 9, 1199–1211
- 600 59 Gibb, R. et al. (2019) Emerging opportunities and challenges for passive acoustics in
- 601 ecological assessment and monitoring. *Methods Ecol. Evol.* 10, 169–185
- 602 60 Sueur, J. et al. (2021) Acoustic biodiversity. Curr. Biol. 31, R1172–R1173
- 603 61 Mankin, R.W. et al. (2011) Perspective and Promise: a Century of Insect Acoustic
- 604 Detection and Monitoring. Am. Entomol. 57, 30–44
- 605 62 Burivalova, Z. et al. (2021) The sound of logging: Tropical forest soundscape before,

during, and after selective timber extraction. *Biol. Conserv.* 254, 108812

607 63 Aide, T.M. et al. (2017) Species Richness (of Insects) Drives the Use of Acoustic Space

- 608 in the Tropics. *Remote Sens.* 9, 1096
- 609 64 Aodha, O.M. *et al.* (2018) Bat detective—Deep learning tools for bat acoustic signal
- 610 detection. PLOS Comput. Biol. 14, e1005995
- 611 65 Chapman, J.W. *et al.* (2011) Recent Insights from Radar Studies of Insect Flight. *Annu.*612 *Rev. Entomol.* 56, 337–356
- 66 Rhodes, M.W. *et al.* (2022) Recent advances in the remote sensing of insects. *Biol. Rev.*614 97, 343–360
- 615 67 Stepanian, P.M. et al. (2020) Declines in an abundant aquatic insect, the burrowing
- 616 mayfly, across major North American waterways. Proc. Natl. Acad. Sci. 117, 2987–2992
- 617 68 Drake, V.A. et al. (2017) Ventral-aspect radar cross sections and polarization patterns of
- 618 insects at X band and their relation to size and form. *Int. J. Remote Sens.* 38, 5022–5044
- 619 69 Mirkovic, D. et al. (2019) Characterizing animal anatomy and internal composition for
- 620 electromagnetic modelling in radar entomology. *Remote Sens. Ecol. Conserv.* 5, 169–179
- 621 70 Gebru, A. et al. (2018) Multiband modulation spectroscopy for the determination of sex
- and species of mosquitoes in flight. J. Biophotonics 11, e201800014
- 623 71 Kirkeby, C. et al. (2021) Advances in automatic identification of flying insects using
- 624 optical sensors and machine learning. Sci. Rep. 11, 1555
- 625 72 Srivathsan, A. et al. (2021) ONTbarcoder and MinION barcodes aid biodiversity
- discovery and identification by everyone, for everyone. *BMC Biol.* 19, 217
- 627 73 Batovska, J. *et al.* (2021) Developing a non-destructive metabarcoding protocol for
- detection of pest insects in bulk trap catches. *Sci. Rep.* 11, 7946
- 629 74 Clare, E.L. *et al.* (2019) Approaches to integrating genetic data into ecological networks.
- 630 Mol. Ecol. 28, 503–519
- 531 75 Ji, Y. et al. (2020) SPIKEPIPE: A metagenomic pipeline for the accurate quantification of

- 632 eukaryotic species occurrences and intraspecific abundance change using DNA barcodes or
- 633 mitogenomes. Mol. Ecol. Resour. 20, 256–267
- 634 76 Elbrecht, V. *et al.* (2018) Estimating intraspecific genetic diversity from community DNA
 635 metabarcoding data. *PeerJ* 6, e4644
- 636 77 Zizka, V.M.A. et al. (2020) Can metabarcoding resolve intraspecific genetic diversity
- 637 changes to environmental stressors? A test case using river macrozoobenthos.
- 638 *Metabarcoding Metagenomics* 4, e51925
- 639 78 Hebert, P.D.N. *et al.* (2003) Biological identifications through DNA barcodes. *Proc. R.*
- 640 Soc. Lond. B Biol. Sci. 270, 313–321
- 641 79 Marquina, D. et al. (2019) Establishing arthropod community composition using
- 642 metabarcoding: Surprising inconsistencies between soil samples and preservative ethanol
- and homogenate from Malaise trap catches. *Mol. Ecol. Resour.* 19, 1516–1530
- 80 Roslin, T. et al. (2022) A molecular-based identification resource for the arthropods of
- 645 Finland. *Mol. Ecol. Resour.* 22, 803–822
- 646 81 Noguerales, V. et al. Community metabarcoding reveals the relative role of
- 647 environmental filtering and spatial processes in metacommunity dynamics of soil
- 648 microarthropods across a mosaic of montane forests. *Mol. Ecol.* n/a,
- 649 82 Roger, F. et al. Airborne environmental DNA metabarcoding for the monitoring of
- terrestrial insects a proof of concept. . 27-Jul-(2021) , bioRxiv, 2021.07.26.453860
- 651 83 Cristescu, M.E. (2019) Can Environmental RNA Revolutionize Biodiversity Science?
- 652 Trends Ecol. Evol. 34, 694–697
- Elbrecht, V. et al. (2019) Validation of COI metabarcoding primers for terrestrial
- arthropods. *PeerJ* 7, e7745
- 655 85 Marquina, D. et al. (2019) New mitochondrial primers for metabarcoding of insects,
- designed and evaluated using in silico methods. Mol. Ecol. Resour. 19, 90–104
- 657 86 Elbrecht, V. et al. (2021) Pooling size sorted Malaise trap fractions to maximize taxon

- 658 recovery with metabarcoding. *PeerJ* 9, e12177
- 659 87 Krehenwinkel, H. et al. (2017) Estimating and mitigating amplification bias in qualitative
- and quantitative arthropod metabarcoding. Sci. Rep. 7, 17668
- 661 88 Greenfield, P. et al. (2019) Kelpie: generating full-length 'amplicons' from whole-
- 662 metagenome datasets. *PeerJ* 6, e6174
- 663 89 Cordier, T. et al. (2021) Ecosystems monitoring powered by environmental genomics: A
- review of current strategies with an implementation roadmap. *Mol. Ecol.* 30, 2937–2958
- 665 90 Ratnasingham, S. and Hebert, P.D.N. (2013) A DNA-Based Registry for All Animal
- 666 Species: The Barcode Index Number (BIN) System. *PLOS ONE* 8, e66213
- 667 91 Morinière, J. et al. (2019) A DNA barcode library for 5,200 German flies and midges
- 668 (Insecta: Diptera) and its implications for metabarcoding-based biomonitoring. *Mol. Ecol.*
- 669 *Resour.* 19, 900–928
- 670 92 Hogeweg, L. et al. (2019) Smart Insect Cameras. Biodivers. Inf. Sci. Stand. 3, e39241
- Bas, Y. *et al.* (2017) Tadarida: A Toolbox for Animal Detection on Acoustic Recordings.
- 672 *J. Open Res. Softw.* 5, 6
- 673 94 Svenningsen, C.S. et al. (2021) Detecting flying insects using car nets and DNA
- 674 metabarcoding. *Biol. Lett.* 17,
- 675 95 Schmeller, D.S. et al. (2009) Advantages of Volunteer-Based Biodiversity Monitoring in
- 676 Europe. *Conserv. Biol.* 23, 307–316
- 677 96 Hayhow, D.B. et al. Oct-(2019), State of nature 2019. . [Online]. Available:
- 678 https://nbn.org.uk/stateofnature2019/reports/. [Accessed: 31-Jan-2022]
- 679 97 WWF (2020) *Living Planet Report Nederland. Natuur en landbouw verbonden*, Wereld
 680 Natuur Fonds.
- 681 98 Gardiner, M.M. and Roy, H.E. (2022) The Role of Community Science in Entomology.
- 682 Annu. Rev. Entomol. 67, 437–456
- Berg, T.B. et al. (2021) The Role and Value of Out-of-School Environments in Science

- 684 Education for 21st Century Skills. Front. Educ. 6,
- 685 100 Mäder, P. et al. (2021) The Flora Incognita app Interactive plant species identification.
- 686 Methods Ecol. Evol. 12, 1335–1342
- 687 101 Wilkinson, M.D. et al. (2016) The FAIR Guiding Principles for scientific data
- 688 management and stewardship. *Sci. Data* 3, 160018
- 689 102 Andersson, A. et al. (2020) Publishing DNA-derived data through biodiversity data
- 690 *platforms. v1.0*, GBIF Secretariat.
- 691 103 Jones, M. *et al.* (2019) *Ecological Metadata Language (EML) version 2.2.0*, KNB Data
 692 Repository.
- 693 104 Wieczorek, J. et al. (2012) Darwin Core: An Evolving Community-Developed Biodiversity
- 694 Data Standard. *PLOS ONE* 7, e29715
- 695 105 Arribas, P. et al. (2021) Connecting high-throughput biodiversity inventories:
- Opportunities for a site-based genomic framework for global integration and synthesis. *Mol. Ecol.* 30, 1120–1135
- 698 106 Zaiko, A. et al. (2022) Towards reproducible metabarcoding data: Lessons from an
- 699 international cross-laboratory experiment. *Mol. Ecol. Resour.* 22, 519–538
- 107 Mousavi-Derazmahalleh, M. et al. (2021) eDNAFlow, an automated, reproducible and
- scalable workflow for analysis of environmental DNA sequences exploiting Nextflow and
- 702 Singularity. Mol. Ecol. Resour. 21, 1697–1704
- 108 Pereira, C.L. et al. (2021) Fine-tuning biodiversity assessments: A framework to pair
- eDNA metabarcoding and morphological approaches. *Methods Ecol. Evol.* 12, 2397–2409
- 109 Bista, I. et al. (2018) Performance of amplicon and shotgun sequencing for accurate
- biomass estimation in invertebrate community samples. *Mol. Ecol. Resour.* 18, 1020–1034
- 110 Wührl, L. et al. DiversityScanner: Robotic handling of small invertebrates with machine
- 708 learning methods. Mol. Ecol. Resour. n/a,
- 709 111 Bianchi, F.M. and Gonçalves, L.T. Getting science priorities straight: how to increase the

reliability of specimen identification? *Biol. Lett.* 17, 20200874

Jinbo, U. *et al.* (2011) Current progress in DNA barcoding and future implications for
entomology. *Entomol. Sci.* 14, 107–124

714

715

716

717 Figure captions

718

Figure 1. Workflows, from data collection to end product, of each of the four coveredtechnologies.

721

Figure 2. Current and potential future scope of the four technologies A non-exhaustive list of

723 current, in development and expected future possibilities for insect ecology and monitoring

vsing the four technological developments discussed in this paper. Colours refer to different

aspects of each technology: taxonomic precision and groups (orange), the metrics for

biodiversity that can be obtained (light blue), the size, scale and type of samples that can be

processed (gold) and the technological challenges for data processing (dark blue). Terms that

transgress the borders between technologies are applicable to both.

729

Highlights

- We appraise four emerging tools and technologies (computer vision, acoustic monitoring, radar and molecular methods) that provide unprecedented opportunities for insect monitoring and the study of insect ecology.
- These technologies have various benefits over traditional insect monitoring methods, including increased resolution of data collection across space and time, and a broader taxonomic coverage.
- At the same time, each technology has its limitations, some of which can be overcome with further methodological developments.
- Key issues regarding open science and international standards need to be addressed.
- While technology can never replace the knowledge of entomological specialists, we expect that integration of data across technologies, along with expert knowledge, will become commonplace in the future.

Outstanding questions

- What standards are needed to ensure interoperability of data and reproducibility of methods using each technology? How should these standards be fostered by the scientific and biodiversity informatics community?
- How do these new technologies compare to traditional methods in accuracy and informativeness? Are new technologies just quicker and cheaper? When should we still use traditional taxonomy and sampling, and how can new technologies and traditional sampling best complement each other?
- What are the sampling biases of new technologies, in terms of both taxonomic accuracy (which groups can be monitored by each technology) as well as species traits (which kinds of species might be missed by each technology)?
- How should the technologies be optimised to maximize both data quantity and data quality, bearing in mind that many of the new technologies may rapidly increase the rate of data collection, outpacing storage capacities and/or the rate at which the data can be validated, processed and analysed?
- How should data aggregation services (such as GBIF and national records) define and label data from novel identification techniques, including the uncertainty in species identity and non-validated insect records, so that these uncertainties are transparent to the end-user?
- How can new technologies overcome common biases associated with the effects of insect activity on monitoring indices and provide unbiased density estimates?
- What is the best sampling design to upscale and integrate different technologies for large-scale and long-term insect monitoring?
- How can new technologies facilitate and enhance engagement with society, promote experience of and learning about insects, and foster meaningful and innovative citizen science?
- How can we ensure that the outputs of these technologies align with policyrelevant indicators for ecological states and trends at relevant spatial and temporal scales?

