

Distribution of bat species in Western Asia: Occurrence records from the Western Asia Bat Research Network (WAB-Net) project

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Academic editor: Miguel Camacho Sanchez

Received: 16 Jul 2024 | Accepted: 09 Aug 2024 | Published: 21 Aug 2024

Citation: Phelps K, Al Abdulasalam Z, Al-Hmoud N, Ali S, Alrwashdeh M, Attaullah, Bilgin R, Ghazaryan A, Hamel L, Hasanov N, Natradze I, Papov G, Sidamonidze K, Spalton A, Urushadze L, Olival K (2024) Distribution of bat species in Western Asia: Occurrence records from the Western Asia Bat Research Network (WAB-Net) project. Biodiversity Data Journal 12: e132199. <u>https://doi.org/10.3897/BDJ.12.e132199</u>

Abstract

Background

Western Asia represents a mixing pot of diverse bat species with distributions spanning across other geographic regions. Yet, relative to other regions, there is a significant gap in coordinated bat research in Western Asia, thereby resulting in a relatively limited number of curated occurrence records.



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New information

The Western Asia Bat Research Network (WAB-Net) project was created to strengthen research capacity and knowledge of the diversity and distribution of bat species in a littlestudied region, as well as to collect data to characterise the diversity and prevalence of coronaviruses in diverse bat species. Over a four-year period (2018–2022), we documented 4,278 individual records for 41 bat species using a cross-sectional survey approach at 50 sites in seven Western Asian countries, specifically Armenia, Azerbaijan, Georgia, Jordan, Oman, Pakistan and Turkiye. At each site, we captured, on average, 90 individual bats (range: 9-131) over multiple consecutive nights and used standardised methods to collect demographic and morphological data from captured individuals. We additionally completed a systematic evaluation of environmental characterisation and human-bat interactions at all 50 sites. Here, we report individual occurrence records and site conditions from this multi-country, multi-year sampling effort.

Keywords

Chiroptera, Middle East, West Asia, biodiversity, mammal

Introduction

Western Asia serves as a convergence point for bat species originating from diverse geographic regions, including North Africa, South Asia and Europe, with the majority of these species distributed in more than one region. However, research on bat species in Western Asia is limited in comparison to other geographic regions (Phelps et al. 2019) and primarily concentrated on documenting bat diversity and distribution in a single country (Shehab et al. 2007, Benda et al. 2011, Benda and Uhrin 2019, Abdulrahman et al. 2021) or across neighbouring countries, such as the Arabian Peninsula (Harrison and Bates 1991).

Systematic bat survey data are invaluable to support both conservation and zoonotic disease research efforts (Phelps et al. 2019). In Western Asia, much of the bat locality data are based on historical surveys or museum collections, which were often collected opportunistically and within a more limited geographic scope. The collection of occurrence data from ongoing field surveys using standardised methods is critical to providing a more comprehensive picture of bat species distribution, abundance and diversity. This is especially true today given active and growing threats to bats in most parts of the world (Frick et al. 2020), including factors like climate change that are leading to range shifts and contractions for many species (Festa et al. 2022). At the same time, bats are important reservoir hosts for zoonotic diseases, including several emerging human and livestock viruses. Field-collected occurrence data can improve our understanding of bat-microbial (e.g. viruses, bacteria, protozoa) dynamics and provide

information for surveillance efforts and disease risk mitigation (Olival et al. 2017, Sánchez et al. 2022).

General description

Purpose: The Western Asia Bat Research Network (WAB-Net) project was created to strengthen research capacity and knowledge of the diversity and distribution of bat species in a little-studied region, as well as to collect data to characterise bat-associated coronaviruses (Phelps et al. 2019). More details on the WAB-Net project are available at <u>www.wabnet.org</u>.

Additional information: Over a four-year period (2018-2022), we conducted standardised, cross-sectional sampling of bat populations at 50 sites in seven Western Asian countries, specifically Armenia, Azerbaijan, Georgia, Jordan, Oman, Pakistan and Turkiye.

Sampling methods

Sampling description: Bats were captured primarily using harp traps and mist nets set in flyways or at the entrance to caves or anthropogenic structures in which bats were roosting. On rare occasions, bats were extracted from crevices with hand-held hoop nets. Trapping began approximately 30 minutes prior to sunset, with trapping duration dependent on capture rate (i.e. to prevent capture of more individuals than can be safely processed in a single night), but on average, nightly trapping duration was 4.8 hours. Captured bats were placed individually in a porous cloth bag and hung in a quiet, dry location away from predators. Standard morphological measurements (i.e. forearm length (mm) and body mass (g)) and demographic information (i.e. sex, age class and reproductive status) of each captured bat were recorded. Expert opinion was relied upon for species identification in the field and, when needed, additional morphological measurements, such as ear length (mm), tail length (mm), hind foot length (mm) and head and body length (mm) along with acoustic recordings, were used to aid in identification. Species identification was later confirmed for a subset of individuals of each species per site from each country via barcoding the cytochrome b gene using previously published methods (Townsen et al. 2008). Bats were immediately released at the original site of capture.

Field team members adhered to biosafety protocols established by the PREDICT Consortium (PREDICT Consortium 2020), including the use of personal protective equipment (i.e. field-dedicated clothing and close-toed shoes that could be disinfected, an N95 respirator mask, double-layer nitrile gloves and eye protection) during the capture, handling and sampling of bats. All methods were approved by the Tufts University Institutional Animal Care and Use Committee (protocols #G2018-01 and #G2020-127) and the USAMRDC Animal Care and Use Review Office (protocol #CT-2017-12).

Geographic coverage

Description: We captured bats at 50 sites in seven Western Asian countries, specifically Armenia, Azerbaijan, Georgia, Jordan, Oman, Pakistan and Turkiye (Fig. 1). The map was created in R version 4.3.2 (R Core Team 2023) using the following packages: "ggrepel" (Slowikowski 2024), "ggspatial" (Dunnington 2023), "rnaturalearth" (Massicotte and South 2023) and "tidyverse" (Wickham et al. 2019).

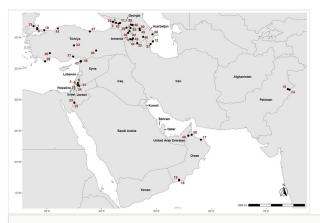


Figure 1.

Bats were sampled at 50 sites in seven countries across Western Asia, including Armenia (n = 6 sites), Azerbaijan (n = 6), Georgia (n = 10), Jordan (n = 8), Oman (n = 5), Pakistan (n = 2) and Turkiye (n = 13).

Locality information and sampling dates for each site are presented in Table 1, with more detailed information about site conditions presented in Suppl. material 3.

Ta	b	le	1	

Locality information and sampling dates for sites included in this study. The map code corresponds to sites in Fig. 1. See Suppl. material 3 for more detailed information about each site.

Map code	Country	Site	Latitude (North)	Longitude (East)	Sampling start date (dd/mm/yyyy)	Sampling end date (dd/mm/yyyy)
1	Turkiye	Cilingoz Cave	41.52	28.22	21/08/2018	24/08/2018
2	Georgia	Ghliana Cave	42.37	42.60	10/09/2018	12/09/2018
3	Jordan	Pella Cave	32.44	35.62	04/10/2018	06/10/2018
4	Georgia	Letsurtsume Cave	42.54	42.11	07/06/2019	10/06/2019
5	Armenia	Areni 1 Cave	39.73	45.20	12/06/2019	16/06/2019
6	Jordan	Al-Himmah Cave	32.71	35.68	07/07/2019	09/07/2019

Map code	Country	Site	Latitude (North)	Longitude (East)	Sampling start date (dd/mm/yyyy)	Sampling end date (dd/mm/yyyy)
7	Georgia	Tetri Senakebi Cave	41.54	45.26	06/08/2019	09/08/2019
8	Turkiye	Yaylacik Cave	41.36	28.22	16/08/2019	18/08/2019
9	Jordan	Baoun Cave	32.39	35.73	22/08/2019	24/08/2019
10	Turkiye	Sofular Cave	41.18	29.51	23/08/2019	27/08/2019
11	Georgia	Samertskhle Klde	42.34	43.35	03/09/2019	05/09/2019
12	Azerbaijan	Fish Reproduction Plant	39.39	49.36	07/09/2019	09/09/2019
13	Turkiye	Dupnisa Cave	41.84	27.56	08/09/2019	11/09/2019
14	Pakistan	Noorjahan & Jahangir Tombs	31.62	74.29	15/10/2019	16/10/2019
15	Pakistan	Sheikhpura Fort	31.71	73.99	18/10/2019	19/10/2019
16	Jordan	Khirbat Al Wahadinah	32.32	35.62	26/10/2019	01/11/2019
17	Oman	Muscat Al Khoud Fort	23.57	58.12	03/11/2019	04/11/2019
18	Oman	Alhaseela Farm	17.01	54.13	01/03/2020	02/03/2020
19	Oman	Suhoor Cave	17.13	54.12	03/03/2020	05/03/2020
20	Georgia	Gremi	42.00	45.66	13/07/2020	16/07/2020
21	Turkiye	Saykoy Cave	36.95	34.79	03/08/2020	05/08/2020
22	Georgia	Becho Cave	42.54	42.00	07/08/2020	12/08/2020
23	Turkiye	Sarihidir Tunnel	38.74	34.94	08/08/2020	10/08/2020
24	Armenia	Arakelots Vanq	41.03	45.07	11/08/2020	13/08/2020
25	Armenia	Shikahogh Mine	39.09	46.48	15/08/2020	17/08/2020
26	Turkiye	Sefer Yitigi Cave	37.48	30.53	17/08/2020	20/08/2020
27	Azerbaijan	Baligchilar Village	38.95	48.92	20/08/2020	22/08/2020
28	Turkiye	Hidirellez Cave	36.17	29.64	22/08/2020	25/08/2020
29	Georgia	Algeti Water Reservoir	41.59	44.53	28/08/2020	02/09/2020
30	Azerbaijan	Gakh	41.43	46.91	09/09/2020	11/09/2020
31	Jordan	King Talal Dam	32.22	35.88	01/10/2020	01/10/2020
32	Jordan	Aqaba Castle	29.52	35.00	12/10/2020	12/10/2020
33	Jordan	Saraya-Aqaba	29.54	34.99	13/10/2020	13/10/2020
34	Jordan	Kufranjah Cave	32.30	35.70	21/10/2020	22/10/2020
35	Georgia	Pichkhovani Church	42.08	45.33	21/06/2021	22/06/2021

Map code	Country	Site	Latitude (North)	Longitude (East)	Sampling start date (dd/mm/yyyy)	Sampling end date (dd/mm/yyyy)
36	Turkiye	Firat	37.91	38.99	07/07/2021	08/07/2021
37	Turkiye	Kahyaoglu	40.99	37.83	13/07/2021	14/07/2021
38	Azerbaijan	Jangi Gobustan	40.51	49.28	15/07/2021	17/07/2021
39	Armenia	Dilijan	40.75	44.82	29/07/2021	31/07/2021
40	Georgia	Grdzeli Chala	42.02	45.62	31/07/2021	07/08/2021
41	Armenia	Getashen	39.70	45.56	03/08/2021	05/08/2021
42	Georgia	Vardigora	42.25	43.09	03/08/2021	09/08/2021
43	Armenia	Jermuk	39.86	45.70	06/08/2021	09/08/2021
44	Turkiye	Cayirkoy Cave	41.46	31.99	10/08/2021	11/08/2021
45	Azerbaijan	Shaki	41.18	46.98	23/08/2021	25/08/2021
46	Azerbaijan	Aghjabadi	40.12	47.57	10/09/2021	12/09/2021
47	Turkiye	Harbiye Cave	36.14	36.14	22/09/2021	26/09/2021
48	Turkiye	Narlica Cave	36.22	36.20	27/09/2021	28/09/2021
49	Oman	Saarani Falaj	24.22	55.81	16/01/2022	18/01/2021
50	Oman	Baidha Tunnel	24.36	56.42	19/01/2022	20/01/2022

Taxonomic coverage

Description: We captured 4,278 individual bats of 41 species belonging to nine families. which represents 43.2% of the 95 bat species distributed across the seven focal countries and 35.6% of the 115 bat species documented in Western Asia (Upham et al. 2024). See Suppl. material 1 for detailed information about individual demographic and morphological traits and Suppl. material 2 for final species determination, based on multiple lines of evidence, including DNA barcode results with associated NCBI GenBank accession numbers. It is important to emphasise that, while the cytochrome b gene is extensively used to confirm species identifications, it captures only the maternal elements of the genome (Ballard and Whitlock 2003). Therefore, conclusions drawn solely from short fragments of the mitochondrial DNA may not be reliable enough to correctly differentiate closely related species (Ruedi et al. 2023). Such is the case for two sibling bat species, Myotis blythii and M. myotis (Furman et al. 2012), which are thought to undergo cryptic hybridisation in geographic areas of sympatry in Western Asia (i.e. Turkiye) (Berthier et al. 2006). Therefore, in addition to DNA barcoding results, which were not informative in some instances, data on morphology, geographic distribution, and/or acoustic recordings was also used to confirm species identifications.

Below, scientific names followed by "*" or "**" indicate a conservation status of "Near Threatened" or "Vulnerable", respectively, as designated by the International Union for Conservation of Nature (IUCN) Red List of Threatened Species (IUCN 2024).

Rank	Scientific Name	Common Name
order	Chiroptera	Bats
family	Emballonuridae	Sheath-tailed Bats
species	Taphozous nudiventris	Naked-rumped Tomb Bat
species	Taphozous perforatus	Egyptian Tomb Bat
family	Hipposideridae	Old World leaf-nosed Bats
species	Asellia arabica	Arabian Trident Leaf-nosed Bat
species	Asellia tridens	Geoffroy's Trident Leaf-nosed Bat
family	Miniopteridae	Bent-winged Bats
species	Miniopterus pallidus*	Pallid Long-fingered Bat
species	Miniopterus schreibersii**	Schreibers' Long-fingered Bat
family	Nycteridae	Slit-faced Bats
species	Nycteris thebaica	Egyptian Slit-faced Bat
family	Pteropodidae	Old World Fruit Bats
species	Rousettus aegyptiacus	Egyptian Rousette
species	Rousettus leschenaultii*	Leschenault's Rousette
family	Rhinolophidae	Horseshoe Bats
species	Rhinolophus blasii	Blasius' Horseshoe Bat
species	Rhinolophus clivosus	Geoffroy's Horseshoe Bat
species	Rhinolophus euryale*	Mediterranean Horseshoe Bat
species	Rhinolophus ferrumequinum	Greater Horseshoe Bat
species	Rhinolophus hipposideros	Lesser Horseshoe Bat
species	Rhinolophus lepidus	Blyth's Horseshoe Bat
species	Rhinolophus mehelyi**	Mehely's Horseshoe Bat
family	Rhinonycteridae	Trident Bats
species	Triaenops persicus	Persian Trident Bat
family	Rhinopomatidae	Mouse-tailed Bats
species	Rhinopoma cystops	Arabian Mouse-tailed Bat

Taxa included:

species	Rhinopoma microphyllum	Greater Mouse-tailed Bat
species	Rhinopoma muscatellum	Muscat Mouse-tailed Bat
family	Vespertilionidae	Vesper Bats
species	Barbastella caspica	Caspian Barbastelle
species	Cnephaeus (Eptesicus) serotinus	Eurasian Serotine
species	Hypsugo savii	Savi's Pipistrelle
species	Myotis alcathoe	Alcathoe Whiskered Myotis
species	Myotis blythii	Lesser Myotis
species	Myotis capaccinii**	Long-fingered Myotis
species	Myotis daubentonii	Daubenton's Myotis
species	Myotis davidii	David's Myotis
species	Myotis emarginatus	Geoffroy's Myotis
species	Myotis myotis	Greater Myotis
species	Myotis nattereri	Natterer's Myotis
species	Myotis tschuliensis	Tschuli <i>Myotis</i>
species	Pipistrellus javanicus	Javan Pipistrelle
species	Pipistrellus kuhlii	Kuhl's Pipistrelle
species	Pipistrellus nathusii	Nathusius's Pipistrelle
species	Pipistrellus pipistrellus	Common Pipistrelle
species	Pipistrellus pygmaeus	Soprano Pipistrelle
species	Plecotus auritus	Brown Long-eared Bat
species	Plecotus macrobullaris	Alpine Long-eared Bat
species	Scotophilus heathii	Greater Asian Yellow Bat
species	Scotophilus kuhlii	Lesser Asian Yellow Bat

Temporal coverage

Notes: August 2018 - January 2022

Usage licence

Usage licence: Other

IP rights notes: Usage of reported data is licensed under a Creative Commons Attribution-Non-Commercial 4.0 License (CC BY-NC 4.0).

Data resources

Data package title: The Western Asia Bat Research Network (WAB-Net) project

Resource link: https://www.gbif.org/dataset/7c56c0cb-66e3-4e8f-acb9-db6370c87451

Number of data sets: 1

Data set name: The Western Asia Bat Research Network (WAB-Net) project

Data format: Darwin Core Format (https://dwc.tdwg.org/terms/)

Description: Occurrence records for 4,278 individual bats from 41 species captured across 50 sites in seven Western Asian countries (Armenia, Azerbaijan, Georgia, Jordan, Oman, Pakistan and Turkiye) between 2018-2022.

Column label	Column description
occurrenceID	An identifier for the dwc:Occurrence (as opposed to a particular digital record of the dwc:Occurrence).
basisOfRecord	The specific nature of the data record.
eventDate	The date-time or interval during which a dwc:Event occurred. For occurrences, this is the date-time when the dwc:Event was recorded.
scientificName	The full scientific name, with authorship and date information if known. When forming part of a dwc:Identification, this should be the name in lowest level taxonomic rank that can be determined. This term should not contain identification qualifications, which should instead be supplied in the dwc:identificationQualifier term.
higherClassification	A list (concatenated and separated) of taxon names terminating at the rank immediately superior to the referenced dwc:Taxon.
kingdom	The full scientific name of the kingdom in which the dwc:Taxon is classified.
phylum	The full scientific name of the phylum or division in which the dwc:Taxon is classified.
class	The full scientific name of the class in which the dwc:Taxon is classified.
order	The full scientific name of the order in which the dwc:Taxon is classified.
family	The full scientific name of the family in which the dwc:Taxon is classified.
genus	The full scientific name of the genus in which the dwc:Taxon is classified.
specificEpithet	The name of the first or species epithet of the dwc:scientificName.
taxonRank	The taxonomic rank of the most specific name in the dwc:scientificName.

lifeStage	The age class or life stage of the dwc:Organism(s) at the time the dwc:Occurrence was recorded.
sex	The sex of the biological individual(s) represented in the dwc:Occurrence.
reproductiveCondition	Categorisation of individuals for eusocial species (including some mammals and arthropods).
identifiedBy	A list (concatenated and separated) of names of people, groups or organisations who assigned the dwc:Taxon to the subject.
dateIdentified	The date on which the subject was determined as representing the dwc:Taxon.
decimalLatitude	The geographic latitude (in decimal degrees, using the spatial reference system given in dwc:geodeticDatum) of the geographic centre of a dcterms:Location. Positive values are north of the Equator, negative values are south of it. Legal values lie between -90 and 90, inclusive.
decimalLongitude	The geographic longitude (in decimal degrees, using the spatial reference system given in dwc:geodeticDatum) of the geographic centre of a dcterms:Location. Positive values are east of the Greenwich Meridian, negative values are west of it. Legal values lie between -180 and 180, inclusive.
geodeticDatum	The ellipsoid, geodetic datum or spatial reference system (SRS), upon which the geographic coordinates given in dwc:decimalLatitude and dwc:decimalLongitude are based.
coordinateUncertaintyInMetres	The horizontal distance (in metres) from the given dwc:decimalLatitude and dwc:decimalLongitude describing the smallest circle containing the whole of the dcterms:Location. Leave the value empty if the uncertainty is unknown, cannot be estimated or is not applicable (because there are no coordinates). Zero is not a valid value for this term.
dataGeneralisations	Actions taken to make the shared data less specific or complete than in its original form. Suggests that alternative data of higher quality may be available on request.
georeferencedDate	The date on which the dcterms:Location was georeferenced.
georeferenceSources	A list (concatenated and separated) of maps, gazetteers or other resources used to georeference the dcterms:Location, described specifically enough to allow anyone in the future to use the same resources.
georeferenceVerificationStatus	A categorical description of the extent to which the georeference has been verified to represent the best possible spatial description for the dcterms:Location of the dwc:Occurrence.
higherGeography	A list (concatenated and separated) of geographic names less specific than the information captured in the dwc:locality term.
continent	The name of the continent in which the dcterms:Location occurs.

country	The name of the country or major administrative unit in which the dcterms:Location occurs.
countryCode	The standard code for the country in which the dcterms:Location occurs.
stateProvince	The name of the next smaller administrative region than country (state, province, canton, department, region etc.) in which the dcterms:Location occurs.
county	The full, unabbreviated name of the next smaller administrative region than stateProvince (county, shire, department etc.) in which the dcterms:Location occurs.
municipality	The full, unabbreviated name of the next smaller administrative region than county (city, municipality etc.) in which the dcterms:Location occurs. Do not use this term for a nearby named place that does not contain the actual dcterms:Location.
locality	The specific description of the place.
language	A language of the resource.
licence	A legal document giving official permission to do something with the resource.
institutionID	An identifier for the institution having custody of the object(s) or information referred to in the record.
institutionCode	The name (or acronym) in use by the institution having custody of the object(s) or information referred to in the record.
collectionCode	The name, acronym, coden or initialism identifying the collection or dataset from which the record was derived.
catalogNumber	An identifier (preferably unique) for the record within the dataset or collection.
recordedBy	A list (concatenated and separated) of names of people, groups or organiations responsible for recording the original dwc:Occurrence. The primary collector or observer, especially one who applies a personal identifier (dwc:recordNumber), should be listed first.
preparations	A list (concatenated and separated) of preparations and preservation methods for a dwc:MaterialEntity.
otherCatalogNumbers	A list (concatenated and separated) of previous or alternative fully qualified catalogue numbers or other human-used identifiers for the same dwc:Occurrence, whether in the current or any other dataset or collection.
previousIdentifications	A list (concatenated and separated) of previous assignments of names to the dwc:Organism.
associatedSequences	A list (concatenated and separated) of identifiers (publication, global unique identifier, URI) of genetic sequence information associated with the dwc:MaterialEntity.

Acknowledgements

We thank the following agencies for permission to conduct this research project: In Armenia (Ministry of Nature Protection, Permit No. 03/577; Ministry of Environment, Permit No. 2/10.2.7/3457); in Azerbaijan (Ministry of Ecology and Natural Resources, Permit No. 21/304); in Georgia (Ministry of Environmental Protection and Agriculture, Permit No. 4894-01-2-201806081223 and Permit No. 3030-01-2-202103261654); in Jordan (Royal Society for Conservation of Nature, Permit No. 2/238/1/10); in Oman (Ministry of Environment and Climate Affairs, Permit No. 6210/10/97 and Permit No. 6210/10/125); in Pakistan (Wildilfe and Parks Department of Punjab Province, Permit No. 4292); in Turkiye (Ministry of Agriculture and Forestry, Permit No. 72784983-488.04 and Permit No. E-21264211-288.04-1581138).

We extend our gratitude to the following field team members: In Armenia (Seda Adamyan, Vanuhi Hambardzumyan, Tatevik Harutyunyan, Tigran Hayrapetyan, Aram Pirumyan, Anna Tadevosyan, Ofik Yelabekyan); in Azerbaijan (Leyla Aliyeva, Sabina Bunyatova, Gulnar Guliyeva, Agil Hakhiyev, Ilaha Kerimli, Sevin Sarukhanova); in Georgia (Maia Chigladze, Rusudan Kodiashvili, Davit Putkaradze, Manana Shanidze, Lamzira Tshkvaradze); in Jordan (Ehab Abu-Basha, Abdallh Al Dwekat, Sief Al Hanandeh, Qusay Al-Khateeb, Zaidoun Hijazeen, Ismail Malkawi, Hani Talafha); in Oman (Ahmed Al Amri, Waheed Al Fazari, Khalifa Al Ghisni, Khalid Al Hikmani, Zakariya Al Kindi, Atika Al Mandari, Abdullah Al Manji, Said Zabanoot); in Pakistan (Abdul Malik, Sajida Noureen, Muhammad Rizwan); in Turkiye (Ekin Azbazdar, Kaan Bayir, Ziya Caylarbasi, Erdem Danyer, Fadime Gencer, Sabri Hacioglu, Evrim Kalkan, Aygun Karacay, Zeynep Kocer, Ayse Mergenci, Selahattin Unsal, Ceylan Yucel); as well as the laboratory team members: in Georgia (Gvantsa Brachveli, Meri Pantsulaia, Ana Papkiauri, Tea Tevdoradze, Giorgi Tomashvilli, Davit Tsaguria); in Jordan (Amany Abdien, Ghadeer Alzghoul); in Turkiye (Fadime Gencer, Aygun Karacay, Yagmur Tarhana).

Lastly, we extend our gratitude to Drs. Paul Bates, Jonathan Epstein, Tigga Kingston, Vincent Munster and Paul Racey for providing general oversight and technical expertise as Scientic Advisory Board members of the WAB-Net.

This research was sponsored by the Department of Defense, Defense Threat Reduction Agency (HDTRA11710064). The content of the information does not necessarily reflect the position or the policy of the federal government and no official endorsement should be inferred.

References

- Abdulrahman M, Gardner A, Yamaguchi N (2021) The occurrence and distribution of bats in Qatar. Journal of Arid Environments 185: 104379. <u>https://doi.org/10.1016/j.jaridenv.</u> 2020.104379
- Ballard JWO, Whitlock M (2003) The incomplete natural history of mitochondria. Molecular Ecology 13 (4): 729-744. <u>https://doi.org/10.1046/j.1365-294x.2003.02063.x</u>

- Benda P, Al-Jumaily MM, Reiter A, Nasher AK (2011) Noteworthy records of bats from Yemen with description of a new species from Socotra. Hystrix 22: 23-56. <u>https://doi.org/</u> <u>10.4404/hystrix-22.1-4473</u>
- Benda P, Uhrin M (2019) On the bat fauna of the Bahrain Archipelago (Chiroptera). Lynx 50: 5-27.
- Berthier P, Excoffier L, Ruedi M (2006) Recurrent replacement of mtDNA and cryptic hybridization between two sibling bat species *Myotis myotis* and *Myotis blythii*.
 Proceedings of the Royal Society B: Biological Sciences 273 (1605): 3101-3109. <u>https:// doi.org/10.1098/rspb.2006.3680</u>
- Dunnington D (2023) ggspatial: Spatial Data Framework for ggplot2. R package version
 1.1.9. URL: <u>https://CRAN.R-project.org/package=ggspatial</u>
- Festa F, Ancillotto L, Santini L, Pacifici M, Rocha R, Toshkova N, Amorim F, Benítez-López A, Domer A, Hamidović D, Kramer-Schadt S, Mathews F, Radchuk V, Rebelo H, Ruczynski I, Solem E, Tsoar A, Russo D, Razgour O (2022) Bat responses to climate change: a systematic review. Biological reviews of the Cambridge Philosophical Society 98 (1): 19-33. <u>https://doi.org/10.1111/brv.12893</u>
- Frick W, Kingston T, Flanders J (2020) A review of the major threats and challenges to global bat conservation. Annals of the New York Academy of Sciences 1469 (1): 5-25. <u>https://doi.org/10.1111/nyas.14045</u>
- Furman A, Çoraman E, Nagy Z, Postawa T, Bilgin R, Gajewska M, Bogdanowicz W (2012) Phylogeography of the large *Myotis* bats (Chiroptera: Vespertilionidae) in Europe, Asia Minor, and Transcaucasia. Biological Journal of the Linnean Society 108 (1): 189-209. https://doi.org/10.1111/j.1095-8312.2012.01994.x
- Harrison DL, Bates PJJ (1991) The mammals of Arabia. 2nd Edition. Harrison Zoological Museum, Kent, UK, 354 pp.
- IUCN (2024) The IUCN Red List of Threatened Species. Version 2024-1. <u>https://</u> www.iucnredlist.org. Accessed on: 2024-7-29.
- Massicotte P, South A (2023) rnaturalearth: World Map Data from Natural Earth. R
 package version 1.0.1. URL: <u>https://CRAN.R-project.org/package=rnaturalearth</u>
- Olival K, Hosseini P, Zambrana-Torrelio C, Ross N, Bogich T, Daszak P (2017) Host and viral traits predict zoonotic spillover from mammals. Nature 546 (7660): 646-650. <u>https:// doi.org/10.1038/nature22975</u>
- Phelps K, Hamel L, Alhmoud N, Ali S, Bilgin R, Sidamonidze K, Urushadze L, Karesh W, Olival K (2019) Bat rsearch networks and viral surveillance: Gaps and opportunities in Western Asia. Viruses 11 (3). <u>https://doi.org/10.3390/v11030240</u>
- PREDICT Consortium (2020) Biosafety and Personal Protective Equipment (PPE) use.
 PREDICT standard operating procedures for one health surveillance. One Health
 Institute, University of California, Davis URL: <u>https://ohi.vetmed.ucdavis.edu/sites/g/files/
 dgvnsk5251/files/inline-files/PREDICT%202%20SOP%20Book_v3.%20Aug2020.pdf</u>
- R Core Team (2023) R: A Language and environment for statistical computing. Vienna, Austria. URL: <u>https://www.R-project.org/</u>
- Ruedi M, Manzinalli J, Dietrich A, Vinciguerra L (2023) Shortcomings of DNA barcodes: a
 perspective from the mammal fauna of Switzerland. Hystrix, the Italian Journal of
 Mammalogy 34 (1): 54-61. <u>https://doi.org/10.4404/hystrix-00628-2023</u>
- Sánchez C, Li H, Phelps K, Zambrana-Torrelio C, Wang L, Olival K, Daszak P (2022) A strategy to assess spillover risk of bat SARS-related coronaviruses in Southeast Asia. Nature Communications 13: 4380. <u>https://doi.org/10.1038/s41467-022-31860-w</u>

- Shehab A, Karataş AH, Amr Z, Mamkhair IB, Sözen MU (2007) The distribution of bats (Mammalia: Chiroptera) in Syria. Vertebrate Zoology 57: 103-123. <u>https://doi.org/10.3897/ vz.57.e30888</u>
- Slowikowski K (2024) ggrepel: Automatically Position Non-Overlapping Text Labels with 'ggplot2'. R package version 0.9.5. URL: <u>https://CRAN.R-project.org/package=ggrepel</u>
- Townsen JS, Brower AV, Judd DD (2008) Identification of mosquito bloodmeals using mitochondrial cytochrome oxidase subunit I and cytochrome b gene sequences. Medical and Veterinary Entomology 22 (4): 386-393. <u>https://doi.org/10.1111/j. 1365-2915.2008.00760.x</u>
- Upham N, Burgin C, Widness J, Liphardt S, Parker C, Becker M, Rochon I, Huckaby D, Zijlstra J (2024) Mammal Diversity Database. <u>https://doi.org/10.5281/zenodo.12738010</u>. Accessed on: 2024-7-20.
- Wickham H, Averick M, Bryan J, Chang W, McGowan LD, Francios R, Grolemund G, Hayes A, Henry L, Hester J, Kuhn M, Pedersen TL, Miller E, Bache SM, Muller K, Ooms J, Robinson D, Seidel DP, Spinu V, Takahashi K, Vaughan D, Wilke C, Woo K, Yutani H (2019) Welcome to the tidyverse. Journal of Open Source Software 4 (43). <u>https://doi.org/ 10.21105/joss.01686</u>

Supplementary materials

Suppl. material 1: Species occurrence, including demographic and morphological data for each captured individual bat

Authors: Kendra Phelps, Zahran Al Abdulasalam, Nisreen Al-Hmoud, Shahzad Ali, Mumen Alrwashdeh, Attaullah, Rasit Bilgin, Astghik Ghazaryan, Luke Hamel, Nijat Hasanov, Ioseb Natradze, George Papov, Ketevan Sidamonidze, Andrew Spalton, Lela Urushadze, Kevin J. Olival **Data type:** occurrences, morphology, demography

Brief description: Detailed information about each of the 4,278 individual bats captured during the WAB-Net project, including taxonomic, demographic and morphologic information. Download file (572.81 kb)

Suppl. material 2: Updated species identifications based on multiple lines of evidence

Authors: Kendra Phelps, Zahran Al Abdulasalam, Nisreen Al-Hmoud, Shahzad Ali, Mumen Alrwashdeh, Attaullah, Rasit Bilgin, Astghik Ghazaryan, Luke Hamel, Nijat Hasanov, Ioseb Natradze, George Papov, Ketevan Sidamonidze, Andrew Spalton, Lela Urushadze, Kevin J. Olival **Data type:** cytB barcoding, species identifications

Brief description: Update of field species identification based on CytB barcoding results, including

associated NCBI GenBank accession numbers, geographic distribution, morphology and/or acoustic signatures.

Download file (527.98 kb)

Suppl. material 3: Environmental conditions and potential bat-human contact interfaces

Authors: Kendra Phelps, Zahran Al Abdulasalam, Nisreen Al-Hmoud, Shahzad Ali, Mumen Alrwashdeh, Attaullah, Rasit Bilgin, Astghik Ghazaryan, Luke Hamel, Nijat Hasanov, Ioseb Natradze, George Papov, Ketevan Sidamonidze, Andrew Spalton, Lela Urushadze, Kevin J. Olival **Data type:** site characteristics

Brief description: We assessed site-level factors within a 1 km² radius at each of the 50 sites in which bats were sampled, ranging from site type (e.g. forested habitat, wetland, caves and/or rock crevices), potential bat-human interfaces (e.g. tourism, wildlife market) and evidence of human disturbance (e.g. guano harvesting, bat hunting, fire remnants). Download file (10.56 kb)