This data article informs about Chiropteran diversity, new records, ecosystem services and possible pathogen carriers in fragmented forests (sub-divided by utility corridors, man-made structures, untouched and secondary plantations) within districts Setiu (Setiu Research Station), Hulu Terengganu (Saok and Lasir waterfalls) and Besut (Gunung Tebu Forest Reserve) of state Terengganu, Peninsular Malaysia. These bats were captured using harp traps and mist nets that were set 10 m apart across flyways, streams and less cluttered trees in the 50 m transect zones (identified at each site). All animals were distinguished by morphology and gender before their release at the site of capture. The data comprise of five bat family groups Hipposideridae, Megadermatidae, Pteropodidae, Rhinolophidae and Vespertilionidae. It is interesting to note that untouched Saok Waterfalls is home to wide variety of bats listed (68.8%), followed by secondary forests of Gunung Tebu Forest Reserve (24.8%), untouched Lasir Waterfalls (4.8%) and lastly, Setiu Research Station as least favored (1.6%). Chiroptera like *Cynopterus brachyotis* (*n* = 23, 37.7%), *Hipposideros bicolor* (*n* = 6, 9.8%) and *Scotophilus kuhli* (*n* = 6, 9.8%) were most dominant in the checklist whereas *Hipposideros armiger*,...
Murina suilla and Scotophilus kuhlii are new data records in the fragmented forests of Terengganu. The data were interpreted into Shannon, Simpson, Margalef, Menhinik and Evenness indices to individually or collectively distinguish chiropteran variety in Terengganu State whereas weight-forearm length (W/FA) informs about chiropteran Body Condition Index (-0.25 to 0.25). The function of bats were also identified to distinguish service providers (pollination and forests regeneration) and zoonotic pathogen carriers (in particular to Leptospira bacteria, Nipah virus and Sindbis virus).

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### Specifications table

| Subject area       | Biology                      |
|--------------------|------------------------------|
| More specific subject area | Bioscience and Biodiversity |
| Type of data       | Tables                       |
| How data was acquired | Four bank harp trap (4.2 m²) and mist nets (Height = 2.6 m and Width = 10 m), Vernier caliper (sensitivity 0.1 cm), measuring tape, analytical balance (sensitivity 0.1 kg) and Paleontological Statistics Software Package (PAST) v.3 |
| Data format        | Raw and Analyzed             |
| Experimental factors | Harp trap and net placement at flyways, streams and less cluttered trees and, the 10 m spacing between the traps and nets. |
| Experimental features | Descriptive abundance by pool, ecosystem services, pathogen-carrying species, guild types, roost, climb limit and range. Interpretations include body condition index and diversity estimations using Shannon, Simpson, Evenness, Margalef and Menhinick Indices. |
| Data source location | 1. Setiu District, Terengganu, East Peninsular Malaysia Setiu Wetland Research Centre: N 5.6771°; E 102.7102°  
  2. Hulu Terengganu District (Terengganu Nature Park), Terengganu, East Peninsular Malaysia Lasir Waterfalls: N 4.9655°; E 102.8396°  
  Saok Waterfalls: N 5.0832°; E 102.7784°  
  3. Besut District, Terengganu, Peninsular Malaysia Gunung Tebu Forest Reserve: N 5.5915°; E 102.6164° |
| Data accessibility | All raw data are available within this article |
| Related research article | Unpublished data |

### Value of the data

- This data is constructed from chiropteran diversity and abundance in fragmented forests of Terengganu with some new records that offer research opportunities and perhaps collaboration to address the subject matter.
- The chiroptera discovered in the present study are ecological service providers that maintain tropical rainforest splendor, influence the genetic vigor of fruits, flora and timber, become pest controllers, function as pathogen reservoirs and indirectly assist agriculture industry through crop pollination, seed dispersal and provide nutrient-rich feces as manure. These form of valuable
information provides insights into forest and wildlife management, serve as benchmark for fragmented forests and as indicator for climate change, assist with food web plotting in different environments (caves, trees, artificial structures and fragmented forests) and also assist with bio-resource studies (like biotechnology and phenology).

- Chiropteran morphometric data translated into body condition index changes the research perspective because wildlife stress, forest carrying capacity and forest regeneration are measurable. These measurements can be incorporated into prediction models used for sustainability and conservation studies.

1. Data

This data article is constructed using survey results that indicate bat diversity and their abundance in fragmented forests within Terengganu (Table 1). From these, additional associations such as new records, ecosystem services, possible pathogen carriers and their lifestyle were added from available literature for forest regeneration impacts, influence towards agriculture activities, trophic level and food web constructs, natural pest control, precursor for diseases, conservation assert (if needed) as well as their geographical tolerance and adaptations (Table 2). Chiropteran morphometric measurements are translated into weight-length data to scale their health in the wild (Table 3). These indicate food resources, its availability and its adaptation to forest fragmentations. The data is also interpreted using Mathematical equations (Shannon, Simpson, Eveness, Margalef and Menhinick) to derive diversity and species richness values which are useful for chiropteran interpolations and assessments within a geographical boundary (Table 4).

2. Experimental design, materials, and methods

Field visits to districts Hulu Terengganu (Saok and Lasir Waterfalls), Setiu (Setiu Research Station) and Besut (Gunung Tebu Forest Reserve) were carried out in June 2017 and spanned six days at each

| Table 1 |
|---|
| Chiropteran abundance from sites assessed in the districts Setiu, Besut and Hulu Terengganu, Peninsular Malaysia. |
| Family | Species | Common name | Abundance (No.) |
|---|---|---|---|
|  |  |  | SW | LW | GTFR | SRS |
| Hipposideridae | *Hipposideros armiger* | Great Leaf-nosed Bat | 0 | 0 | 2 | 0 |
|  | Hipposideros bicolor | Bicoloured Leaf-nosed Bat | 5 | 0 | 1 | 0 |
|  | Hipposideros cervinus | Fawn Leaf-nosed Bat | 2 | 1 | 0 | 0 |
|  | Hipposideros diadema | Diadem Leaf-nosed Bat | 0 | 0 | 1 | 0 |
| Megadermatidae | Megaderma lyra | Greater False Vampire Bat | 0 | 0 | 1 | 0 |
|  | Balionycteris maculata | Spotted-winged Fruit Bat | 0 | 0 | 1 | 0 |
| Pteropodidae | *Cynopterus brachyotis* | Lesser Dog-faced Fruit Bat | 21 | 1 | 0 | 1 |
|  | Eonycteris spelaea | Cave Nectar Bat | 3 | 0 | 0 | 0 |
|  | MacroGLOSSUS sobrinus | Long-tongued Fruit Bat | 3 | 0 | 0 | 0 |
|  | Megerops caudatus | Tailless Fruit Bat | 1 | 0 | 0 | 0 |
|  | Penthetor lucasi | Dusky Fruit Bat | 2 | 1 | 0 | 0 |
| Rhinolophidae | *Rhinolophus affinis* | Intermediate Horseshoe Bat | 1 | 0 | 3 | 0 |
|  | Rhinolophus lepidus | Blyth’s Horseshoe Bat | 0 | 2 | 0 | 0 |
| Vespertilionidae | *Marina suilla* | Brown Tube-nosed Bat | 1 | 0 | 0 | 0 |
|  | *Scotophilus kuhlii* | Lesser Asiatic Yellow House Bat | 0 | 0 | 6 | 0 |
| Total | | | 42 | 3 | 15 | 1 |

Note: The status of all bats are ‘Least Concern’ unless marked with ‘*’ to indicate it as ‘Not Evaluated’ in the IUCN Red List. New records are shown with ‘N’ for their absence even in the most recent study by Pounsin et al. [1]. Sites for field visits are abbreviated as SW = Saok Waterfalls, LW = Lasir Waterfalls, GTFR = Gunung Tebu Forest Reserve and SRS = Setiu Research Station. Abundance of bats are denote with ‘no.’ to represent number of individuals.
The geographical range of bats are classified into Primary Forests (Pr), Regenerated Forests (R) and Secondary forests (S) (personal observations). Pathogens that associate with Chiroptera include Carey Island Virus, Juga virus (J), Japanese Encephalitis virus (JE), Nipah virus (N), Phnom Penh Bat virus (PP), Sindbis virus (Sb), Issyk-Kul (Keterah Virus), Leptospira bacteria and SARS coronavirus (Sc). Chiroptera guilds are denote with AI = Aerial Insectivore, C = Carnivore, F = Frugivore, I = Insectivore and N = Nectarivore whereas their roost are denote with B = buildings, Cv = Cave, Fs = Forests, P = Plantations and T = Trees. Chiroptera climb limits were recorded as maximum climb height. The geographical range of bats are classified into E1 = Endemic to Indomalayan Realm; E2 = Endemic to Indonesia & Malaysia; E3 = Endemic to Malaysia and N = Native.

### Table 2
Mean forearm length and weight of voucher Chiroptera used to construct the weight to length ratio and the comparisons used to construct the mass index.

| Species                  | Forearm length (mm) | Weight (g) | W/FA (%) | Repository | BC Index |
|--------------------------|---------------------|------------|----------|------------|----------|
|                          | Male               | Female     | Male     | Female     | W/FA     | Male     | Female     |
| Hipposideros armiger     | 91.2 ± 0.0         | 89.6 ± 0.0 | 68.0 ± 0.0 | 49.1 ± 0.0 | 74.5      | 54.8     | 48.0       |
| Hipposideros bicolor     | 43.4 ± 1.3         | 46.0 ± 0.0 | 8.1 ± 0.9 | 5.8 ± 0.9  | 18.6      | 12.5     | 20.8 -0.1 |
| Hipposideros cervinus    | 47.0 ± 0.0         | 50.1 ± 0.0 | 9.4 ± 0.0 | 10.9 ± 0.0 | 20.0      | 21.7     | 20.0 0.0H |
| Hipposideros diadema     | 83.6 ± 0.0         | 79.1 ± 0.0 | 25.7 ± 0.0 | 52.4 ± 0.0 | 30.7      | 54.7     | -0.4U -0.4H|
| Megaderma lyra           | -                  | -          | 79.1 ± 0.0 | -          | 66.2      | 56.8     | -0.2H -0.4U|
| Balonycteris maculata    | -                  | 68.0 ± 0.0 | -          | -          | 65.0      | 79.3     | -0.2H     |
| Macroglossus sobrinus    | 45.3 ± 0.6         | -          | 16.2 ± 0.6 | -          | 35.7      | 44.3     | -0.2H     |
| Megaerops ecaudatus      | 54.0 ± 0.0         | -          | 24.1 ± 0.0 | -          | 44.6      | 44.3     | 0.0H -0.0H|
| Penthetor lucasi         | 54.0 ± 0.0         | 56.1 ± 0.0 | 26.7 ± 0.0 | 27.9 ± 0.0 | 49.4      | 49.7     | 71.0 -0.3H|
| Rhinolophus affinis      | 51.0 ± 3.6         | 52.2 ± 0.0 | 13.5 ± 1.8 | 14.1 ± 0.0 | 26.5      | 27.0     | 27.7 0.0H|
| Rhinolophus lepidus      | 42.5 ± 0.0         | -          | 15.9 ± 0.0 | -          | 37.4      | -13.4    | 18.0 -     |
| Murina suilla            | -                  | -          | 31.5 ± 0.0 | -          | 4.0       | -12.7    | -14.0 -0.1H|
| Scotophilus kuhlii        | 50.8 ± 0.0         | 51.0 ± 1.4 | 19.6 ± 0.0 | 20.3 ± 2.7 | 38.6      | 39.9     | 42.3 -0.1H|

Note: Weight to forearm length (W/FA) measurements were recorded from voucher chiropteran and compared with known measurements (W/FA*) to construct the Body Condition Index (BCI). Values in this index were scaled by 0.5 unit differences into U = Underweight (-0.75 to -0.25), H = Ideal (-0.25 to 0.25), E = Overweight (0.25 to 0.75) and B = Obese (> 0.75). Symbol – indicates absence of samples.

### Table 3
Chiroptera ecological services, possible pathogen associations and limitations in the fragmented forests of Terengganu.

| Species                  | Service | Location | Possible pathogen associations | Guild | Roost | Climb Limit (× 10³ m) | Range   |
|--------------------------|---------|----------|--------------------------------|-------|-------|-----------------------|---------|
| Hipposideros armiger     | /       | /        | /                              | Al    | Cv    | 2.0                   | N       |
| Hipposideros bicolor     | /       | /        | /                              | Al    | Cv    | 0.6                   | E²      |
| Hipposideros cervinus    | /       | /        | /                              | Al    | Cv, T | 1.4                   | N       |
| Hipposideros diadema     | /       | /        | /                              | Al    | Cv, T | 1.2                   | N       |
| Megaderma lyra           | /       | /        | /                              | C, I  | B, Cv | 1.0                   | E¹      |
| Balonycteris maculata    | /       | /        | /                              | F, N  | Cv    | 1.5                   | E³      |
| Cytopterus brachyotos     | /       | /        | /                              | F, N  | Cv, T | 1.6                   | N       |
| Cytopterus horsfieldii    | /       | /        | /                              | F, N  | Cv, T | > 1.0                 | N       |
| Eonycteris spelaea       | /       | /        | /                              | N     | Cv    | 1.0                   | N       |
| Macroglossus sobrinus    | /       | /        | /                              | N     | T     | 2.0                   | N       |
| Megaerops ecaudatus      | /       | /        | /                              | F, N  | Fs    | 3.0                   | E²      |
| Penthetor lucasi         | /       | /        | /                              | F, N  | Cv    | 0.6                   | E²      |
| Rhinolophus affinis      | /       | /        | /                              | Al    | Cv    | 2.0                   | N       |
| Rhinolophus lepidus      | /       | /        | /                              | Al    | Cv    | 2.3                   | E¹      |
| Murina suilla            | /       | /        | /                              | Al    | Fs, P | 1.5                   | N       |
| Scotophilus kuhlii        | /       | /        | /                              | Al    | B     | 1.1                   | E¹      |

Note: Chiroptera service as pollinator is indicated with (P). The forest types are divided into Primary Forests (Pr), Regenerated Forests (R) and Secondary forests (S) (personal observations). Pathogens that associate with Chiroptera include Carey Island virus (CI), Juga virus (J), Japanese Encephalitis virus (JE), Nipah virus (N), Phnom Penh Bat virus (PP), Sindbis virus (Sb), Issyk-Kul (Keterah Virus), Leptospira bacteria and SARS coronavirus (Sc). Chiroptera guilds are denote with AI = Aerial Insectivore, C = Carnivore, F = Frugivore, I = Insectivore and N = Nectarivore whereas their roost are denote with B = buildings, Cv = Cave, Fs = Forests, P = Plantations and T = Trees. Chiroptera climb limits were recorded as maximum climb height. The geographical range of bats are classified into E¹ = Endemic to Indomalayan Realm; E² = Endemic to Indonesia & Malaysia; E³ = Endemic to Malaysia and N = Native.
While four-bank harp and mist nets were used to trap the bats (chiropterans), these devices were constructed in open spaces between trees after adapting with procedures of Jayaraj et al. \[2\]. In consideration, the trees should have ten meter distances, a vital criteria to construct the \(50 \times 50\ (2500\ \text{m}^2)\) transect. Entanglement and capture of bats were carried out hourly every day after sunset and before sunrise (between 1830 and 0630). All bats were measured for size and weight using Vernier caliper and portable analytical balance (sensitivity \(\pm 0.01\ \text{kg}\)). Specific measurements of bats such as ear, head-body, tail, forearm, tarsus and hind foot were used to distinguish the bat identity. Descriptive measurements of bats such as forearm length and weight were used to construct the Body Condition Index adopted from Suba et al. \[3\]. In the presence of negative values, the body mass scale was adapted with the values and separated by 0.5 differential margins that give rise to underweight, ideal, overweight and obese. Bat identity were associated with services, dwelling locations, possible pathogen associations, guilds and roosts, climb limits and range \[4,5\]. All chiropteran were released at the site of capture after data acquisition. Recapture was prevented by excluding bats with same gender and size that have hair shavings on their hind foot. The data was interpret into diversity values of Shannon, Simpson, Evenness, Menhinick and Margalef using Paleontological Statistics Software Package (PAST) v.3 because these constructs accurately profile bats by abundance, diversity and richness at each site.

### Acknowledgments

Authors in this data article extend their appreciation to Mr. Hasrulzaman Hassan Basri, Mr. Azuan Roslan and Mr. Syamsul Aznain Ariffin for assistance and guidance at the field and also to authorities from Department of Wildlife and National Parks, Forestry Department and Central Terengganu Development Authority (KETENGAH) for permits and assistance. This work was supported by Trans-Disciplinary Research Grant Scheme (TRGS-59373, 2015), Niche Research Grant Scheme (NRGS-5313-286, 2015) and Geopark Kenyir Development Fund (KGDR-53167-3, 2015).

### Transparency document. Supplementary material

Transparency document associated with this article can be found in the online version at [https://doi.org/10.1016/j.dib.2018.11.058](https://doi.org/10.1016/j.dib.2018.11.058).
References

[1] G. Pounsin, N.S. Wahab, A. Roslan, M.A. Zahidin, E. Pesiu, N.A.M. Tamrin, M.T. Abdullah, Diversity of bats in contrasting habitats of Hulu Terengganu Dipterocarp Forest and Setiu Wetland BRIS forest with a note on preliminary study of vertical stratification of Pteropodid Bats, Trop. Life Sci. Res. 29 (2018) 51–69.

[2] V.K. Jayaraj, N.F.D.A. Tahir, N.A. Udin, N.F.K. Baharin, S.K. Ismail, S.N.A. Zakaria, Species diversity of small mammals at Gunung Stong State Park, Kelantan, Malaysia, J. Threat. Taxa 4 (2012) 2617–2628.

[3] J. Suba, V. Vintulis, G. Petersons, Body weight provides insights into the feeding strategy of swarming bats, Hystrix Ital. J. Mammal. 22 (2010) 179–181.

[4] A.C. Hughes, C. Satasook, P.J. Bates, S. Bumrungsri, G. Jones, The projected effects of climatic and vegetation changes on the distribution and diversity of Southeast Asian bats, Glob. Change Biol. 18 (2012) 1854–1865.

[5] IUCN (International Union for Conservation of Nature), The IUCN Red List of Threatened Species, Version 2017-3, 2017. (http://www.iucnredlist.org) (Accessed 3 July 2018).