



# Confirmation of Skywalker Hoolock Gibbon (*Hoolock tianxing*) in Myanmar Extends Known Geographic Range of an Endangered Primate

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## Abstract

Characterizing genetically distinct populations of primates is important for protecting biodiversity and effectively allocating conservation resources. Skywalker gibbons (*Hoolock tianxing*) were first described in 2017, with the only confirmed population consisting of 150 individuals in Mt. Gaoligong, Yunnan Province, China. Based on river geography, the distribution of the skywalker gibbon has been hypothesized to extend into Myanmar between the N'Mai Kha and Ayeyarwaddy Rivers to the west, and the Salween River (named the Thanlwin River in Myanmar and Nujiang River in China) to the east. We conducted acoustic point-count sampling surveys, collected noninvasive samples for molecular mitochondrial cytochrome *b* gene identification, and took photographs for morphological identification at six sites in Kachin State and three sites in Shan State to determine the presence of skywalker gibbons in predicted suitable forest areas in Myanmar. We also conducted 50 semistructured interviews with members of communities surrounding gibbon range forests to understand potential threats. In Kachin State, we audio-recorded 23 gibbon groups with group densities ranging between 0.57 and 3.6 group/km<sup>2</sup>. In Shan State, we audio-recorded 21 gibbon groups with group densities ranging between 0.134 and 1.0 group/km<sup>2</sup>. Based on genetic data obtained from skin and saliva samples, the gibbons were identified as skywalker gibbons (99.54–100% identity). Although these findings increase the species' known population size and confirmed distribution, skywalker gibbons in Myanmar are threatened by local habitat loss, degradation, and fragmentation.

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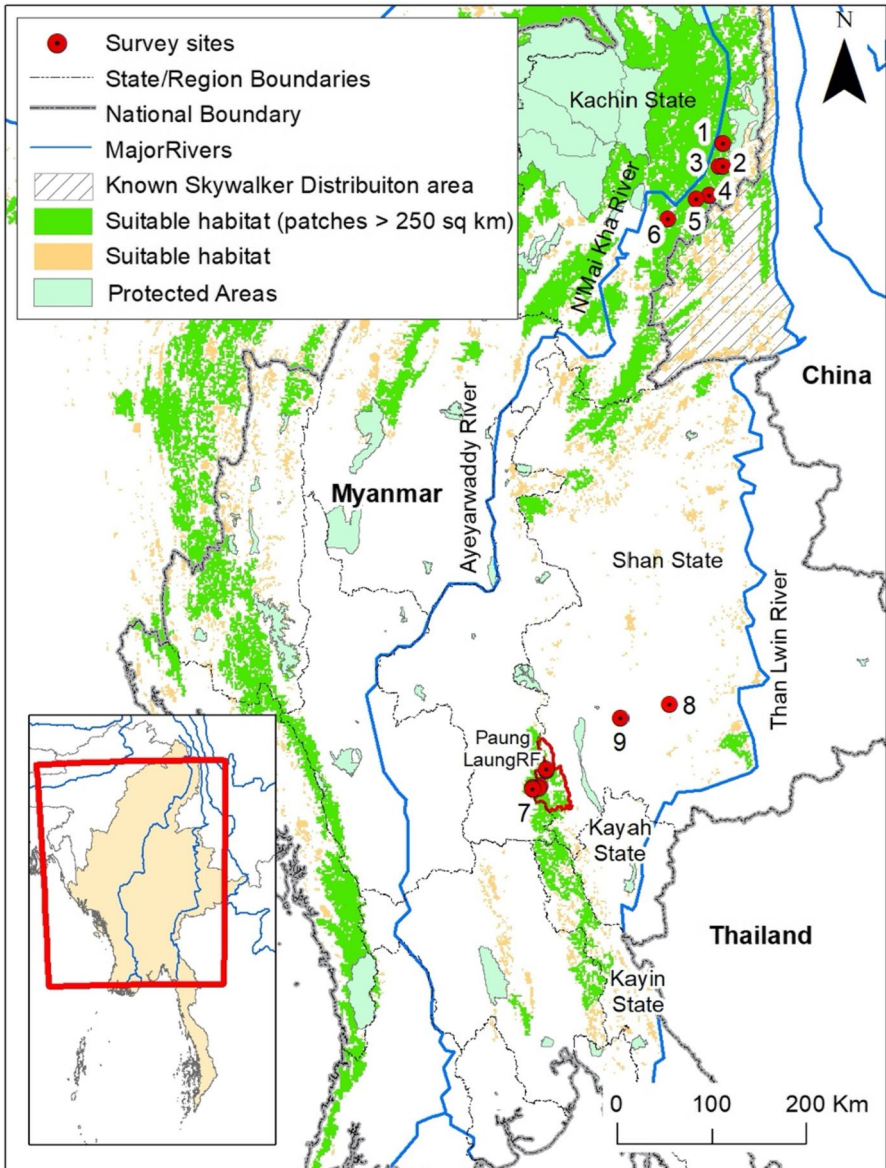
Most of the skywalker gibbon population in Myanmar exists outside protected areas. Therefore, the IUCN Red List status of the skywalker gibbon should remain as Endangered.

**Keywords** Hylobatidae · Primate · Species range · Endangered · Conservation · IUCN Red List

## Introduction

Effective designation and management of conservation areas for wildlife is increasingly important in a world of limited resources and intensifying anthropogenic pressure (Torres-Romero *et al.*, 2023). Basic estimates of animal populations, such as presence/absence and density, are essential to select appropriate conservation areas and inform wildlife management and international conservation policy. The International Union for Conservation of Nature (IUCN) uses estimates of the population size and distribution of a species as essential criteria for defining conservation status (IUCN, 2022). Such estimates often are difficult to obtain, particularly with cryptic or highly morphologically similar species. Animals with similar morphology can represent genetically distinct and divergent lineages, despite their similar appearance. To protect biodiversity, particularly among Hylobatidae, which have extraordinarily high genome plasticity (Carbone *et al.*, 2014), it is important to identify genetically distinct populations and their distributions.

Skywalker hoolock gibbons (*Hoolock tianxing*) (hereafter referred to as skywalker gibbons) were first described in 2017, with the only confirmed population of 150 individuals occurring in Mt. Gaoligong, Yunnan Province, China (Fan *et al.*, 2017). Based on gibbon ecology and the biogeographic patterns of large rivers acting as dispersal barriers that demarcate speciation boundaries in gibbons (Groves, 1967; Thinh *et al.*, 2010a, b), the distribution of the skywalker gibbon is hypothesized to extend into Myanmar between the N'Mai Kha and Ayeyarwaddy Rivers to the west and the Salween River (named the Thanlwin River in Myanmar and the Nujiang River in China) to the east (Fan *et al.*, 2017) (Fig. 1). A historical gibbon specimen at the Natural History Museum, London, collected in 1913 from the Goteik region of Shan State, Myanmar, which is located between these two riverine boundaries, was morphologically identified as a skywalker gibbon (Fan *et al.*, 2017). However, no live skywalker gibbons have been confirmed as occurring in Myanmar over the past century. According to historical records and a limited population survey conducted in 2010, before the recognition of the skywalker gibbon as a distinct species, eastern hoolock gibbons (*Hoolock leuconedys*) were described as present in Kachin, Shan, Kayah, and Kayin States in Myanmar, which are now known to be within the potential range of the skywalker gibbon (Geissmann *et al.*, 2013). During the 2010 assessment, the estimated gibbon population numbers were 89,000 in the Northern Kachin Forest complex, 150,000 in Southern Kachin-Northern Sagaing forests, 50,000 in Shan state subtropical forests, and 16,000 in Kayah-Kayin Montane rainforests (Geissmann *et al.*, 2013).



**Fig. 1** Survey site locations for acoustic gibbon monitoring conducted between December 2021 and March 2023. Site locations were located between the Ayeyarwaddy/N'Mai Kha and Salween (named Thanlwin in Myanmar and Nujiang in China) rivers. Habitat suitability determined according to Ngwe Lwin *et al.* (2021).

Primate surveys have historically been limited in Myanmar given its complex history of civil ethnic conflict concentrated near protected areas and reserve forests. Many of the historical hoolock gibbon records for Myanmar were collected by the Bombay Natural History Society's "Mammal Survey of India, Burma and

Ceylon” conducted between 1913 and 1915 (Wroughton, 1915), the Vernay-Hopwood Chindwin expedition of 1935 (Carter, 1943), and the Vernay-Cutting expedition conducted between 1938 and 1939 (Anthony, 1941). There is a considerable gap in records between 1940 and the 1980s. The most recent assessments in 2010 were limited in several geographic regions due to security issues. The accuracy of gibbon population estimates generated in 2010 is unknown, with the recognition of the skywalker gibbon as a separate species from the eastern hoolock gibbon further complicating the interpretation of these population estimates.

Our overarching goal is to provide accurate information for the skywalker gibbon species distribution and threats to guide updating of the IUCN Red List status. Our objectives were to: (1) conduct acoustic population surveys of gibbons in potential skywalker gibbon range; (2) conduct morphological and DNA identification of any individuals found; and (3) assess threats to new gibbon populations identified in Myanmar through the use of key informant interviews.

## Methods

### Evaluation of Predicted Suitable Habitat and Selection of Gibbon Population Survey Sites

To select survey sites between the Ayeyarwaddy and N’Mai Kha Rivers to the west and Salween River to the east, we collated current and historical records of hoolock gibbons from peer reviewed publications (Fan *et al.*, 2011, 2017, Lwin *et al.*, 2021, 2022) and published reports (Geissmann *et al.*, 2013; Htoo & Grindley, 2010). In a previous study, we used 414 independent locations with documented gibbon presence in 1914–2019 to model the predicted range of hoolock species (Lwin *et al.*, 2021). We identified nine forest areas with predicted hoolock species presence that fell within the estimated range of the skywalker gibbon in Myanmar. The largest forest areas within the theoretical skywalker gibbon range were in Kachin, Shan, Kayah, and Kayin States, including Paung Laung Reserve Forest (Fig. 1) (Lwin *et al.*, 2021).

To improve the chance of locating gibbons in selected survey areas, we gathered additional information through interviews with local wildlife conservation organizations (Wapalaw Wildlife Watch and Friends of Wildlife) and members of the Myanmar Timber Enterprise, which conducts elephant logging operations throughout the proposed skywalker gibbon habitat. We telephoned local wildlife researchers, bird guides, and forestry staff to ask them questions about gibbon presence and hunting pressures. We selected sites to survey if they had: 1) a high likelihood of gibbon presence based on occupancy models; 2) a confirmed recent report of a gibbon sighting based on telephone interviews; and 3) were in close proximity to known populations of hoolock gibbon species and/or collection localities for historical museum specimens of skywalker gibbons. We also tried to select locations as far west and south as feasible to determine the western and southern-most borders of the range.

**Table 1** Number of groups and density estimates for gibbons identified in Kachin and Shan State, Myanmar, resulting from acoustic population surveys conducted between December 2021 and March 2023

Survey location	No. groups	Density estimate (groups / km <sup>2</sup> )	2.5% CI	97.5% CI	No. listening post transects	Forest size (km <sup>2</sup> )	Elevation (m)	Predominant forest type	Anthropogenic activity*	PA status	Land use status
1. Jid Lwai	3	0.79	0.04	1.6	1	~10	1500	Broadleaved evergreen forest	Forest conversion for subsistence rice farming and commercial cardamom; hunting; NTFP collection	NPA	NP (adjacent to Imawbum National Park)
2. Lang Yang	3	3.6	0.6	7.8	1	~50	1700	Broadleaved evergreen forest	Forest conversion for subsistence rice farming; hunting; NTFP collection	NPA	NP (South of Imawbum National Park)
3. Kyauk Phyu	12	2.7	1.1	4.3	1	~50	2000	Broadleaved evergreen forest	Stone mining	NPA	NP (South of Imawbum National Park)
4. Tamu kone	3	2.6	0	5.3	1	~20	2000	Broadleaved evergreen forest	Shifting cultivation for maize and cardamom cultivation; hunting	NPA	NP
5. Chaung Maw	1	2.3	0	5	1	~20	2000	Broadleaved evergreen forest	Shifting cultivation	NPA	NP
6. Sei Law	1	0.57	0	1.5	1	~40	600	Mixed evergreen forest	Forest conversion for agriculture	NPA	CMF
7. Paung Laung	19	0.85	0.41	1.56	4	~1,700	300 – 1,700	Mixed deciduous forest	Dam construction, logging	NPA	RF

Table 1 (continued)

Survey location	No. groups	Density estimate (groups / km <sup>2</sup> )	2.5% CI	97.5% CI	No. listening post transects	Forest size (km <sup>2</sup> )	Elevation (m)	Predominant forest type	Anthropogenic activity*	PA status	Land use status
8. Sat Taw	0	0	0	0	1	~100	1000	Mixed deciduous forest	Shifting cultivation	NPA	RF
9. Mae Nei Taung	2	0.134	0.076	0.344	1	~3	1,500 – 2,200	Mixed coniferous evergreen forest	Crop cultivation in mountain area	NPA	CMF

\*PA = protected area under the Myanmar Protected Area System; *NTPP* = nontimber forest product; *NPA* = nonprotected area under the Myanmar Protected Area System; *RF* = reserved forest is the best quality and higher commercial value forest, where the public has no harvesting rights (Source: Myanmar Forest Department 2020); *CMF* = community managed forests involve all sustainable forest management and utilization activities, in which the local community itself is involved, including establishing new plantations and managing existing forests, to create employment and income opportunities from subsistence to commercial purpose, to generate food, to stabilize ecosystems and to improve environmental conditions (Source: Community Forestry Instructions Myanmar 8 May 2019); *NP* = no protections documented under Myanmar Forest Law, Wildlife Law or Vacant, Fallow and Virgin Land Law

We ultimately conducted acoustic population monitoring in nine forest blocks between the Ayeyarwaddy/N'Mai Kai and Salween rivers in Kachin and Shan States (Table 1). During the study period, sites in Kayah and Kayin States were inaccessible due to conflict (Fig. 1).

1. Jid Lwai (26°11' N, 98°19'), Myitkyina District, Kachin State, in a forest block adjacent to Imawbum National Park. The forest consists of broadleaved evergreen forest surrounded by shifting cultivation, predominantly cardamon.
2. Lang Yang and 3. Kyauk Phyu Taung (25°58' N, 98°19' and 25°57' N, 98°17'), Hsaw Law Township, Myitkyina District, Kachin State. The forest consists of mountainous broadleaved evergreen forest south of Imawbum National Park.
3. Tamu Kone (25°41' N, 98°11'), Chibwe Township, Myitkyina District, Kachin State. The forest consists of broadleaved evergreen forest with shifting cultivation for corn and cardamon plantations.
4. Chaung Maw (25°39' N, 98°02'), Wine Maw Township, Myitkyina District, Kachin State. The forest is located east of Sha Ngaw River, a tributary of the N'Mai Kha River, and is classified as remnant broadleaved evergreen forest surrounded by shifting cultivation.
5. Sei Law (25°27' N, 97°45'), Wine Maw Township, Myitkyina District, Kachin State. The forest is located in community managed forest near the N'Mai Kha River, and is a small mixed evergreen forest block on a north-south mountain range surrounded by a plantation area.
6. Paung Laung Reserve Forest (19°45'–20°28' N, 96°22'–96°49'), Pinlaung Township, Taunggyi District, Southern Shan State. The forest is classified as mixed deciduous in lower elevations and broadleaved evergreen forest in higher elevations (Kitahara *et al.*, 2019; Mon *et al.*, 2009).
7. Sat Taw Reserve Forest (20°36'–20°43' N, 97°45'–97°52'), Mongnei Township, Langkho District, Southern Shan State. The forest is classified as mixed deciduous forest, with some conifers present at higher elevations, with shifting cultivation.
8. Mae Nei Taung Mountain Range (20°38'–20°48' N, 97° 18'–97°23'), Hopong Township, Taunggyi District, Southern Shan State. The forest is classified as vacant land, with the remaining forested areas managed by local communities using traditional practices. The area is hilly and classified as evergreen forest, with mixed coniferous forests, hill savannas and Rhododendron forests found at higher elevations along the mountain range. Crops are grown in the mountain area, predominantly Assyrian plum (*Cordia myxa*), tea (*Camellia sinensis*), opium poppy (*Papaver somniferum*), and coffee (*Coffea arabica*).

## Acoustic Population Survey Methods

We surveyed three forest blocks in Shan State between December 2021 and March 2023 and six forest blocks in Kachin State between March and May 2022. We used acoustic point-count sampling methods (i.e., triangulation), using established

methods (Brockelman & Srikosamatara, 1993; Buckley *et al.*, 2006; Cheyne *et al.*, 2008; Brockelman *et al.*, 2020; Hankinson *et al.*, 2021), to investigate the presence of gibbons and to estimate gibbon density within each area (Cheyne *et al.*, 2008). We established three listening posts in a triangular array, 300–600 m apart, within each site within forest blocks. We established sites 1–2 km apart to form an array. To avoid sampling bias due to elevation, we set up listening posts on hills/ridges on both sides of a mountain/valley. We avoided listening posts at the edge of forests or large rivers, where less gibbons are expected to be present. Gibbons require continuous canopy so degraded areas, rivers, and areas with highly fragmented canopy are not ideal habitats for gibbons and will yield an inaccurate estimate of the population (Cheyne *et al.*, 2016; Hamard *et al.*, 2010). We visited each array of three listening posts on 4–5 consecutive days from 06h00 to 11h00 am and recorded all song types at 3-min intervals. Because of local regulations and safety concerns, field teams could not enter forest areas long before or after dark. For each song heard, we recorded bearing, distance, song type (e.g., solo, duet, great calls (female vocalization)), and start and end time.

### Other Field Sampling Methods

Following morning acoustic monitoring periods, we attempted to visually locate individual gibbons to take photographs of them and collect noninvasive biospecimens. We collected chewed plant samples ( $n = 11$ ) to recover saliva from 11 individual gibbons observed feeding in Paung Laung using methods described previously (Smiley Evans *et al.*, 2016). Specifically, we collected samples of dropped plants or fruits when a gibbon moved to a new foraging tree, cut out the portion of the discarded plant or fruit material that was most masticated using disposable sterile scalpel blades, and placed plant material in DNA/RNA Shield (Zymogen, Inc). We also opportunistically obtained a dry skin sample from a hunter in Lon Than village in Tamu Kone area. We preserved the dry specimen in a plastic bag at ambient temperature. We also obtained a single piece of skin from a museum specimen from the Natural History Museum of the United Kingdom (NHMUK 1933.7.29.15) collected in 1913 from Gokteik, northern Shan State (22°21'N, 96°55'E) that was provisionally classified as a skywalker gibbon based on morphology (Fan *et al.*, 2017).

### Photographic Morphologic Assessment

We examined photographs from individual gibbons from Shan State. We examined photographs visually for discrete external characteristics that distinguish between skywalker gibbons and other hoolock species based on descriptions from Mt. Gao-ligong, China (Fan *et al.*, 2017). We examined the following characteristics that are unique to *H. tianxing* and distinguish the species from eastern hoolock gibbons (*Hoolock leuconedys*): 1) the eyebrow streaks are thinner and separated by a large gap; 2) the beard is completely black or brown instead of white; 3) white hair is absent in the suborbital area; 4) the genital tuft is black, brown, or dark gray instead



of whitish; and 5) adult females have incomplete white face rings, with only sparse white hairs present on the lateral orbital and suborbital regions (Fan *et al.*, 2017).

## Molecular Analyses

We used saliva, skin, and museum samples to molecularly verify species identity. We extracted total genomic DNA from saliva and the skin sample using Qiagen DNeasy kits (Qiagen, Inc.). From the saliva we amplified a 284-bp fragment of the mitochondrial cytochrome b (cytb) gene using primers 5'-ATGATACGAAAACCATCGTTG-3' and 5'-GCGAATAATTCAACCATAGTTC-3'. We performed amplification for 45 cycles, each with a denaturation step at 94 °C for 30 s, annealing at 52 °C for 30 s, and extension at 72 °C for 30 s, followed by a final extension step at 72 °C for 5 min. From the skin sample, we amplified the complete cytb gene by using previously described protocols (Thinh *et al.*, 2010a, b). We checked PCR amplifications on 1.5% agarose gels. Subsequently, we cleaned PCR products with Spin-Prep PCR Clean-up kits (Milipore) and sent them for Sanger-sequencing at Eurofins Genomics using the corresponding amplification primers. We checked sequence electropherograms with 4Peaks 1.8 and aligned sequences in SeaView 5.0.4 (Gouy *et al.*, 2010). We also extracted the complete cytb sequence from an unpublished (C. Roos) complete mitochondrial genome of an adult male gibbon specimen in the NHMUK. We deposited this and newly generated sequences in GenBank, which are available under accession numbers OR159231-OR159233. We performed species identification using the online Basic Local Alignment Search Tool (BLAST) of the National Center for Biotechnology Information (NCBI) using standard settings.

## Threat Assessment Surveys

We conducted key informant interviews in each site to assess overall anthropogenic activities present. Because there are no protected areas in Shan State and wildlife threat assessments have not previously been performed in this region, we conducted these key informant interviews to determine whether similar threats may impact gibbons in this region compared with others in Myanmar. We conducted interviews in 12 villages in Shan State between December 2021 and July 2022 to gather local ecological knowledge about gibbons and determine potential anthropogenic threats impacting any existing populations. We asked respondents about their religious affiliation, because it potentially had bearing on the level of hunting threat posed. We randomly selected villages based on proximity to two forest areas (Paung Laung and Mae Nei Taung Forest Reserves) where skywalker gibbon populations were confirmed to occur based on the combined results of our acoustic population surveys and DNA analyses. Villages ranged from 1 to 4 km from the edge of the forest. Within each village, we prioritized interviews with hunters or former hunters to obtain the most accurate information on gibbon threats. We invited individuals to be interviewed if they were aged 18 years or older and if they were likely to be engaged in activities posing threats to gibbons (e.g., hunting, logging, or forest disturbance).

Given the sensitive nature of hunting, we identified potential respondents within a household by asking less direct questions, such as “who spends the most time in the forest?” and “who has the most contact with animals?” We conducted only one interview per household to ensure independence of responses and limit biases. All respondents gave informed oral consent to partake in the survey after being informed of the purpose and description of the study. Participation was voluntary and respondents were able to terminate their participation in the study at any time.

We created a standard Burmese-language questionnaire for all interviews, consisting of four sections comprised of 39 individual questions in total (Supplementary Material). Section one gathered basic demographic respondent information; section two focused on occupation and potential hunting activities in the forest; section three gathered information on gibbon knowledge, including respondents’ awareness and personal encounters with gibbons and potential threats to gibbon populations; and section four asked questions on forest cover changes and the potential reasons for such changes. We used Burmese research assistants with extensive experience working in hunting and agriculturally based communities to conduct questionnaires. We selected research assistants based on their ability to empathize with the human desire for hunting to increase relatability with respondents.

## Statistical Methods

To estimate gibbon group density, we used the Acoustic Spatial Capture-Recapture (ASCR) R package developed for the IUCN SSC Primate Specialist Group Section on Small Apes (<http://gibbons.asia/tag/ascr>) (Stevenson, 2014). This package fits spatially explicit capture-recapture (SECR) models to acoustic data and has been used as an efficient density estimation method in previous gibbon studies (Kidney *et al.*, 2016). SECR models require the use of a mask—a fine grid of latitude and longitude coordinates around each trap (array of listening posts)—to provide a set of plausible locations for each detected call. We set the maximum distance between each mask point and the closest trap to 2,000 m. We set the distance between adjacent mask points to 200 m. We determined the most appropriate detection function based on the Akaike Information Criterion (AIC). We ultimately used the half-normal detection function to model density:  $g(d;g_0,\sigma) = g_0\exp\left(\frac{-d^2}{2\sigma^2}\right)$ , where  $d$  is the density of groups that call on a single day. Further division by the probability that a group calls on a given day (estimated from external data) yields an estimate of gibbon group density (in groups per hectare).  $g_0$  indicates the intercept for either the halfnormal or hazard-rate detection function, and  $\sigma$  is the scale parameter for either the halfnormal or hazard-rate detection function.  $g_0$  was fixed as 1 for the single occasion surveys and  $d$  at 0 m (Kidney *et al.*, 2016). We conducted analyses separately for each survey site (four sites in Paung Laung and one site in each of the eight remaining locations).

To calculate suitable skywalker gibbon habitat existing within and outside protected areas, we based overall habitat suitability estimates on earlier hoolock gibbon occupancy models that included skywalker gibbon data (misclassified as

eastern hoolock gibbons, because the species had not yet been identified in Myanmar) (Lwin *et al.* 2021) overlapped with the new skywalker gibbon range proposed through this study according to the IUCN Red List Assessment Guidelines (<https://www.iucnredlist.org/assessment/process>). We calculated km<sup>2</sup> of suitable habitat both within and outside of protected areas by overlaying shape files from Lwin *et al.* (2021) with Protected Area System boundaries provided by the Myanmar Forest Department.

To better understand dynamics contributing to threats in Skywalker gibbon forest sites, we assessed whether respondent sociodemographic characteristics, hunting activities, gibbon knowledge, and forest cover change varied between Paung Laung and Mae Nei Taung Forest sites. Interviews collected data on respondents' age, gender, ethnicity, religion, occupation, education level, and frequency of visiting the local forest, all of which have been shown to influence knowledge and awareness of gibbons and other biodiversity within indigenous communities in China (Turvey *et al.*, 2017; Zhang *et al.*, 2020), and data on local interactions with gibbons and associated environmental change (perceptions of local gibbon population decline, reported hunting of gibbons, taboos against hunting gibbons, perceptions of local forest loss). To investigate variation in these responses between survey sites, we created generalized linear models (GLMs) with Gaussian or binomial error structures in R Version 4.2.2 and RStudio. However, most of the variables showed little variation across the overall respondent sample, showed collinearity with survey site (e.g., the two sites contained different ethnic and religious groups, meaning that the relative contribution of these factors to variation in interview responses could not be investigated independently of survey site), or had no responses from one of the two survey sites (e.g., no respondents reported gibbon hunting from Mae Nei Taung). As such, differences between the two survey sites were only investigated statistically for respondent age, whether respondents visited the forest at least once per week, reported gibbon declines, taboos against hunting gibbons, and perceptions of local forest loss. Given the relatively small respondent sample size ( $n = 24$  in Paung Laung and  $n = 26$  in Mae Nei Taung), we investigated relationships between each response variable and survey site using a series of separate models containing a single response variable and survey site as a single predictor variable to maximize statistical power within each model.

## Ethical Note

All study procedures involving gibbons were non-invasive in nature. All gibbon survey procedures were approved by the Myanmar Forest Department and the Shan State Forest Department. All human surveys were conducted anonymously and no personal identifying information was collected. All human survey procedures were approved by the Institutional Review Board of the University of

California, Davis (IRB #1863876-1) and the Myanmar Forest Department. The authors declare that they have no conflict of interest.

**Data Availability** The datasets gathered and analyzed during the current study are available from the corresponding author on reasonable request.

## Results

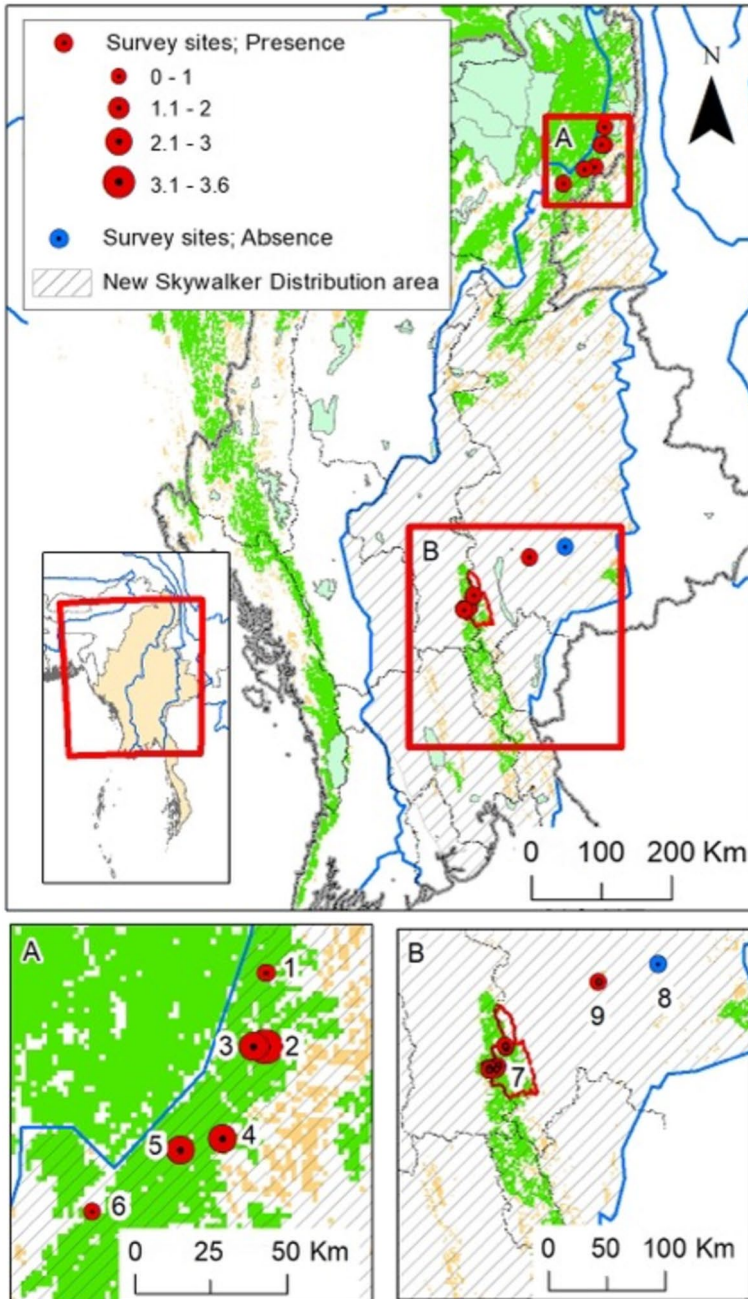
### Population Surveys

In Kachin State, we recorded 23 gibbon groups at six survey sites: 12 groups at Kyauk Phyu, three groups each at Jid Lwai, Lang Yang and Tamu Kone, and one group each at Chaung Maw and Sei Law. Group densities estimated using the ASCR method varied between 0.57 and 3.6 group/km<sup>2</sup> (Table 1). In Shan State, we recorded 21 gibbon groups at two survey sites: 19 groups in Paung Laung, and two groups in Mae Nei Taung. We did not locate any gibbons in Sat Taw Reserve Forest. Group densities estimated using the ASCR method varied between 0.134 and 0.85 group/km<sup>2</sup> (Table 1; Fig. 2).

The group density estimates for non-protected forest areas where gibbons were present (Jid Lwai, Lang Yang, Kyauk Phyu, Tamu Kone, Chaung Maw, Sei Law, Paung Laung, Mae Nei Taung) were not substantially different from the group density estimates for two of the more recently designated protected areas in Myanmar containing eastern hoolock gibbons, the Indawgyi Biosphere Reserve (0.71 group/km<sup>2</sup>, 95% confidence interval [CI] 0.4–1.27) and Mahamyaing Wildlife Sanctuary (0.97 group/km<sup>2</sup>, 95% CI 0.71–1.32) (Lwin *et al.*, 2022, Tun *et al.*, 2023). The main forest area in Shan State that we surveyed, Paung Laung Forest Reserve, was similar in size (1,337 km<sup>2</sup>) to Indawgyi Biosphere Reserve (1,700 km<sup>2</sup>) and Mahamyaing Wildlife Sanctuary (1,180 km<sup>2</sup>) but has a much lower level of legal conservation protection.

### Morphological Descriptions

We evaluated 60 photographs of five individual gibbons (15 photographs of two adult females and one adult male from Paung Laung Reserve Forest, and 45 photographs of two adult males in Mae Nei Taung). All five individuals were very similar and could be clearly identified as skywalker gibbons based on distinct morphological traits. In the adult males, the eyebrows are well separated and white-colored hair is scant or absent around the eyes, chin, and genital tuft, making these areas appear brown or black (close to the color of the rest of the body). These features distinguish them from eastern hoolocks, which have many white hairs around their eyes and a white chin and genital tuft. In the adult females, there is less white fur around the eyes and the white fur does not form the conspicuous white face ring (connecting the chin and eyebrows) that is seen in the eastern hoolock gibbon (Fig. 3).



**Fig. 2** Proposed new skywalker gibbon (*Hoolock tianxing*) distribution area in Myanmar. Locations of survey sites in Kachin State (A) and Shan State (B), showing presence or absence of gibbons as assessed between December 2021 and March 2023. Size of circles in insets A and B represent gibbon densities.



**Fig. 3** Morphological comparison of hoolock gibbon species and subspecies. Photographs of *Hoolock hoolock*, *Hoolock hoolock mishmiensis*, *Hoolock leuconedys*, and *Hoolock tianxing* (China) sourced and adapted from Fig. 2 in “Description of a new species of Hoolock gibbon (Primates: Hylobatidae) based on integrative taxonomy” (Fan *et al.*, 2017). *H. tianxing* (Myanmar) photographs by Nature Conservation Society Myanmar and additional *H. tianxing* (China) photographs by P-F Fan. Top row: Male gibbons. Male *H. tianxing* have less convex tops of head, straighter white brows, and less conspicuous white beards compared with *H. hoolock*, *H. h. mishmiensis*, and *H. leuconedys*. Bottom row: Female gibbons. Female *H. tianxing* have fainter white face rings compared with *H. hoolock*, *H. h. mishmiensis*, and *H. leuconedys*.

## Molecular Identification

We obtained a partial (217 bp) cytb sequence from one chewed plant sample collected in Paung Laung and the complete (1140 bp) cytb sequence from the skin sample from Tamu Kone. We also obtained the complete cytb sequence from the mitochondrial genome of the skin museum specimen from Gokteik. Ten chewed plant specimens collected did not contain sufficient DNA to detect gibbon genetic material. All three sequences obtained were identical in the 217 bp partial cytb sequence, whereas the two full-length cytb sequences differed in three positions (0.26%). Using BLAST search, we identified all three sequences unambiguously as skywalker gibbons. For the partial cytb sequence from Paung Laung we revealed 99.54–100% (0–1 differences) identity with skywalker gibbon, 97.70–98.62% (3–5 differences) identity with eastern hoolock gibbon and 97.24–97.70% (5–6 differences) identity with western hoolock gibbon (*Hoolock hoolock*) (100% query coverage). For the complete cytb sequence from Tamu Kone, we obtained 99.74–99.91% (1–3 differences) identity with skywalker gibbon, 97.70–98.42% (11–18 differences) identity with eastern hoolock gibbon and 96.84–97.19% (32–36 differences) identity with western hoolock gibbon (100% query coverage). For the complete cytb sequence of the museum specimen from Gokteik, BLAST results revealed 99.65–99.82% identity (2–4 differences) with skywalker gibbon, 98.33–98.95% (12–19 differences) identity with eastern hoolock gibbon, and 96.93–97.28% (31–35 differences) identity with western hoolock gibbon (100% query coverage).

## Threat Assessments

We interviewed 50 people in Paung Laung (five villages, 24 respondents; mostly Karen ethnicity) and Mae Nei Taung (seven villages, 26 respondents; mostly Pa'O ethnicity). People in Paung Laung were equally divided between Buddhist and Christian religious affiliation and people in Mae Nei Taung were all Buddhist. Respondents varied in age from 19 to 85 years, 96% ( $n = 48/50$ ) were male, and 94% ( $n = 47/50$ ) practiced farming as their primary occupation. Respondents had considerable experience with local forests; 88% ( $n = 44/50$ ) reported that they had visited the forest at least once per month, and 44% ( $n = 22/50$ ) visited at least once per week for farming, collection of fuelwood and nontimber forest products, timber extraction, and hunting (for subsistence and sale). Hunting was reported by 82% ( $n = 41/50$ ) of respondents. There was no statistically significant difference between the two forest areas in respondent age (GLM with Gaussian error structure: estimate =  $-0.56$ , standard error [SE] =  $4.16$ ,  $t = -0.13$ ,  $p = 0.894$ ) or whether respondents visited the forest at least once per week (GLM with binomial error structure: estimate =  $0.09$ , SE =  $0.60$ ,  $z = 0.15$ ,  $p = 0.884$ ). These were the only two sociodemographic indices that showed substantial variation across the respondent sample, indicating that responses can be compared directly between the two areas without needing to control for additional potential confounding effects of respondent characteristics.

Respondents had a high level of awareness and experience of skywalker gibbons. In total, 82% ( $n = 41/50$ ) could identify a skywalker gibbon from a photograph, and 98% ( $n = 49/50$ ) could identify it from a playback of its call. Overall, only one respondent could not recognize the species at all. In total, 82% ( $n = 41/50$ ) had seen and/or heard gibbons in the forest close to their village. Only 34% ( $n = 17/50$ ) of respondents reported the date when they had last seen a gibbon in the local forest with dates ranging from within the past year to 30 years earlier.

Most respondents (70%;  $n = 35/50$ ) felt that gibbons had declined, and there was no statistically significant difference in numbers of respondents reporting gibbon declines between the two forest areas (GLM with binomial error structure: estimate =  $-0.69$ , SE =  $0.63$ ,  $z = -1.10$ ,  $p = 0.270$ ). When asked about local hunting of gibbons, 46% ( $n = 11/24$ ) reported that people hunted gibbons around Paung Laung; two respondents specifically stated that juvenile gibbons were hunted. In contrast, no respondents around Mae Nei Taung reported local hunting of gibbons. Awareness of local hunting in Paung Laung was reported by both Buddhists ( $n = 7/12$ ) and Christians ( $n = 4/12$ ), and from all five villages surveyed around this forest area. Considerably more respondents reported the existence of taboos against hunting gibbons around Mae Nei Taung ( $n = 17/26$ ) compared with Paung Laung ( $n = 9/24$ ); however this difference was not statistically significant (GLM with binomial error structure: estimate =  $-1.15$ , SE =  $0.59$ ,  $z = -1.95$ ,  $p = 0.052$ ).

Of the 34 respondents who described their perceptions for causes of gibbon decline, 85% ( $n = 29/34$ ) of respondents considered that forest cover had decreased locally, with local forest loss reported to have occurred because of expansion of agriculture (including shifting cultivation) and agroforestry, logging, forest burning, and fuelwood collection. More respondents considered that forest cover had

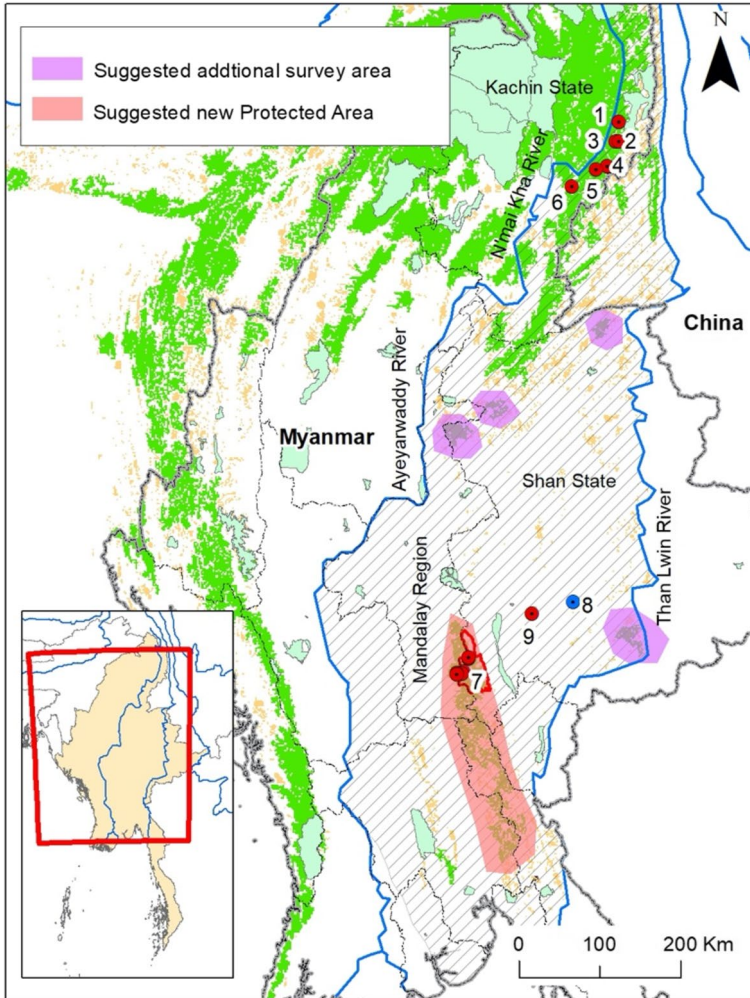
decreased in Paung Laung ( $n = 23/24$ ) compared with Mae Nei Taung ( $n = 19/26$ ); however, this difference was not statistically significant (GLM with binomial error structure: estimate = 2.14, SE = 1.11,  $z = 1.92$ ,  $p = 0.055$ ). Only two respondents thought that gibbon declines could be attributed to hunting.

## Discussion

We report confirmation of skywalker gibbon presence in Myanmar, extending the species' current range beyond China, and further supporting the hypothesis that the species range is demarcated by the N'Mai Kha and Ayeyarwaddy Rivers to the west and Salween River (Thanlwin in Myanmar and Nujiang in China) to the east. Our findings confirm that skywalker gibbons occur as far south as Shan State. Surveys in Kayin and Kayah States were not possible at the time of this study (due to security issues); however, it is likely that gibbon populations previously identified as eastern hoolock gibbons in Kayin and Kayah States are in fact skywalker gibbons. Based on these findings, we conclude that most of the world's skywalker gibbon populations occur in Myanmar.

We confirmed skywalker gibbon presence noninvasively without access to voucher specimens or captive individuals, using chewed, naturally discarded plants. Recent gibbon voucher specimens are not preserved anywhere in Myanmar and historical specimens in international museums come with a degree of uncertainty regarding specific collection sites. Similarly, gibbons coming from zoological institutions within Myanmar may have inaccurate reported locations of origin. Fecal samples have long been the most commonly targeted noninvasive sample type for the identification of gibbon DNA (Oka & Takenaka, 2001; Bradley *et al.*, 2001). However, feces often are a challenging specimen type to collect, given the frequency gibbons defecate in the presence of humans or other predatory animals. Identifying the origin of fecal samples in densely forested locations also can be challenging. Scat detection dogs have recently improved the likelihood of detection of gibbon fecal samples (Orkin *et al.*, 2016); however, this resource was not available in Myanmar. We therefore used chewed, naturally discarded plants for DNA identification (Smiley Evans *et al.*, 2016). We attempted to collect fecal samples during our surveys, but we did not find any. We did recover gibbon DNA from one out of 11 chewed plant specimens. This was a lower rate of DNA detection from chewed plants compared with similar studies performed using these methods with habituated mountain gorillas (*Gorilla beringei beringei*) and golden monkeys (*Cercopithecus mitis kandti*) (Smiley Evans *et al.*, 2016). Our lower rate of detection was likely due to the greater distance at which wild gibbons had to be observed compared to habituated individuals from other species. Prioritizing collection of longer chewed plant specimens is easier the closer they can be observed (Smiley Evans *et al.*, 2016). While we had a low rate of DNA detection, given the abundance of chewed plant samples available, we found this to be a viable method—showing that saliva recovered from this specimen type can be used for DNA identification of wild, nonhabituated gibbons.





**Fig. 4** Proposed new protected area(s) to conserve the southern population of skywalker gibbons in Myanmar.

We used both molecular identification and knowledge of gibbon ecology to determine the new proposed species range. We selected site locations for gibbon surveys located between the major north-south rivers of the Ayeyarwaddy/N<sup>o</sup>Mai Hka and Salween rivers and did not conduct gibbon surveys outside of this river-bound landscape. We made this decision based on other gibbon speciation records that show large rivers represent barriers for gibbon dispersal (Groves, 1967; Thinh *et al.*, 2010a, b). Gibbons do not swim and are largely restricted to the forest canopy making river valleys generally unfavorable for gibbons (Groves, 1967; Thinh *et al.*, 2010a, b). Other gibbon species in Myanmar have species distributions bounded by rivers: the Chindwin River is the boundary between western hoolock and eastern

hoolock gibbons (Groves, 1967), and the Lohit River may act as the boundary between western hoolock and Mishmi Hills hoolock gibbon (*Hoolock hoolock mishmiensis*) (Choudhury, 2013). Based on this information, we did not conduct surveys west of the Ayerwaddy or east of the Thanlwin; however, for further confirmation of the species boundary, this survey effort could be attempted in future.

We found skywalker gibbons in a wide range of habitat types, extending from high elevational seasonal montane forests in the Mt. Gaoligong region of China (Fei *et al.*, 2019) to lower elevation mixed deciduous forests of Shan State, Myanmar, including disturbed forest areas, demonstrating the adaptability of the species. We document an elevational range of 600–2000 meters above mean sea level (MAMSL) on the Myanmar side of the border. There are lower elevation forests in Myanmar within the hypothesized skywalker gibbon range, such as the Bago Yoma. None of these forests, however, were surveyed, because they had no recent gibbon sightings, a criterion for our selection of acoustic population monitoring locations. Skywalker gibbons have been documented between 1596–2685 MAMSL in China (Fan *et al.*, 2011). This wide elevational range is similar to other hoolock gibbons: the western hoolock gibbon has been documented between 15 MAMSL (Ahsan, 2000) and 2300 MAMSL (Geissmann *et al.*, 2013). The highest recorded elevation of a gibbon species is recorded in China, where *Nomascus concolor* has been documented above 2600 MAMSL (Fan & Jiang, 2010; Haimoff *et al.*, 1987).

As further efforts are put toward revising the protected area system in Myanmar, we recommend that areas that support confirmed skywalker gibbon populations be included under protection to curb extremely high hunting and land conversion pressures from surrounding communities. We propose better protection for skywalker gibbons in the southernmost, lower elevation region of the newly defined skywalker gibbon range (Fig. 3). For example, in Shan State, where there are no protected areas, we identified gibbon groups in forested areas that are heavily degraded by logging and other anthropogenic activities. This finding indicates the extreme conservation urgency for this region, as skywalker gibbons remain but in low-quality habitats with high hunting pressures. Much less focus has been placed on protecting remaining forests in this region compared with more northern regions of Myanmar, where large areas of undisturbed forest landscape still exist. Restoration of habitats is a critical component of a multidimensional approach to primate conservation (Chapman *et al.*, 2020) and identifying skywalker gibbons within degraded forests in Myanmar is an opportunity to help support conservation and reforestation in some of these areas. We therefore recommend that a portion of the southernmost habitat for skywalker gibbons also be prioritized for inclusion in the Myanmar protected area system (Fig. 4). Protected areas are positively associated with safeguarding primate species diversity globally (Torres-Romero *et al.*, 2023) and should be incorporated into skywalker gibbon conservation action planning in Myanmar.

Community-based conservation efforts are also urgently needed and might be enacted on a much shorter time frame than that required to establish and manage new government organized protected areas. Based on discussions with communities who live in the survey areas in Kachin State, two communities (Jid Lwai, Site 1; and Tamu Kone, Site 4) stated their willingness to establish community protected areas to protect gibbons and to prevent land grabs for mining. In addition, one community (Sei Law,

Site 6) wanted to start a gibbon conservation awareness program for their community to prevent hunting of gibbons. Shan State communities were enthusiastic for the conservation of gibbons and communities preferred to encourage indigenous and community conserved areas rather than the designation of a government operated protected area given current concerns around indigenous people's rights in Shan State. Forested habitat within indigenous peoples' lands (Garnett *et al.*, 2018) is positively associated with safeguarding primate species diversity globally (Torres-Romero *et al.*, 2023) and has had proven success in conserving primates in other parts of Myanmar (Thompson *et al.*, 2023). Community-managed forests are one option to integrate forest management into national poverty reduction programs in rural areas. These forests have lower and less variable annual deforestation rates than protected ones (Porter-Bolland *et al.*, 2012). Increasingly, a collective effort of stakeholders, including government-led protected areas as well as communities and indigenous peoples' groups, is being recognized as the only effective way to protect and save our closest living relatives (Estrada *et al.*, 2017; Fan, 2017; Pan *et al.*, 2016; Li *et al.*, 2018).

We also identified variation in local gibbon hunting, taboos, and forest loss across different social-ecological landscapes in Myanmar that highlight important local-scale differences in attitudes and interactions with gibbons and their habitat, indicating the need for region-specific community outreach and forest protection as part of future conservation efforts. These findings suggest that the skywalker gibbon population at Paung Laung may be more threatened by both hunting and habitat loss in comparison to other areas, and further research of this nature should be conducted as a matter of urgency to assess the status and viability of this population. More widely, our interview findings also demonstrate the high potential for using an interview survey approach to collect indigenous knowledge about the status of local gibbon populations more widely across Myanmar in the future as part of the conservation toolkit for strengthening the evidence-base for this poorly known species. These methods have been successful in measuring local ecological knowledge to detect cryptic threatened primates, such as the Hainan gibbon (Turvey *et al.*, 2017), and could be expanded in Myanmar.

Skywalker gibbons are currently considered by the IUCN in its Red List assessment as Endangered A4cd (IUCN, 2022) because of the suspected population decline caused by significant and accelerating rates of deforestation and overexploitation for both food and medicinal purposes (Fan *et al.*, 2020). Having confirmed the existence of skywalker gibbon populations in Myanmar, we propose the expansion of the species' range to the Ayerwaddy River as the western demarcation and the Salween River as the eastern demarcation of the range (Fig. 4). Based on this updated distribution range, there is currently an estimated 7,248 km<sup>2</sup> of suitable habitat in China and 31,703 km<sup>2</sup> of suitable habitat in Myanmar (38,951 km<sup>2</sup> in total) for skywalker gibbons. Of the suitable skywalker gibbon habitat within the Myanmar and China Protected Area Systems (areas with official conservation protection status), we determined that there are 1,012 km<sup>2</sup> in China and 1,368 km<sup>2</sup> in Myanmar. Overall, a very small fraction of the suitable habitat for skywalker gibbons in Myanmar exists within the protected area system (~4%).

While skywalker gibbons have now been confirmed to occur in Myanmar, thus increasing the species' global population size and confirmed distribution, threats to the survival of the species across the region are extensive, including local habitat loss,

degradation and fragmentation, and hunting (both for meat and for live-trade) in both China and Myanmar. Nearly half of the remnant population in China lives in landscapes used for agroforestry (Fei *et al.*, 2017), and based on our data, greater than 95% of the population in Myanmar exists outside protected areas. A suspected population decline caused by significant rates of deforestation in Myanmar (estimated at 0.94% annually between 2002 and 2014, totaling >2 million ha forest loss) is projected to continue and potentially accelerate (Bhagwat *et al.*, 2017). A population loss equaling or exceeding 50% can thus be anticipated over three generations (between 2000 and 2045). Therefore, the conservation status of the skywalker gibbon should remain as Endangered A4cd.

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## Declarations

**Inclusion and Diversity Statement** Twelve authors (including joint first, second, and third authors) are contributors from the location where the research was conducted, and participated in study conception, study design, data collection, analysis, and interpretation of the findings.

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