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A new species of *Theloderma* (Amphibia: Anura: Rhacophoridae) from Vietnam

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Abstract

We describe a new species of Theloderma from northwestern Vietnam based on morphological differences and molecular divergence. The loderma annae sp. nov. is distinguishable from its congeners on the basis of a combination of the following characters: Size small, SVL 27.1–28.5 mm in males, 30.3–32.6 mm in females; head longer than wide; vomerine teeth absent; snout long (SL/SVL 0.16–0.19); spines on upper eyelid absent; tibiotarsal projection absent; dorsal skin smooth; dermal fringes on forearm and tarsus absent; dorsal surface gravish green; and throat and ventral surface of arms and thighs brown with white spots.

Key words: Theloderma annae sp. nov., karst forest, molecular phylogeny, taxonomy, Hoa Binh Province

Introduction

The genus Theloderma Tschudi, 1838 is considered as the most poorly known group of tree frogs due to their cryptic habits and rarity of representatives (Nguyen et al. 2014). This genus currently comprises 25 species, of which one-third have been described in the last ten years (Frost 2016). Vietnam contains the greatest number of species in the genus, with 13 species recorded from the country, six of which having been described since 2009, viz. T. bambusicola Orlov, Poyarkov, Vassilieva, Ananjeva, Nguyen, Nguyen & Geissler, T. chuyangsinensis Orlov, Poyarkov, Vassilieva, Ananjeva, Nguyen, Nguyen & Geissler, T. lateriticum Bain, Nguyen & Doan, T. nebulosum Rowley, Le, Hoang, Dau & Cao, T. palliatum Rowley, Le, Hoang, Dau & Cao, and T. vietnamense Poyarkov, Orlov, Moiseeva, Pawangkhanant, Ruangsuwan, Vassilieva, Galoyan, Nguyen, and Gogoleva (Bain et al. 2009; Nguyen et al. 2009; Orlov et al. 2006, 2012; Rowley et al. 2011; Poyarkov et al. 2015). However, Poyarkov et al. (2015) subsequently synonymized T. bambusicola with T. laeve and T. chuvangsinensis with T. palliatum. In addition, Nguyen et al. (2014) recently recorded T. petilum (Stuart & Heatwole) for the first time from Vietnam.

During recent field work in northwestern Vietnam, specimens of a small tree frog species were collected in the karst forest of Ngoc Son - Ngo Luong Nature Reserve in Hoa Binh Province. Molecular comparisons revealed this taxon being nested within the genus Theloderma. Closer morphological examination showed that the specimens from Hoa Binh are clearly distinguished from other known members of Theloderma by a combination of morphological features. The phylogenetic analyses revealed this taxon to be clustered within the *Theloderma* truongsonense group with a strong support value but being distinctly separated from congeners. Due to morphological and molecular differences of the newly-collected specimens to all known species in the genus, we herein describe the newly discovered *Theloderma* population from Hoa Binh as a new species.

Materials and methods

Sampling. Field surveys were conducted in April 2014 by C.T. Pham, H.T. An, H.N. Ngo (hereafter C.T. Pham *et al.*), in October 2014 and in April 2015 by T.Q. Nguyen, C.T. Pham, M.D. Le, H.N. Ngo (hereafter T.Q. Nguyen *et al.*) in Ngoc Son – Ngo Luong NR, Lac Son District, Hoa Binh Province, northwestern Vietnam. Specimens were collected between 19:00 and 23:30 h. After taking photographs, specimens were anaesthetized with ethyl acetate, fixed in 80% ethanol for 4–6 hours, and then later transferred to 70% ethanol for permanent storage. Tissue samples were preserved separately in 95% ethanol. Preserved specimens were deposited in the collection of the Institute of Ecology and Biological Resources (IEBR), Hanoi; Vietnam National Museum of Nature (VNMN), Hanoi, Vietnam; and the Zoologisches Forschungsmuseum Alexander Koenig (ZFMK), Bonn, Germany. Other specimens referred to in this paper were deposited at the American Museum of Natural History (AMNH), Australian Museum (AMS), Chengdu Institute of Biology (CIB), Hanoi National University of Education (HNUE), Graduate School of Human and Environmental Studies, Kyoto University (KUHE), Field ID of Zoological Museum, Moscow University (NAP).

Molecular data and phylogenetic analyses. We used the protocols of Kuraishi *et al.* (2013), modified by Nguyen *et al.* (2015), for DNA extraction, amplification, and sequencing. Fragments of three mitochondrial DNA genes 12S rRNA, tRNA^{val}, and 16S rRNA were amplified using the primers following Kuraishi *et al.* (2013). *Nyctixalus pictus* (Peters) and *Liuixalus romeri* (Smith) were selected as outgroups according to Nguyen *et al.* (2015) and Li *et al.* (2016) (Table 1).

Chromas Pro software (Technelysium Pty Ltd., Tewantin, Australia) was used to edit the sequences, which were aligned using MAFFT version 7 (Katoh & Standley 2013) with default settings. We then checked the initial alignments by eye and adjusted slightly. Phylogenetic trees were constructed by using maximum likelihood (ML) and Bayesian inference (BI). Prior to ML and Bayesian analyses, we chose the optimum substitution models for entire sequences using by Kakusan 4 (Tanabe 2011) based on the Akaike information criterion (AIC). We compared both partitioned and non-partitioned models on our data set, and the latter was selected based upon AIC. The best model selected for ML was the general time reversible model (GTR: Tavaré 1986) with a gamma shape parameter (G: 0.319 in ML and 0.339 in BI). The BI summarized two independent runs of four Markov Chains for 10,000,000 generations. A tree was sampled every 100 generations and a consensus topology was calculated for 70,000 trees after discarding the first 30,001 trees (burn-in = 3,000,000). We checked parameter estimates and convergence using Tracer version 1.5 (Rambaut & Drummond 2009). The strength of nodal support in the ML tree was analyzed using non-parametric bootstrapping (MLBS) with 1000 replicates. We regarded tree nodes in the ML tree with bootstrap values of 70% or greater as sufficiently resolved (Huelsenbeck & Hillis 1993), and nodes with a BPP of 95% or greater as significant in the BI analysis (Leaché & Reeder 2002). Pairwise comparisons of uncorrected sequence divergences (p-distance) were calculated for 16S rRNA fragments only between species of the genus Theloderma.

Morphological characters. Measurements were taken with a digital caliper to the nearest 0.1 mm. Abbreviations are as follows: SVL: Snout-vent length, HL: Head length (from the back of mandible to the tip of snout), HW: Maximum head width (across angle of jaws), SNL: Snout length (from anterior corner of eye to the tip of snout), NS: Distance from nostril to the tip of snout, EN: Distance from anterior corner of the eye to the nostril, IN: Internarial distance, IOD: Interorbital distance, ED: Eye diameter, UEW: Maximum width of upper eyelid, DAE: Distance between anterior corner of eyes, DPE: Distance between posterior corners of eyes, MAE: Distance between angle of jaws and anterior corner of the eye, MPE: Distance between angle of jaws and posterior corner of the eye, MN: Distance from the back of mandible to the nostril, TYD: Tympanum diameter, TYE: Distance from anterior margin of tympanum to posterior corner of the eye, FLL: Forelimb length (from axilla to elbow), HAL: Hand length (from elbow to the tip of third finger), fd1-4: Width of discs of fingers I-IV, fw1-4: Width of fingers I-IV, TFL: Third finger length, OPT: Outer palmar tubercle length, FeL: Femur length (from vent to knee), TbL: Tibia length (from knee to tarsus), TbW: Maximum tibia width, FoL: Foot length (from tarsus to the tip of fourth toe), FTL: Fourth toe length, IMT: Inner metatarsal tubercle length, OMT: Outer metatarsal tubercle length, td1-4: Width of discs of toes I-IV, tw1-4: Width of toes I-IV. Terminology for describing eye coloration in life and webbing formula followed those of Glaw & Vences (1997, 2007). Sex was determined by the presence of nuptial pads and gonadal inspection.

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No	Species	Voucher	Locality	Genbank number	Keterences
1.	Theloderma sp. nov.	IEBR 3732	Vietnam: Hoa Binh	LC168170	This study
2.	Theloderma sp. nov.	IEBR 3733	Vietnam: Hoa Binh	LC168171	This study
Э.	Theloderma sp. nov.	IEBR 3734	Vietnam: Hoa Binh	LC168172	This study
4.	T . asperum	Pet trade	Malaysia: Perak	KT461929	Poyarkov et al. (2015)
5.	T. albopunctatum	VNMN J291	Vietnam: Vinh Phuc: Tam Dao	KJ802914	Nguyen et al. (2015)
6.	T. albopunctatum	VNMN 3540	Vietnam: Lao Cai: Sa Pa	KJ802913	Nguyen et al. (2014)
7.	$T.\ albopunctatum$	VNMN 4404	Vietnam: Kon Tum: Ngoc Linh	LC012854	Nguyen et al. (2015)
8.	T. albopunctatum	VNMN 4406	Vietnam: Thanh Hoa: Xuan Lien	LC012856	Nguyen et al. (2015)
9.	$T.\ albopunctatum$	VNMN PAE 262	Vietnam: Son La: Ta Xua	LC012857	Nguyen et al. (2015)
10.	T. bicolor	VNMN 3536	Vietnam: Lao Cai: Sa Pa	KJ802915	Nguyen et al. (2014)
11.	T. corticale	VNMN J2892	Vietnam: Vinh Phuc: Tam Dao	KJ802916	Nguyen et al. (2014)
12.	T. corticale	VNMN J2932	Vietnam: Tuyen Quang: Na Hang	KJ802917	Nguyen et al. (2014)
13.	T. gordoni	KUHE 32447	Laos: Houaphan	KJ802919	Nguyen et al. (2014)
14.	T. gordoni	VNMN PAE217	Vietnam: Son La: Ta Sua	KJ802918	Nguyen et al. (2014)
15.	T. gordoni	VNMN 4407	Vietnam: Kon Tum: Ngoc Linh	LC012852	Nguyen et al. (2015)
16.	T. laeve	VNMN 4403	Vietnam: Gia Lai: Mang Yang	LC012846	Nguyen et al. (2015)
17.	T. lateriticum	VNMN 1215	Vietnam: Bac Giang: Tay Yen Tu	LC012850	Nguyen et al. (2015)
18.	T. lateriticum	VNMN PAE 226	Vietnam: Son La: Ta Sua	LC012849	Nguyen et al. (2015)
19.	T. lateriticum	AMNH 168757	Vietnam: Lao Cai: Sa Pa	LC012848	Nguyen et al. (2015)
00	T longramm	/IEBR A.0860 k11HE 52581	Malaweia: Narari Samhilan: Kanahoi	A B8/7178	Materii <i>at al (</i> 2014)
-07 07	1. teporosum			AD04/120	Maisul et $at. (2014)$
21.	T. licin	KUHE 19426	Thailand: Nakon Sri Tamarat	LC012859	Nguyen <i>et al.</i> (2015)
22.	T. licin	KUHE 52599	Malaysia: Selangor	KJ802920	Nguyen et al. (2014)
23.	T. nebulosum	AMS R 173409	Vietnam: Kon Tum	JN688168	Rowley et al. (2011)
24.	T. nebulosum	AMS R 173877	Vietnam: Kon Tum	JN688168	Rowley et al. (2011)
25.	T. nebulosum	ROM 39588	Vietnam: Kon Tum	LC012845	Nguyen et al. (2015)
26.	T. palliatum	NAP 2735	Vietnam: Dak Lak	LC012843	Nguyen et al. (2015)
27.	T. palliatum	NAP 2736	Vietnam, Dak Lak	LC012844	Nguyen et al. (2015)
28.	T. palliatum	AMS R173130	Vietnam: Lam Dong	JN688172	Rowley et al. (2011)
29.	T. palliatum	AMS R173131	Vietnam: Lam Dong	JN688173	Rowley et al. (2011)
30.	T. petilum	HNUE MNA.2012.0001	Vietnam: Dien Bien: Muong Nhe	KJ802925	Nguyen et al. (2014)
31.	T. rhododiscus	CIB GX200807017	China: Guangxi	LC012842	Nguyen et al. (2015)
32.	T. ryabovi	VNMN 3924	Vietnam: Kon Tum: Mang Canh	LC012860	Nguyen et al. (2015)
33.	T. stellatum	stellatum-1	Thailand: Chanthaburi: Phliu	KT461918	Poyarkov et al. (2015)
34.	T. truongsonense	VNMN 4402	Vietnam: Khanh Hoa: Hon Ba	LC012847	Nguyen et al. (2015)
35.	T. vietnamense	VNMN 3687	Vietnam: Phu Yen: Krong Trai	KJ802923	Nguyen <i>et al.</i> (2014)
36.	T. vietnamense	VNMN 3686	Vietnam: Phu Yen: Krong Trai	KJ802922	Nguyen <i>et al.</i> (2014)
37.	Nyctixalus pictus	KUHE 53517	Borneo: Sarawak: Bario	LC012863	Nguyen et al. (2015)
38.	Liuixalus romeri	CIB 20080048	China: Hong Kong	AB871412	Nguyen et al. (2014)



FIGURE 1. Maximum likelihood (ML) tree based on partial sequence of mitochondrial genes (12S rRNA, tRNAval, and 16S rRNA) for samples of *Theloderma* and referenced species. Numbers above and below branches represent bootstrap support for ML and Bayesian posterior probabilities (MLBS/ BPP).

Species	1	2	Э	4	5	9	L	8	6	10	11	12	13	14	15	16	17	18
1. Theloderma annae sp. nov.																		
2. T. albopunctatum	15.9	ı																
3. T. asperum	14.8	11.7	ı															
4. T. bicolor	10.5	11.1	12.1															
5. T. corticale	11.0	13.8	13.6	4.8														
6. T. gordoni	14.8	16.0	16.8	11.4	12.8	ı												
7. T. laeve	10.7	13.0	13.8	10.2	12.7	15.6	ı											
8. T. lateriticum	11.7	15.2	16.4	8.4	11.0	14.7	11.6	ı										
9. T. leporosum	12.3	13.6	13.9	6.5	8.7	10.3	11.5	10.2	ı									
10. T. licin	17.3	11.3	11.3	13.3	15.4	17.9	17.0	17.5	14.6									
11. T. nebulosum	7.4	12.3	12.1	7.7	9.0	12.0	7.2	11.2	8.9	13.9	ı							
12. T. palliatum	13.1	14.2	15.1	7.2	8.9	13.0	13.3	11.2	10.4	16.5	11.7	ı						
13. T. petilum	14.8	8.5	10.8	10.6	14.7	16.4	14.2	14.5	12.7	11.4	12.7	14.0	ı					
14. T. ryabovi	13.3	13.7	12.6	10.1	10.7	14.3	10.0	13.0	11.0	15.1	10.7	11.7	13.3	ı				
15. T. rhododiscus	13.9	12.4	12.5	6.9	9.5	13.0	11.0	11.4	10.1	15.5	10.2	9.1	11.8	10.2	ı			
16. T. stellatum	18.1	17.9	21.1	15.2	18.3	21.3	18.6	17.7	19.3	19.3	16.3	17.7	16.4	21.3	16.7	·		
17. T. truongsonense	9.7	13.4	13.4	10.0	12.7	16.7	8.9	11.3	13.0	14.9	7.4	12.8	13.4	13.4	11.8	17.7	ı	
18. T. vietnamense	18.5	18.7	20.5	14.7	16.5	20.1	18.3	17.2	18.2	19.1	17.3	17.4	18.2	18.5	16.8	8.8	18.0	ı

Results

Phylogenetic analyses. Aligned, combined sequences of 12S rRNA, tRNA^{val}, and 16S rRNA yielded a total of 1628 bp (104 bp of 12S, 70 bp of tRNA^{val} and 1454 bp of 16S). The data set also included nine shorter sequences (ca. 530 bp) obtained from three samples of *Theloderma* **sp. nov.** from Hoa Binh, one sample of *T. asperum* from Genbank (KT461929), two samples of *T. nebulosum* from GenBank (JN688168, JN688169), two samples of *T. palliatum* from GenBank (JN688172, JN688173), and one sample of *T. stellatum* from Genbank (KT461918). Of 1628 nucleotide sites, 799 were variable and 685 were parsimony informative within the ingroup. The ML and Bayesian analyses produced topologies with -lnL = 14948.102 and 15000.616, respectively.

Phylogenetic analyses employing ML and BI methods yielded slightly different topologies only among referenced species, and only the ML tree is presented in Fig. 1. The unidentified species of *Theloderma* from Hoa Binh was placed in the clade containing *T. laeve* (Smith), *T. nebulosum*, and *T. truongsonense* (Orlov & Ho) with strong support (MLBS = 100%, BPP = 1.00) (Fig. 1).

The interspecific uncorrected genetic p-distances at the fragment of 16S rRNA gene examined between the unnamed *Theloderma* species from Hoa Binh and all congeners analysed varied from approximately 7.4% (compared with *T. nebulosum*) to 18.5% (compared with *T. vietnamense*) (Table 2).

Theloderma annae sp. nov.

(Figs. 2, 3)

Holotype. IEBR 3732 (Field number HB 2014.185), adult male, collected by T. Q. Nguyen *et al.* on 9 October 2010 in karst forest near Cho Village (20°24.909'N, 105°19.102'E, at an elevation of 343 m), Tu Do Commune, Lac Son District, Hoa Binh Province, Vietnam.

Paratypes. Seven specimens collected from Ngoc Son – Ngo Luong Nature Reserve in Lac Son District, Hoa Binh Province, Vietnam: IEBR 3733–3735 (Field numbers HB 2014.92–2014.94), adult males, collected by C. T. Pham *et al.* on 19 April 2014 in the karst forest near Khu Village (20°27.976'N, 105°18.421'E, at an elevation of 650 m) Ngoc Son Commune; ZFMK 97289 (Field number HB 2014.121), adult female, collected by T.Q. Nguyen *et al.* on 2 October 2014 in the karst forest near Khu Village (20°26.862'N, 105°20.144'E, at an elevation of 553 m) Ngoc Son Commune; ZFMK 97290 (Field number HB 2014.147), adult male, collected by T.Q. Nguyen *et al.* on 6 October 2014 in the karst forest near Khang Village (20°23.778'N, 105°20.847'E, Vietnam, at an elevation of 219 m) Tu Do Commune; IEBR 3736 (Field number HB 2014.186), adult female, collected by T.Q. Nguyen *et al.* on 9 October 2010 in the karst forest near Cho Village (20°24.964'N, 106°19.116'E, at an elevation of 279 m), Tu Do Commune; VNMN A.2016.5 (Field number HB 2014.211), adult male, collected by T.Q. Nguyen *et al.* on 10 October 2014 in the karst forest near Khu Village (20°26.517'N, 105°20.147'E, at an elevation of 570 m) Ngoc Son Commune; VNMN A.2016.5 (Field number HB 2014.211), adult male, collected by T.Q. Nguyen *et al.* on 10 October 2014 in the karst forest near Khu Village (20°26.517'N, 105°20.147'E, at an elevation of 570 m) Ngoc Son Commune; VNMN A.2016.5 (Field number HB 2014.211), adult male, collected by T.Q. Nguyen *et al.* on 10 October 2014 in the karst forest near Khu Village (20°26.517'N, 105°20.147'E, at an elevation of 570 m) Ngoc Son Commune.

Diagnosis. Morphologically, the frog specimens from Hoa Binh Province showed diagnostic characters of the genus *Theloderma*, for instance a distinct tympanum, round canthus rostralis, bony ridges from canthus rostralis to occiput absent, and skin of head not co-ossified to the skull (see Liem 1970; McLeod & Norhayati 2007; Rowley *et al.* 2011). However, because no morphological synapomorphy is known for the genus *Theloderma*, and its monophyly is not certain (Bain *et al.* 2009; Li *et al.* 2009; Rowley *et al.* 2011) we used molecular evidence to provide independent support of our generic placement.

The new species is distinguished from its congeners and other small rhacophorid species by a combination of the following characters: 1) Size small (SVL 27.1–28.5 mm in males, 30.3–32.6 mm in females); 2) head longer than wide; 3) vomerine teeth absent; 4) snout long (SNL/SVL 0.16–0.19); 5) spines on upper eyelid absent; 6) tibiotarsal projection absent; 7) dorsal skin smooth; 8) dermal fringes on forearm and tarsus absent; 9) dorsal surface greyish green; and 10) throat and ventral surfaces of arms and thighs brown with white spots.

Description of holotype. Adult male, body robust (SVL 27.4 mm), dorsoventrally compressed. Head slightly longer than wide (HL 11.7 mm, HW 10.8 mm), convex above; snout round in dorsal view, slightly protruding, its length longer than horizontal diameter of eye (SNL 5.1 mm, ED 4.1 mm); canthus rostralis round, loreal region oblique, concave; interorbital distance wider than internarial distance and upper eyelid (IOD 3.6 mm, IN 3.3 mm, UEW 2.9 mm); distance between anterior corner of eyes (DAE 6.2 mm) about 68% distance between posterior

corner of eyes (DPE 9.2 mm); nostril round, without a lateral flap of skin, closer to tip of snout than to the eye (NS 2.4 mm, EN 3.4 mm); pupil oval, horizontal; tympanum distinct (TYD 2.7 mm), round, smaller than eye diameter (ED 4.1 mm), greater than the distance between tympanum and eye (TYE 1.1 mm); pineal ocellus absent; spinules on upper eyelid absent; vomerine teeth absent; choanae small, oval; tongue cordate, deeply notched posteriorly; vocal sac absent; supratympanic fold distinct, extending from behind the eye to beyond level of axilla.

Forelimbs: Arm short, about half of hand length (FLL 5.8 mm, HAL 14.2 mm), dermal fringe along outer side of forearm absent; fingers free of webbing, relative lengths of fingers I \leq II \leq V \leq III; tips of fingers with enlarged discs with distinct circummarginal grooves; disc of finger III approximately two times of width of finger III (fd3/ fw3 1.99) but smaller than tympanum diameter (fd3/TYD 0.58); subarticular tubercles distinct, blunt, round, formula 1, 1, 2, 2; nuptial pads prominent, oval; outer palmar tubercle divided into two.

Hindlimbs: Heels overlapping when held at right angles to the body; tibia length about five times greater than tibia width (TbL 16.7 mm, TbW 3.5 mm), longer than thigh length (FeL 14.7 mm) but shorter than foot length (FoL 20.8 mm); relative length of toes I<II<III \leq V<IV; tips of toes with enlarged discs with distinct circummarginal grooves, discs slightly smaller than those of fingers; webbing formula 11-1½ II1-2III1-2IV2-1V; subarticular tubercles distinct, blunt, round, formula 1, 1, 2, 3, 2; inner metatarsal tubercle small (IMT 1.7 mm); dermal ridge along outer side of tibia and tarsal fold absent; projection at tibiotarsal articulation absent.

Skin texture in life: Dorsal surface of head and body smooth; posterior part of dorsum, flanks, dorsal surface and lateral sides of limbs with small granules; dorsolateral folds absent; throat and chest smooth, belly and ventral surface of thighs granular; dermal appendage at vent absent.

Coloration in life: Iris greyish green, pupil round; background of dorsal surface greyish green; head with a mossy green blotch between eyes, in triangular shape; tympanum brown; dorsum and upper part of flanks with dark markings and blotches, forming a network; dorsal surface of forelimbs yellowish green with some mossy green spots and bars; throat and chest white with brown marbling; belly immaculate white. The coloration is brighter at night.

Coloration in preservative: Dorsal surface of head, dorsum, and upper part of flanks greyish brown with dark markings and blotches; upper lip and tympanum light brown; forelimb, dorsal surface of thigh, tibia and foot grey with dark bands, posterior part of thigh below the vent greyish brown with small white spots; chin and throat brown with small white spots; belly immaculate cream to white; ventral part of forelimbs white to grey with small white spots; tibia and webbing dark brown.

Variation and sexual dimorphism. Measurements and morphological characters of the type series are provided in Table 3. The dorsal markings are indistinct in the specimen IEBR 3735, more prominent in the specimens IEBR 3734, 3736, ZFMK 97289, 97290 and consisting of three dark blotches in the individual IEBR 3733. The males are smaller in size than the females (SVL 27.1–28.6 mm, n = 6 vs. 30.3–32.6 mm, n = 2, respectively).

Etymology. We name this new species in honor of our colleague and friend, Anna Rauhaus, section animal keeper of the terrarium division of the Cologne Zoo, Germany, in recognition of her support of our research and conservation work in Vietnam. As common names we suggest Anna's Mossy Frog (English), Éch cây sần an-na (Vietnamese), and Annas Moosfrosch (German).

Ecological notes. *Theloderma annae* **sp. nov.** appears to be closely associated with karstic environment. Specimens were found at night between 19:00–23:30 h near cave entrances and in valleys surrounded by limestone cliffs, relatively far from water sources (Fig. 4). Advertisement calls, eggs and tadpoles of the species have not been recorded during our field surveys. The main habitat at the type locality is secondary karst forest of medium and small hardwoods mixed with shrubs and vines. Most of specimens were found on leaves, about 0.3–1.2 m above the ground but some specimens were collected on a limestone cliff. The air temperatures at the times of collection ranged from 20.4–30.2°C and relative humidity from 59–87%. In Ngoc Son – Ngo Luong Nature Reserve, several species of *Theloderma* have been recorded, viz. *T. albopunctatum* (Liu & Hu), *T. corticale* (Boulenger), and *T. lateriticum*, but they were found in tree holes filled with water or on tree leaves near streams in the valleys. Other species of amphibians and reptiles found at the site were *Rhacophorus* sp., *Raorchestes parvulus* (Boulenger), *Cyrtodactylus bobrovi* Nguyen, Le, Ngo, Pham, Hoang, Pham & Ziegler, and *Gekko* cf. *palmatus* Boulenger.

Distribution. *Theloderma annae* **sp. nov.** is currently known only from the type locality in Ngoc Son – Ngo Luong Nature Reserve, Hoa Binh Province, Vietnam. Several field surveys were conducted in the adjacent karst forest of Hang Kia – Pa Co Nature Reserve in Hoa Binh Province but no records of this species have been made so far.



FIGURE 2. Dorsal and ventral views of the holotype (IEBR 3732, adult male) of *Theloderma annae* sp. nov. from Hoa Binh Province, Vietnam.



FIGURE 3. Ventral side of hand (A) and foot (B) of the holotype (IEBR 3732) of *Theloderma annae* sp. nov. in preservative.

	IEBR 3732	VNMN A.2016.5	IEBR 3733	IEBR 3734	IEBR 3735	ZFMK 97290			ZFMK 97289	IEBR 3736
Sex	3	3	3	3	3	3	Min-Max	Mean±SD	Ŷ	Ŷ.
Type status	Н	Р	Р	Р	Р	Р	(n=6)	(n=6)	Р	Р
SVL	27.4	27.7	28.0	28.5	27.8	27.1	27.1–28.5	27.7±0.34	30.3	32.6
HW	10.8	10.2	10.2	10.9	10.4	10.3	10.2-10.9	10.5±0.26	11.8	12.7
HL	11.7	11.2	11.1	11.2	11.6	11.6	11.1–11.7	11.4±0.23	12.4	13.0
MN	2.2	2.3	2.1	2.3	2.0	2.1	2.0-2.3	2.2 ± 0.08	2.0	2.5
MFE	5.2	4.8	4.9	4.8	4.8	4.6	4.6–5.2	4.8±0.13	4.2	5.2
MBE	7.8	8.0	8.4	8.6	9.7	8.7	7.8–9.7	8.4±0.27	8.2	8.9
SNL	5.1	4.9	4.6	5.0	4.6	5.0	4.6–5.1	4.9±0.16	5.0	5.4
ED	4.1	4.2	4.2	4.5	4.2	4.4	4.1-4.5	4.3±0.11	4.8	4.9
UEW	2.9	2.8	2.9	2.8	2.7	2.6	2.6-2.9	2.8±0.07	2.9	3.2
IN	3.3	3.4	3.1	3.2	3.2	3.0	3.0-3.4	3.1±0.18	3.4	3.5
IOD	3.6	3.8	3.8	3.4	3.7	3.7	3.4–3.8	3.7±0.09	3.8	4.0
DAE	6.2	6.1	5.8	5.8	6.3	6.1	5.8-6.2	6.0±0.16	6.4	6.6
DPE	9.2	8.8	8.5	8.9	8.8	8.6	8.5–9.2	8.8±0.16	9.4	10.2
NS	2.4	2.1	2.2	2.1	2.2	2.3	2.1–2.4	2.2±0.08	2.4	2.5
EN	3.4	3.3	3.1	3.2	3.2	3.0	3.0-3.4	3.2±0.11	3.1	3.4
TYD	2.9	2.6	2.4	2.5	2.6	2.7	2.4–2.9	2.6±0.12	3.1	3.2
TYE	1.1	1.2	0.9	0.9	1.0	1.1	0.9–1.2	01.0 ± 0.10	0.9	1.0
FLL	5.8	5.1	4.4	4.8	4.7	4.	4.4–5.8	4.9±0.36	5.5	6.3
HAL	14.2	13.3	13.6	13.9	13.8	13.4	13.3–14.2	$13.7{\pm}~0.26$	14.3	15.8
TFL	6.2	5.6	5.7	5.8	5.8	5.3	5.3-6.2	5.7±0.20	5.9	6.3
fd3	1.7	1.5	1.6	1.6	1.7	1.5	1.5–1.7	1.6 ± 0.06	1.5	1.6
FeL	14.7	15.1	13.8	14.8	14.1	13.8	13.8–15.1	14.4 ± 0.48	15.0	15.8
TbL	16.7	16.2	15.8	15.7	16.4	17.0	15.7-17.0	16.3±0.38	17.4	18.9
TbW	3.5	3.0	3.1	3.0	3.1	2.9	2.9–3.5	3.1 ± 0.14	3.1	3.5
FoL	20.8	21.1	19.7	20.2	20.0	20.7	19.7–20.8	$20.3{\pm}~0.33$	22.0	23.4
FTL	12.9	12.3	12.1	11.7	10.7	11.0	10.7–12.9	$11.8\pm\!0.63$	13.5	13.3
SNL/SVL	0.19	0.18	0.18	0.17	0.17	0.18	0.17–0.19	$0.18{\pm}~0.01$	0.16	0.17
ED/SNL	0.80	0.87	0.91	0.90	0.90	0.88	0.80-0.91	$0.88{\pm}0.03$	0.97	0.90
TYE/TYD	0.40	0.45	0.38	0.34	0.37	0.39	0.34–0.45	0.30 ± 0.03	0.33	0.30
fd3/TYD	0.58	0.60	0.66	0.62	0.67	0.55	0.55–0.67	$0.61{\pm}~0.04$	0.52	0.48

TABLE 3. Measurements (in mm) and proportions of the type series of *Theloderma annae* **sp. nov.** (H = holotype, P = paratype, other abbreviations defined in text).

Comparisons. We compared the new species with other members of the genus *Theloderma* based on morphological specimen examination (see Appendix I) and data obtained from the literature (e.g., Ahl 1927, 1931; Bain & Nguyen 2004; Bain *et al.* 2009; Boulenger 1903; Bourret 1937, 1942; Chanda 1994, Chan-ard 2003; Chan & Norhayati 2009; Fei *et al.* 2009, 2012; Inger *et al.* 1999; Jiang *et al.* 2009; Kunz *et al.* 2010; Liu & Hu 1962; McLeod & Norhayati 2007; Nguyen *et al.* 2014; Orlov *et al.* 2005, 2006, 2012; Smith 1924; Stuart & Heatwole 2004; Taylor 1962; Tschudi 1838; Rowley *et al.* 2011).

The loderma annae **sp. nov.** differs from *T. bicolor* (Bourret), *T. corticale*, *T. gordoni* Taylor, *T. (Stelladerma* in Poyarkov et al. 2015) horridum (Boulenger), *T. kwangsiense* (Liu & Hu), *T. leporosum* Tschudi, *T. moloch* (Annandale), *T. nagalandense* Orlov, Dutta, Ghate & Kent, and *T. phrynoderma* (Ahl) by having a smaller size (SVL 27.1–28.6 mm in males and 30.3–32.6 mm in females vs. SVL \geq 40 mm in the other species) and smooth dorsal skin (vs. with large warts in other species).



FIGURE 4. *Theloderma annae* **sp. nov.** (IEBR 3734) *in situ* on vegetation in Ngoc Son – Ngo Luong Nature Reserve, Hoa Binh Province, Vietnam.

The new species differs from T. andersoni (Ahl) by having smooth dorsal skin (vs. with small tubercles in T. andersoni), in dorsal color pattern (greyish green vs. dark olive with two yellow spots on middle of flanks and a large black spot on groin in T. andersoni) and ventral color (white vs. yellow with fine brown spots anteriorly in T. andersoni); from T. albopunctatum (Liu & Hu) and T. asperum (Boulenger) by having smooth dorsal skin (vs. with small tubercles in latter species), in dorsal color pattern (greyish green vs. greyish brown with large conspicuous white pattern in latter species), in ventral color pattern (white vs. marbled black and bluish grey in latter species), and iris color (greyish green vs. reddish brown in latter species); from T. baibengense Jiang, Fei & Huang by having smooth dorsal skin (vs. with small tubercles in *T. baibengense*), in dorsal color pattern (greyish green vs. back with large conspicuous white pattern in T. baibengense), and ventral color pattern (white vs. dark with white pattern in T. baibengense); from T. lateriticum in dorsal color pattern (greyish green vs. brick red with single black mid-dorsal spots in T. lateriticum), in ventral color pattern (white vs. uniform grey-brown with white spots in T. lateriticum), and iris color (greyish green vs. brick-red in T. lateriticum); from T. laeve (Smith) in dorsal color pattern (greyish green vs. golden beige to light brown above with small dark brown spots and blotches forming an x-shaped figure, and dark longitudinal streaks on flanks in T. laeve), in ventral color pattern (white vs. light greyish pink to violet), and iris color (greyish green vs. grey above and dark brown below in T. laeve); from T. licin McLeod & Ahmad by having dorsal skin smooth (vs. with small tubercles in T. licin), in dorsal color pattern (grevish green vs. white to pale brown in *T. licin*), in ventral color pattern (white vs. white with brown reticulations in T. licin), and iris color (greyish green vs. red in T. licin); from T. nebulosum by having smooth dorsal skin (vs. with small tubercles in T. nebulosum), in dorsal color pattern (greyish green vs. brown with dark marking in T. nebulosum), in ventral color pattern (white vs. dark brownish black with pale blue in T. nebulosum), and iris color (greyish green vs. pale gold above, reddish brown below in *T. nebulosum*); from *T. palliatum* by having smooth



FIGURE 5. Type locality (black square) of *Theloderma annae* sp. nov. in Hoa Binh Province, Vietnam.

dorsal skin (vs. with small tubercles in T. palliatum), in dorsal color pattern (greyish green vs. pale brown or pale yellow to light straw brown with dark blotches forming an x-shaped figure in *T. palliatum*), in ventral color pattern (white vs. dark brownish black with pale blue or greyish with light bluish and dark spots and blotches in T. palliatum), and iris color (greyish green vs. pale gold above, reddish brown below in T. palliatum); from T. petilum (Stuart & Heatwole) in dorsal color pattern (greyish green vs. light brown with dark brown reticulations and scattered black spots in T. petilum), and the iris color (greyish green vs. upper part reddish brown, lower part grey in T. petilum); from T. rhododiscus (Liu & Hu) in dorsal color pattern (greyish green vs. brown in T. rhododiscus), in ventral color pattern (white vs. brown black scattered with grey white network in T. rhododiscus), and iris color (greyish green vs. red brown in T. rhododiscus); from T. ryabovi Orlov, Dutta, Ghate & Kent by having a smaller size (SVL 27-33 mm vs. 44 mm in T. ryabovi), in dorsal color pattern (greyish green vs. light beige with lilac pattern and black spots in T. ryabovi), in ventral color pattern (white vs. dark grey in T. ryabovi), and iris color (greyish green vs. dark brown in T. ryabovi); from T. (Stelladerma in Poyarkov et al. 2015) stellatum Taylor and T. (Stelladerma in Poyarkov et al. 2015) vietnamense in dorsal color pattern (greyish green vs. brown with light markings in the latter), in ventral color pattern (white vs. dark brown reticulated with cream in in the latter), and iris color (greyish green vs. brown with black reticulations in the latter); and from T. truongsonense (Orlov & Ho) in dorsal color pattern (greyish green vs. pale brown with small, distinct darker brown markings in T. truongsonense), in ventral color pattern (white vs. dark grey with black round speckles in *T. truongsonense*), and iris color (greyish green vs. pale gold above, reddish brown below in T. truongsonense).

In terms of dorsal color pattern *Theloderma annae* **sp. nov.** is similar to *Gracixalus quyeti* (Nguyen, Hendrix, Böhme, Vu & Ziegler) from Quang Binh Province and *G waza* Nguyen, Le, Pham, Nguyen, Bonkowski & Ziegler from Cao Bang Province. In addition, *Theloderma annae* **sp. nov.** and the two afore mentioned *Gracixalus* species also have similar life histories, all inhabiting limestone karst forest far from water sources. However, *Theloderma annae* differs from *G quyeti* by having dorsal skin smooth (vs. with small sharp tubercles in *G quyeti*), the absence of a dark inverse Y-marking on dorsum (vs. present in *G quyeti*), and the outer palmar tubercle prominent (vs. indistinct in *G quyeti*). *Theloderma annae* **sp. nov.** differs from *G waza* in the ratio of TYD/TDE (0.39 in males and 0.32 in females vs. 0.87 in males and 0.98 in females in *G waza*), and the absence of a dark inverse Y-marking on dorsum (vs. present in *G waza*).

Discussion

In our phylogenetic analysis *Theloderma annae* is assigned to the *T. truongsonense* group (or the *T. laeve* group in Poyarkov *et al.* 2015) with significant values of nodal support. The mean genetic distance (16S gene) between the new species and other members of this species group varied from 7.4% (compared with *T. nebulosum*) to 10.7% (compared with *T. laeve*) (Table 2). Morphologically, the new species can be clearly distinguished from members of the *T. truongsonense* group in having smooth dorsal skin and the dorsal surface being greyish green with mossy markings. The greenish brown or moss-green coloration on the dorsal surface of the new species, which can blend remarkably well into the background of stones covered with lichens or tree leaves, seems to be an adaptation to the life mode associated with the karst environment (see also Nguyen *et al.* 2012). However, biodiversity of karst forests in Ngoc Son – Ngo Luong Nature Reserve as well as in northwestern Vietnam is currently threatened due to the effects of road construction, expanding agriculture, and illegal timber logging (T. Nguyen pers. obs.).

The discovery of additional new species of *Theloderma* in Vietnam suggests that the current species richness of the genus remains underestimated. More studies using an integrative approach, i.e., combining morphological and molecular data, will help to reveal the extent of species richness of *Theloderma* in the poorly studied regions of northern Vietnam.

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APPENDIX I. Examined specimens.

- Gracixalus quyeti (2): Vietnam: Quang Binh Province: ZFMK 82999 (holotype), VNUH 160706 (paratype).
- *G. waza* (6): Vietnam: Cao Bang Province: Ha Lang District: IEBR A.2012.2 (holotype), IEBR A.2012.3, VNMN A.2012.2, A.2012.3, ZFMK 93666, 93667 (paratypes).
- *Theloderma albopunctatum* (8): Vietnam: Bac Giang Province: Son Dong District: IEBR A. 2013.82, A.2013.83; Vietnam: Hoa Binh Province: Mai Chau District: IEBR 3760, 3761; Vietnam: Hoa Binh Province: Lac Son District: IEBR 3762–3765.
- T. bicolor (2): Vietnam: Lao Cai Province: Sa Pa District: IEBR 3740, 3741.
- *T. corticale* (4): Vietnam: Bac Giang Province: Son Dong District: IEBR A. 2013.84, A.2013.85; Vietnam: Hoa Binh Province: Lac Son District: IEBR 3758, 3759.
- *T. gordoni* (3): Vietnam: Hoa Binh Province: Lac Son District: IEBR 3737; Vietnam: Vinh Phuc Province: Tam Dao NP: IEBR 3738, 3738.
- *T. lateriticum* (8): Vietnam: Lao Cai Province: Sa Pa District: IEBR 3745–3747; Vietnam: Hoa Binh Province: Lac Son District: IEBR 3748–3753.
- T. truongsonense (4): Vietnam: Quang Binh Province: Le Thuy District: IEBR 3754-3757.
- T. vietnamense (3): Vietnam: Kien Giang Province: Phu Quoc District: IEBR 3742-3744.