



# Thermal Sensitivity of Heat Sensor TRPA1 Correlates With Temperatures Inducing Heat Avoidance Behavior in Terrestrial Ectotherms

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Temperature is an essential environmental factor that controls an organism's performances. As ectothermic animals largely rely on external heat sources for adjusting their body temperature, thermal perception is a primary process of behavioral thermoregulation. Transient receptor potential ankyrin 1 (TRPA1) is a heat sensitive ion channel in most non-mammalian species, and its heat activation has been suggested to induce heat avoidance behaviors in ectothermic animals. However, associations between TRPA1 and ecologically relevant temperatures have not been investigated, and the analyses including diverse taxa will provide robust support for understanding the associations. Here, I conducted extensive literature review, and assembled published data on thermal threshold of TRPA1 and three physiological parameters: the experimental voluntary maximum (EVM), which is body temperatures when heat avoidance behaviors are induced; the critical thermal maximum (CT<sub>max</sub>), which is a point in temperature beyond which an organism becomes incapacitated; and average body temperature (T<sub>mean</sub>) recorded in the field. Then, I examined the relationships between thermal threshold of TRPA1 and each of the three physiological parameters. As phylogenetically closely related species tend to show similar trait values among species, I conducted the regression analyses by accounting for phylogenetic distances among species. This study supports previous research by affirming that thermal threshold of TRPA1 is substantially correlated with body temperature that the animals escaped from the heat source, represented here as EVM. Nevertheless, thermal threshold of TRPA1 showed a statistically insignificant correlation with CT<sub>max</sub> and T<sub>mean</sub>. The results suggest that although thermal threshold of TRPA1 is evolutionarily labile, its associations with EVM is highly conserved among diverse terrestrial ectotherms. Therefore, thermal threshold of TRPA1 could be a useful parameter to evaluate species vulnerability to thermal stress particularly in the recent climate warming scenario.

Keywords: thermal biology, terrestrial ectotherms, thermoregulation, thermal sensitivity, transient receptor potential ankyrin 1, heat avoidance behavior, phylogenetic generalized least squares

# INTRODUCTION

Temperature is one of the most influential environmental factors that affects the organisms' performances. As ectothermic animals largely rely on external heat sources for adjusting their body temperature, thermal perception is an essential process of behavioral thermoregulation (Tattersall et al., 2012). Animals perceive temperature through sensory neurons in the peripheral nervous system, in which a subset of temperature-sensitive ion channels is expressed (Tominaga and Caterina, 2004; Dhaka et al., 2006). The transient receptor potential (TRP) ion channel superfamily consists of Ca<sup>2+</sup> permeable non-selective cation channels, which function as polymodal receptors detecting mechanical, chemical, and thermal stimuli (Saito and Tominaga, 2017; Hoffstaetter et al., 2018). Among the TRP superfamily, a subset of TRPs (called thermoTRPs) is temperature sensitive, and known to participate in thermal perception in animals (Patapoutian et al., 2003; Venkatachalam and Montell, 2007). Recent studies reported that the activation temperatures of each thermoTRP are diverse among species, suggesting that the functions of thermoTRPs evolve with change in thermal physiology (Gracheva and Bagriantsev, 2015).

In electrophysiological analysis of the heterologous expression of thermoTRPs, thermal thresholds could be estimated from the Arrhenius plot that represents the logarithmic rate of inward currents against inverse temperatures in Kelvin (Voets et al., 2005). The intersection of lines fitted to baseline and linearly increasing currents of the Arrhenius plot indicates the initial temperature which activates thermoTRPs and it represents the thermal threshold of thermoTRPs (Boukalova et al., 2010; Gracheva et al., 2010; Vriens et al., 2014). The use of thermal thresholds for describing and comparing thermoTRPs has been discouraged because thermoTRPs are activated gradually rather than at a specific threshold and also vary unless the experimental conditions are kept constant across studies (Voets et al., 2005; Vriens et al., 2014). Although thermal thresholds of thermoTRPs should still be carefully discussed, they have been one of indices for describing ranges of thermal sensitivity (Gracheva and Bagriantsev, 2015). Studies on thermoTRPs have, in fact, revealed that each thermoTRP possesses channelspecific activation temperatures covering temperature ranges that animals normally experience (Patapoutian et al., 2003; Dhaka et al., 2006).

TRP ankyrin 1 (TRPA1) is a member of thermoTRPs and is among those which have been studied in a high number of species. Its thermal sensitivity has been debated in mammalian TRPA1, as cold activates TRPA1 in rodent species but not in primate species (Caspani and Heppenstall, 2009; Chen et al., 2013). In contrast, heat consistently activates TRPA1 in non-mammalian species except *Caenorhabditis elegans* (Chatzigeorgiou et al., 2010; Saito et al., 2014). In *Drosophila melanogaster*, four TRPA1 subtypes have been reported, and mutant analyses in larval or adult flies have demonstrated that the ortholog of vertebrate TRPA1 (dTRPA1) and Painless are requisite for avoiding noxious temperatures (Rosenzweig et al., 2005; Hamada et al., 2008; Neely et al., 2011; Peng et al., 2015). Recently, TRPA1 has been analyzed in conjunction with heat avoidance behaviors across various taxa, such as mosquitoes, *Xenopus* frogs, and *Anolis* lizards (Saito et al., 2016; Akashi et al., 2018; Li et al., 2019). These studies revealed a trend that species with higher thermal thresholds of TRPA1 showed heat avoidance behaviors induced at higher temperatures. These mutant and comparative analyses suggested that animals perceive noxious heat through the activation of TRPA1, which induces heat avoidance behavior.

Temperature-dependent performance of ectothermic animals is often graphically described using thermal reaction norms or thermal performance curves. Animals achieve their highest performance at the peak of the curve but become incapable of keeping right posture and performing at both extreme ends of the curve, which represent critical thermal maximum and minimum (CT<sub>max</sub> and CT<sub>min</sub>) (Cowles and Bogert, 1944; Huey and Stevenson, 1979; Lutterschmidt and Hutchison, 1997). The curve is typically asymmetric and left-skewed so that a deviation of body temperature from the optimum will reduce the performance greater toward the higher thermal limit than the lower thermal limit of the curve. Therefore, detection of temperature above the optimum is crucial. Animals accept the temperature rise or fall until their body temperature reaches the limit that they voluntarily tolerate. The highest and lowest voluntarily tolerable temperatures are defined as maximum and minimum voluntary tolerances, respectively, and the activity of animals takes place within thermal-safety margins bounded by the two voluntary tolerance temperatures (Cowles and Bogert, 1944; Vitt and Caldwell, 2008). The highest voluntary tolerance temperature is experimentally measured as experimental voluntary maximum (EVM) at which animals show the first attempt to escape from the heat sources (Ruibal, 1961; Hertz, 1981). Tmean is the mean body temperature recorded in the field and is not necessarily the temperature that maximizes performance (i.e., the peak of the thermal performance curve) as ectothermic animals regulate their body temperatures presumably through a compromise between costs and benefits associated with thermoregulatory behaviors (Huey and Slatkin, 1976; Huey and Stevenson, 1979).

Ectothermic animals, particularly thermoregulators, sense ambient temperatures to keep T<sub>mean</sub> within thermal-safety margins distant from CT<sub>max</sub> and CT<sub>min</sub>. TRPA1 is an excellent candidate molecule for the detection of temperature. Previous comparative studies have focused on the relationship between thermal threshold of TRPA1 and temperature that induces thermal avoidance behaviors (Saito et al., 2016; Akashi et al., 2018; Li et al., 2019). However, these studies have not considered whether thermal threshold of TRPA1 is associated with other ecologically relevant parameters, such as CT<sub>max</sub> and T<sub>mean</sub>. In addition, ecological or physiological similarities among species would arise either from adaptation to similar environments or from close phylogenetic relationships (Cavender-Bares et al., 2004; Losos, 2008; Cooper et al., 2011; Kellermann et al., 2012), thus, comparative analyses should consider the phylogenetic distances among species (Felsenstein, 1985; Hansen and Orzack, 2005). The previous studies might have overlooked the possibility of correlations between the physiological properties of TRPA1 and other physiological parameters.

In this study, I assembled published data regarding thermal threshold of TRPA1 and the physiological traits EVM,  $CT_{max}$ , and  $T_{mean}$ , in ectothermic animals. Then, I analyzed whether thermal threshold of TRPA1 correlates with any of the relevant physiological traits by conducting ordinary least squares (OLS) and phylogenetic generalized least squares (PGLS) analyses to control the effects of phylogenetic distances. Here, I found that thermal threshold of TRPA1 correlates specifically with EVM even after considering phylogenetic distances among species, but not with  $CT_{max}$  nor  $T_{mean}$ . Therefore, the current study suggests that the association between TRPA1 and EVM is highly conserved among diverse terrestrial ectotherms, and TRPA1 contributes in part to setting the upper limit of species thermal-safety margins.

# MATERIALS AND METHODS

# Literature Review and Data Acquisition

I searched National Center for Biotechnology Information (NCBI)<sup>1</sup> for studies reported complete coding sequences of and physiological properties of TRPA1, and also Google Scholar<sup>2</sup> and PubMed<sup>3</sup> for studies on TRPA1 which included its thermal threshold as an index of thermal sensitivity. The literature was searched for studies published before February 28, 2021. The query term for NCBI was "TRPA1." The query terms for Google Scholar and PubMed were the following: trp, "temperature threshold" OR "thermal threshold," and "Arrhenius plot." The technical terms were double quoted. Two phrases were linked with "OR" and gueried if the phrases were equally used across studies. For species whose thermal thresholds of TRPA1 have been reported, I conducted further literature surveys with the following terms to identify studies reporting (1) EVM, (2)  $CT_{max}$ , and (3)  $T_{mean}$ . The query terms were the following: (1) "species," "voluntary maximum" OR "maximum voluntary" OR nocifensive response behavior; (2) "species," "critical maximum" OR "critical thermal maximum" OR CT<sub>max</sub>; (3) "species" field average body temperature. Plural forms were also queried. Ectothermic species were included for subsequent analyses if thermal threshold of TRPA1 and/or at least one of the three physiological parameters had been reported.

I summarized the reported values by calculating a single grand mean and grand standard error (SE) for each trait and species. This calculation requires means, errors (SE or standard deviations, SD), and sample sizes to be reported in the original papers. However, several studies presented the data in figures rather than as numerical values. Thus, depending on the reported data set, I defined the grand mean and grand SE as follows: (a) For data reported by multiple and comprehensive studies, I summarized the data by calculating the grand means and grand SE using total data sets within each trait for each species. The data with missing SE or SD and sample size were omitted from the grand means and grand SE. (b) For data reported by multiple but incomprehensive studies (i.e., lack of SE or SD and sample size in all the reported studies), I calculated the mean and SE of the reported means as the grand mean and grand SE, respectively. (c) For data reported by a single study, the mean and SE were treated as the grand mean and grand SE, respectively. If the data lacked SE or SD and sample size was not indicated, their mean was used as the grand mean without an error.

As thermal threshold of TRPA1 was reported from two different electrophysiological approaches using HEK293 cells or *Xenopus* oocytes, the data obtained from HEK293 cells and *Xenopus* oocytes and those obtained only from *Xenopus* oocytes were separately summarized. If different isoforms were investigated in the electrophysiological analyses, the data for the longer isoform was adopted for the current study. All four parameters reviewed here should not be taken as fixed values for the species as each value varied across or within studies to some extent, depending on experimental protocols. All statistical analyses were performed using the R, version 3.5.1 (R Development Core Team, 2018).

# **Phylogenetic Analysis**

Coding sequences of TRPA1 for 18 species that passed the criteria of the literature review were obtained from GenBank<sup>4</sup>. The accession numbers of these sequences are shown in Figure 1. I translated the nucleotide sequences to amino acid sequences and constructed the multiple protein sequence alignments with MUSCLE (Edgar, 2004), using default penalties for a gap-opening (-2.9), a gap-extension (0), and hydrophobicity multiplier (1.2). For phylogenetic analysis, the aligned amino acid sequences were converted back to nucleotide sequences. Then, I inferred a phylogenetic tree from the aligned nucleotide sequences by using the maximum likelihood method with 1,000 bootstrap replicates in MEGA X (Kumar et al., 2018; Stecher et al., 2020). The best nucleotide substitution model for maximum likelihood method was determined using the "find best DNA/protein models" tool, and the Kimura 2-parameter model of sequence evolution under a gamma distribution of rates with invariant sites was adopted based on the lowest Bayesian information criterion scores yielded from the analysis (Kimura, 1980). All positions with less than 95% site coverage were eliminated, resulting in a total of 3,177 positions in the final dataset for the analysis. TRPA1 of Patiria pectinifera was used as the outgroup.

# Evaluating Phylogenetic Signal and Correlations Between Physiological Parameters

I assessed whether phylogenetic distances explain the similarity and divergence in thermal threshold of TRPA1 and the physiological parameters (EVM,  $CT_{max}$ , and  $T_{mean}$ ) by means of Pagel's lambda ( $\lambda$ ) (Pagel, 1999). Lambda scales the internal branches relative to the tip branches of the phylogenetic tree from 1 to 0, inferring the extent of the phylogenetic signal. Lambda equal to 1 indicates that traits evolve under a Brownian motion model and closely related species are expected to have similar trait values. Lambda equal to 0 indicates that traits

<sup>&</sup>lt;sup>1</sup>https://www.ncbi.nlm.nih.gov/

<sup>&</sup>lt;sup>2</sup>https://scholar.google.com/

<sup>&</sup>lt;sup>3</sup>https://pubmed.ncbi.nlm.nih.gov/

<sup>&</sup>lt;sup>4</sup>http://www.ncbi.nlm.nih.gov/genbank/



The tree was estimated from the maximum likelihood (ML) analysis of TRPA1 coding sequences. The numbers at each node indicate bootstrap values obtained from 1,000 replicates using the ML method. Scale bar refers to a phylogenetic distance of 0.2 nucleotide substitutions per site. Accession numbers are shown in parentheses.

evolve independently of the phylogeny and signifies the absence of phylogenetic signal. In this analysis, the inferred maximum likelihood tree was first imported into the R by using the R package "ape" (Paradis et al., 2004). Lambda was then calculated from the grand mean and grand SE for each trait by using the *phylosig* function in the R package "phytools" (Revell, 2012). For thermal threshold of TRPA1, lambda was calculated using the species with inferred phylogeny. The studies reporting thermal threshold of TRPA1 did not have a complete set of physiological parameters (i.e., EVM,  $CT_{max}$ , and  $T_{mean}$ ). Species without those parameters were pruned from the inferred phylogeny, and lambda was calculated for EVM,  $CT_{max}$ , and  $T_{mean}$  using the remaining species with the corresponding pruned phylogeny.

I investigated the relationship between the grand means of thermal threshold of TRPA1 and the physiological parameters ( $CT_{max}$ , EVM, and  $T_{mean}$ ) by conducting OLS and PGLS regression analyses. As the activation of TRPA1 was assumed to regulate the physiological parameters, thermal threshold of TRPA1 was selected as the sole explanatory variable, and the three physiological parameters were analyzed as response variables. I

first fitted the OLS regression to the data by using the *lm* function in R so that the fitted regression line indicated how well an explanatory variable was associated with the response variables. Residual error in OLS regression model needs to be independent among observations. However, the error will no longer be independent particularly in the case of interspecific comparisons. PGLS analysis considers the phylogenetic relationship based on an assumption that the residual error in the regression model is proportional to the internal branch length of the phylogenetic tree (Felsenstein, 1985; Revell, 2010). Thus, I secondly conducted three independent PGLS analyses on thermal threshold of TRPA1 paired with one of the three parameters and each of the corresponding pruned phylogenies by using the pgls function in the R package "caper" (Orme et al., 2018). The PGLS analysis estimates the maximum likelihood values of lambda in the residual error simultaneously with the regression parameters.

### RESULTS

The literature review showed that thermal threshold of TRPA1 has been estimated from the Arrhenius plot in 18 ectothermic species (Supplementary Table 1). Among those species, the physiological parameters (EVM, CT<sub>max</sub>, and T<sub>mean</sub>) were reported for 9, 7, and 6 species, respectively (Supplementary Table 2). The grand mean and grand SE from both HEK293 cells and Xenopus oocytes and those only from Xenopus oocytes are summarized in Table 1 and Supplementary Table 3, respectively. In the former case, Pagel's lambda for thermal threshold of TRPA1 was 0.786, and the null hypothesis of no phylogenetic signal (i.e.,  $\lambda = 0$ ) was rejected (P = 0.010). EVM also showed a high value of lambda ( $\lambda = 0.826$ ) but with insignificant *P*-value (P = 0.192). Both CT<sub>max</sub> and T<sub>mean</sub> showed lambda to be 0 with P-value of 1.0. In addition to estimating lambda for each trait, OLS and PGLS analyses were conducted for each of the three combinations of the two parameters (Figure 2 and Table 2). OLS analysis showed a significant positive correlation between thermal threshold of TRPA1 and EVM (slope = 0.603, P = 0.014) (Figure 2A). However, thermal threshold of TRPA1 was not correlated with CT<sub>max</sub> nor T<sub>mean</sub>  $(CT_{max}: slope = -0.464, P = 0.187; T_{mean}: slope = 0.166,$ P = 0.524) (Figures 2B,C). Although PGLS analysis detected strong phylogenetic signal in the model of thermal threshold of TRPA1 and EVM ( $_{ML}\lambda$  = 0.904), thermal threshold of TRPA1 was positively correlated with EVM as observed in OLS analysis (slope = 0.458, P = 0.026) (Figure 2A). PGLS analysis did not show a significant correlation between thermal threshold of TRPA1 and  $CT_{max}$  nor  $T_{mean}$  ( $CT_{max}$ : slope = -0.461, P = 0.184;  $T_{mean}$ : slope = 0.212, P = 0.511) with no phylogenetic signal in these traits (Figures 2B,C). The OLS and PGLS analyses conducted on data obtained only from Xenopus oocytes showed the same trend described above (Supplementary Table 4).

# DISCUSSION

Here, I investigated the relationship between thermal threshold of TRPA1 and the physiological traits (EVM,  $CT_{max}$ , and  $T_{mean}$ )

**TABLE 1** Summary of the grand mean  $\pm$  grand standard error for each trait used in the analyses.

Species	TRPA1	EVM	CT <sub>max</sub>	T <sub>mean</sub>	References		
Crotalus atrox	27.6 ± 0.9 (6) <sup>c</sup>	34 (–) <sup>c</sup>	39 (–) <sup>c</sup>	27.4 (–) <sup>c</sup>	Brattstrom, 1965; Gracheva et al., 2010		
Elaphe obsoleta	$37.9 \pm 0.4 \ (18)^a$	38 (–) <sup>c</sup>	38 (–) <sup>c</sup>	$27 \pm 0.4 \ (158)^a$	Brattstrom, 1965; Blouin-Demers and Weatherhead, 2001; Gracheva et al., 2010; Kurganov et al., 2017		
Python regius	$32.7 \pm 0.5 \ (6)^{c}$	-	-	-	Gracheva et al., 2010		
Corallus hortulanus	$29.6 \pm 0.3 \ (6)^{c}$	-	-	-	Gracheva et al., 2010		
Anolis carolinensis	$35.2 \pm 0.5  (35)^a$	$36.0 \pm 0.7 \ (15)^a$	$41.7 \pm 0.2(49)^{a}$	$29.8 \pm 0.3  (39)^a$	Brattstrom, 1965; Corn, 1971; Wilson and Echternach 1990; Lailvaux and Irschick, 2007; Saito et al., 2012; Kurganov et al., 2014		
Anolis allogus	$33.5 \pm 0.7 \ (26)^{c}$	$32.3 \pm 0.2  (47)^{a}$	-	$28.3 \pm 0.1 \ (248)^{a}$	Ruibal, 1961; Schettino et al., 2010; Akashi et al., 2018		
Anolis homolechis	$36.4 \pm 0.8 \ (18)^{c}$	$34.5 \pm 0.3  (53)^{a}$	-	$29.4 \pm 0.1 \ (276)^{a}$	Ruibal, 1961; Schettino et al., 2010; Akashi et al., 2018		
Anolis sagrei	$36.1 \pm 0.8 \ (22)^{c}$	$34.1 \pm 0.3 \ (9)^{c}$	$39.4 \pm 0.1 \ (102)^a$	32.0 ± 0.2 (295) <sup>a</sup>	Ruibal, 1961; Corn, 1971; Schettino et al., 2010; Akashi et al., 2018; Gunderson et al., 2020		
Xenopus tropicalis	$39.8 \pm 0.4 \ (44)^a$	40 (4) <sup>c</sup>	34 (18) <sup>c</sup>	-	Herrel and Bonneaud, 2012; Saito et al., 2012; Saito et al., 2016		
Xenopus laevis	$37.5 \pm 0.4 \ (51)^a$	38 (4) <sup>c</sup>	$31.6 \pm 0.1 \ (51)^a$	-	Sherman and Levitis, 2003; Saito et al., 2016; Araspin et al., 2020		
Ambystoma mexicanum	$39.7 \pm 1.0 \ (6)^{c}$	-	-	-	Oda et al., 2019		
Drosophila melanogaster	$29.7\pm0.3~\text{(-)}^{c}$	$30.2 \pm 0.6$ (–) <sup>b</sup>	$41.0 \pm 0.1 \ (12)^a$	-	Folk et al., 2007; Wang et al., 2008; Oswald et al., 2011; Kang et al., 2012		
Bombyx mori	21.6 (–) <sup>c</sup>	_	-	-	Sato et al., 2014		
Anopheles gambiae	$28.5 \pm 0.7 \ (38)^{c}$	_	-	-	Li et al., 2019		
Anopheles stephensi	$30.3 \pm 0.9 \ (17)^{c}$	-	-	-	Li et al., 2019		
Aedes aegypti	$32.0 \pm 0.8 \ (18)^{c}$	_	-	-	Li et al., 2019		
Culex pipiens pallens	$21.8 \pm 0.7$ (20) <sup>c</sup>	-	-	-	Li et al., 2019		
Patiria pectinifera	$34.8 \pm 0.5 \ (23)^{c}$	-	-	-	Saito et al., 2017		

TRPA1, thermal threshold of TRPA1; EVM, experimental voluntary maximum; CT<sub>max</sub>, critical thermal maximum; T<sub>mean</sub>, field average body temperature; Hyphens, data not available. Letters in upper case denotes the criteria (a, b, and c) used for the data summary as described in method.

The values are given in °C, and numbers in parentheses denote the sample sizes for each trait. The values for TRPA1 derived from the electrophysiological analysis using HEK293 cells and Xenopus oocytes.



**FIGURE 2** Relationships between thermal threshold of TRPA1 and three physiological parameters. **(A–C)** Experimental voluntary maximum (EVM), critical thermal maximum (CT<sub>max</sub>), and average body temperature (T<sub>mean</sub>) were plotted against thermal thresholds of TRPA1. The dashed and solid lines represent regression lines produced from OLS and PGLS analyses, respectively. The numbers near the dots indicate the species studied: (1) *Crotalus atrox* (western diamondback rattlesnake), (2) *Elaphe obsoleta* (Texas rat snake), (3) *Anolis carolinensis* (green anole), (4) *Anolis allogus*, (5) *Anolis homolechis*, (6) *Anolis sagrei* (brown anole), (7) *Xenopus tropicalis* (western clawed frog), (8) *Xenopus laevis* (African clawed frog), and (9) *Drosophila melanogaster* (fruit fly).

in terrestrial ectotherms by conducting OLS and PGLS analyses. Previous studies on individual species or sets of closely related species suggested that the activation of TRPA1 induces heat avoidance behaviors (Rosenzweig et al., 2005; Hamada et al., 2008; Neely et al., 2011; Saito et al., 2016; Akashi et al., 2018; Li et al., 2019). The regression analyses also supported the previous findings even with the inclusion of diverse taxa, and revealed that thermal threshold of TRPA1 was specifically related

Model	Framework	Intercept		Slope			
		Estimate ± SE	Р	Estimate ± SE	Р	R <sup>2</sup>	MLλ
EVM ~ TRPA1	OLS	14.21 ± 6.48	0.065	0.6 ± 0.18	0.014	0.6	n/a
	PGLS	$18.23 \pm 5.88$	0.017	$0.46 \pm 0.16$	0.026	0.53	0.904
$\text{CT}_{\text{max}} \sim \text{TRPA1}$	OLS	$54.0 \pm 10.66$	0.004	$-0.46 \pm 0.30$	0.187	0.32	n/a
	PGLS	$54.0 \pm 10.42$	0.004	$-0.46 \pm 0.30$	0.184	0.32	0
$T_{mean} \sim TRPA1$	OLS	$23.28 \pm 8.2$	0.047	$0.17 \pm 0.24$	0.524	0.11	n/a
	PGLS	$21.96 \pm 10.3$	0.1	$0.21\pm0.29$	0.51	0.11	0

TABLE 2 | Results of ordinary least squares (OLS) and phylogenetic generalized least squares (PGLS) analyses performed on the data obtained from HEK293 cells and Xenopus oocytes.

Significant correlations are shown in boldface; n/a, not applicable.

to EVM rather than to  $CT_{max}$  and  $T_{mean}$ . The results suggest that although thermal threshold of TRPA1 is evolutionarily labile, its association with EVM is highly conserved among diverse terrestrial ectotherms. Animals stay active within thermal range bounded by maximum and minimum voluntarily tolerable temperatures (Vitt and Caldwell, 2008); thus, TRPA1 may be involved in defining the upper limit of species thermal-safety margins.

EVM is, by definition, lower than  $CT_{max}$  as the animals should retreat from heat sources before they become incapable of keeping right posture due to the heat stress. Therefore, thermal threshold of TRPA1 may have been kept lower than CTmax through adaptive evolution, allowing the animals to perceive stressful temperatures before experiencing lethal temperatures. The insignificant correlation between thermal threshold of TRPA1 and CT<sub>max</sub>, otherwise, suggests that these two parameters evolve independent of one another, such that the other sensory molecules are responsible for perceiving the lethal temperatures. Field body temperature of ectothermic animals tends to fluctuate throughout a day, season, and also along a thermal environmental gradient and thus should ideally be assessed by a large number of observations, e.g., observations on at least 10 individuals in the case of lizards (Hertz et al., 2013). In the current study, each of the grand mean for T<sub>mean</sub> derived from extensive field observations except for C. atrox (Table 1). Nevertheless, Tmean did not yield any significant correlation with thermal threshold of TRPA1. As ectothermic animals adjust their body temperatures depending on costs and benefits associated with thermoregulation, such as intra- and/or interspecific competition (Huey and Slatkin, 1976; Huey and Stevenson, 1979), the ectothermic animals may not be able to achieve or adjust their body temperatures, depending solely on temperature perceived through TRPA1.

While thermal threshold of TRPA1 and EVM were significantly correlated one another, that of TRPA1 in some species deviated from EVM in a few Celsius degrees. In other words, heat avoidance behaviors seem not to be induced exactly at thermal threshold of TRPA1. This could be true but alternate explanations are also possible. Thermal thresholds of TRPA1 assessed in the current study derived from the electrophysiological analysis conducted on heterologously expressed TRPA1 using *Xenopus* oocytes or HEK293 cells. Two-electrode voltage clamp or patch-clamp recordings are particularly useful for analyzing the physiological properties

of ion channels although cell physiology, such as the ion compositions across cell membranes and/or the membrane potentials, varies depending upon the experimental protocols. The artificial cell physiology could result in the deviation of thermal thresholds of TRPA1 from those in vivo. Based on Ca<sup>2+</sup>imaging experiments, for example, a TRPA1 agonist-responsive neurons in dorsal root ganglions (DRG) increased intracellular calcium concentration at a threshold a few degrees lower than thermal thresholds of TRPA1 estimated from heterologous expression systems in chickens (Saito et al., 2014). In addition, the involvement of the other molecules may account for an offset of few Celsius degrees between the two parameters. In DRG neurons, the other thermoTRP channel, TRP vanilloid 1 (TRPV1), is often co-expressed with TRPA1 and has also been suggested as a heat sensitive channel in native DRG neurons as well as TRPA1 (Story et al., 2003; Prober et al., 2008; Saito et al., 2012, 2014). In the case of fruit flies, four TRP channels belong to the TRPA subfamily, and Painless in addition to TRPA1 was also reported to be involved in avoidance of noxious heat (Neely et al., 2011). Thus, the integration of these heat-sensitive channels may better explain thermal sensitivities at whole-animal levels.

Although thermal thresholds of TRPA1 were analyzed in the current study, other physiological properties of thermoTRPs could also be important parameters. For instance, a member of thermoTRP channels, TRP melastatin 8 (TRPM8), showed that its half-maximal activation temperatures, rather than its thermal threshold, were correlated with core body temperatures in chicken, rat, and two *Xenopus* species, *X. laevis* and *X. tropicalis* (Myers et al., 2009). In addition, the thermal sensitivities of TRPA1, as characterized by normalized current sizes, were less sensitive to heat in the presumed hot-adapted species, *X. tropicalis* and *X. borealis*, compared with the presumed cold-adapted species, *X. laevis* and *X. muelleri* (Saito et al., 2019). As such, other physiological properties of thermoTRPs may also provide keys to account for species thermal biology.

In this study, the regression analyses on diverse taxa showed that the values of EVM can be explained as a function of thermal threshold of TRPA1. Therefore, the study on thermal physiological properties of TRPA1 may help evaluate species' vulnerability to thermal stress that is not always easy to measure from behavioral experiments due to limitations of animal handling, such as substantially large or small body size, conditions under captivity, or the conservation status. In these cases, the thermal physiological properties of TRPA1 could be useful particularly in the recent climate warming scenario, since electrophysiological analysis can be conducted if the nucleotide sequences of target thermoTRPs have been reported. Further studies on thermoTRPs both from physiological and ecological perspectives are expected and will lead to a better understanding of thermoTRPs' functional evolution regarding species' thermal ecology.

### DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found in the article/ **Supplementary Material**.

# **AUTHOR CONTRIBUTIONS**

The author confirms the sole responsibility for the following: study conception and design, data collection, analysis and interpretation of the results, and manuscript preparation.

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# SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fevo.2021. 583837/full#supplementary-material

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**Conflict of Interest:** The author declares that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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