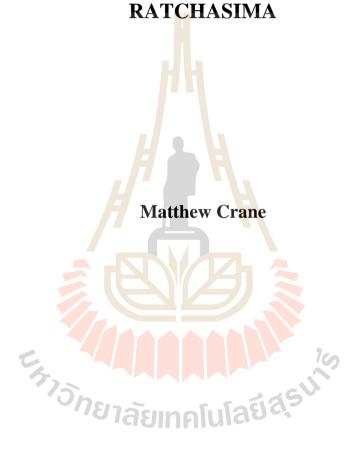
EFFECT OF LAND-USE AND HUMAN DISTURBANCE ON HERPETOFAUNAL COMMUNITIES IN THE SAKAERAT BIOSPHERE RESERVE, NAKHON



A Thesis Submitted in Partial Fulfillment of the Requirements for the

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ผลกระทบจากการใช้ที่ดินและการรบกวนของมนุษย์ต่อชุมชนของ สัตว์เลื้อยคลานและสัตว์สะเทินน้ำสะเทินบก ในเขตสงวนชีวมณฑลสะแกราช จังหวัดนครราชสีมา



วิทยานิพนธ์นี้เป็นส่วนหนึ่งของการศึกษาตามหลักสูตรปริญญาวิทยาศาสตรมหาบัณฑิต สาขาวิชาชีววิทยาสิ่งแวดล้อม มหาวิทยาลัยเทคโนโลยีสุรนารี ปีการศึกษา 2558

EFFECT OF LAND-USE AND HUMAN DISTURBANCE ON HERPETOFAUNAL COMMUNITIES IN THE SAKAERAT BIOSPHERE RESERVE, NAKHON RATCHASIMA

Suranaree University of Technology has approved this thesis submitted in partial fulfillment of the requirements for a Master's Degree.

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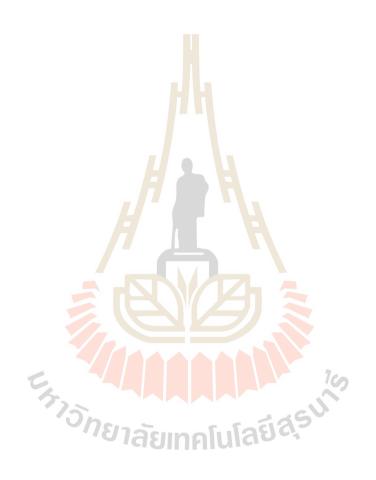
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แมททิวเครน: ผลกระทบจากการใช้ที่ดินและการรบกวนของมนุษย์ ต่อชุมชนของ สัตว์เลื้อยคลานและสัตว์สะเทินน้ำสะเทินบก ในเขตสงวนชีวมณฑลสะแกราชจังหวัด นกรราชสีมา (EFFECT OF LAND-USE AND HUMAN DISTURBANCE ON HERPETOFAUNAL COMMUNITIES IN THE SAKAERAT BIOSPHERE RESERVE, NAKHON RATCHASIMA) อาจารย์ที่ปรึกษา: ผู้ช่วยศาสตราจารย์ ดร.พงศ์ เทพ สุวรรณวารี, 82 หน้า.

พื้นที่อนุรักษ์เพียงอย่างเดียวไม่สามารถช่วยรักษาความหลากหลายทางชีวภาพไว้ได้อย่าง สมบรูณ์ ดังนั้นการศึกษาพื้นที่ที่ถูกมนุษย์รบควนเพื่ออนุรักษ์ความหลากหลายทางชีวภาพของโลก จึงเป็นเรื่องที่จำเป็น ในแถบเอเชียตะวันออกเฉียงใต้ซึ่งป่าไม้ถูกทำลายไปอย่างมากมายแต่ยังขาด การวิจัยด้านสังคมของสัตว์เลื้อยคลานและสัตว์สะเทินน้ำสะเทินบกในพื้นที่ถูกมนุษย์รบกวน งานวิจัยนี้มุ่งศึกษาชุมชนของสัตว์เลื้อยคลานและสัตว์สะเทินน้ำสะเทินบกในป่าเต็งรัง ป่าที่ถูก รบกวน และสวนป่ายูลาลิปตัส การสำรวจดำเนินการในเดือนพฤษภาคม มิถุนายนและกันยายน พ.ศ. 2558 โดยการวางกับดักตาข่ายจำนวน 11 แห่ง แต่ละแห่งประกอบด้วยกับดักกรวยสองชั้น จำนวน 12 กับดักและหลุมกับดักขนาด 40 ลิตรจำนวน 3 หลุม และเก็บข้อมูลลักษณะแหล่งที่อยู่ อาศัยแต่ละแห่งจากแปลงขนาด 1 x 1 ตารางเมตร จำนวน 6 แปลง ข้อมูลด้านภูมิประเทศ เช่น ความสูงจากระดับน้ำพะเลและขนาดของพื้นที่ศึกษา ถูกกำหนด โดยโปรแกรม ArcGIS การ วิเคราะห์ความสอดคล้องเชิงพหุ (Multiple Correspondence Analysis: MCA) พบว่าพื้นที่ศึกษามี ความแตกต่างกันอย่างเป็นอิสระ

ผลการศึกษาพบสัตว์เลื้อยคลานและสัตว์สะเทินน้ำสะเทินบกจำนวน 861 ตัว จำแนกได้ 40 ชนิด โดยป่ายูคาลิปตัส (361 ตัว, 37 ชนิด) และป่าที่ถูกรบกวน (373 ตัว, 29 ชนิด) มีความชุกชุม และจำนวนชนิดมากกว่าป่าเต็งรัง (127 ตัว, 29 ชนิด) สำหรับสัตว์สะเทินน้ำสะเทินบกพบความชุก ชุมในป่ายูคาลิปตัสและป่าที่ถูกรบกวนมากกว่าในป่าเต็งรัง แต่ในสัตว์เลื้อยคลานกลับพบความชุก ชุมในป่ายูกาลิปตัสมากกว่าป่าทั้งสองประเภท

ความมากชนิดของสัตว์เลื้อยกลานและสัตว์สะเทินน้ำสะเทินบกไม่ได้ผันแปรตาม ประเภทของป่า แต่เมื่อเปรียบเทียบเฉพาะความมากชนิดของสัตว์สะเทินน้ำสะเทินบกเท่านั้นพบว่า ในป่าเต็งรังมีพบความหลากหลายทางจำนวนชนิดของสัตว์สะเทินน้ำสะเทินน้อยกว่าในป่าที่ถูก รบกวนและป่ายูกาลิปตัส ผู้วิจัยใช้เทคนิก non-metric multidimensional scaling และทดสอบด้วย PERMANOVA เพื่อศึกษา หาความแตกต่างโครงสร้างของชุมชนสัตว์เลื้อยคลานและสัตว์สะเทินน้ำสะเทินบกไม่พบความ แตกต่างอย่างมีนัยสำคัญระหว่างประเภทของป่ากับโครงสร้างของชุมชน



สาขาวิชาชีววิทยา ปีการศึกษา 2558 MATTHEW CRANE: EFFECT OF LAND-USE AND HUMAN
DISTURBANCE ON HERPETOFAUNAL COMMUNITIES IN THE
SAKAERAT BIOSPHERE RESERVE, NAKHON RATCHASIMA.
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82 PP.

DIVERSITY/SPECIES RICHNESS/COMMUNITY STRUCTURE/HABITATS

Protected areas cannot completely sustain biodiversity, thus understanding the role of human-disturbed areas in conserving the world's diversity is critical. Despite intensive deforestation, Southeast Asia is underrepresented in studies investigating faunal communities in human-modified landscapes. This project assessed the herpetofaunal community in dry dipterocarp forest, secondary disturbed forest, and eucalyptus plantations in the Sakaerat Biosphere Reserve. In May, June, and September of 2015, we surveyed using 11 passive trapping arrays. Each array consisted of 12 double funnel traps and three 40 L pitfall traps. We assessed habitat characteristics at each site collected from six 1 m x 1 m quadrats at each site. To collect landscape variables such as elevation and patch size, we used ArcGIS mapping software. Analysis of the habitat characters using multiple correspondence analysis (MCA), did not reveal any clustering of the sites.

The results showed 861 individuals were captured representing 40 species.

The eucalyptus plantations (361 individuals, 37 species) and highly disturbed forest

sites (373 individuals, 29 species) hosted higher herpetofauna abundance and richness than protected areas of dry dipterocarp forest (127 individuals, 29 species).

Both the PLE and HDF (224 and 275 individuals respectively) had higher amphibian abundance than the DDF (57 individuals)Reptiles in contrast showed the highest abundance in the PLE sites compared to the other two forest types, with 100 captures compared to 54 in the DDF and 57 in the HDF. Based on sample based rarefaction herpetofaunal species richness did not vary between forest types, but did vary significantly between the DDF compared HDF and PLE sites when comparing amphibian species. We did not find any significant differences in community structure among forest types based on non-metric multidimentional scaling and PERMANOVA.



School of Biology Academic Year 2015 Student's Signature Motor

Advisor's Signature P. Suran

Co-advisor's Signature

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CONTENTS

	Page
ABSTRACT IN THAI	I
ABSTRACT IN ENGLISH	III
ACKNOWLEDGEMENTS	V
CONTENTS	VII
LIST OF TABLES.	X
LIST OF FIGURES.	XII
CHAPTER	
I INTRODUCTION	1
1.1 Introduction	1
1.2 Research objectives	4
1.3 Scope and limitations	4
II LITERATURE REVIEW	6
III METHODS	13
3.1 Study site	13
3.2 Assessment of forest types	15

CONTENTS (Continued)

		Page
	3.3 Captures and biometrics	21
	3.4 Data analysis	22
	3.4.1 Abundance, richness, and diversity	22
	3.4.2 Community structure	24
	3.4.3 Body condition	25
IV	RESULTS AND DISCUSSION	26
	4.1 Results	26
	4.1.1 Habitat assessment	26
	4.1.2 Plot captures.	27
	4.1.3 Mortalities	31
	4.1.4 Comparison of funnel and pitfall traps	31
	4.1.5 Abundances	32
	4.1.6 Species richness	37
	4.1.7 Diversity	39
	4.1.8 Community structure	40
	4.1.9 Analysis of biometrics	47

CONTENTS (Continued)

	Page
4.2 Discussion	49
4.2.1 Abundance, species richness, and diversity	49
4.2.2 Community structure	50
4.2.3 Biometrics	51
V CONCLUSION AND RECOMMENDATIONS	52
REFERENCES	54
APPENDICES	
APPENDIX A SITE HABITAT AND ENVIRONMENTAL DAT	A63
APPENDIX B RICHNESS, ABUNDANCE, AND COMMUNITY	Y DATA70
APPENDIX C BIOMETRICS DATA	79
CURRICULUM VITAE	82
"ยาลยเทคโนโลยด์"	

LIST OF TABLES

Tab	ole	Page
2.1	The impacts of human landscapes on herpetofaunal communities in the	
	tropics	7
3.1	Landscape characteristics for each plot site used in the study, negative	
	values are used to indicate that site is located inside the SBR core area	16
3.2	Each category that was assessed in the study and which data and statistical	
	methods were used	22
4.1	Contribution to the variance between sites from eigenvalues resulting from	
	multiple correspondence analysis of site habitats	27
4.2	Comparison of sampling effort as the number of trap nights for each forest	
	type	27
4.3	Amphibian captures by species from three forest types DDF, HDF, and	
	PLE 160	29
4.4	Reptile captures by species from three forest types DDF, HDF, and PLE	30
4.5	Reptile and amphibian trap mortalities from each forest type and for each	
	month sampled	31
4.6	The number of amphibians and reptiles captured in the SBR from each forest	
	type and the capture efficiency of each trap night	32
4.7	The calculated Shannon-Weiner and Simpson diversity indexes for each	
	plot	40

LIST OF TABLES (Continued)

Tabl	le	Page
4.8	PERMANOVA results modeling reptile, amphibian, and total captures	
	based on forest type and other identified environmental factors	46
4.9	Significant results from indicator species analysis using abundance and	
	presence absence community data	47
4.10	Results from the ANOVA of BCI by forest type for E. macularia and L.	
	capucinus	48
4.11	Processing information for all snakes captured during the study	48



LIST OF FIGURES

Figu	ure	Page
3.1	Map of the Sakaerat Biosphere Reserve delineating core, buffer, and	
	transition areas	14
3.2	All three forest types that were sampled in the Sakaerat Biosphere Reserve	
	(Highly disturbed forest, dry dipterocarp forest, and plantation	
	forest)	17
3.3	Map of the study area showing steambeds and composite of Landsat 8	
	images to clearly determine water sources	18
3.4	Passive trapping array showing the line, wings, and traps	19
3.5	Plot layout showing the locations used to assess habitats and an example	
	quadrat used to estimate ground cover in the study	21
4.1	Plotting of sites based on multiple correspondence analysis of habitat	
	variables	26
4.2		
	amphibians, reptiles, and all herpetofauna	33
4.3	Whittaker plots for complete amphibian captures and reptiles for DDF, PLE,	
	and HDF	35
4.4	Seasonal rank abundance curves from reptile and amphibians species from	
	all forest types combined	36
4.5	Plot of amphibian abundance by distance to the edge of SBR, distance to	
	water patch size and elevation	37

LIST OF FIGURES (Continued)

Figu	ire	Page
4.6	Sample based and individual based rarefaction for each forest type based on	
	all captured herpetofauna	38
4.7	Sample and individual based rarefaction for amphibians and for reptiles	39
4.8	Comparison of heirarchic clustering of samples sites based on the recorded	
	abundances of reptiles and amphibians	41
4.9	Comparison of heirarchic clustering between abundance and presence	
	absence data for all captured herpetofauna	42
4.10	Non-metric multidimensional scaling for sites based on observed	
	amphibians, reptiles, and herpetofauna with significant environmental	
	variables	44

CHAPTER I

INTRODUCTION

1.1 Introduction

Global biodiversity stands on the brink of a mass collapse. Human activities threaten species across all taxa and regions, but tropical regions in particular are a major battle ground for global biodiversity. The tropics hold over 60% percent of the Earth's species, yet human population growth has directly conflicted with biodiversity conservation causing widespread deforestation in many tropical regions (Dirzo and Raven, 2003). Conservation measures generally focus on developing and expanding the global protected area network as refuges for biodiversity.

The global protected area network protects 460 million ha (~12.5% of total forest area) (FAO, 2010) of forest cover from deforestation; however many reserves are becoming isolated from other large tracts of undisturbed landscapes (Sánchez-Azofeifa et al., 1999; DeFries and Hansen, 2005). Isolation could mean extinction for a multitude of species as global climate change puts additional pressure on populations by shifting suitable habitat ranges (Bickford et al., 2010). Additionally, many reserves may not effectively protect biodiversity as illegal poaching can lead to the large scale defaunation (Harrison, 2011). Rodrigues et al. (2004) calculated a conservative estimate of gap species, those not represented in protected reserves, by overlaying species distributions and protected area maps, identifying 1,424 species not protected under the current reserve system. The

limitations for the current reserve system for biodiversity conservation have led to an increase in research focusing on the role of human-disturbed areas in sustaining the world's diversity.

Deforestation and the accompanied loss of biodiversity is a particular challenge for sustainable development in Southeast Asia (Fox and Vogler, 2005). Southeast Asia has the highest rates of annual deforestation of all tropical regions, which has been projected to result in the loss of 13-42% of all regionally endemic species in the next century (Brooks et al., 2002). Thailand in particular has been victim to extensive loss of forest cover as land-use changes over time. From 1961 to 1998 estimates indicated that the forest cover declined from 53% to 25% in the 37 year period (Charupat, 2000). A landscape change assessment for the Northeast of Thailand showed an increase in the isolation for a protected forest along with an increase in agriculture and urban development in the area from 1980 to 2010 (Sutthivanich and Ongsomwang, 2015).

Deforestation and other anthropogenic impacts such as urbanization and poaching are causing declines across all taxa; however, amphibians are the most threatened of all terrestrial vertebrates (Sodhi et al., 2010). Despite very different morphological and life history traits reptile species are imperiled by many of the same anthropogenic influences (Gibbon et al., 2000). A recent assessment on the global conservation status of reptiles revealed that over 18% of the 9,084 described species are threatened (Böhm et al., 2013). The status of amphibians mirrors that of reptiles with 7.4% of the 5,743 known species listed as Critically Endangered on the IUCN red list (Stuart et al., 2004). Along with lizards and amphibians snake populations are thought to be in decline globally (Reading and Luiselli, 2010), highlighting the need for herpetofaunal

community and population studies to monitor the observed trends. Reptile populations face a wide array of extinction threats from anthropogenic activities. Species affected by anthropogenic habitat loss were particularly prevalent in Southeast Asia (Böhm et al., 2013). The conservation value of human-modified landscapes must be assessed to inform conservation action plans in the region.

Despite the high rate of deforestation, Southeast Asia is generally underrepresented in studies on faunal community response to habitat loss and response to human-modified landscapes (Trimble and Aarde, 2012). Additionally, herpetofauna is globally underrepresented in community studies and in their response to anthropogenic disturbance (Voris, 2006; Trimble and Aarde, 2012). Thailand is home to more than 142 species of amphibians and over 218 species of reptiles (IUCN, 2014), and has the smallest area of remnant forest cover in Southeast Asia (Sodhi et al., 2010). The herpetofaunal diversity and the level of human disruption make Thailand an ideal site to investigate the impacts of land-use change on tropical amphibian and reptile communities.

The United Nations Educational, Scientific, and Cultural Organization (UNESCO) Man and the Biosphere Programme was developed to establish sustainable landscapes; balancing biodiversity conservation and sustainable human development. The Sakaerat Biosphere Reserve is home to 41% of Thailand's reptile species and 22% of the amphibian species listed on the IUCN red list (IUCN, 2014; SERS, 2014). With a diverse landscape of protected forest and mixed agricultural land it is an ideal location to study herpetofauna and the impacts of land-use on communities. Recent studies in the area have led to a deeper understanding of specific amphibian and reptile species such as hatching plasticity in the frogs (Poo and Bickford, 2014), the

spatial ecology and behavior of king cobras (Strine et al., 2014), and the discovery of a new frog species (*Limnonectes megastomias*) (McLeod, 2008). However only one study from the area has assessed a herpetofaunal community, which was focused only on semi-aquatic snake species (Genus *Enhydris*) (Karns et al., 2010). The proposed project will be the first full herpetofaunal assessment of the disturbed landscapes in the Sakaerat Biosphere Reserve.

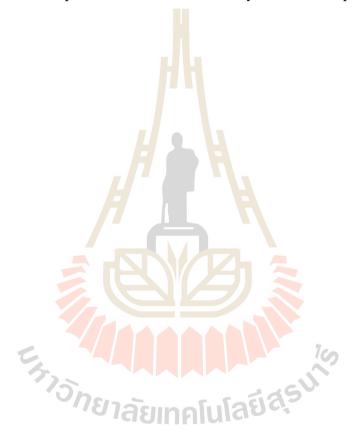
1.2 Research objectives

- 1) To determine abundance, richness, and diversity of reptile (excluding birds) and amphibian species in the (a) secondary forests, (b) plantation forests, and (c) protected areas of the Sakaerat Biosphere Reserve.
- 2) To evaluate the structure of amphibian and reptile communities in three different land use types
- 3) To determine the impact of land-use on body condition of highly abundant species

1.3 Scope and limitations

Starting in May 2015 herpetofaunal community sampling took place in the core area and the transition area of the Sakaerat Biosphere Reserve. The study area was confined to the Northeast including the majority of the dry dipterocarp forest in the core area and a patchwork of mixed agricultural and small plantation. We sampled three land use types: heterogeneous disturbed forest (HDF), eucalyptus plantation forest (PLE), and primary dry dipterocarp forest (DDF). Dry evergreen forest which is present within the Southwest of the Biosphere Reserve were not sampled, as the

disturbed areas were most likely dry dipterocarp forest before conversion by human activities. Plantations forests were considered monoculture forest stands of eucalyptus. Heterogeneous disturbed forests are fragments of natural forests with high levels of anthropogenic disturbance embedded in an agricultural matrix. The survey methods were limited to passive trapping arrays (12) using both double-funnel and pitfalls traps. We sampled once each month in May, June, and September.



CHAPTER II

LITERATURE REVIEW

2.1 Herpetofaunal diversity in human-dominated landscapes

Research projects investigating community response to human-disturbance are typically heavily biased towards specific taxa and regions. Most studies come out of the Neotropical forests and focus on mammals, plants, or invertebrates (Trimble and Aarde, 2012). However, several studies have assessed both the methods of studying herpetofauna in human-disturbed landscapes and the impact on community assemblages (Urbina-Cardona et al., 2006; Gardner et al., 2007b; Ribeiro-Júnior et al., 2008) (Table 2.1). The ability of plantation forests to maintain herpetofaunal diversity is often compared to pristine forests to elucidate similarities in community structure impacts from land-use change.

รักยาลัยเทคโนโลยีสุรมใช้

Table 2.1 The impacts of human landscapes on herpetofaunal communities in the tropics.

Habitat types	Places	Amphibian		Reptile		- References	
Habitat types	1 faces	Species	Individuals	Species	Individuals	References	
Remnant forest bordered with palmetto		18	142	10	58		
Remnant forest bordered with pasture	Near La Selva Biological Station, Costa	17	225	10	61	Kurz et al.	
Palmetto plantation	Rica	11	68	2	2	(2014)	
Pasture		9	46	12	66		
Mature primary rainforest		22	515	25	681		
14-19 years secondary forest	Northeastern Brazilian Amazonia, Brazil	14	510	15	87	Gardner et al. (2007b)	
4–5 years <i>Eucalyptus</i> plantation		5	714	14	757	ai. (20070)	
40-400 years secondary forest	Hong Vone Chine	6	52	7	237	Sung et al.	
30-60 years <i>Lophostemon confertus</i> plantation	Hong Kong, China	4	16	8	134	(2012)	
Primary rainforest		7	46	2	2		
Selective logged rainforest	Kibale National Park, Uganda	8	94	4	17	Vonesh (2001)	
Pine plantation		8	102	4	4	(====)	
Primary rainforest		8	236	6	49		
Pentaclethra macroloba plantation	La Selva Biological Station, Costa Rica	5	95	4	13	Folt and Reider	
Virola koschnyi plantation	La Selva Biologicai Station, Costa Rica	7	117	3	25	(2013)	
Vochysia guatemalensis plantation	6. 4.	8	128	5	48		
Fragmented tropical rainforest	725	13	588	22	258	Urbina-	
Edge of fragmented tropical rainforest	Los Tuxtlas Biosphere Reserve, Mexico	14	570	25	208	Cardona et	
Pasture	<i>''ชา</i> ลยเท	12 3	98	11	157	al. (2006)	
Tropical dry forest	Chamela Biosphere Reserve, Mexico	4-13	9-46	22-28	178-276	Suazo- Ortuño et	
Disturbed tropical dry forest		6-8	23-40	19-29	238-304	al. (2008)	

While some results did not agree, most studies found significant differences in the abundance, richness, and community structure responses for amphibians and reptiles (Gardner et al., 2007b; Wanger et al., 2010, Kurz et al., 2014;). Typically amphibian communities were more sensitive to human disturbance than reptiles, which makes ecological sense as amphibian morphology (for example semi-permeable skin) means that environmental factors can have a greater impact (Cushman, 2006). However, the literature does not support a unanimous conclusion about the response of amphibian communities to human disturbance gradients.

One study from La Selva Biological Station in Costa Rica tested the influence of land use along the forest edge on herpetofaunal communities sampling palmetto, pasture, and natural forests (Kurz et al., 2014). They used generalized linear mixed models and determined the best habitat classification model for describing the observed species richness and abundance. The best fit model for amphibians grouped both disturbance areas (open pasture and palmetto plantation) as a single type compared to forest. The results indicate that regardless of the land-use the amphibian communities were sensitive to human habitat disturbance. A second study from the same site aimed at answering a different question related to human disturbance impacts.

Folt and Reider (2013) examined plantation forests of three separate species (*Pentaclethra macroloba*, *Virola koschnyi*, and *Vochysia guatemalensis*) compared to natural growth forest. The study found a significant difference in amphibian species richness between only *one* plantation type (*P. macroloba*) and the reference forest at the 95% confidence interval. The two other plantation forests under review contained comparable species richness to the reference forest. However, the authors also note

that community assemblage differed significantly between both *P. macroloba* and *V. guatemalensis* when compared to the primary forest amphibian community structure. Frog abundances were also highest in one of the plantation forests (*V. koschnyi*) than any other forest type. The study indicates that based on species richness, native monoculture plantation stands can support biodiversity conservation. The two studies from La Selva Biological Station illustrate the complexity of community response to human disturbance and the need for further research to isolate all of the factors in disturbance impacts.

Work from the Brazilian Amazonia tropical forests found similar results to both of the La Selva Station studies on the influence of human disturbance schemes on amphibian communities. Gardner et al. (2007a) sought to standardize the comparison of primary, secondary, and plantation forests in a Brazilian rainforest. Instead of investigating a reference primary forest in comparison to native species mono-culture plantations, the study sampled a gradient of disturbed forests: primary, secondary, and non-native *Eucalyptus* plantations. Results from the study found that amphibian abundances were similar across all forest types while the community structure varied between primary forest and secondary and plantation forests. The amphibian species found in the two disturbed forest types (secondary and plantation) were a subset of the primary forest amphibian community.

Similar to Folt and Reider (2013) who determined that plantation forests can support some species diversity, however Gardner et al. (2007b) did not find any disturbed forest that was comparable to the reference forest. Gardner et al. (2007b) additionally determined that secondary forests held significantly more amphibian species than plantation forests.

Additional evidence exists that secondary forests can contain relatively higher amphibian abundances than non-native plantation forests, even if species composition does not differ (Sung et al., 2012). Additionally, work in a Neotropical dry forest supported the conclusions that amphibian abundances are similar across disturbed and primary landscapes, while species richness and community structure decrease (Suazo-Ortuño et al., 2008). In Southeast Asia these results also are confirmed from the only study conducted in the region in Indonesia. Amphibian species richness decreased significantly along the human disturbance gradient (Wanger et al., 2010).

Despite many studies arriving at similar conclusions some notable studies found that disturbed areas had either a neutral impact or even a positive impact on amphibian communities. Vonesh (2001) investigated the differences between an undisturbed primary forest, a similar forest with historical selective logging, and a pine plantation forest in Kibale National Park, Uganda. He determined that the logged forest had higher herpetofaunal species richness and abundance than the undisturbed forest, but that the pine plantation forest had the highest overall species richness and abundance. Fredricksen and Fredricksen (2004) also arrived at very different conclusions in terms of amphibian species richness and abundance. The researchers also studied the impact of selective logging on amphibian communities, but found no significant differences between logged and unlogged habitats in Bolivia. The conflicting results demands more research to assess either the methodological shortcomings that caused the results or identify critical factors that can lead to such drastically different conclusions.

Reptile communities show different trends than amphibians in response to human habitat conversion. In general, reptile communities were more resilient to forest

change and human disturbance, and in some cases species richness and abundance increased along disturbance gradients. For example, the previously discussed Folt and Reider (2013) study determined that reptile community assemblages between the primary forest and two of the plantation forests were indistinguishable (*V. koschnyi* and *V. guatemalensis*) as opposed to a single plantation type for amphibians. Kurz et al. (2014) did not find that the studied disturbed forest type (palmetto) held higher reptile species richness or abundance compared to primary forest, but did determine that palmetto was less dissimilar to primary forest than heavily disturbed pasture land. Gardner et al. (2007b) found that primary forests held higher species diversity than secondary or non-native plantation forests, but that the species richness between secondary and plantation forests did not differ significantly for reptiles in direct contrast to amphibian species richness. Additionally, several unique reptile species were found only in plantation forests. While none of these studies found that disturbance increased reptile species richness or abundance, all of them illustrated that reptile communities are relatively less sensitive to change than amphibians.

Two studies on herpetofaunal communities did discover that reptile species richness increased. In an agricultural matrix around a Neotropical dry forest, lizard species richness, diversity, and abundance was higher than in the remnant comparison forest (Suazo-Ortuño et al., 2008). However, from the same study, turtles were highly sensitive to habitat conversion, and no snake species was sensitive to disturbance, indicating that treating all reptiles as a group may lead to misinterpreting the results. In the study by Wanger et al. (2010) out of Indonesia, reptile species richness and abundance peaked in a natural shade-cacao plantation forest. The results from all the

studies show positive implications for the availability of plantation and regrowth forest regimes in maintaining reptile diversity and abundance.

In addition to investigating the effect of human disturbance on reptile species richness, abundance, and community structure one study also attempted to assess the secondary impacts of disturbance on individual health. Sung et al. compared secondary forests and plantation forests in Hong Kong selecting the most abundant species of reptile (*Sphenomorphous indicus*) creating a body condition index from a linear regression of mass and SVL and counting the number of individuals exhibiting tail loss. Secondary forest individuals exhibited no significant difference in body condition; however, plantation individuals were significantly more likely to show tail loss, which the authors attributed to fewer shelter sites in plantation forests.



CHAPTER III

METHODS

3.1 Study site

The study was conducted within the Sakaerat Biosphere Reserve located in Nakhon Ratchasima Province, Thailand (14.44–14.55°N, 101.88–101.95°E). The reserve has an 80 km² core area and a combined 360 km² making up the buffer and transitional zones, which consist mostly of agricultural and settlement areas (Figure 3.1). The core area predominately consists of primary growth dry evergreen forest (60%), dry dipterocarp forest (18%), and secondary plantation forest (<18%) (Tongyai, 1980). The dry dipterocarp forest is endemic to South East Asia and is characterized with thick *Vietnamosasa pusilla* ground cover and dipterocarp trees such has *Shorea siamensis* and *Shorea obtusa* (Lamotte et al., 1998). The transition zone of SBR comprises nearly 82% of the total area and is characterized by isolated forest fragments in a patchwork of agricultural fields, small plantation forests, and human settlements.

The Sakaerat Biosphere Reserve hosts 93 reptile and 29 sp amphibian species. Four reptiles and three amphibian species are categorized as Threatened or Near Threatened on the IUCN Red List (IUCN, 2014; Sakaerat Environmental Research Station, 2014). However, the exact number of species is still unknown as populations of known species in the area may in fact be separate species (Voris, 2006). The estimate is also low as several species are Data Deficient or not present on the list.

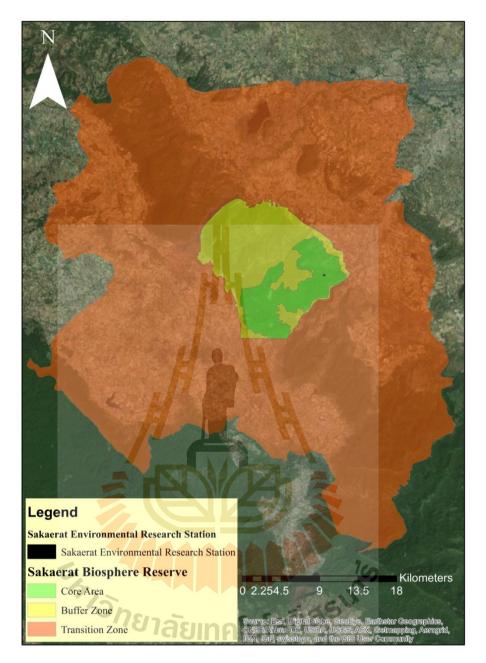


Figure 3.1 Map of the Sakaerat Biosphere Reserve delineating core, buffer, and transition areas.

3.2 Assessment of forest types

I assessed the herpetofaunal community assemblages across a gradient of human disturbance, specifically in remnant dry dipterocarp forest (DDF), highly disturbed forest (HDF), and eucalyptus plantation forest (PLE). For the purpose of the study disturbed forests were considered as secondary forests embedded in an agricultural matrix, characterized by high levels of anthropogenic change, but still similar to remnant forests in dominate vegetative cover type. Cycads and bamboo grass are typically present, along with species of Dipterocarpacae. We considered, eucalyptus plantations as forest stands planted with *Eucalyptus camaldulensis* for economic production. Within SBR eucalyptus stands are typically harvest every 2-5 years, at which point the trees are cut while the base is allowed to regrow. The eucalyptus trees are planted in straight lines with roughly two to five meters between rows.

Current land use maps provided by the Thai government were inaccurate with 7 out of 10 randomly selected points not matching the indicated classification. I used satellite imagery from Google Earth to create polygons by hand of each identifiable plantation and secondary forest within the study area. To test the accuracy of visual identification, I ground-truthed the newly created areas, which resulted in 9 out of 10 randomly selected sites falling in appropriate land-use type. This initial assessment indicated that the hand generated plantation and disturbed forest areas are sufficient to randomly select sites; however, further analyses of the area was required to accurately determine land use composition of the transition area.

We randomly selected twelve sites (4 in each forest type) using ArcMap 10.1. Plots were set no closer than 450 meters from each other to control for spatial autocorrelation. The total area of the disturbed forest patches (127.13 ha) and the

plantation forests (110.97 ha) were roughly equal; however, the plantation forests were much more fragmented and smaller (Figure 3.2).

Table 3.1 Landscape characteristics for each plot site used in the study, negative values are used to indicate that site is located inside the SBR core area.

Site	Elevation (m)	Patch size (ha)	Slope (°)	Distance from edge of core area (m)	Distance to water (m)	Water type
DDF - 1	368	527.7	7.0	-1298	475	Stream
DDF - 2	386	179.2	24.2	-324	950	Stream
DDF - 3	396	527.7	7.5	-1125	680	Stream
DDF - 4	294	527.7	12.1	-218	185	Stream
HDF - 1	254	33.1	4.5	1081	150	Pond
HDF - 2	272	57.9	7.1	3252	150	Pond
HDF - 3	265	33.1	1.5	979	310	Pond
HDF - 4	251	7.3	2.4	3150	46	Pond
PLE - 1	246	8.3	2.5	3988	160	Pond
PLE - 2	253	1.1	3.1	328	12	Stream
PLE - 3	274	1.3	9.6	2430	123	Pond

I investigated landscape factors for each plot site, including distance to water, patch size, slope, elevation, and distance to the edge of the protected area (Table 3.1). To determine distance to water I used a 32-day composite of Landsat 8 data from April 7th to May 9th to identify sources of water within the study area. Additionally I used a digital elevation model of the study area to determine stream beds using the Hydrology toolbox in ArcGIS 10.1 (Figure 3.3).

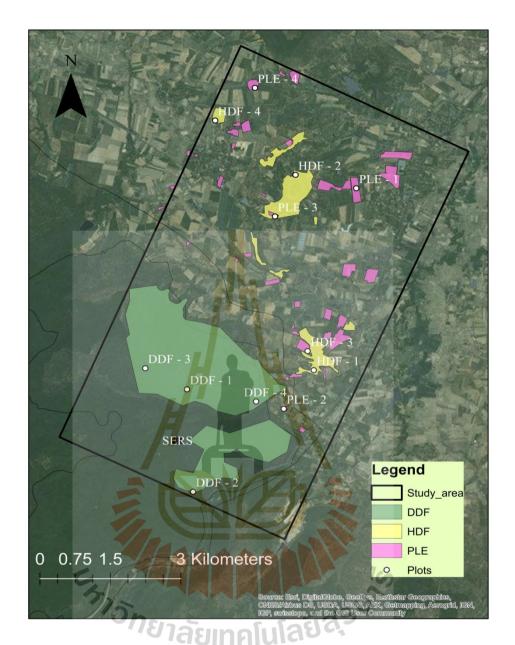


Figure 3.2 All three forest types that were sampled in the Sakaerat Biosphere Reserve (Highly disturbed forest, dry dipterocarp forest, and plantation forest).

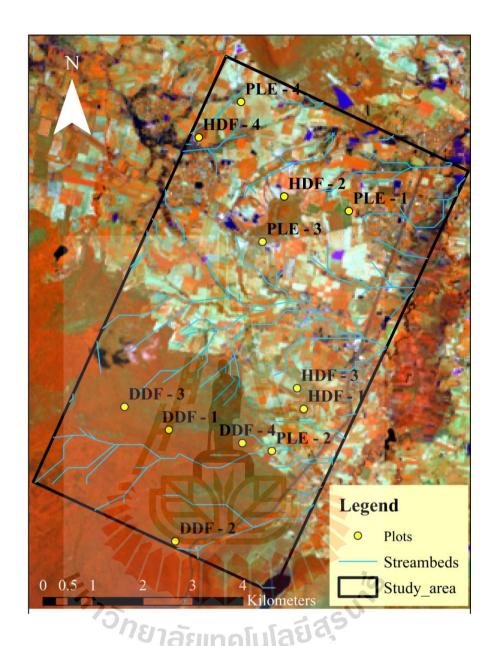


Figure 3.3 Map of the study area showing steambeds and composite of Landsat 8 images to clearly determine water sources.

To effectively sample the forest areas, we built Y-shaped drift-fence arrays with double funnel traps and 40 L pitfall traps. We attached two double funnel traps measuring 2 m x 0.5m x 0.3 m at the end of each line. A 40 L pitfall trap was placed at the midpoint of each line for a total of three pitfalls per array. Six double funnel traps were affixed to the center of each array for a total of 12 funnel traps and 3 pitfall traps (Figure 3.4). Each arm of the plot was 15 m in length with ground vegetation cleared half a meter on both sides. Each plot was oriented with 1 line running North-South with the two other lines angled 120° from the North-South line.



Figure 3.4 Passive trapping array showing the line (A), wings (B), and traps (C and D).

Each site was sampled for three days in May, June, and September. We sampled half of the plots, representing an equal number for each forest type, for 3 days and then switched to the second set. Prior to sampling each site was visited and repairs made to ensure that all plots were equivalent.

To account for habitat variations between sites, measured several environmental variables at each plot including: percent canopy cover, percent groundcover, leaf litter depth, leaf litter ground coverage, and vegetative density at the groundstory(<1 m), understory (1-3), midstory (3-5 m), and abovestory (> 5 m). Each variable was assessed from a 1 m x 1 m quadrat at 6 sites per plot, three set 7.5 m from the center away from each line, and three set 3 m away from the end of each line (Figure 3.5). To measure canopy density, I used hand-made densiometer created from a PVC pipe with fishing line creating four equal areas. Additionally, I assessed ground cover within each quadrat including percent coverage of leaf litter, rocks, grass, dead vegetation, fallen logs, trees/saplings, and bare ground. When leaf litter was present, I used a rigid ruler to measure depth to the closest1-mm. Ground cover and vegetation density were divided into 7 classifications (None, very light, light, medium, heavy, very heavy, complete).

Both May (8.55 mm) and June (79.25 mm), had lower precipitation than September (259.85 mm). Despite the differences in precipitation in 2015, the average rainfall in May (107.8 mm) and June (90.8 mm) over the past 4 years are comparable. Due to differences in average temperature, relative humidity, and rain fall May and June were categorized as dry season samples and September represented a single rainy season month (see Appendix 3 and 7).

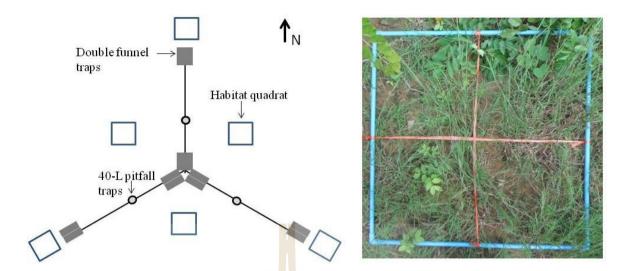


Figure 3.5 Plot layout showing the locations used to assess habitats (Left) and an example quadrat used to estimate ground cover in the study (Right).

3.3 Captures and biometrics

All snakes captured through the study were brought to the lab for processing to increase measurement precision. Individuals were marked with a field cauterizing unit on the ventral scales for mark and recapture analysis (Winne et al., 2006). Isoflorane was used to anesthetize captured snakes to determine accurate biometrics, which is proven to be a more accurate method for measuring individuals (Blouin-Demers, 2003; Setser, 2007).

We collected biometrics for an additional five species: *Kaloula pulchra, Kaloula mediolineata, Dixonious siamensis, Eutropis macularia,* and *Leiolepis reevesii rubritaeniata*. Captured individuals were processed in the field, at the time and site of capture. We collected snout-to-vent lengths (SVL) as well as mass, using a digital caliper and digital scale respectively. Mass and SVL were used to construct a body condition index for abundant species (Schulte-Hostedde et al., 2005; Peig and Green, 2009). Lizards were marked with xylene free permanent markers on the left and right

posterior jaw. While not a long term marking method, the technique allowed us to determine recaptures within sampling sessions.

3.4 Data analysis

3.4.1 Abundance, richness, and diversity

Table 3.2 Each category that was assessed in the study and which data and statistical methods were used.

Category	Data used	Methods
Environment	Habitat from site	Multiple correspondence analysis
Abundance	Captures	ANOVA, Kruskal-Wallis
Species richness	Captures	Sample based rarefaction
Diversity	Captures	Kruskal-Wallis
Community structure	Captures; landscape data; habitat data	NMDS, PERMANOVA, hierarchic clustering
Body condition	Biometric data	ANOVA, general least squares regression

We analyzed amphibian and reptile communities separately for all methods as life history traits and response to human disturbance can vary drastically between these two groups. Additionally we conducted the same analyses for all herpetofauna captures to determine whether any observed trends were consistent for both groups as a whole. Additionally, we compared differences in abundance, richness, and diversity for each month that was sampled.

Abundance was plotted against patch size, elevation, and distance to water to determine whether any potential correlations may exist. However, since the sample

size was limited to 11 sites and we select sites based on forest type and not environmental variables further statistical tests to verify differences were not possible. I used the categorical variables in a multiple correspondence analysis to investigate which environmental variables if any contributed to separation of the three forest types. Multiple correspondence analysis is similar to principle component analysis, but incorporates categorical rather than numeric inputs (Abdi and Valentin, 2007).

We compared both reptile and amphibian abundances between forest types using the accumulated total for each plot. Additionally to control for unequal sampling between seasons, we also tested the difference in abundance between the forest types for each month. For any that was non-parametric, we conducted a Kruskal-Wallis signed rank test followed by a pairwise comparison using a Wilcoxon signed rank test for any significant results at the 95% confidence interval. For data that fit the assumptions of parametric testing we used ANOVA with Tukey's post-hoc testing to determine significant differences at the 95% confidence interval. Additionally, we created Whittaker plots for each forest type to display the species dominance patterns for the entire landscape and for each individual forest type.

To analyze species richness we first created sample-based rarefaction curves using each day that a plot was open as a sampling unit. Rarefaction curves were created for each forest type and then assessed for significance at the 95% confidence interval. To assess sampling completeness we divided the observed species richness at each site by the estimated richness calculated using the Chao 1 estimate. Chao1 is a non-parametric estimator that provides a lower bound for species richness (Chao, 1984).

We compared diversity between forest types using two different indices. Firstly we used the standard Shannon-Wiener index which incorporates species richness and evenness to calculate diversity.

$$H' = -\sum_{i=1}^{s} p_i \ln (p_i)$$

Additionally we investigated differences between diversity using the Simpson index which incorporates species dominance to define diversity.

$$\lambda = \sum_{i=1}^{R} p_i^2$$

3.4.2 Community structure

Prior to any testing I transformed the community data for each group. I applied a total relativization to amphibian captures as several sites had a much higher number of individuals. Both reptiles and total herpetofauna were transformed using a log(x+1) to reduce the impact of highly abundant species (McCune and Grace, 2002). First to explore community similarity between sample sites I applied hierarchic clustering analysis based on the Bray-Curtis dissimilarity matrix for all plot sites. We used the complete linkage method to create the clusters, as our aim was to visualize the most compact groups with minimal within group spread (Manning et al., 2008). Next we tested for spatial auto-correlation between sites using the Mantel test. To remove any spatial correlations from later analysis we also created groups based on geographic distances using hierarchic clustering and then used that as an additional grouping variable.

We used non-metric multidimensional scaling (NMDS) as an ordination technique to visually represent the dissimilarity between sites. NMDS is a non-

parametric ordination technique that requires less assumptions from the data, and was more appropriate for our dataset which contained a high number of zeros (Zuur et al., 2007). We applied environmental fitting to the NMDS to determine which variables provided significant influence on site separation. Any variables that were in the 90% confidence were used in the formula for the follow up hypothesis testing. To test the hypothesis that community structure varied between forest types, we ran a PERMANOVA test with 999 permutations (Anderson and Walsh, 2013). We also used indicator species analysis to determine which species contributed to differences between forest types. All analyses were done in R Studio using packages "vegan", "MASS", and "BiodiversityR" (Venable and Ripley, 2002; Kindt and Coe, 2005; Husson et al., 2016; Oksansen et al., 2016; R Core Development Team, 2016).

3.4.3 Body condition

I created a body condition index using the residuals from a ordinary least squares regression of SVL to mass. Only species with measurements of at least 2 individuals from each forest type were select to create a body condition index. Two species, *D. siamensis* and *E. macularia*, fit the sample size requirements, but only *E. macularia* had data that was normally distributed and homogenous allowing for a linear regression.

We also analyzed whether forest type influenced whether reptile individuals had lost their tails as an indicator for fitness. Tail loss indicates potential predation pressure, and requires energy reserves to re-grow the dropped appendage. We used a chi-square test for independence, to determine whether forest type influenced tail loss for all captured reptiles and for the two most abundant species: *D. siamensis* and *E. macularia*

CHAPTER IV

RESULTS AND DISCUSSION

4.1 Results

4.1.1 Habitat assessment

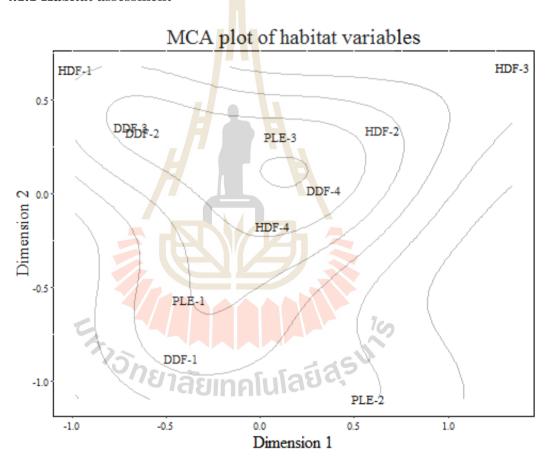


Figure 4.1 Plotting of sites based on multiple correspondence analysis of habitat variables.

The results of the multiple correspondence analysis did not show clear clustering of forest types (Figure 4.1). Additionally, the MCA results showed that no measured variables contributed strongly to the variance between each site (Table 4.1).

Table 4.1 Contribution to the variance between sites from eigenvalues resulting from multiple correspondence analysis of site habitats.

Dimension	1	2	3	4	5	6	7	8	9	10
Variance %	0.41	0.34	0.32	0.25	0.22	0.17	0.14	0.12	0.07	0.04
% Variance explained	19.4 6	16.4 7	15.3 0	12.0 9	10.6 0	8.32	6.67	5.57	3.57	1.93
Cumulative % variance explained	19.4 6	35.9 4	51.2	63.3	73.9	82.2 6	88.9	94.5	98.0 7	100

4.1.2 Plot captures

We sampled for a total of 1,977 trap nights over the course of the study. Two out of the 12 plots were stolen during the course of the study (1 in eucalyptus plantation and 1 in heterogeneous disturbed forest) resulting in uneven sampling between forest types (Table 4.2).

Table 4.2 Comparison of sampling effort as the number of trap nights for each forest type.

Forest type	Veral Funnel U. a.	Pitfall	Total
DDF	576	144	720
HDF	540	135	675
PLE	468	114	582
Grand Total	1584	393	1977

Throughout the study we recorded a total of 861 individuals comprising 40 recognized species. From the 861 captures 110 individuals (12.8% of total) were not completely identified to species, including 48 reptiles and 62 amphibians (Table 4.3

and Table 4.4). Geckos contributed to the majority of the unidentified reptiles (38 individuals), primarily due to the close similarity between the species *Hemidactylus* frenatus and Hemidactylus garnottii. For analysis we considered both Hemidactylus spp. as a single species.

Over half of the unidentified amphibians (35 individuals) came from the Microhylidae family and specifically the genus microhyla. Additionally, several individuals were not identifiable to genus due to ants destroying the specimen. For the analyses in the study these individuals were removed as we assessed richness and diversity at the species level.



Table 4.3 Amphibians captures by species from three forest types DDF, HDF, and PLE.

Family	Species	DDF	HDF	PLE	Total		
Bufonidae	Duttaphrynus melanostictu	S	7	5	12		
Microhylidae	Calluella guttulata		1	10	11		
	Glyphoglossus molossus	18	3	28	49		
	Kaloula mediolineata	20	27	11	58		
	Kaloula pulchra	13	6	13	32		
	Microhyla butleri	1	13	19	33		
	Microhyla heymonsi	1	20	28	49		
	Microhyla fissipes	1	78	51	130		
	Microhyla pulchra	1	53	8	62		
	Microhyla sp.	Н	12	23	35		
	Micryletta in <mark>o</mark> r <mark>n</mark> ata			11	11		
Ranidae	Hylarana e <mark>ryth</mark> raea			1	1		
	Hylarana <mark>macrodactyla</mark>		2	3	5		
Dicroglossidae	Fejervarya li <mark>mnocharis</mark>	2	66	31	99		
	Occ <mark>ido</mark> zyga lima			5	5		
Unknown	Unknown		23	4	27		
Species	7////	8	11	14	14		
Grand total		57	311	251	619		
⁷⁷ จักยาลัยเทคโนโลยีสุรูป							

Table 4.4 Reptile captures by species from three forest types DDF, HDF, and PLE.

Family	Species	DDF	HDF	PLE	Total
Agamidae	Calotes sp.			1	1
	Calotes versicolor	3		1	4
	Leiolepis reevesii	1	5	8	14
Colubridae	Boiga multimaculata	2			2
	Boiga siamensis	1			1
	Chrysopelea ornata	1		1	2
	Coelognathus radiatus	1			1
	Dendrelaphis suborcula <mark>ris</mark>			1	1
	Enhydris plumbea			1	1
	Lycodon capucinus	7	4	4	15
	Lycodon laoensis		1		1
	Oligodon fasciolatus	1		2	3
	Oligodon pseudotaen <mark>i</mark> atus	2	1		3
	Oligodon taeniatu s			1	1
	Psammodynates				
	pulverulentus _	1			1
	Rhabdophis c <mark>hry</mark> sargus			1	1
Gekkonidae	Boiga multi <mark>mac</mark> ulata		1		1
	Dixoneus s <mark>i</mark> amensis	14	9	17	40
	Gehyra la <mark>c</mark> erata	4	3	2	9
	Hemidactylus sp.		6	14	20
Elapidae	Bung <mark>aru</mark> s candidus		1	3	4
	Calliophis maculiceps	1			1
	Naja s <mark>iamensi</mark> s	1	1		2
Scincidae	Eutropis <mark>macularia</mark>	10	16	39	65
	Lygosoma bowringii	3	7	4	14
Typhlopidae	Ramphotyphlops albiceps	1 4	1		1
	Ramphotyphlops braminus	IJakia.	1		2
Viperidae	Calloselasma rhodostoma	Ulcio		1	1
Unknown	Unknown	16	5	9	30
Species		17	14	17	26
Grand total		70	62	110	242

4.1.3 Mortalities

Over the course of the study 12.9% (111 individuals) died either in the trap or from processing. However, morality rates fluctuated between forest type, trap type, and sample month (Table 4.5).

Table 4.5 Reptile and amphibian trap mortalities from each forest type and for each month sampled.

Forest type	Month	Reptiles	Amphibians
DDF	May	1	0
	June	1	0
	September	0	0
	Total	2	0
HDF	May	3	23
	June	2	22
	Septem <mark>ber</mark>	0	3
	Total	5	48
PLE	May	2	2
	June	8	39
	September		4
	Total	11	45
Grand Total		18	93

HDF and PLE site had higher mortalities rates (14.2% and 15.5% respectively), when compared to the DDF sites (1.6%). Amphibians were more sensitive to trap related mortalities losing 93 individuals (15.0% of total), compared to just 18 reptiles (7.4% of total). Trap type also effected mortality rates with funnel traps proving more dangerous to animal safety with 57 mortalities compared to just eight.

4.1.4 Comparison of funnel and pitfall traps

Due to the plot design funnel traps accounted for a higher amount of trapping effort, and as a result caught a higher number of individuals. When

comparing trap efficiency (captured individuals/trap night), pitfall and funnel traps were similar for all herpetofauna; however when dividing reptile and amphibian captures, the results suggest that pitfall traps did not perform as well for reptiles than for amphibians (Table 4.6). Despite similar efficiencies in capturing individuals, funnel traps captured more amphibian and reptiles species, 14 and 24 species respectively, than pitfall traps, 6 and 12 species respectively. However, both trap types captured at least 1 unique species. I analyzed abundance, species richness, and diversity using the combined captures from both trap types to cover any biases in either method.

Table 4.6 The number of amphibians and reptiles captured in the SBR from each forest type and the capture efficiency of each trap night.

Trap	Forest _	Amp	hibians	Re	eptiles	7	Гotal
Туре	Type	Count	Efficiency	Count	Efficiency	Count	Efficiency
Турс	Турс	Count	(%)	Count	(%)	Count	(%)
Funnel	DDF	46	8.0	64	_11.1	110	19.1
	HDF	242	44.8	55	10.2	297	55.0
	PLE	202	43.2	99	21.2	301	64.3
То	otal	490	30.9	218	13.8	708	44.7
Pitfall	DDF	11/6	1a7.6na	16a	4.2	17	11.8
	HDF	69	51.1	7	5.2	76	56.3
	PLE	49	43.0	11	9.6	60	52.6
То	otal	129	32.8%	24	6.1%	153	38.9%
Grand To	otal	619	31.3	242	12.2	861	43.6

4.1.5 Abundances

Observed abundances varied across the three forest types under investigation for both amphibians and reptiles (Figure 4.2). Total herpetofaunal

abundances did not significantly vary by forest type (F-value = 3.146, df = 2, p-value = 0.0982). Additionally, complete amphibian abundances did not differ between forest types (F - value = 2.352, df = 2, p-value = 0.157), while reptile abundances did show a significant difference (χ^2 = 6.0829, df = 2, p-value = 0.0478). Pairwise comparison results found that only HDF and PLE reptile abundances were significantly different (W = 0, p-value = 0.04975). 2.352 0.157

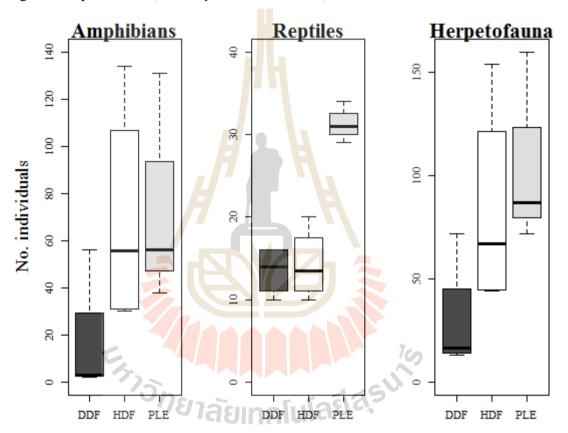


Figure 4.2 A comparison of the observed abundances in each forest type for amphibians, reptiles, and all herpetofauna.

Reptile abundances differed significantly between forest types in the dry season ($\chi^2 = 6.385$, df = 2, p-value = 0.0411) and in the wet season ($\chi^2 = 6.409$, df = 2, p-value = 0.0406). Pairwise post hoc analysis revealed that over the dry season the dry dipterocarp did not differ significantly from the heterogeneous disturbed forest

(W = 6, p-value = 0.653), but both DDF and HDF significantly differed from eucalyptus plantation (W = 0, p-value = 0.0477 and W = 12, p-value = 0.0498 respectively). The pairwise comparison for reptile abundances in the wet season revealed that only DDF and PLE were significantly different (W = 0, p-value = 0.0477), as HDF abundances did not differ significantly from DDF (W = 8, p-value = 0.559) nor PLE (W = 0, p-value = 0.0722).

Amphibian abundances in the dry season varied significantly between forest types ($\chi^2 = 7.616$, df = 2, p-value = 0.0222), but did not differ significantly in the wet season ($\chi^2 = 2.917$, df = 2, p-value = 0.233). Dry season amphibian abundances were different between the DDF and both HDF and PLE (W = 0, p-value = 0.0265 and W = 0, p-value = 0.0436 respectively); however abundances were not different between HDF sites and PLE sites (W = 9, p-value = 0.4).

The Whittaker plots generated for each forest type indicated that reptiles show high unevenness, with little difference in evenness between forest types (Figure 4.3). The two most abundant reptile species, *D. siamensis* and *E. macularia*, remained the same between forest types; however while in the HDF and PLE the *E. macularia* was the most common species the *D. siamensis* was more abundant in the DDF. The third most common species (or species complex) in both the PLE and HDF, *Hemidactlyus* spp., did not occur at all within the DDF.

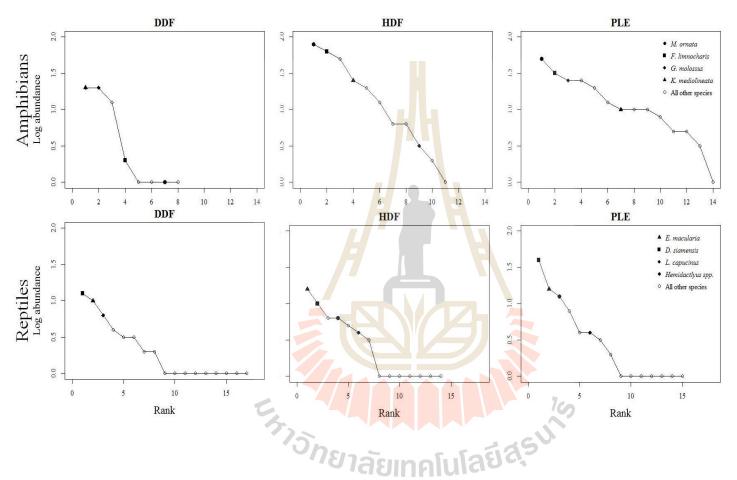


Figure 4.3 Whittaker plots for complete amphibian captures (top row) and reptiles (bottom row) for DDF, PLE, and HDF

Amphibians in contrast show variation in the between DDF and both HDF and PLE. The two most abundant amphibian species, *M. fissipes* and *F. limnocharis*, in the PLE and HDF sites were identical, but two completely different species, *K. mediolineata* and *G.molossus*, dominated the DDF. Amphibians rank abundance changed seasonally, with the *F. limnocharis* moving from the top spot during the dry season to 7th in the wet season, while reptile rank abundances did not drastically vary between the two observed seasons (Figure 4.4).

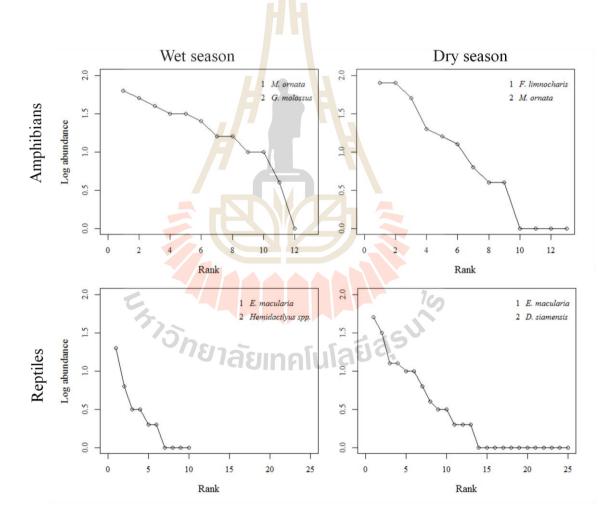


Figure 4.4 Seasonal rank abundance curves from reptile and amphibians species from all forest types combined.

Plotting amphibian abundance by different landscape factors did not reveal any potential correlations with the possible exception of distance to water (Figure 4.5). Reptile abundance did not show any potential trends when plotted by the same factors (Appendix 9)

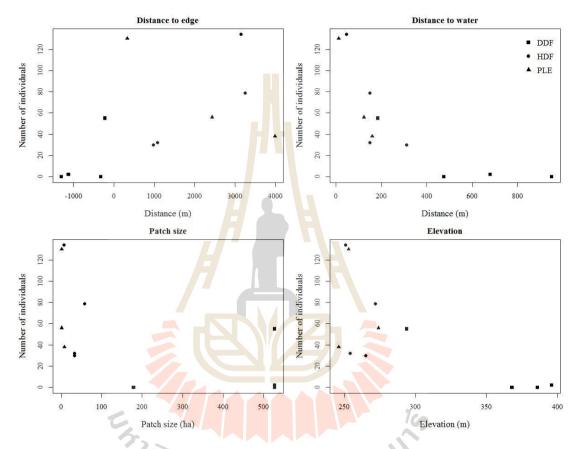


Figure 4.5 Plot of amphibian abundance by distance to the edge of SBR, distance to water, patch size, and elevation.

4.1.6 Species richness

Sampling completeness varied widely between sites for both amphibians and reptiles meaning that within forest types there was high variation between sites (Appendix 4). As DDF amphibian captures were primarily limited to a single site (DDF-4), the estimated richness was actually lower than the observed richness. The lower estimated richness occurred because we used sampling with replacement to

accurately determine confidence intervals. Additionally, sampling completeness also varied by season with more reptile species observed in May and June than in September. The trend shifted for amphibians in the DDF, with almost all species only observed in September.

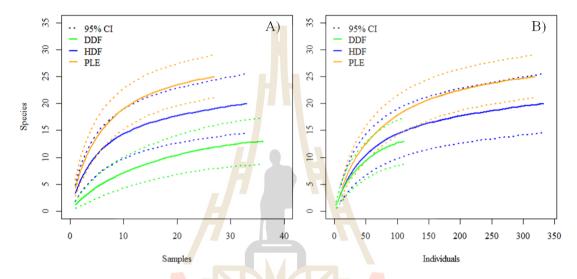


Figure 4.6 Sample and individual based rarefaction for amphibians (A and B) and for reptiles (C and D).

The extrapolated rarefaction curves for each forest type showed a significant difference in total herpetofauna species between PLE and DDF; however, HDF fell within the 95% confidence interval for both of the other two forest types (Figure 4.6). However, when correcting for the number of individuals using individual based rarefaction the significant differences were no longer apparent.

When comparing reptile species richness between forest types, no significant differences appeared; however the curves never reached an asymptote. For amphibian species richness the PLE showed significantly higher richness than the HDF, even when controlling for differences in the number of captured individuals (Figure 4.7).

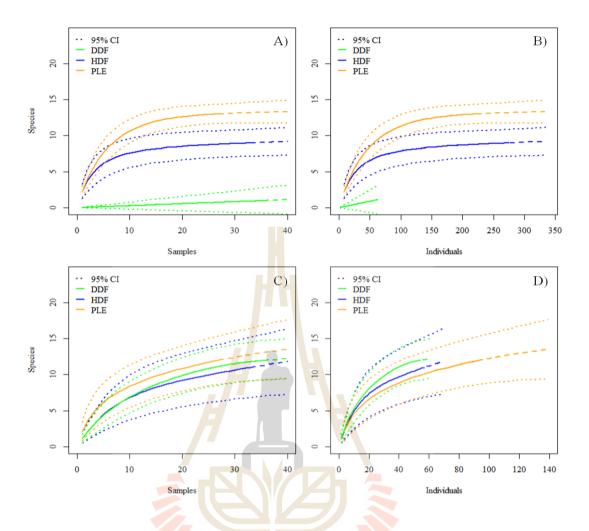


Figure 4.7 Sample and individual based rarefaction for amphibians (A and B) and for reptiles (C and D).

4.1.7 Diversity

Diversity based on the Shannon-Weiner index differed significantly between forest types when considering all herpetofauna ($\chi^2 = 6.4091$, df = 2, p-value = 0.04058) and amphibians ($\chi^2 = 8.2648$, df = 2, p-value = 0.01604); however reptile diversity did not significantly vary between forest types ($\chi^2 = 1.5455$, df = 2, p-value = 0.4618) (Table 4.7). Pairwise comparison using Wilcoxon's rank sum test for all herpetofauna revealed that no significant differences existed between forests at the 95% confidence interval. When comparing amphibian diversity only DDF and PLE

were significantly different (W = 0, p-value = 0.04975). Comparing the Simpson index between forest types showed only a significant difference when using all herpetofauna ($\chi^2 = 6.1818$, df = 2, and p-value = 0.04546).

Table 4.7 The calculated Shannon-Weiner and Simpson diversity indexes for each plot.

	Amph	ibians	F	Reptiles	7	Total
_	Shannon-		Shannon-		Shannon-	
Plot	Weiner	Simpson	Weiner	Simpson	Weiner	Simpson
DDF - 1	0.00	1.00	1.72	0.78	1.72	0.78
DDF - 2	0.00	1.00	1.82	0.82	1.82	0.82
DDF - 3	0.69	0.50	1.42	0.72	1.75	0.79
DDF - 4	1.43	0.71	1.57	0.73	2.00	0.81
HDF - 1	1.66	0.80	1.63	0.78	2.24	0.88
HDF - 2	1.34	0.70	1.50	0.76	1.71	0.76
HDF - 3	1.47	0.67	2.08	0.86	2.31	0.84
HDF - 4	1.59	0.75	0.75	0.45	1.87	0.80
PLE - 1	1.89	0.79	1.79	0.78	2.53	0.89
PLE - 2	2.18	0.86	1.73	0.74	2.57	0.90
PLE - 3	1.91	0.82	1.74	0.76	2.50	0.89

4.1.8 Community structure

The Mantel test results showed that amphibian and total herpetofauna assemblages were spatially auto-correlated by plot (R = 0.5307, p-value = 0.008 and R = 0.4728, p-value = 0.002 respectively). Reptile communities however were not spatially auto-correlated (R = 0.1744, p-value = 0.093). Spatial autocorrelation of amphibian communities could either indicate that the sampling sites were set to close together. To account for spatial autocorrelation, I created a new categorical variable called distance group. Distance groups were based on geographic distance between sites divided into 3 groups: far, edge, and forest.

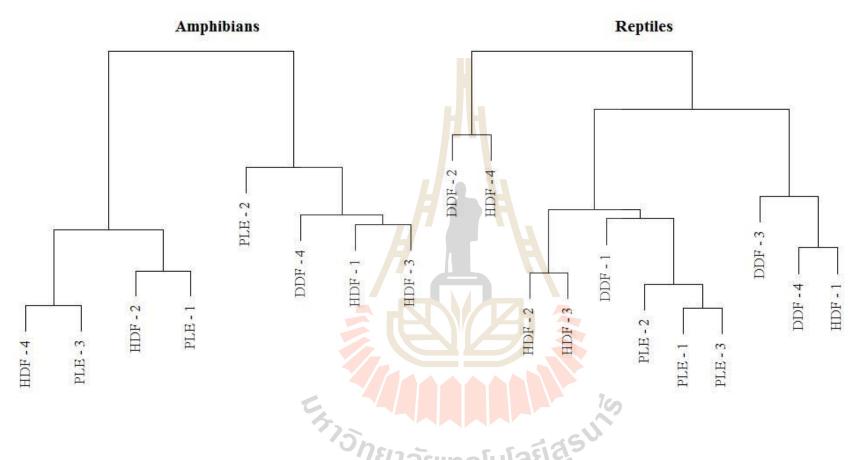


Figure 4.8 Comparison of hierarchic clustering of samples sites based on the recorded abundances of reptiles (right) and amphibians (left).

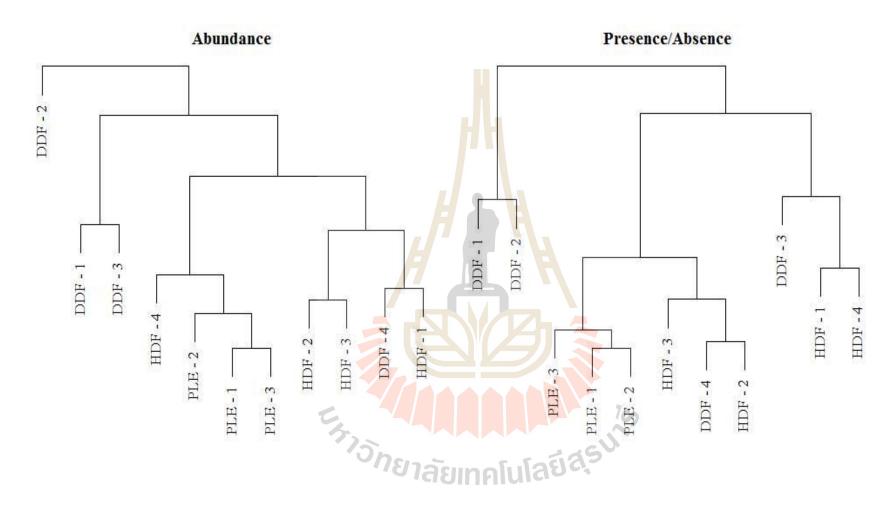


Figure 4.9 Comparison of hierarchic clustering between abundance and presence absence data for all captured herpetofauna

Hierarchic clustering based on complete linkages showed that sites did not cluster based solely by forest type, and different patterns emerged when considering amphibians and reptiles separately (Figure 4.8). Additionally, clusters changed when using the abundance data and presence-absence data (Figure 4.9), which means that species abundances varied between sites and not just presence.

The results of NMDS on the abundance data did not show clustering of sites based on forest type. For both total herpetofaunal and reptile assemblages only the PLE sites showed a pattern of clustering on the NMDS. Since several sites had no recorded amphibian captures, only the PLE and HDF sites were used in the NMDS. Sites showed no clustering based on amphibian captures. The stress values for each NMDS plot all fell within the generally accepted range for good representation, meaning that the NMDS plots accurately displayed the resulting ordination in two dimensions.

The environmental fitting for each NMDS found that elevation, distance to water, and patch size were significant variables when analyzing all herpetofauna. Distance to SBR was the only significant variable for amphibians, while no variables fit for reptiles (Appendix 5). Only the environmental fitting for all herpetofauna captures showed that forest type was a significant factor in the NMDS, in addition to canopy cover (Appendix 6).

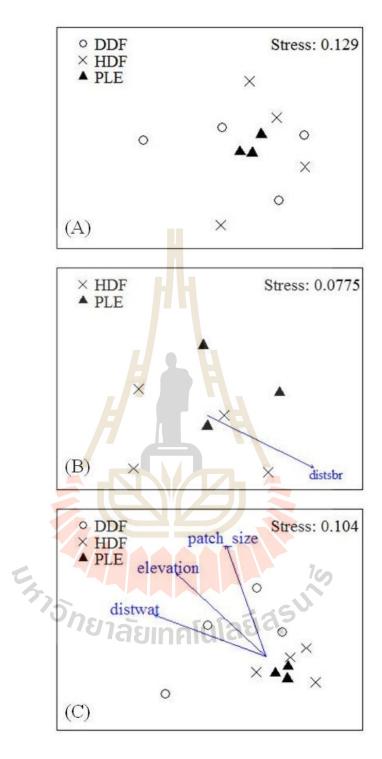


Figure 4.10 Non-metric multidimensional scaling (NMDS) for sites based on observed A) amphibians, B) reptiles, and C) herpetofauna with significant environmental variables.

Analysis of the beta dispersion for total, reptile, and amphibian captures, showed that only the total captures were significant different between forest types (F = 5.5083 and *p*-value = 0.03132). The results from the PERMANOVA found no significant difference between forest types, and no significance for any of the environmental variables used in the model with the exception of amphibians and distance grouping (Table 4.8). Additionally, the PERMANOVA for the presence absence data did not find any significant differences (Appendix 8).



Table 4.8 PERMANOVA results modeling reptile, amphibian, and total captures based on forest type and other identified environmental factors.

	Variable	Df	Sum of Squares	Mean Squares	F- Model	R	<i>p</i> -value
Total	Distance group	2	0.99	0.50	2.27	0.44	0.11
	Forest type	2	0.20	0.10	0.46	0.09	0.91
	Distance to water	1	0.22	0.22	1.02	0.10	0.46
	Distance to SBR	1	0.06	0.06	0.25	0.02	0.96
	Canopy cover	1	0.28	0.28	1.26	0.12	0.37
	Elevation	1	0.21	0.21	0.96	0.09	0.55
	Patch size	1-	0.08	0.08	0.36	0.04	0.92
	Residuals	1	0.22	0.22	0.10		
	Total	10	2.26	1.00			
Reptile	Distance group	2	0.38	0.19	0.82	0.19	0.65
	Forest type	2	0.20	0.10	0.43	0.10	0.95
	Residuals	6	1.40	0.23	0.71		
	Total	10	1.99	1.00			
Amphibian	Distance group	1	0.36	0.36	4.91	0.40	0.02
	Forest type	1	0.15	0.15	2.12	0.17	0.17
	Distance to water	าลุยเ	0.13	0.13	1.80	0.15	0.22
	Distance to SBR	1	0.07	0.07	0.92	0.08	0.55
	Litter depth	1	0.10	0.10	1.43	0.12	0.32
	Residuals	1	0.07	0.07	0.08		
	Total	6	0.89	1.00			

The indicator species analysis based on abundance data identified five species as significantly contributing to forest type differences (Table 4.9). When using

only species presence absence, we found that only two species, *Bungarus canditus* and *Callouela guttalata*, were significant indicators. For all indicator species using both methods the test found them as indicators for the PLE forests.

Table 4.9 Significant results from indicator species analysis using abundance and presence absence community data.

Data	Species	F	orest type	Indica valu	n-value
Abundance	Callouela gutta <mark>lat</mark> a		PLE	0.93	3 0.029
	Bungarus can <mark>ditus</mark>		PLE	0.80	0.041
	Hemidactlyus spp.		PLE	0.73	3 0.035
	Micryletta in <mark>o</mark> rnata		PLE	0.6	7 0.039
	Eutropis macularia		PLE	0.6	7 0.042
Presence/Absence	Callouela <mark>gu</mark> ttalata		PLE	0.80	0.027
	Bunga <mark>rus</mark> c <mark>a</mark> nditus		PLE	0.80	0.028
	Micry <mark>lett</mark> a inornata		PLE	0.6	7 0.050

4.1.9 Analysis of biometrics

From the 861 total captures we collected biometrics on 86 individuals, including 40 snakes and 48 individuals from the five species *D. siamensis*, *E. macularia*, *K. pulchra*, *K. mediolineata*, and *L. reevesii rubritaeniata*. We selected the three most abundant reptiles to create a BCI from the residuals of an ordinary least squares regression, *D. siamensis* (n = 15), *E. macularia* (n = 14) and *Lycodon capucinus* (n = 10). The SVL measurements for the *D. siamensis* was not normal (W = 0.5984, *p*-value = 2.463e-05) and thus violated the assumptions for an ordinary least squares regression; however *E. macularia* and *L. capucinus* measurements fit the assumptions and were used to create a BCI. The results of an ANOVA on BCI to forest type found no significant difference for either *E. macularia* or *L. capuncinus* (Table 4.10).

Table 4.10 Results from the ANOVA of BCI by forest type for *E. macularia* and *L. capucinus*.

Species		Df	Sum Sq	Mean Sq	F-value	<i>p</i> -value
E. macularia	forest	2	0.6715	0.3358	1.344	0.301
	Residuals	11	2.7487	0.2499		
L. capucinus	forest	2	53.18	26.59	0.868	0.461
	Residuals	7	214.49	30.64		

We also analyzed whether forest type influenced fitness by assessing whether reptile individuals had lost their tails. The chi-square analysis for all reptiles showed that forest type did not influence tail loss ($\chi^2 = 1.5353$, df = 2, p-value = 0.4641). Looking at the two most abundant species. D. siamensis and E. macularia, also show that forest type and tail loss were independent ($\chi^2 = 5.2724$, df = 2, p-value = 0.07163 and $\chi^2 = 0.2171$, df = 2, p-value = 0.8971 respectively).

As snout-to-vent length and mass can vary widely between species, we did not conduct any tests on these metrics (Table 4.11 and Appendix C). Form the 40 snakes processed, 30 were male compared to only 6 females, with 4 individuals too young to safely determine sex. The high male to female capture ratio pattern was observed across all forest types.

Table 4.11 Processing information for all snakes captured during the study.

Forest type	Individuals	Species	Male:Female	Max SVL	Min SVL	Avg SVL
DDF	18	11	14:3	1266	84	445.5
HDF	8	4	7:1	1036	232	433.0
PLE	14	10	9:2	1070	144	415.2

4.2 Discussion

4.2.1 Abundance, species richness, and diversity

A single plot was responsible for over 90% of the amphibian captures in the DDF, which may be confounding the results for both the total abundance comparison and when comparing abundances specifically in the wet season. The low abundance of amphibians in the DDF, agree with similar results from several other studies such as Vonesh (2001) in which amphibian abundance was higher in disturbed forests and pine plantations than in primary forest. Additionally, Gardner et al. (2007b) found that amphibian abundances were actually highest in 4-5 year Eucalyptus plantations compared to primary and secondary forests in the Neotropics. However, several other studies arrived at different conclusions showing that plantations do not necessarily house higher numbers of amphibians (Folt and Reider, 2013; Kurz et al., 2014). However, open habitats, i.e. pastures, appear to show the greatest negative impact on amphibian abundance and diversity, which were not investigate in this study (Urbina-Cardona et al., 2006; Kurz et al., 2014). The species that were found in the DDF were also primarily burrowing frogs (G. molossus and K. mediolineata). As burrowing frogs, they spend a large portion of their time underground and potentially face less risk of desiccation in dry habitats such as the DDF (Suazo-Ortuño et al., 2008). All three of the species G. molossus, K. pulchra, and K. mediolineata abundances showed that they peaked in sites at the edge of the protected area and dropped in sites farther away from the edge. In contrast the most abundant amphibian species found in the HDF and PLE sites, F. limnocharis and M. fissipes, displayed the opposite trend.

The similarity in reptile species richness across all forest types supports the hypothesis that reptiles are not as sensitive to fragmentation as other taxa, and that

some species even thrive in partially disturbed habitats (Wanger et al., 2010). Canopy heterogeneity provides reptiles with a plethora of basking sites, and some studies suggests that at least insect prey communities are positive impacted in disturbed habitats (Heliölä et al., 2001; Suazo-Ortuño et al., 2008). However, our results also show that the slopes of the individual based rarefaction for reptiles did not reach an asymptote and that both the DDF and HDF had higher species richness when accounting for the number of individuals. More samples could provide a better comparison between forest types as the accumulation curve levels off.

The lack of significant difference in pairwise comparison of diversity between forest types potentially derives from an issue in sample size. The variation between sites in the same forest type was high which could be overcome with a larger sample size; however passive trapping arrays are time intensive and expensive to create.

The results from the MCA of habitat variables does not necessarily reflect that all forest types comprised similar habitats. One potential issue was the correlation between variables which violates the assumptions of MCA. Additionally since we the collected the data as categorical values, small variations between forest types may have gone unnoticed.

4.2.2 Community structure

The drastic difference in amphibians between the majority of the DDF sites and the other sites created separation when comparing the total herpetofaunal community. The lack of any significant environmental variables for reptiles suggests that other factors are contributing to the difference or that no true differences exist on the limited spatial and/or temporal scale. However, the NMDS shows that the PLE

sites formed a tight grouping, but may overlap with the other forest types in a way that obscures the clustering.

Clustering for amphibians appears to occur based on the spatial distance between sites. As observed by Kurz et al. (2014) the habitat type along the edge of a forest did not influence herpetofaunal communities with the exception of forest to pasture. The sites near the edge, may house more species as individuals can still benefit from the primary forest. Further study in the area is needed to isolate the potential edge effect and the effective distance that supports amphibian communities.

4.2.3 Biometrics

While tail loss did not show a significant result at the 95% it did at the 90% indicating that a larger sample size may reveal a true significant difference. Sung et al. results support that reptiles in disturbed habitats may face increased predation pressure as evidenced through tail loss. Additionally comparing a BCI for the most abundant species does not necessarily reflect how habitat influences body condition as highly abundant species are typically generalists and are not as sensitive to habitat degradation. An assessment of rare species would require a longer time scale to achieve an acceptable sample size, but could provide contrasting information.

CHAPTER V

CONCLUSION

As a preliminary study on the effects disturbance on herpetofauna, this study provides useful results for establishing continued study in the Sakaerat Biosphere Reserve. The high mortality rates throughout the dry season suggest that sampling with passive trapping may not be ethical in all seasons. Disturbed habitats that experience high temperatures and low humidity may require different sampling procedures, such as only opening traps at night.

Our results also confirmed the assertion that reptiles and amphibians do not show the same patterns with regard to disturbance. Abundance, species richness, and diversity comparisons clearly showed that these two groups should not be combined when considering their response to an environmental gradient. Reptiles appeared less sensitive to habitat differences, while amphibians showed a high degree of difference.

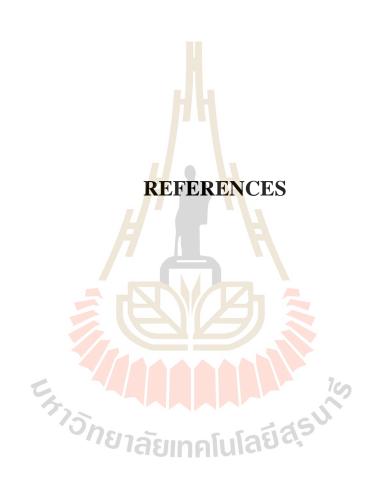
The study also documented that amphibian abundance and richness was very low in the protected forest, which contrasts with findings in other studies. More research is required to determine what environmental factors may influence the low number of amphibians in the DDF, and whether seasonal shifts occur.

The biometrics of several abundant species did not reveal any significant differences between forest types. However, studying the most abundant species does not necessarily provide a clear indicator for habitat quality. Species that are abundant across all forest types, are generalists and should not show any evidence in reduced

body condition along a disturbance gradient. Future studies should investigate body condition for species that are present, but not necessarily common in all forest types to assess the impact of disturbance.

Our study did not find any significant differences in community structure between forest types using ordination (non-metric multidimensional scaling) nor with PERMANOVA. Additionally, continued sampling over several years and season would improve detection of rare species. Addressing both the temporal and spatial aspects will aid in more accurate ordination and multivariate hypothesis testing.

While some of the results suggest that there are significant differences between the studied forest types, the study did not provide enough conclusive evidence to completely explain how herpetofauna communities are effected by habitat disturbance in the SBR. One major issue that we encountered was that the forest type categories we created do not necessarily work as a disturbance gradient. We suggest that future studies focus on a single forest type and identify different factors, such as patch size, management activities, and distance to water, that influence herpetofauna communities.



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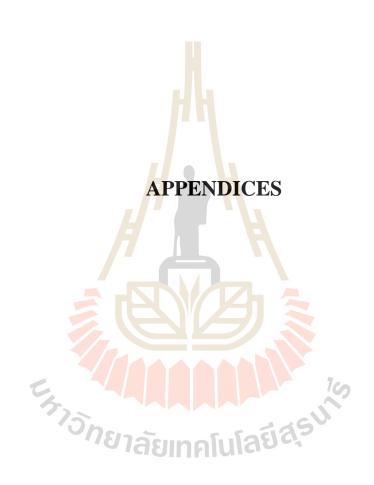
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APPENDIX A SITE HABITAT AND ENVIRONMENTAL DATA





Appendix A-1 Photographs of each dry dipterocarp forest site.





Appendix A-2 Photographs of each highly disturbed forest site.





Appendix A-3 Photographs of each eucalyptus plantation site.



Appendix A-4 Daily weather conditions for the Sakaerat Biosphere Reserve in May, June, and September of 2015 for the all days sampled including the day traps were opened.

	Max	Min	Average	Relative	ъ.
Date	temperature	temperature	temperature	Humidity	Rain
	(°C)	(°C)	(°C)	(%)	(mm)
15/05/2015	35.25	24.75	25.35	75	0
16/05/2015	36	25.35	25.95	74.5	0
17/05/2015	35.6	26.05	26.65	73	0
18/05/2015	36.55	25.3	25.9	74	0
25/05/2015	35.75	26.4	27	72	0
26/05/2015	35.75	25.95	26.55	72.5	0
27/05/2015	35.55	25.8	26.4	75	0
28/05/2015	36.8	25.8	26.4	73.5	0
17/06/2015	31.35	23.5	24.1	83	0
18/06/2015	31.4	23.85	24.45	80	0
19/06/2015	33.75	24.1	24.7	77	0
20/06/2015	34.65	24.05	24.65	74.5	0
23/06/2015	30.8	25.05	25.65	76.5	2.35
24/06/2015	31.8	24.8	25.4	78.5	0
25/06/2015	32.05	25.35	25.95	75	0
26/06/2015	29.45	25.05	25.65	74	0
17/09/2015	25.8	38 23.27 U	23.8	92.5	38.45
18/09/2015	26.65	22.65	23.25	94	1.65
19/09/2015	31.2	22.25	22.85	93	0
20/09/2015	32.75	22.45	23.05	91	0
21/09/2015	32.8	23.6	24.2	88	28.55
22/09/2015	30.4	22.7	23.3	91.5	52.4
23/09/2015	30.15	22.6	23.2	91.5	0
24/09/2015	30.8	23.65	24.25	86	0

Appendix A-4 Monthly weather conditions for May, June, and September at the Sakaerat Environmental Research Station.

Month _		Temperature (°C)			Rain
	Max	Min	Average	Humidity (%)	(mm)
May	36.0	25.8	26.4	74.8	8.6
June	34.2	25.0	25.6	75.7	79.3
September	30.9	23.4	24.0	84.3	259.9

Appendix A-5 Ground cover assessment for each site.

	Vegetative ground	• 1	
Plot	cover	Bare ground coverage	Leaf litter coverage
DDF-1	Light (16-25%)	Light (16-25%)	Medium (25-65%)
DDF-2	Medium (25-65%)	Medium (25-65%)	Very light (1-15%)
DDF-3	Medium (25-65%)	Very light (1-15%)	Medium (25-65%)
DDF-4	Light (16-25%)	Medium (25-65%)	Very light (1-15%)
HDF-1	Medium (25-65%)	Very light (1-15%)	Very light (1-15%)
HDF-2	Medium (25-65%)	Medium (25-65%)	Very light (1-15%)
HDF-3	Medium (25-65%)	Medium (25-65%)	None (0%)
HDF-4	Medium (25-65%)	Light (16-25%)	Very light (1-15%)
PLE-1	Medium (25-65%)	Light (16-25%)	Very light (1-15%)
PLE-2	Light (16-25%)	Medium (25-65%)	Medium (25-65%)
PLE-3	Medium (25-65%)	Medium (25-65%)	Very light (1-15%)

Appendix 10 Canopy and vegetation factors for each plot site.

Plot	Canopy cover	Groundstory density	Understory density	Midstory density	Abovestory density
DDF-1	Heavy (66-80%)	Medium (25-65%)	Very light (1-15%)	Very light (1-15%)	Medium (25-65%)
DDF-2	Heavy (66-80%)	Heavy (66-80%)	None (0%)	Very light (1-15%)	Medium (25-65%)
DDF-3	Medium (25-65%)	Heavy (66-80%)	Very light (1-15%)	Very light (1-15%)	Light (16-25%)
DDF-4	Medium (25-65%)	Light (16-25%)	Very light (1-15%)	Very light (1-15%)	Very light (1-15%)
HDF-1	Medium (25-65%)	Heavy (66-80%)	Very light (1-15%)	Very light (1-15%)	Very light (1-15%)
HDF-2	Medium (25-65%)	Light (16-25%)	Light (16-25%)	Medium (25-65%)	Light (16-25%)
HDF-3	Very light (1-15%)	Light (16-25%)	Very light (1-15%)	None (0%)	None (0%)
HDF-4	Medium (25-65%)	Medium (25-65%)	Very light (1-15%)	Medium (25-65%)	Very light (1-15%)
PLE-1	Very light (1-15%)	Medium (25-65%)	Very light (1-15%)	Very light (1-15%)	Very light (1-15%)
PLE-2	Very light (1-15%)	Medium (25-65%)	Light (16-25%)	Very light (1-15%)	None (0%)
PLE-3	Light (16-25%)	Very light (1-15%)	Very light (1-15%)	Very light (1-15%)	Very light (1-15%)

APPENDIX B RICHNESS, ABUNDANCE, AND COMMUNITY DATA



Appendix B-1 Sampling completeness for trapping site including observed richness and estimated richness using Chao1 method.

Family	Forest type	Observed richness	Chao1 richness	Completeness
Amphibians	DDF - 1	0	7.73	0.0%
	DDF - 2	0	7.73	0.0%
	DDF - 3	2	7.73	25.9%
	DDF - 4	8	7.73	103.5%
	HDF - 1	6	10.09	59.5%
	HDF - 2	5	10.09	49.6%
	HDF - 3	7	10.09	69.4%
	HDF - 4	9	10.09	89.2%
	PLE - 1	10	13.49	74.1%
	PLE - 2	13	13.49	96.4%
	PLE - 3	9	13.49	66.7%
Reptiles	DDF - 1	7	16.74	41.8%
	DDF - 2	7	16.74	41.8%
	DDF - 3	5	16.74	29.9%
	DDF - 4		16.74	41.8%
	HDF - 1	6	12.63	47.5%
	HDF - 2	5	12.63	39.6%
7	HDF - 3	9	12.63	71.3%
	HDF - 4	3	12.63	23.8%
	PLE - 1	ลัยเทอโนโล	14.3	62.9%
	PLE - 2	9	14.3	62.9%
	PLE - 3	9	14.3	62.9%

Appendix B-2 Monthly abundance, species, and reptile tail loss at each sample site.

Month	Plot	Amphil	oians	Repti	les	Reptile
MOIIII	FIOL	Abundance	Species	Abundance	Species	tail loss
May	DDF - 1	0	0	4	4	0
-	DDF - 2	0	1	4	3	0
	DDF - 3	0	0	3	2	0
	DDF - 4	0	0	5	3	0
	HDF - 1	5	3	7	4	0
	HDF - 2	3	3	4	2	0
	HDF - 3	0	0	6	4	0
	HDF - 4	55	7	11	3	1
	PLE - 1	6	5	13	6	0
	PLE - 2	0	1	7	4	0
	PLE - 3	4	3	10	5	2
	PLE - 4	0	0	6	4	0
Tot	al	73	11	80	21	3
June	DDF - 1	0	0	9	4	1
	DDF - 2	0	0	7	6	0
	DDF - 3	1	1	6	4	0
	DDF - 4	0	1	6	2	0
	HDF - 1	13	3	4	4	0
	HDF - 2	45	5	5	4	0
	HDF - 3	10	4	7	5	0
	HDF - 4	79	8	9	2	0
	PLE - 1	9	4	10	5	0
	PLE - 2	7	4	14	6	0
	PLE - 3	29	6	13	6	0
Tot	al	193	11	90	17	1
September	DDF - 1	0	0	3	3	2
	DDF - 2	0	0.0	13.31	1	0
	DDF - 3	a iahili	Hulia	1	1	1
	DDF - 4	55	8	5	4	0
	HDF - 1	14	4	1	1	0
	HDF - 2	31	4	1	1	0
	HDF - 3	20	6	2	2	0
	PLE - 1	23	7	11	4	0
	PLE - 2	123	11	8	3	2
	PLE - 3	23	6	8	3	1
Tot	al	290	12	41	10	6
Grand Total		556	14	211	27	10

Appendix B-3 Rank abundance for amphibian species captured in each forest type.

Forest type	Species	Rank	Abundance	Proportion	Log(Abund)
DDF	Kaloula mediolineata	1	20	35.1	1.3
	Glyphoglossus molossus	2	18	31.6	1.3
	Kaloula pulchra	3	13	22.8	1.1
	Fejervarya limnocharis	4	2	3.5	0.3
	Microhyla butleri	5	1	1.8	0
	Microhyla heymonsi	6	1	1.8	0
	Microhyla fissipes	7	1	1.8	0
	Microhyla pulchra	8	1	1.8	0
HDF	Microhyla fissipes	1	78	28.4	1.9
	Fejervarya limnocharis	2	66	24	1.8
	Microhyla pulchra	3	52	18.9	1.7
	Kaloula mediolineata	4	27	9.8	1.4
	Microhyla heymonsi	5	20	7.3	1.3
	Microhyla butleri	6	13	4.7	1.1
	Duttaphrynus melan <mark>osti</mark> ctus	7	7	2.5	0.8
	Kaloula pulchra	8	6	2.2	0.8
	Glyphoglossus mo <mark>l</mark> os <mark>s</mark> us	9	3	1.1	0.5
	Hylarana macro <mark>d</mark> actyla	10	2	0.7	0.3
	Calluella gutt <mark>ula</mark> ta	11	1	0.4	0
PLE	Microhyla f <mark>issip</mark> es	1	51	22.8	1.7
	Fejervarya <mark>limno</mark> charis	2	31	13.8	1.5
	Glyphoglossus molossus	3	28	12.5	1.4
	Microhyla heymonsi	4	28	12.5	1.4
	Microhyla butleri	5	19	8.5	1.3
	Kaloula pulchra	6	13	5.8	1.1
	Kaloula mediolineata	7	-5119	4.9	1
	Micryletta inornata		3011	4.9	1
	Calluella guttulata	9	10	4.5	1
	Microhyla pulchra	10	8	3.6	0.9
	Duttaphrynus melanostictus	11	5	2.2	0.7
	Occidozyga lima	12	5	2.2	0.7
	Hylarana macrodactyla	13	3	1.3	0.5
	Hylarana erythraea	14	1	0.4	0

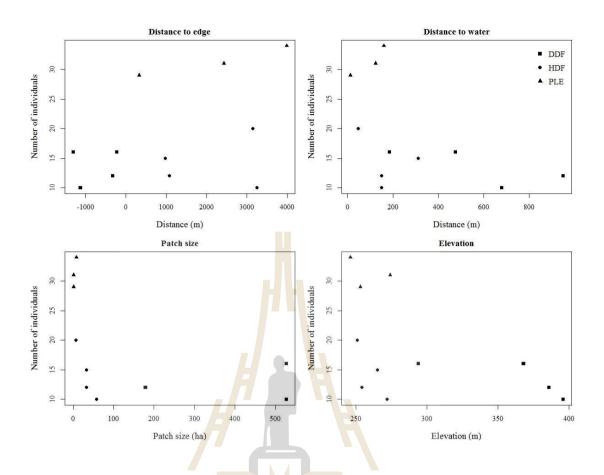
AppendixB-4 Rank abundance for reptile species captured in each forest typ.e

Forest type	Species	Rank	Abundance	Proportion	Log(Abun)
DDF	Dixoneus siamensis	1	14	25.9	1.1
	Eutropis macularia	2	10	18.5	1
	Lycodon capucinus	3	7	13	0.8
	Gehyra lacerata	4	4	7.4	0.6
	Calotes versicolor	5	3	5.6	0.5
	Lygosoma bowringii	6	3	5.6	0.5
	Boiga multomaculata	7	2	3.7	0.3
	Oligodon pseudotaeniatus	8	2	3.7	0.3
	Boiga siamensis	9	1	1.9	0
	Calliophis maculiceps	10	1	1.9	0
	Chrysopelea ornata	11	1	1.9	0
	Coelognatus radiatus	12	1	1.9	0
	Leiolepis reevesii	13	1	1.9	0
	Naja siamensis	14	1	1.9	0
	Oligodon faciolatus	15	1	1.9	0
	Psammodynastes	16	1	1.9	0
	pulverulentus				
	Ramphotyphlops <mark>b</mark> raminus	17	1	1.9	0
HDF	Eutropis macularia	1	16	28.1	1.2
	Dixoneus siamensis	2	9	15.8	1
	Lygosoma b <mark>owri</mark> ngii	3	7	12.3	0.8
	Hemidactylus spp	4	6	10.5	0.8
	Leiolepis reev <mark>esii</mark>	5	5	8.8	0.7
	Lycodon capucinus	6	4 7	5 7	0.6
	Gehyra lacerata	7	3	5.3	0.5
	Boiga multomaculata	8	3,45	1.8	0
	Bungarus candidus		384	1.8	0
	Lycodon laoensis	10	1	1.8	0
	Naja siamensis	11	1	1.8	0
	Oligodon pseudotaeniatus	12	1	1.8	0
	Ramphotyphlops albiceps	13	1	1.8	0
	Ramphotyphlops braminus	14	1	1.8	0

Appendix B-4 (Continued)

Forest type	Species	Rank	Abundance	Proportion	Log(Abun)
PLE	Eutropis macularia	1	39	41.5	1.6
	Dixoneus siamensis	2	15	16	1.2
	Hemidactylus spp.	3	12	12.8	1.1
	Leiolepis reevesii	4	8	8.5	0.9
	Lygosoma bowringii	5	4	4.3	0.6
	Lycodon capucinus	6	4	4.3	0.6
	Bungarus candidus	7	3	3.2	0.5
	Gehyra lacerata	8	2	2.1	0.3
	Calloselasma rhodostoma	9	1	1.1	0
	Calotes versicolor	10	1	1.1	0
	Dendrelaphis subocularis	11	1	1.1	0
	Enhydris plumbea	12	1	1.1	0
	Oligodon faciolatus	13	1	1.1	0
	Oligodon taeniatus	14	1	1.1	0
	Rhabdophis chrysargus	15	1	1.1	0





Appendix B-5 Plot of reptile abundance by distance to the edge of SBR, distance to water, patch size, and elevation.

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Appendix B-6 Results from environmental fitting on NMDS of abundance community data for continuous variables.

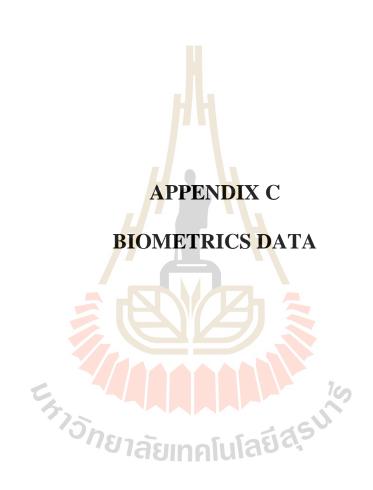
Community analyza	Environmental variable	NMDS1	NMDS2	R^2	<i>p</i> -value
Reptiles	Groundstory	-0.294	-0.956	0.240	0.334
•	Canopy cover	-0.794	-0.608	0.362	0.160
	Grass	0.201	-0.980	0.100	0.676
	Litter depth	-0.349	-0.937	0.117	0.628
	Elevation	-0.975	-0.220	0.159	0.503
	Distance to water	-0.998	0.065	0.310	0.223
	Distance to SBR edge	0.613	-0.790	0.032	0.889
	Patch size	0.911	-0.412	0.016	0.959
	Slope	-0.973	0.231	0.269	0.305
	Aspect	1.000	-0.022	0.013	0.946
Amphibians	Groundstory	0.721	0.693	0.401	0.350
	Canopy cover	0.702	-0.712	0.153	0.714
	Grass	0.515	-0.857	0.334	0.460
	Litter depth	0.432	0.902	0.753	0.064
	Elevation	-0.109	-0.994	0.494	0.291
	Distance to water	-0.401	-0.916	0.606	0.145
	Distance to SBR edge	0.900	-0.437	0.759	0.043
	Patch size	-0.089	-0.996	0.447	0.313
	Slope	0.555	-0.832	0.100	0.820
	Aspect	0.468	-0.884	0.526	0.225
Total	Groundstory	-0.801	-0.599	0.009	0.968
3	Canopy cover	-0.969	0.246	0.488	0.078
	Grass	0.694	-0.720	0.197	0.458
	Litter depth	-0.273	0.962	0.105	0.669
	Elevation	-0.837	0.548	0.861	0.003
	Distance to water	-0.984	0.179	0.851	0.005
	Distance to SBR edge	0.631	-0.776	0.512	0.051
	Patch size	-0.428	0.904	0.768	0.007
	Slope	-0.936	-0.353	0.492	0.110
	Aspect	0.480	-0.877	0.431	0.123

Appendix B-7 Fitting of environmental factors on NMDS of abundance community data.

Community analyzed	Environmental variable	R^2	<i>p</i> -value
Reptiles	Forest type	0.038	0.960
	Distance group	0.209	0.453
	Ground vegetation	0.222	0.705
	Leaf litter coverage	0.201	0.748
Amphibians	Forest type	0.206	0.387
	Distance group	0.441	0.014
	Ground vegetation	0.473	0.516
	Leaf litter coverage	0.611	0.254
Total	Forest type	0.448	0.039
	Distance group	0.591	0.003
	Ground vegetation	0.053	0.997
	Leaf l <mark>it</mark> ter co <mark>v</mark> erage	0.195	0.748

Appendix B-8 PERMANOVA results for all herpetofauna captures, reptile captures, and amphibian captures using presence/absence transformation.

Community analyzed	Variable	Df	Sum of Squares	Mean Squ <mark>are</mark> s	F-Model	\mathbb{R}^2	<i>p</i> -value
Total	Distance group	2	0.69	0.35	1.82	0.37	0.18
	Forest type	2	0.11	0.06	0.30	0.06	0.97
	Distance to water	1	0.14	0.14	0.75	0.08	0.60
	Elevation	1	0.18	0.18	0.93	0.10	0.52
	Canopy cover	1	0.17	0.17	0.90	0.09	0.54
	Patch size	-1	0.17	0.17	0.90	0.09	0.54
	Residuals		0.38	0.19	0.21		
	Total	10	1.85	1.00			
Reptiles	Forest type	2	0.32	0.16	0.84	0.17	0.62
	Canopy cover	1	0.27	0.27	1.40	0.14	0.26
	Residuals	7	1.34	0.19	0.69		
	Total	10	1.93	1.00			
Amphibians	Distance group	1	0.02	0.02	0.34	0.07	0.79
	Forest type	1	0.08	0.08	1.09	0.21	0.45
	Litter depth	1	0.05	0.05	0.71	0.14	0.62
	Residuals	3	0.21	0.07	0.58		
	Total	6	0.36	1.00			



Appendix C-1 Captured snake biometrics for each species by forest type.

Forest	C	C	Average	Average	
type	Species	Count	SVL (mm)	Mass (g)	
DDF	Boiga multomaculata	1	598	22.8	
	Boiga siamensis	1	93.5	74.4	
	Calliophis maculiceps	1	265	4.9	
	Chrysopelea ornata	1	570	36.8	
	Coelognathus radiatus	1	827	165.8	
	Lycodon capucinus	5	360.8	25.4	
	Lycodon laoensis	1	328	14.1	
	Naja siamensis	1	1266	758.5	
	Oligodon fasciolatus	1	616	86.7	
	Oligodon pseudota <mark>eni</mark> atus	1	269	8.7	
	Rhabdophis sub <mark>min</mark> iatus	1	248	5.7	
HDF	Boiga multom <mark>a</mark> culata	1	518	16.8	
	Lycodon capucinus	5	371.2	16.12	
	Lycodon <mark>lao</mark> ensis	1	321	12.5	
PLE	Bungaru <mark>s can</mark> didus	2	700.5	138.6	
	Calloselasma rhodostoma	1	552	36.7	
	Chrysopelea ornata	1	724	726	
	Dendrelaphis subocularis	1	450	22.7	
	Enhydris plumbea	apa	182	1.5	
	Lycodon capucinus	3	337.7	14.1	
	Oligodon fasciolatus	2	376	46.6	
	Oligodon pseudotaeniatus	1	288	13.4	
	Oligodon taeniatus	1	307	15.3	
	Rhabdophis chrysargos	1	144	10.9	

Appendix C-2 Biometric information for species by each forest type examined.

Species	Forest	orest Individuals		SVL		Mass	
Species	type	marviduais	Avg	StdDev	Avg	StdDev	
Dixoneus siamensis	DDF	1	50.00	-	2.25	-	
	HDF	8	49.88	1.89	3.24	0.31	
	PLE	6	43.06	11.53	2.14	1.30	
Eutropis macularia	DDF	4	51.50	7.55	4.13	2.22	
	HDF	5	57.80	2.95	5.07	0.85	
	PLE	5	49.51	3.77	3.47	0.97	
Kaloula	HDF	10	45.99	11.38	11.62	6.03	
mediolineata							
	PLE		22.48	-	1.28	-	
Kaloula pulchra	HDF	2	61.00	1.41	17.34	0.44	
Leiolepis reevesii	HDF	2	55.67	35.82	11.01	8.68	
	PLE	4	73.85	5.66	10.52	2.76	



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