



TEXAS A&M UNIVERSITY
SAN ANTONIO

INSIGHTS INTO SUBTERRANEAN CONNECTIVITY WITHIN THE AQUIFER OF THE YUCATÁN PENINSULA: POPULATION GENETICS AND DISTRIBUTION OF *TYPHLATYA* SPECIES

Gabrielle Vaughn, Lauren Ballou, David Brankovits,
Efrain Chavez-Solis, Fernando Calderón- Gutiérrez,
Tom Iliffe, Nuno Simões, and Elizabeth Borda

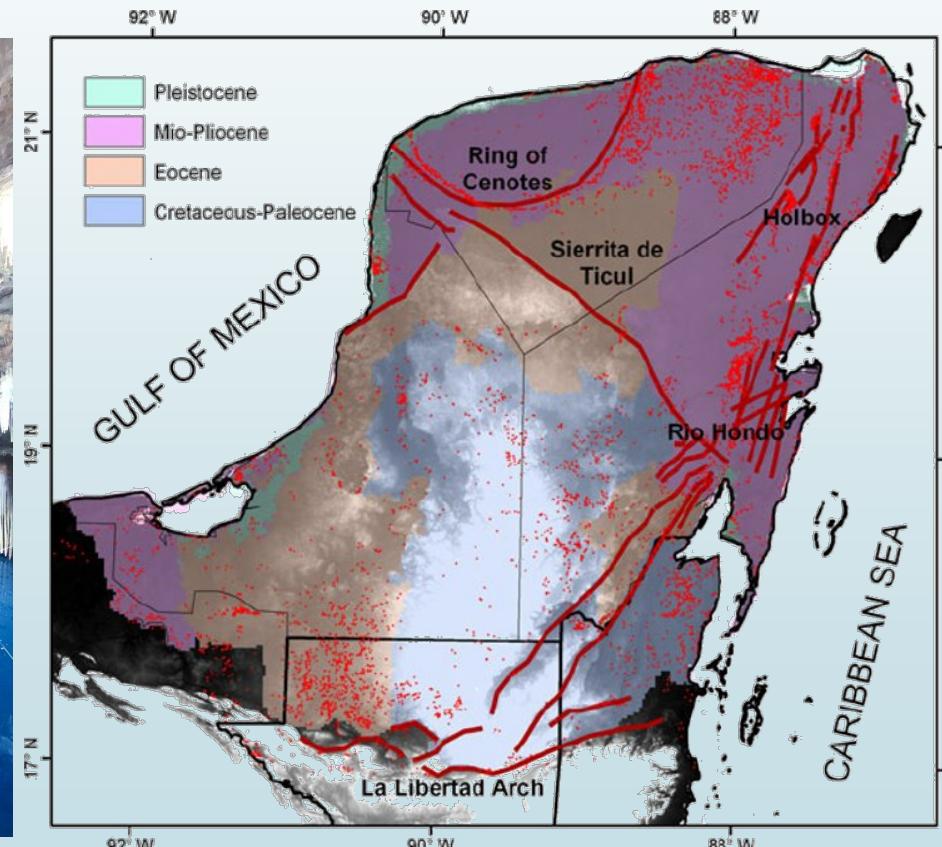
Department of Natural Sciences, Texas A&M
University- San Antonio

Texas Groundwater Invertebrate Forum 2024



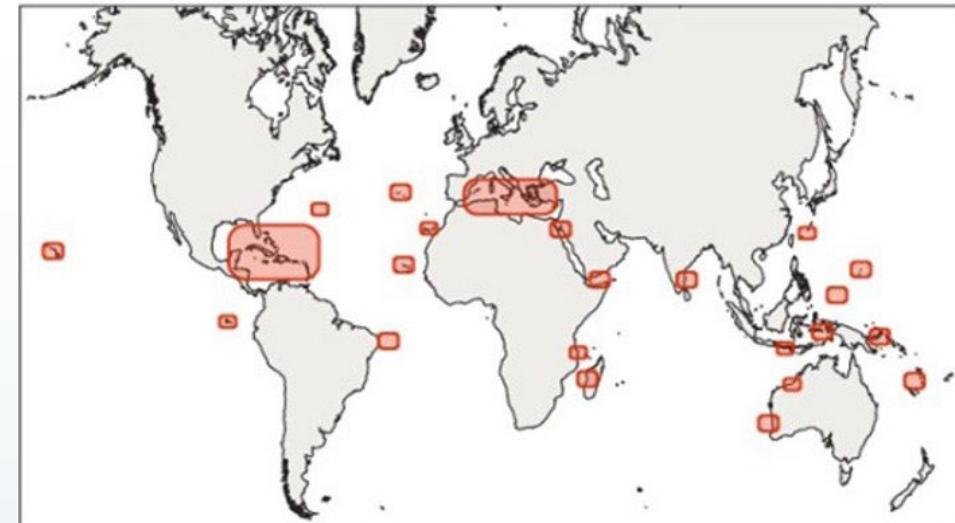
The Yucatán Peninsula

- No major lakes or rivers
- Extensive karst aquifer ecosystems
- ~80 stygobiont species (>70% crustaceans)
 - High endemism
 - Low abundance

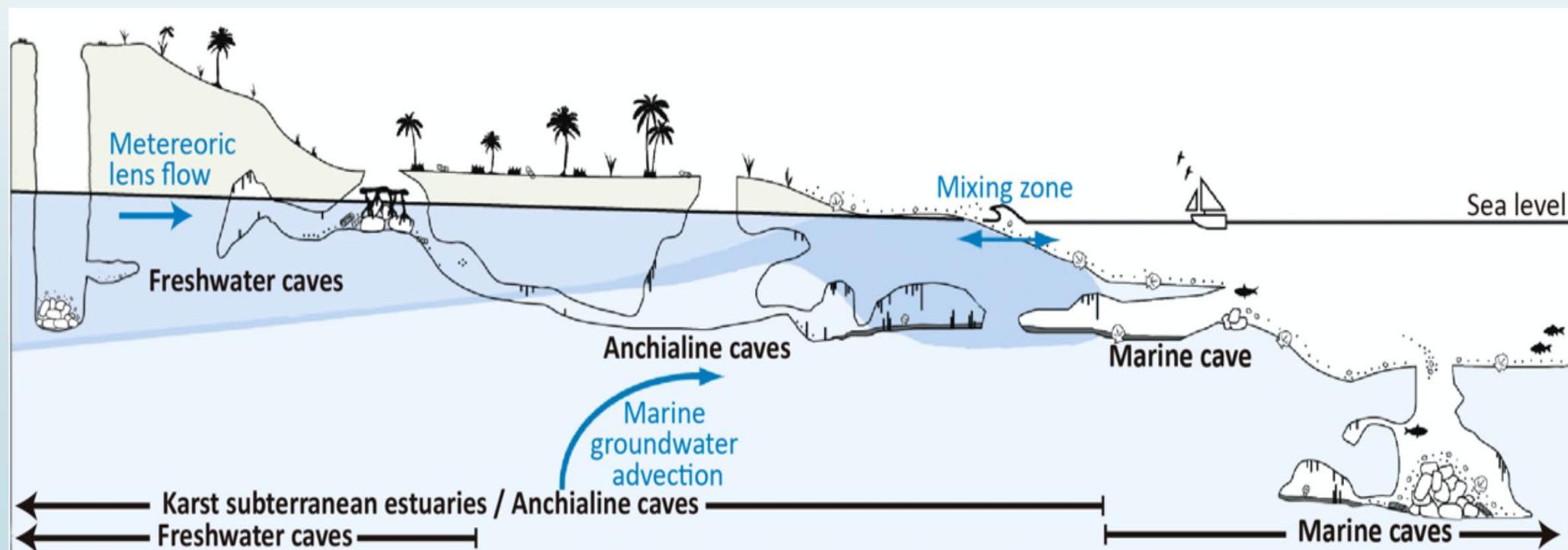


Anchialine Caves

- Karst subterranean estuaries
- Stratified salinity (fresh – saline), separated by a halocline
- Transient ecosystems relative to sea level rise and fall

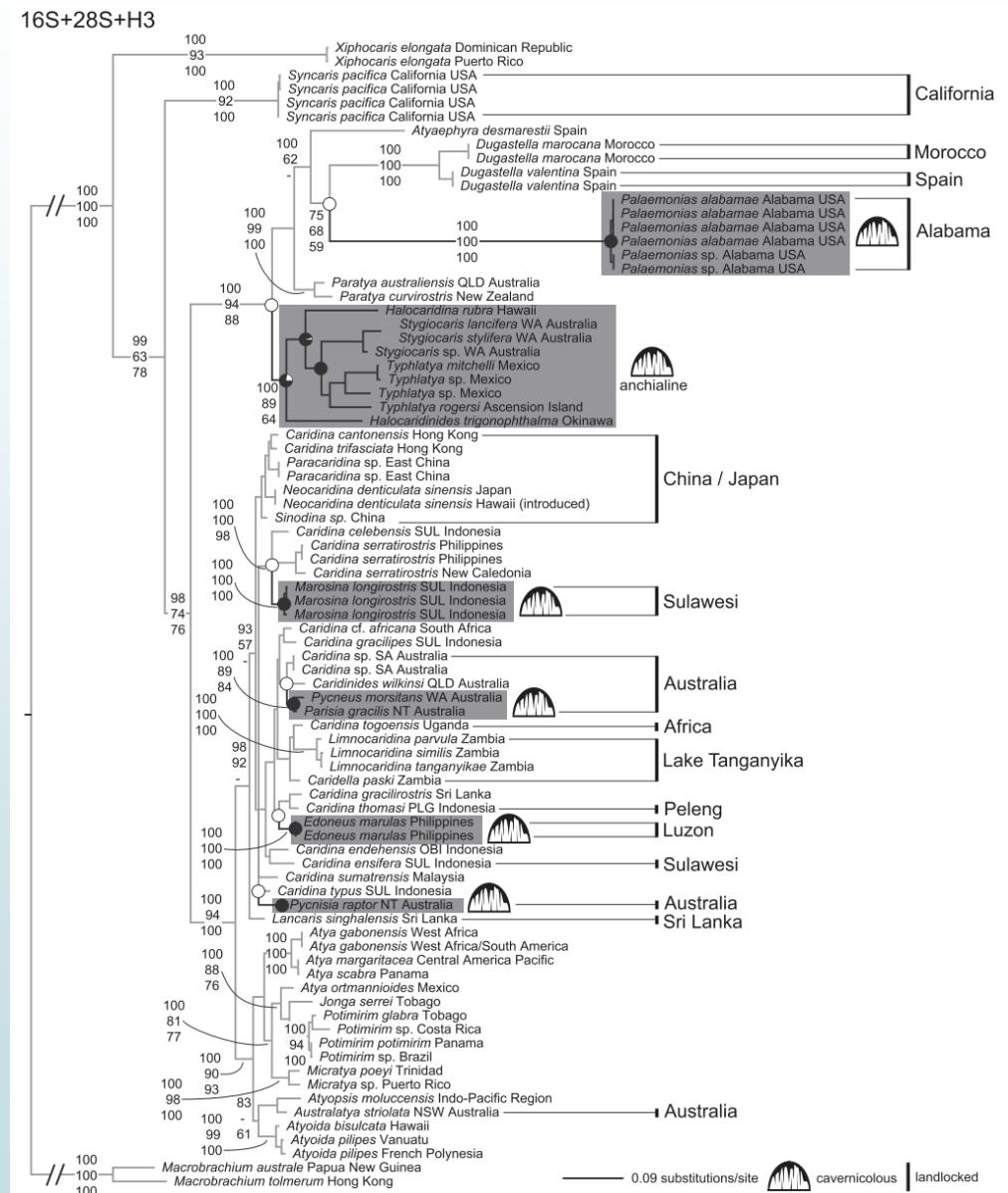


Iliffe and Kornicker 2009, Martinez et al. 2018

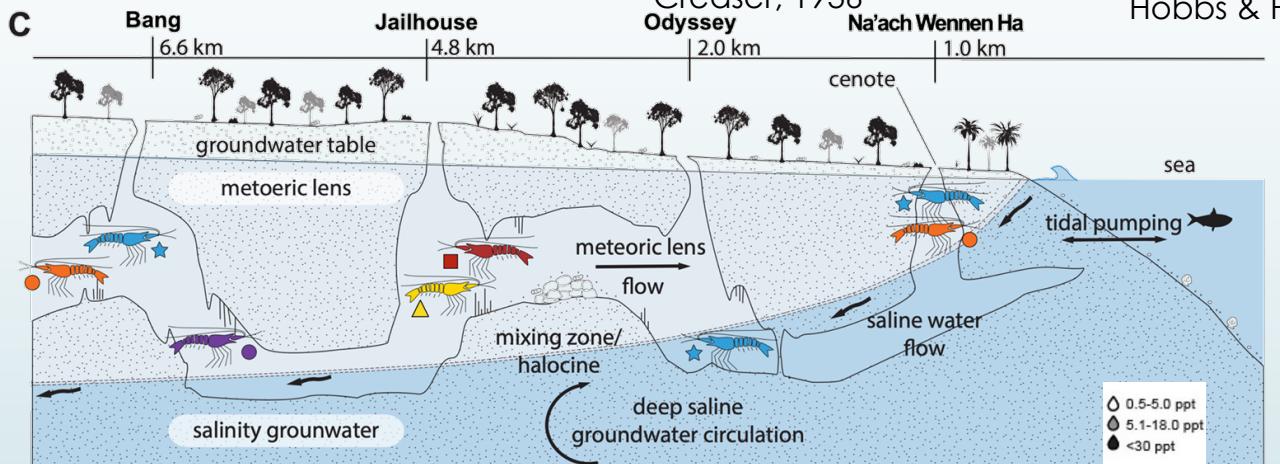
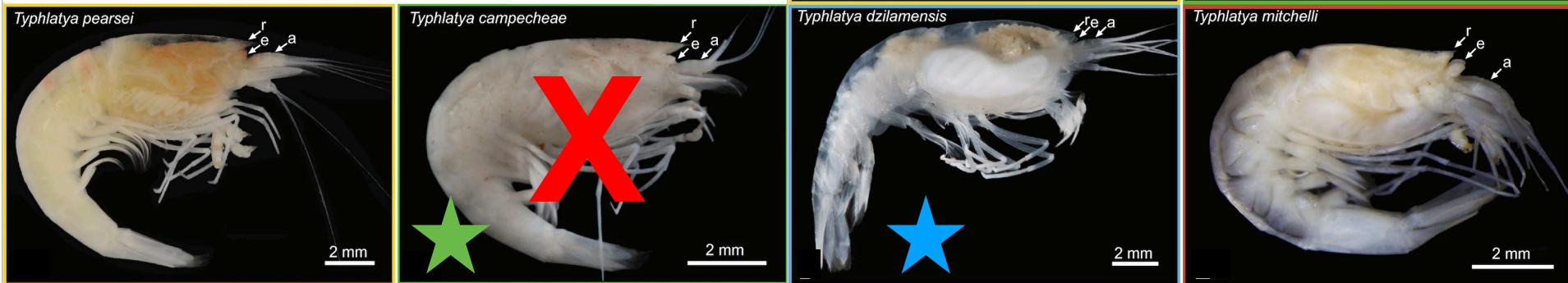


Atyidae

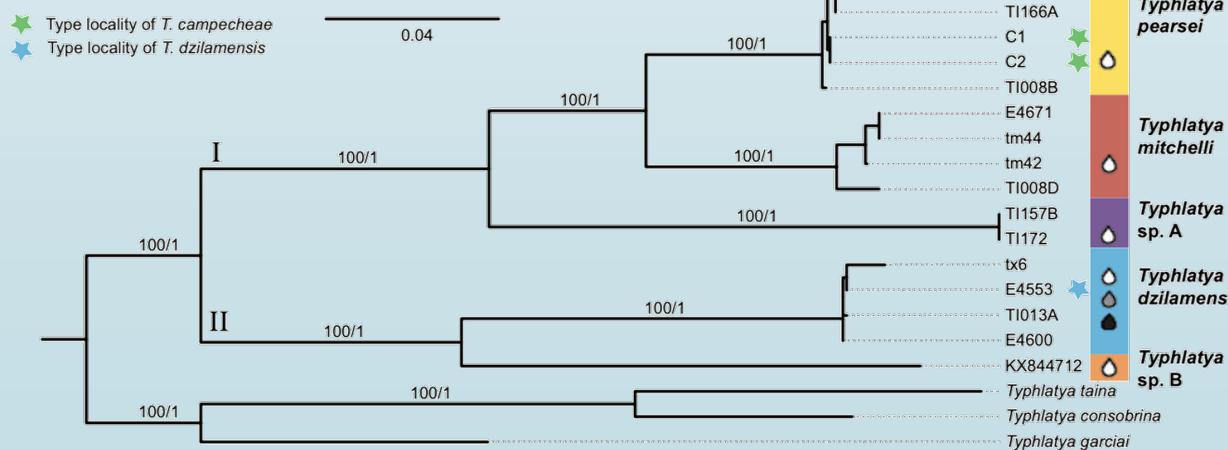
- 450 species, 40 genera
- Freshwater family
- Convergent cave adaptations
- Single anchialine adapted clade
 - Fresh \leftrightarrow Saline
- Not recorded in the open ocean



Typhlatya in the Yucatán Peninsula



16S rRNA, COI, CYTB, 18S rRNA, 28S rRNA, H3



- Endemic
- Evidence of syntopy
- Coastal euryhaline
Typhlatya dzilamensis demonstrate phenotypic plasticity (e.g., salinity tolerance)

Research Questions

- What is the distribution of the *Typhlatya* species within the Yucatán Peninsula?
- Can the distribution and population genetics of *Typhlatya* species provide insights into connectivity between and among cave systems within the Yucatán Peninsula?



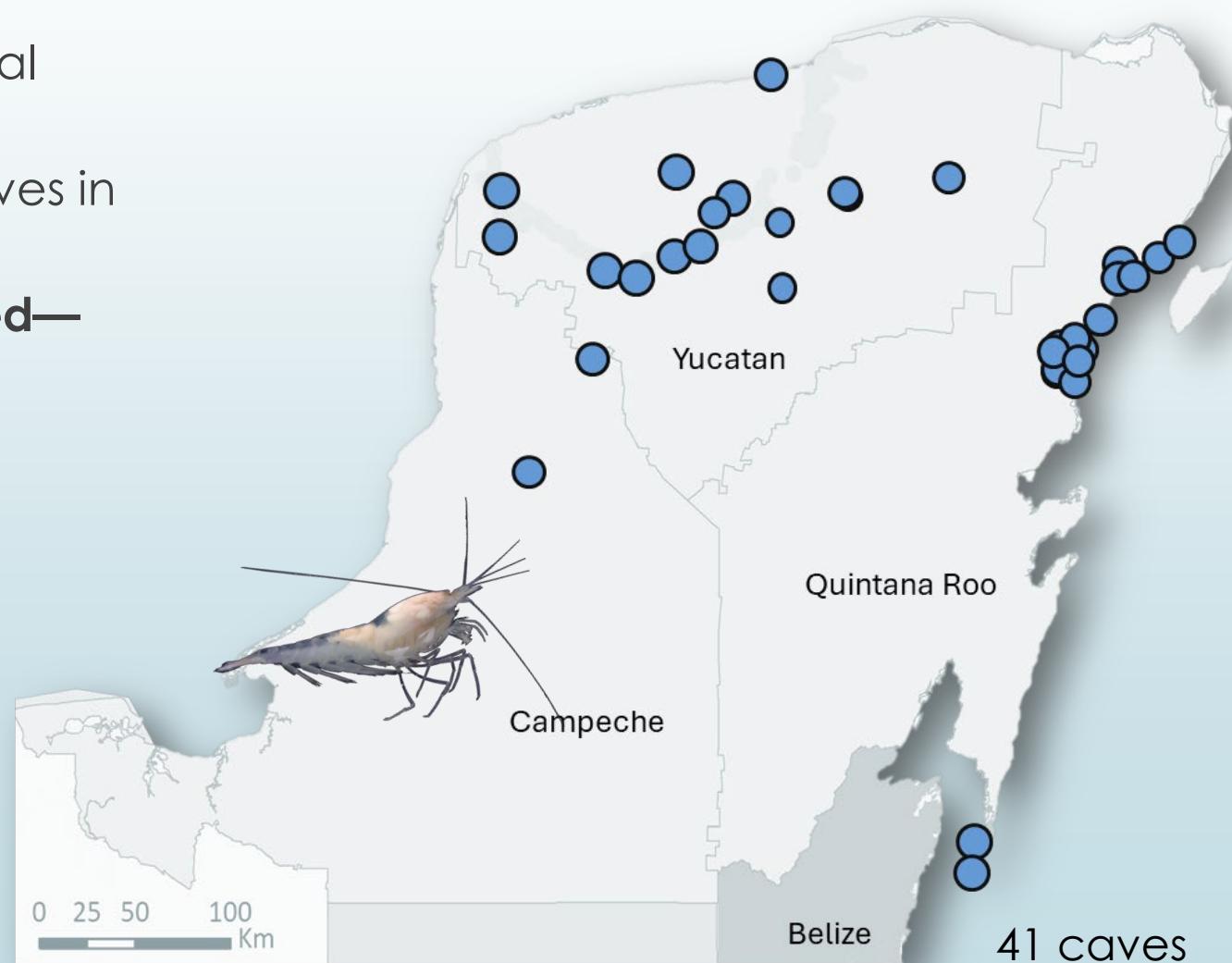
Methods

Sampling—

- ~41 inland and coastal caves
- Includes 2 marine caves in Belize

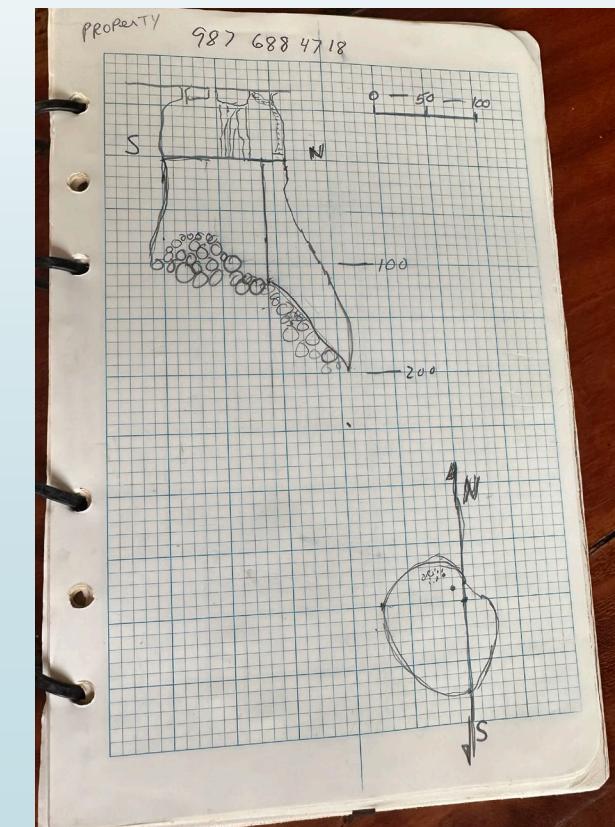
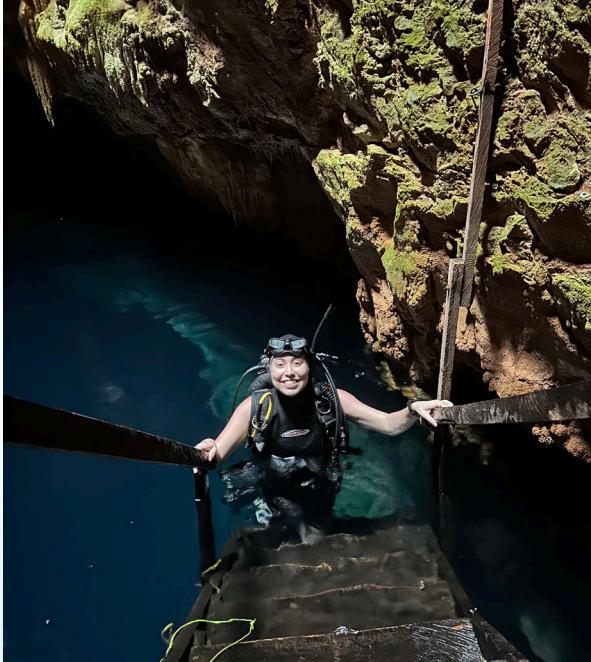
Molecular Data extracted—

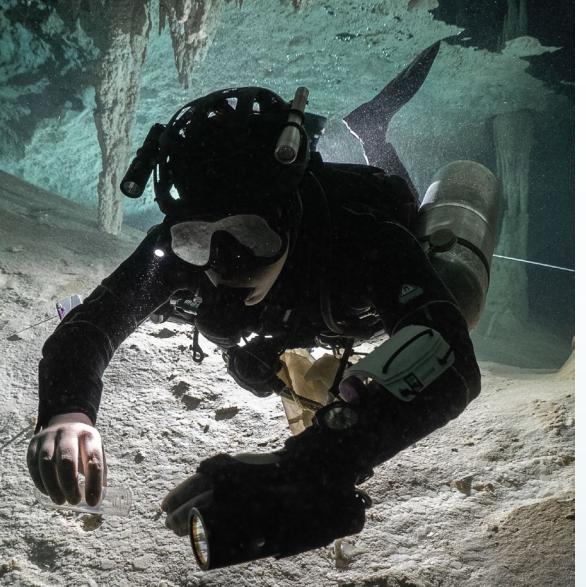
- Mitochondrial genes
 - 16S rRNA
 - CO1
 - CYTB
- Nuclear genes
 - 28S rRNA
 - ITS



Methods— Sampling in SOTUTA, YP

- 10-day fieldwork expedition
- Exploration and mapping of unmapped cenote, 'Tres Rayos'
 - Courtesy of Imelda & Steve Moon
- Specimen collection
 - Only one *T. mitchelli* was collected ☹
 - Low abundance!





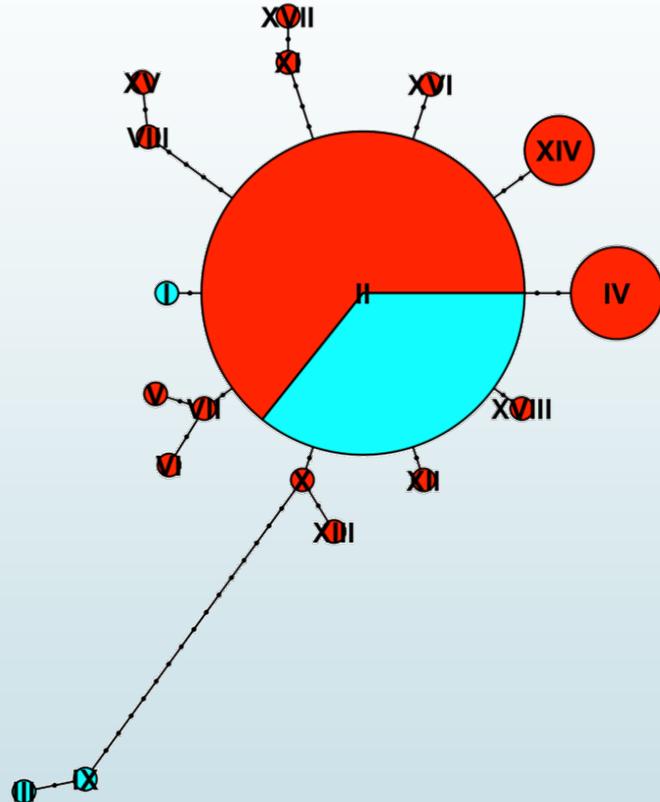
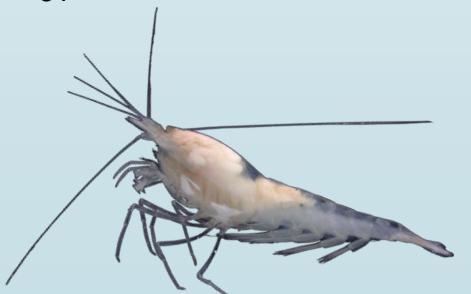
Methods

Population Genetics—

- Haplotype Network Reconstruction
- Population structure
- Connectivity

Statistical Analysis—

- Genetic diversity
- Isolation by distance (IBD)
- F_{ST}



Results



Typhlatya pearsei

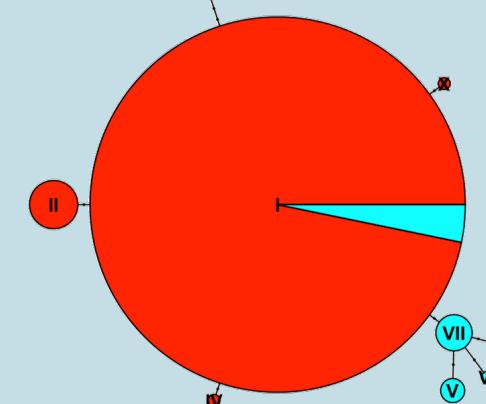
12 Caves



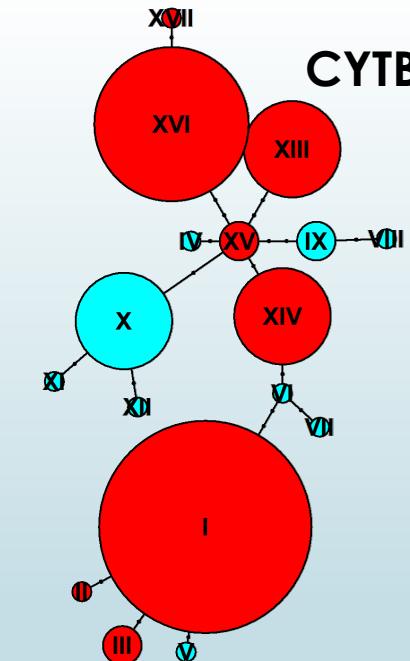
COI



16S



CYTB



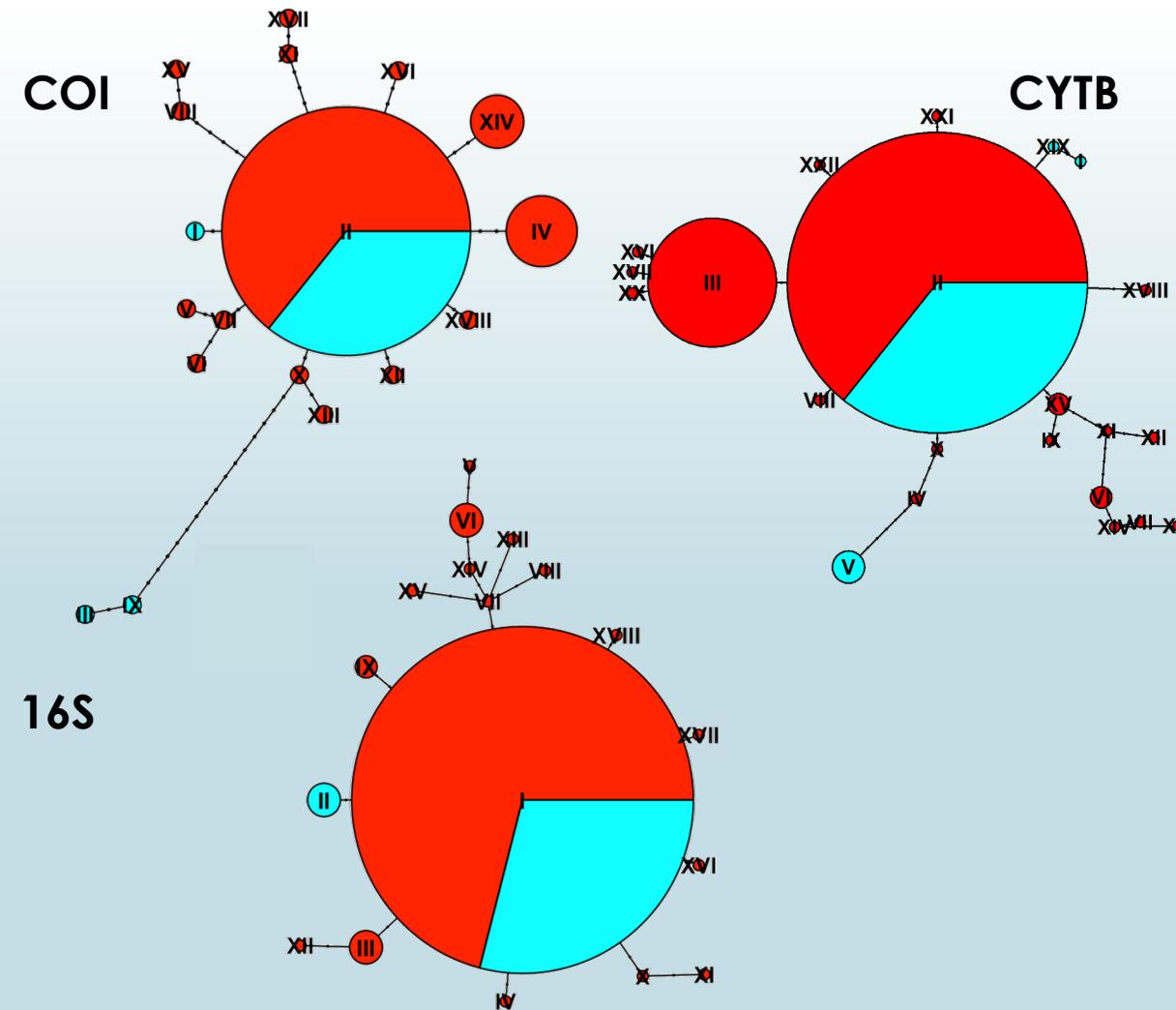
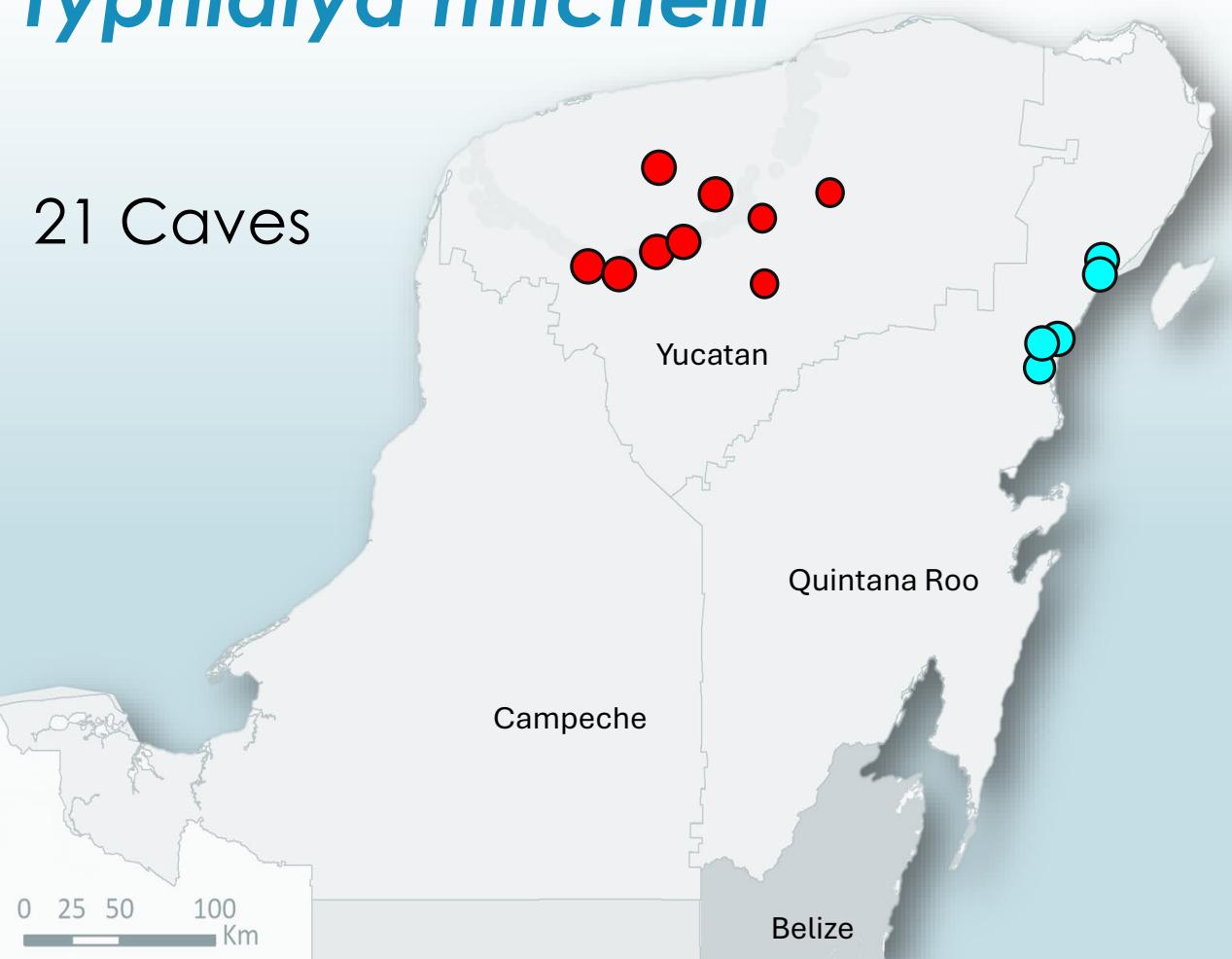
28S

1 haplotype

Gene	n	n (inland)	n (coastal)	F _{ST}	IBD (r; significance)	Hd	nH
COI	34	24	10	0.4234	0.1534; 0.021	0.9857	27
CYTB	49	35	14	0.1677	0.0907; 0.036	0.9014	17
16S	47	36	11	0.2612	0.4095; 0.001	0.5597	10
28S	27	16	11	0	-	0	1

Typhlatya mitchelli

21 Caves



Gene	n	n (inland)	n (coastal)	F _{ST}	IBD (r; significance)	Hd	nH
COI	36	29	7	0.1205	0.4325; 0.013	0.8413	18
CYTB	65	50	15	0.0375	0.2043; 0.014	0.7707	22
16S	56	40	16	0.0451	0.2165; 0.015	0.6714	18
28S	27	22	5	-	-	0	1

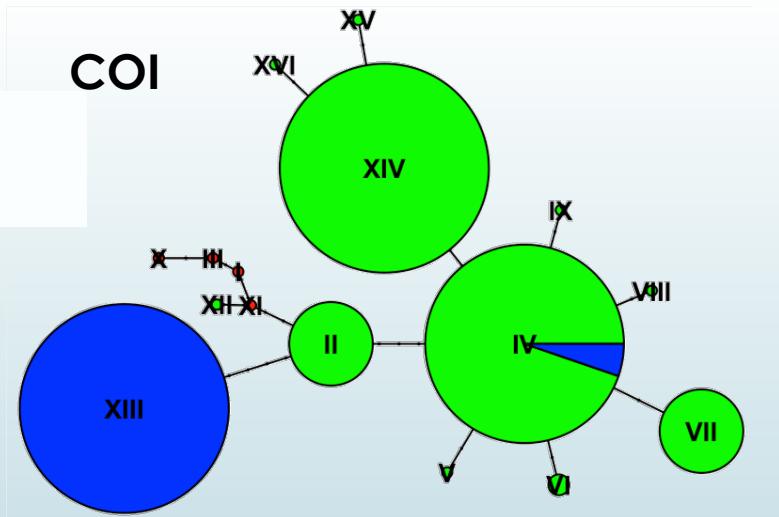
28S
1 haplotype

Typhlatya dzilamensis

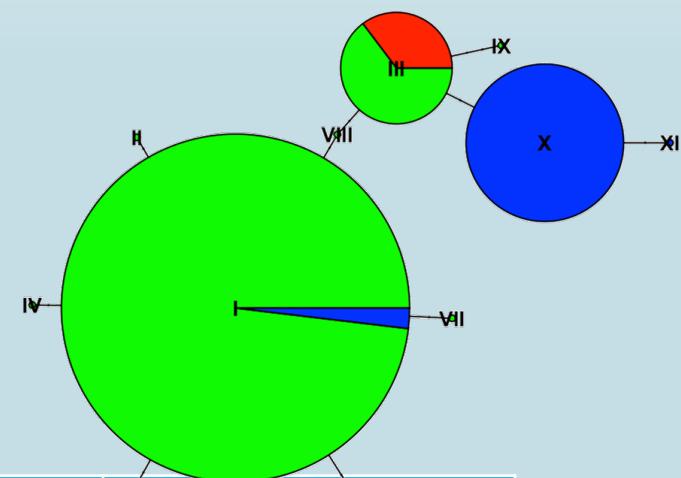
16 Caves



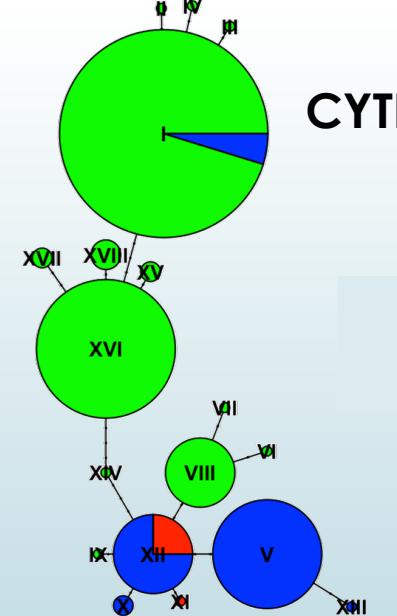
COI



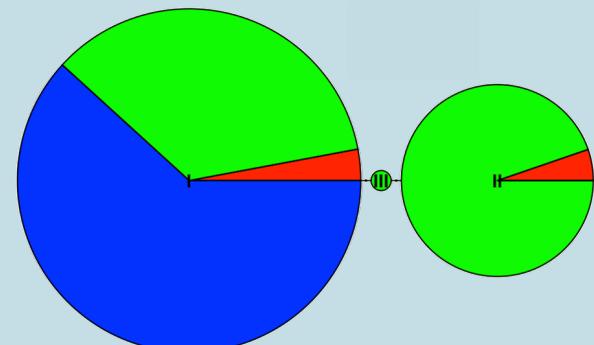
16S



CYTB



28S

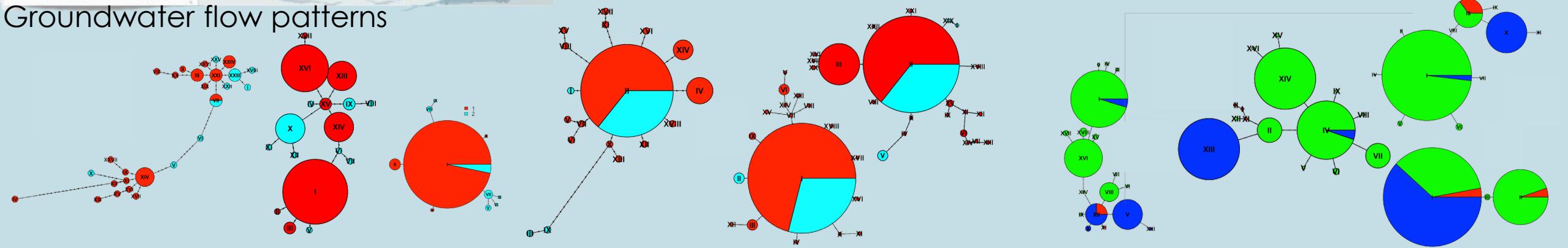


Gene	n	n (Yucatan coast)	n (Caribbean coast)	n (Belize)	F _{ST}	IBD (r; significance= 0.001)	Hd	nH
COI	87	4	62	21	0.6745	0.5861	0.8375	16
CYTB	79	3	21	55	0.2411	0.228	0.8666	18
16S	107	6	75	26	0.6539	0.539	0.6828	11
28S	55	2	33	20	0.6752	0.8523	0.5064	3

Summary

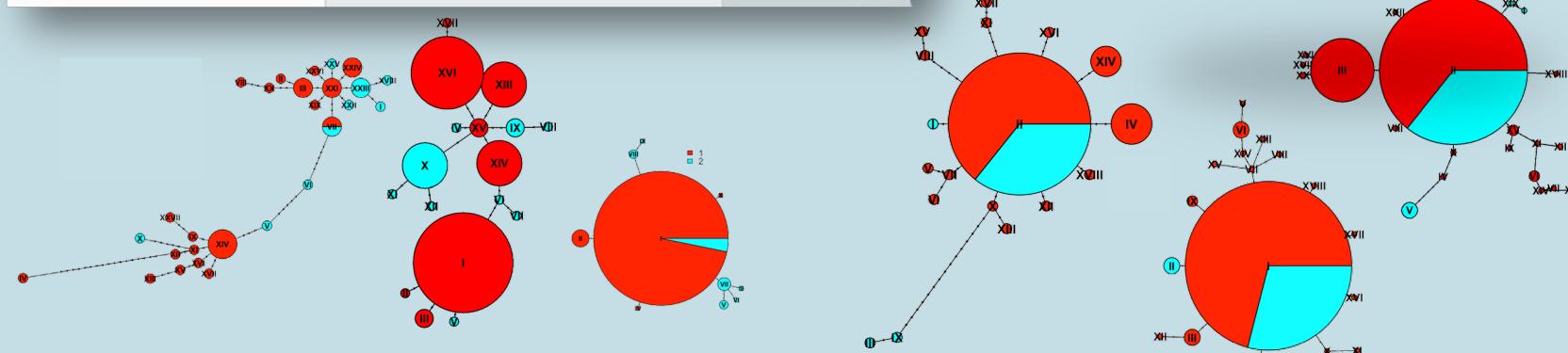
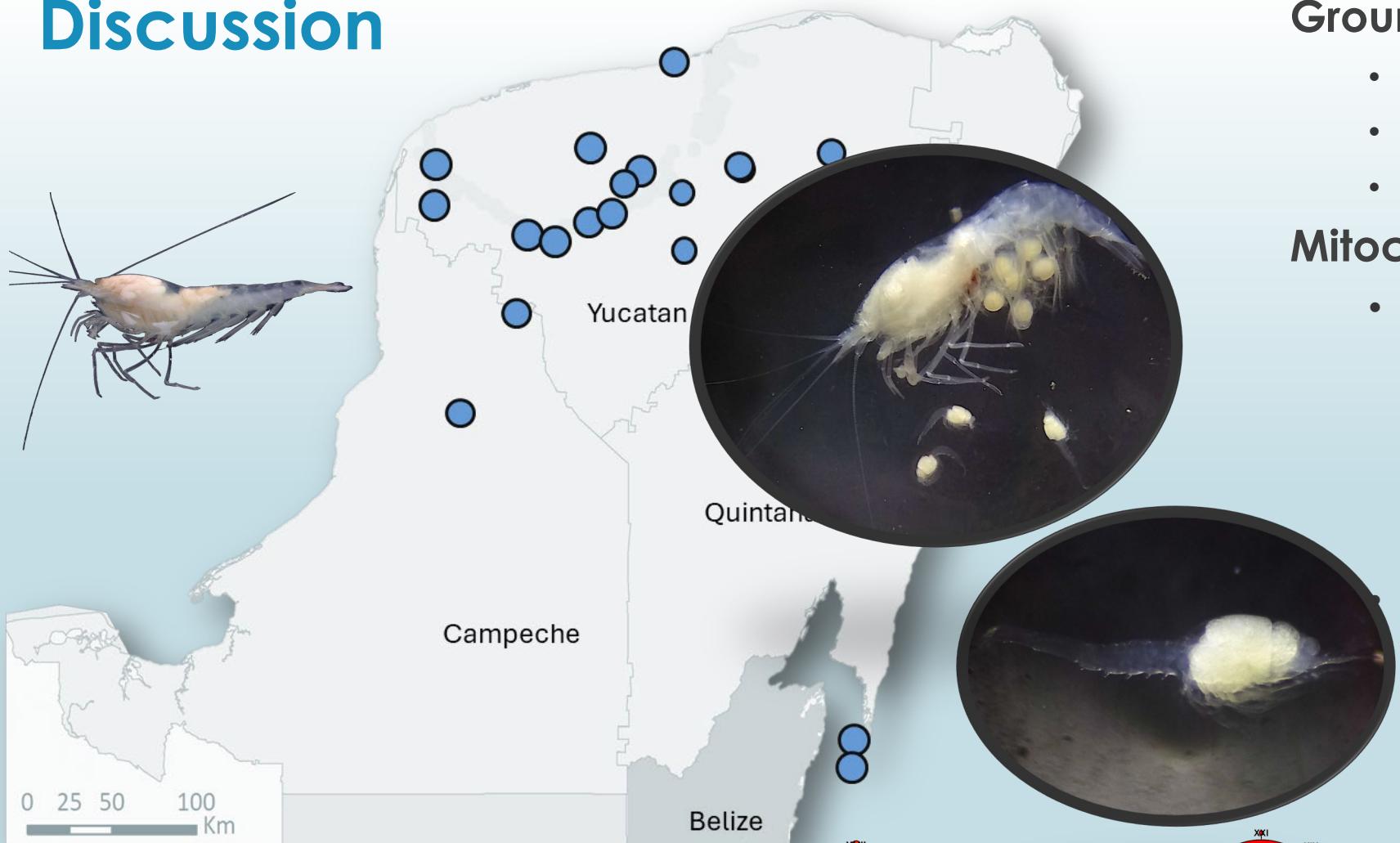


Groundwater flow patterns



- Importance of molecular tools towards more **accurate identification of populations** within Yucatán Peninsula aquifer
- Provide insights into the
 - Potential **connections between / among** inland and coastal cave systems
 - Potential connections between / among proximal and distant cave systems
 - Distributional ranges
- **Subterranean caves are NOT isolated**

Discussion



Groundwater connectivity & barriers

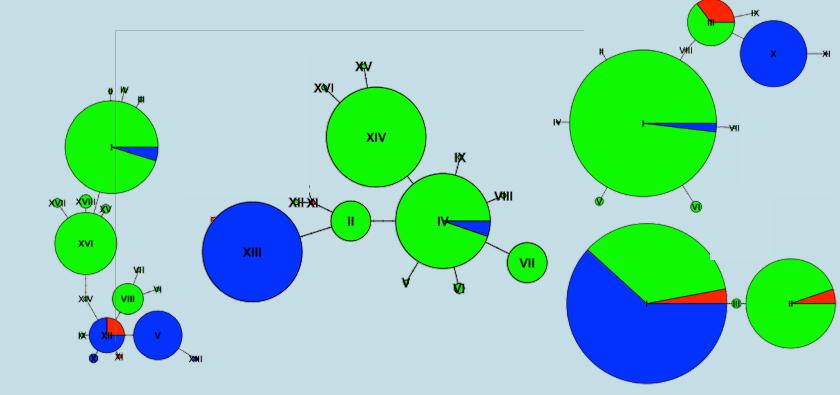
- Inland vs coastal
- Cave passages
- Salinity stratification

Mitochondrial vs. Nuclear diversity

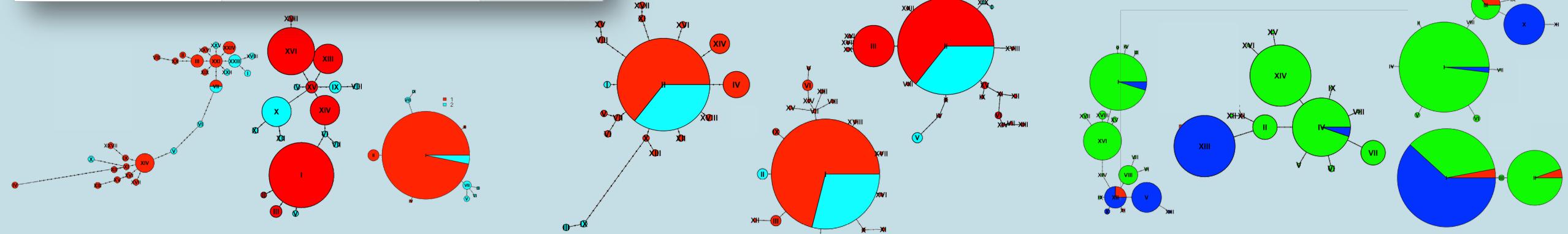
- mtDNA
 - Maternally inherited
 - Brooding females – limited dispersal?
 - Lecitotrophic or direct developing larvae?

nucDNA

- Different patterns
- Moving males



Continuing Work



- Evaluation of the population connectivity by cave
- Linking environmental data with population structure
- Increase sample sizes
- Decrease geographic sampling gaps
- Inclusion of ITS (In progress) will provide added insight among populations of *Typhlatya* species
- Setting the stage for ongoing work to integrate genomic data

Acknowledgements



Advisor/Mentor: Dr. Elizabeth Borda

Borda lab

Co-authors: Lauren Ballou, David Brankovits, Efrain Chavez-Solis, Fernando Calderón- Gutiérrez, Tom Iliffe and Nuno Simões

Imelda & Steve Moon, Danielle Bragg

KISS Rebreather team

Research funding:

Texas A&M University San Antonio Research Council & Summer Research Grants

NSF Biological Oceanography (#2136322)



TEXAS A&M UNIVERSITY
SAN ANTONIO



Universidad Autónoma
de México - Sisal

TEXAS A&M
UNIVERSITY
GALVESTON CAMPUS®





Questions?