# Yuanning Li

**Position:** Postdoc

Address: 165 Prospect Street, New Haven, CT, 06511

**Phone:** 334-844-3223

Email: Yuanning.li@yale.edu

## **Research Interests**

Animal comparative genomics, systematics, and life history evolution; Genome evolution of marine invertebrates; Genomic and symbioses adaptation to deep-sea chemosynthetic (hydrothermal vents, cold seeps or whale bones) habitats, especially in tubeworms; Mitogenomics.; Root of the animal tree of life.

### **Education**

2018-present PostDoc, Yale University Advisor: Casey Dunn

2017-2018 PostDoc Auburn University Advisor: Ken Halanych

2012-2017 Ph.D **Auburn University** (Molecular Systematics)

**Committee**: Drs. Kenneth Halanych (advisor), Scott Santos, Jason Bond, Eric Peatman

**Courses:** Advanced Biological Computing, Invertebrate Biodiversity, Molecular Ecology, Molecular Genetics And Biotech, Biogeography, Evolutionary Biology, Invertebrate Zoology.

2011-2012 M.S. Ocean University of China (Aquatic biology)

2007-2011 B.S. Shanghai Ocean University (Biological Science)

#### **Research Projects**

- Population genetics of wood-eating bivalves Xylophaga and bone-eating worms Osedax
- Whole genome sequencing and annotation of deep-sea tubeworm Lamellibrachia
- Comparative genomics of siboglinid endosymbionts.
- Mitogenomics of wood-eating bivalves Xylophaga

## **Research Cruises**

2013 **BoWLs** - Two weeks long NSF funded deployment of bone/wood landers for

capture of Osedax and Xylophaga in the Pacific Northwest.

Workshop

May 2014 **AU Bioinformatic Bootcamp.** 

October 2014 FAB Fall Taxonomy Workshop Polychaetes part III: the Spionida

### **Publication**

- 13. Li, Y., A.F. Bernardino, K.M. Halanych. (*In prep*). Whole genome sequencing of tubeworm Lamllibrachia reveals adaptation to deep-sea chemosynthetic environments. *PNAS*
- 12. Li, Y., P.G. Galaska, A.F. Bernardino, K.M. Halanych. (*In prep*). Recruitment patterns of deep-sea wood-boring bivalves (Xylophagidae, Mollusca) inferred from SNP data.
- 11. L. Kong, Li. Y, K.M. Kocot, K.M. Halanych. (*In prep*) (2018) Mitogenomics reveals phylogenetic relationships of Arcoida molluscs and multiple independent expansions in mitogenome size. Molecular Phylogenetics and Evolution. (Co-first Author).
- 10. P.G. Galaska, Li, Y., K.M. Kocot, A.R. Mahon, K.M. Halanych. (*In press*) (2018) Conservation of brittle stars (Ophiuroidea, Echinodermata) mitochondrial genome arrangements. *Molecular Phylogenetics and Evolution*
- 9. Li, Y., A.F. Bernardino, K.M. Halanych. (Submitted). Mitogenomics reveal evolutionary relationships and dynamic genomic architecture of deep-sea wood-boring bivalves (Molluscs: Xylophagaidae). Molecular Phylogenetics and Evolution.
- 8. Li, Y., K.M. Kocot, M.G. Tassia, J.T. Cannon, M. Bernt, K.M. Halanych. (*In press*). The case of the codon reassignment: A novel hemichordate mitochondrial genetic code provides insights on deuterostome evolution. *Genome Biology and evolution*. IF=3.940
- 7. Li, Y., M.R. Liles, K.M.Halanch. (*In press*). (2018) Endosymbiont genomes yield clues of tubeworm success. *ISME journal*. **IF=9.520**
- 6. A.F. Bernardino, Li, Y., C.R. Smith, K.M. Halanych. (2017) Multiple introns in a deep-sea Annelid mitochondrial genome. *Scientific Report* 7:4295.
- 5. D.S. Waits, S.R. Santos, D.J. Thornhill, Li, Y., K.M. Halanych. (2016) Evolution of Sulfur Binding in Hemoglobin in Siboglinidae (Annelida) with Special Reference to Bone Eating Worms, Osedax. Journal of Molecular Evolution. 82: 219-229.
- 4. Li, Y., K.M. Kocot, N.V.Whelan, S.R. Santos, D.S. Waits, K.M. Halanych. (2016)
  Phylogenomics of tubeworms (Siboglinidae, Annelida) and comparative
  performance of different reconstruction methods. *Zoologica Scripta*. 46: 200-213.
  IF=3.057
- 3. Li, Y., K.M. Kocot, S.R. Santos, D.J. Thornhill, K.M. Halanych. (2015) Mitogenomics reveals phylogeny and dramatic size variations of control regions in the deep-sea family Siboglinidae (Annelida). *Molecular Phylogenetics and Evolution*. doi:10.1016/j.ympev.2015.02.008. IF=4.412
- 2. Li, Y., P. Ma, P. Liu, Q. Li. The genetic diversity and phylogenetic analysis of ITS1 in mitochondrial DNA of four populations of Portunus Trituberculatus. (2012)

  Oceanologia Et Limnologia Sinica. 2012,43(4).

1. X. Su, G. Wang, J. Feng, J. Li, Y. Genetic diversity of 15 *Macrobrachium nipponense* populations in Poyang Lake based on microsatellite analysis. (2011) *Chinese Journal of Ecology*. 2011,30(9):2007-2013.

#### **Oral Presentations**

- 8. Li, Y. (speaker), K.M. Halanych. Comparative genomics of seep-dwelling tubeworm (Siboglnidae: Annelida) endosymbionts. The Society for Integrative & Comparative Biology (SICB). San Francisco, CA, USA. January 2018
- 7. Li, Y. (speaker), K.M. Halanych. Comparative genomics reveal symbiont-host evolution of deep-sea tubeworms. 6th International Symposium on Chemosynthesis-Based Ecosystems. Woods Hole, Mass. USA. Augustus 2017
- 6. Li, Y. (speaker), A.F. Bernardino, K.M. Halanych. Recruitment patterns of deep-sea wood-boring bivalves (Xylophagidae, Mollusca) inferred from SNP data. *The Society for Integrative & Comparative Biology (SICB)*. New Orleans, LA, USA. January 2017
- **5. Li, Y.** (speaker), K.M. Halanych. **Using genomic tools to understand symbiont-host evolution of deep-sea tubeworms.** *12th International Polychaete Conference (IPC12)*. Cardiff, UK. August 2016
- Li, Y. (speaker), K.M. Halanych. Recruitment of deep-sea wood-boring (Xylophagidae, Mollusca) inferred from RAD sequence Data. Evolution 2016. Austin, TX, USA. May 2016
- 3. Li, Y. (speaker), N.V. Whelan, K.M. Kocot, S.R. Santos, K.M. Halanych. Phylogenomics of tubeworms (Annelida: Siboglinidae) facilitates comparative performance of supermatrix versus species tree phylogenetic approaches. *The Society for Integrative & Comparative Biology (SICB)*. Portland, OR, USA. January 2016
- 2. Li, Y. (speaker), K.M. Kocot, S.R. Santos, K.M. Halanych. Phylogenomics and species-tree analyses reveal deep-sea family Siboglinidae (Annelida) relationships. *The Society for Integrative & Comparative Biology (SICB)*. West Palm Beach, FL, USA. January 2015
- 1. Li, Y. (speaker), K.M. Kocot, S.R. Santos, K.M. Halanych. Mitogenomics and Phylogenomics of the family Siboglinidae (Annelida). Southeastern Ecology and Evolution Conference (SEEC). March 2014

## **Outreach/Service**

2016-present	Manager of the Auburn University Next-generation Sequencing Center.
2013-2017	<b>AU Explore.</b> (outreach event to interest area grade-school students in higher education) 7
2016	Outreach in Drake Middle School 7th grade. (2 classes, 40 students)

#### **Grants and Fellowships**

2017 Ocean University of China-Auburn University (OUC-AU) Grant. Genetic resources for Ark Shell bivalves: An evolutionary perspective on aquaculture resources. (Listed as CO-I) \$78,000

2012-2016 Chinese Government Scholarship. (1600\$/month, 4 years)

**2016** COSAM Travel Award. (\$250)

**2016 CMB Summer GRA.** (\$1683/Mo, \$5049 Total)

2015 Auburn University Graduate School Travel Award. (\$400)

**2014** Biological Sciences Department Travel Award. (\$250)

**2014 COSAM Travel Award.** (\$350)

Awards

2017 Kenneth Ottis Distinguished Graduate Fellowship (For outstanding

achievement in physiology/Molecular Biosciences).

2017 COSAM Outstanding International Graduate Student.

**Teaching Experience** 

2016 **Graduate Teaching Assistant**. Introductory Biology. Auburn University

2017 **Graduate Teaching Assistant**. Introductory Biology. Auburn University

### References

Dr. Kevin Kocot Dr. Scott Santos Dr. Ken Halanych Professor of Biology Assistant Professor of Biology Professor of Biology Auburn University The University of Alabama **Auburn University** 101 Rouse Life Bldg 307 Mary Harmon Bryant Hall 101 Rouse Life Bldg Tuscaloosa, AL 35847 Auburn, AL 36849 Auburn, AL 36849 205-348-4052 334-844-3222 santos@auburn.edu ken@auburn.edu kmkocot@ua.edu