

# Timothy Stephens

## Curriculum vitae

### Personal summary

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I am a computational biologist working on understanding the forces that govern the interactions between organisms in ecologically critical environments, such as coral reefs, and the forces that drive major evolutionary transitions, such as the one that gave rise to the first photosynthetic Eukaryotas.

📍 Department of Biochemistry and Microbiology, Rutgers University, USA  
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🌐 timothystephens.github.io  
🐙 TimothyStephens

### Education

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2012 – 2015	Bachelor of Biotechnology, Institute for Molecular Bioscience, The University of Queensland
2015 – 2015	Honours, Institute for Molecular Bioscience, The University of Queensland
2016 – 2019	PhD, Institute for Molecular Bioscience, The University of Queensland

### Research Experience

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2012 – 2013	Undergraduate Research Project, Institute for Molecular Bioscience, The University of Queensland
2013 – 2013	Paid Research Assistant, School of Biological Sciences, The University of Queensland
2014 – 2014	Undergraduate Researcher Project, School of Biological Sciences, The University of Queensland
2015 – 2015	Honours Project, Institute for Molecular Bioscience, The University of Queensland
2016 – 2019	PhD, Institute for Molecular Bioscience, The University of Queensland
2019 – present	Post Doctoral Associate, Department of Biochemistry and Microbiology, Rutgers University

## Teaching experience

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- 2012 – 2014      **CASPiE Tutor**  
University of Queensland  
Duties included guiding undergraduate chemistry students through an advanced set of research-focused practicals, with the aim of developing their critical and experimental thinking.
- 2013              **Science Mentor**  
University of Queensland  
Duties included organizing and running first year science student introduction and social engagement events designed to inform students about study/research opportunities available at The University of Queensland.
- 2013 – 2014      **Peer Assisted Study Session (PASS) Tutor**

University of  
Queensland  
Duties included  
planning and  
leading multiple  
weekly tutorial  
sessions, each  
comprising 20-  
30+ students.

2016 – 2017

### **Practical Tutor**

University of Queensland

Duties included assisting students with the completion of set questions and marking of assignments.

2018

### **Head Practical Tutor**

University of Queensland

Duties included designing and developing assessment material for the course and organising marking of assignments.

2020 – 2022

### **Guest lecturer**

Rutgers University

Conducted lectures on de

novο next-  
generation  
sequencing,  
genome and  
transcriptome  
sequencing, and  
metagenomics  
in a join  
undergraduate  
and  
postgraduate  
course titled  
“Fundamentals  
of Microbial  
Genomics”.  
Also assisted in  
designing and  
marking student  
exam questions  
and students’  
oral  
presentations.

## Professional development

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- 2017 UQ Idea Hub, University of Queensland
- 2017 Research Commercialisation Workshop, University of Queensland

## Awards and honours

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- 2011 Merit Scholarship for academic achievement, University of Queensland
- 2012 Selected for the Advanced Study Program in Science (ASPinS; based on academic merit),

University  
of  
Queensland

2012	Dean's Commendation for Academic Excellence (achieved a GPA over 6.6), University of Queensland
2016	Research Training Program (RTP) scholarship, University of Queensland
2016	Top poster prize at the IMB Research Higher Degree Student Symposium, University of Queensland
2017	Won best pitch prize at the UQ Idea Hub, University of Queensland
2018	Registration award for Society for Molecular Biology & Evolution (SMBE) 2018, SMBE
2021	2019 Dean's Award for Outstanding Higher Degree by Research Theses, University of Queensland

## Publications

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2022 Davies S. W., Gamache M. H., Howe-Kerr L., Kriefall N.G., Baker A.C., Banaszak A., Bay L., Bellantuono A.J., Chan C. X., Claar D.C., Coffroth M.A., Cuning R., del Campo J., Frommlet J. C., Fuess L. E., Goulet T. L., Hoadley K. D., Hume B. C. C., Kemp D. W., Kitchen S. A., LaJeunesse T. C., Lin S., McIlroy S., McMinds R., Nitschke M. R., Oakley C. A., Peixoto R. S., Prada C., Putnam H. M., Quigley K., Reich H. G., Reimer J. D., Rosales S., Saad O. S., Santos S. R., Shoguchi E., **Stephens T. G.**, Strader M. E., Suggett D. J., Swain T. D., Tran C., Traylor-Knowles N., Voolstra C. R., Weis V., Wright R., Yamashita H., Ziegler M., Correa A. M. S., and Parkinson J. E. Building consensus around the assessment and interpretation of

symbiodiniaceae

diversity. *Under*

*review in PeerJ*,

2022. [[Preprint](#)]

- 2022           Bhattacharya D., Etten J. V., Benites L. F., and **Stephens T. G.** Endosymbiotic ratchet accelerates divergence after organelle origin. *Under review in BioEssays*, 2022.
- 2022           Meng Z., Williams A., Liao P., **Stephens T. G.**, Drury C., Chiles E. N., Su X., Javanmard M., and Bhattacharya D. Development of a portable toolkit to diagnose coral thermal stress. *Scientific Reports*, 12:14398, 2022. [[URL](#)]
- 2022           Benites L. F., **Stephens T. G.**, and Bhattacharya D. Multiple waves of viral invasions in Symbiodiniaceae algal genomes. *Accepted for publication in Virus Evolution*, 2022. [[Preprint](#)]
- 2022           \*Gabr A., \***Stephens T. G.**, and Bhattacharya D. Loss of key endosymbiont genes may facilitate early host control of the chromatophore in *Paulinella*. *iScience*,



25:104974,

2022.

[\[URL\]](#)

\*Co-first  
authorship

- 2022      **Stephens T. G.**, Lee J., Jeong Y., Yoon H. S., Putnam H. M., Majerova E., and Bhattacharya D. High-quality genome assemblies from key Hawaiian coral species. *Accepted for publication in GigaScience*, 2022.
- 2022      Bhattacharya D., **Stephens T. G.**, Tinoco A., Richmond R., and Cleves P. A. Life on the edge: Hawaiian model for coral evolution. *Limnology and Oceanography*, 67:1976-1985, 2022. [\[URL\]](#)
- 2022      \*Calatreva V., \***Stephens T. G.**, Gabr A., Grossman A. R., and Bhattacharya D. Retrotransposition facilitated the establishment of a primary plastid in the thecate amoeba *Paulinella*. *PNAS*, 119:e2121241119, 2022. [\[URL\]](#) \*Co-first authorship
- 2022      \*Gabr A., \***Stephens T. G.**, and Bhattacharya

D. Hypothesis:

*Trans*-splicing  
generates  
evolutionary  
novelty in the  
photosynthetic  
amoeba

*Paulinella*.

*Journal of  
Phycology*,  
58:392-405,  
2022. [[URL](#)]

\*Co-first  
authorship

2022

Dougan K. E., Gonzalez-Pech R. A., **Stephens T. G.**, Shah S., Chen Y., Ragan M. A., Bhattacharya D., and Chan C. X. Genome-powered classification of microbial eukaryotes: focus on coral algal symbionts. *Trends in Microbiology*, 30:831-840, 2022. [[URL](#)]

2022

Gabr A., Zournas A., **Stephens T. G.**, Dismukes G., and Bhattacharya D. Evidence for a robust photosystem II in the photosynthetic amoeba *Paulinella*. *New Phytologist*, 234:934-945, 2022. [[URL](#)]

2021

**Stephens T. G.**, Gabr A., Calatрева V., Grossman A. R., and Bhattacharya

D. Why is  
primary  
endosymbiosis  
so rare?. *New  
Phytologist*,  
231:1693-1699,  
2021. [[URL](#)]

- 2021 Bernard G., **Stephens T. G.**, Gonzalez-Pech R. A., and Chan C. X. Inferring phylogenomic relationship of microbes using scalable alignment-free methods. *Methods in Molecular Biology*, 2242:69-76, 2021. [[URL](#)]
- 2021 Jacobus A. P., **Stephens T. G.**, Youssef P., Gonzalez-Pech R., Ciccotosto-Camp M. M., Dougan K. E., Chen Y., Basso L. C., Frazzon J., Chan C. X., and Gross J. Comparative genomics supports that Brazilian bioethanol *Saccharomyces cerevisiae* comprise a unified group of domesticated strains related to cachaça spirit yeasts. *Frontiers in Microbiology*, 12:644089, 2021. [[Preprint](#)] [[URL](#)]
- 2021 Gonzalez-Pech,

R. A., **Stephens T. G.**, Chen Y., Mohamed A. R., Cheng Y., Shah S., Dougan K. E., Fortuin M. D. A., Lagorce R., Burt D. W., Bhattacharya D., Ragan M. A., and Chan C. X. Comparison of 15 dinoflagellate genomes reveals extensive sequence and structural divergence in family Symbiodiniaceae and genus *Symbiodinium*. *BMC Biology*, 19:73, 2021.

[\[Preprint\]](#) [\[URL\]](#)

- |      |  |
|------|--|
| 2021 | Williams A., Pathmanathan J. S., <b>Stephens T. G.</b> , Su X., Chiles E. N., Conetta D., Putnam H. M., and Bhattacharya D. Multi-omic characterization of the thermal stress phenome in the stony coral <i>Montipora capitata</i> . <i>PeerJ</i> , 9:e12335, 2021. <a href="#">[Preprint]</a> <a href="#">[URL]</a> |
| 2020 | Lhee D., Lee J., Ettahi K., Cho C. H., Ha J., Chan Y., Zelzion U., <b>Stephens T. G.</b> , Price D. C.,  |

Gabr A.,  
 Nowack E. C.  
 M.,  
 Bhattacharya  
 D., and Yoon  
 H. S. Amoeba  
 genome  
 reveals  
 dominant host  
 contribution to  
 plastid  
 endosymbiosis.

*Molecular  
 Biology and  
 Evolution*,  
 38:344-357,  
 2020. [[URL](#)]

2020

**Stephens T. G.**, Gonzalez-Pech R. A., Cheng Y., Mohamed A. R., Burt D. W.,  
 Bhattacharya D., Ragan M. A., and Chan C. X. Genomes of the dinoflagellate  
*Polarella glacialis* encode tandemly repeated single-exon genes with adaptive  
 functions. *BMC Biology*, 18:56, 2020. [[Preprint](#)] [[URL](#)] Featured by [IMB News](#)

2019

Chen Y., Gonzalez-Pech R. A., **Stephens T. G.**, Bhattacharya D., and Chan C. X.  
 Evidence that

inconsistent  
gene  
prediction  
can mislead  
analysis of  
dinoflagellate  
genomes.

*Journal of  
Phycology*,  
56:6-10, 2019.

[\[Preprint\]](#)

[\[URL\]](#)

- 2018      Liu H., **Stephens T. G.**, Gonzalez-Pech R. A., Beltran V. H., Lapeyre B., Bongaerts P., Cooke I., Aranda M., Bourne D. G., Foret S., Miller D. J., van Oppen M. J. H., Voolstra C. R., Ragan M.A., and Chan C.X. *Symbiodinium* genomes reveal adaptive evolution of functions related to coral-dinoflagellate symbiosis. *Communications Biology*, 1:95, 2018. [\[Preprint\]](#) [\[URL\]](#) Featured by multiple outlets: [IMB News](#), [GBRE](#), [Video Feature](#), [IMB 2018 year in review video](#)
- 2018      Lee J, Yang E. C., Graf L.,

Yang J. H.,  
Qiu H.,  
Zelzion U.,  
Chan C. X.,  
**Stephens T.**  
**G.**, Weber A.  
P. M., Boo G.  
H., Boo S.  
M., Kim K.  
M., Shin Y.,  
Jung M., Lee  
S. J., Yim H.  
S., Lee J. Y.,  
Bhattacharya  
D., and Yoon  
H. S.

Analysis of  
the draft  
genome of  
the red  
seaweed  
*Gracilariopsis*  
*chorda*  
provides  
insights into  
genome size  
evolution in  
Rhodophyta.  
*Molecular*  
*Biology and*  
*Evolution*,  
35:1869-1886,  
2018. [[URL](#)]

2018      \*Gonzalez-Pech R. A., \***Stephens T. G.**, and Chan C. X. Commonly misunderstood parameters of NCBI BLAST and important considerations for users. *Bioinformatics*, 35:2697-6998, 2018. [[URL](#)] \*Co-first authorship

[F1000  
recommended]

- 2018      **Stephens T. G.**, Ragan M. A., Bhattacharya D., and Chan C. X. Core genes in diverse dinoflagellate lineages include a wealth of conserved dark genes with unknown functions. *Scientific Reports*, 8:17175, 2018. [[URL](#)]
- 2016      **Stephens T. G.**, Bhattacharya D., Ragan M. A., and Chan C. X. PhySortR: a fast, flexible tool for sorting phylogenetic trees in R. *PeerJ*, 4:e2038, 2016. [[Preprint](#)] [[URL](#)]

## Talks

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- 2017 **Stephens T. G.**, Bhattacharya D., Ragan M. A. and Chan C. X. Insights into coral reef symbiosis from the genome of cold-adapted algae. *EMBL Australia Postgraduate Symposium 2017*, 29th November-1st December 2017, Sydney, Australia.
- 2018 **Stephens T. G.**, Bhattacharya D., Ragan M. A., and Chan C. X. *Polarella* genomics: understanding the evolutionary transition to algal symbiosis and cold adaptation. *Botany Department, Biosciences Institute, University of Sao Paulo*. 14th December 2018, Sao Paulo, Brazil.
- 2018 **Stephens T. G.**, Bhattacharya D., Ragan M. A., and Chan C. X. *Polarella* genomics:

understanding  
the  
evolutionary  
transition to  
algal  
symbiosis and  
cold  
adaptation.

*2nd Bioenergy  
Workshop,  
UNESP-USP-  
UNICAMP  
Integrated  
Postgraduate  
Program in  
Bioenergy,  
Institute for  
Research in  
Bioenergy,  
State  
University of  
Sao Paulo  
(UNESP). 6th  
December  
2018, Sao  
Paulo, Brazil.*

2019

**Stephens T. G.,** Bhattacharya D., Ragan M. A., and Chan C. X. *Polarella* genomics: understanding the evolutionary transition to algal symbiosis and cold adaptation. *Joint Academic Microbiology Seminars (JAMS)*. 9th April 2019, Brisbane, Australia.

2021

**Stephens T. G.,** Calatrava V., Gabr A., Grossman A., and Bhattacharya D. Insights into the evolution of a primary

endosymbiosis  
through  
analysis of the  
*Paulinella*  
genome. *12th*  
*International*  
*Phycological*  
*Congress*. 22-  
26th March  
2021, Chile.

2021

**Stephens T. G.**, Calatrava V., Gabr A., Grossman A., and Bhattacharya D. Insights into the evolution of a primary endosymbiosis through analysis of the *Paulinella* genome. *75th Annual Meeting of the Phycological Society of America*. 13-22nd July 2021, online.

2022

**Stephens T. G.**, Etten J. V., McDermott T., and Bhattacharya D. Analysis of environmental meta-omics data from the extremophilic red algae Cyanidiophyceae. *Joint Aquatic Sciences Meeting*. 14-20th May, 2022, Grand Rapids, USA.

2022

**Stephens T. G.**, Strand E. L., Putnam H. M., and Bhattacharya D. Differences in ploidy and the prevalence of clonal

propagation  
between  
*Montipora*  
*capitata* and  
*Pocillopora*  
*acuta* from  
Kane'ohe  
Bay,  
Hawai'i.  
15th  
International  
Coral Reef  
Symposium.  
3-8th July  
2022,  
Bremen,  
Germany.

## Poster presentations

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- 2018     **Stephens T. G.**, Bhattacharya D., Ragan M. A., and Chan C. X. *Polarella* genomics: understanding cold adaptation and evolutionary transition to symbiosis in dinoflagellates. *Society for Molecular Biology & Evolution (SMBE) annual meeting 2018*, 8-12th July 2018, Yokohama, Japan.
- 2016     **Stephens T. G.**, Chan C. X., and Ragan M. *Polarella* genomics: understanding the

evolutionary  
transition to  
algal  
symbiosis  
and cold  
adaptation,  
*IMB  
Research  
Higher  
Degree  
Student  
Symposium,*  
13th Jul  
2016, The  
University  
of  
Queensland,  
Brisbane,  
Australia.

## Grants

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- 2018 UQ-FAPESP Strategic Research Fund SPRINT (2018/15159-9), \$20,000 (~USD \$14,000)  
Integrated genomic approaches to understand stress tolerance in bioethanol-producing yeasts and coral reef symbionts  
Principal Investigators: Cheong Xin Chan (UQ) and Jeferson Gross (State University of Sao Paulo), Jointly funded by University of Queensland (UQ) and Sao Paulo State Foundation (FAPESP)
- 2022 Core Facility Utilization Application, \$5,000 USD  
Funds for sequencing of coral microbiome samples.  
Principal Investigators: Debashish Bhattacharya (Rutgers University), Rutgers University

2022 Center for Nutrition, Microbiome, and Health Small Grant FY-22, \$2,000 USD  
Characterizing the coral microbiome biogeography across colonies and reefs.  
Principal Investigators: Debashish Bhattacharya (Rutgers University), Center for  
Nutrition, Rutgers University

## Media coverage

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2020 Our paper [Amoeba Genome Reveals Dominant Host Contribution to Plastid Endosymbiosis](#) covered by [Rutgers Today](#)

2021 Our paper [Why is primary endosymbiosis so rare?](#) was covered by [Rutgers Newsroom](#).  
We also produced two animated videos: [Video 1](#), [Video 2](#)

2022 Our paper [Retrotransposition facilitated the establishment of a primary plastid in the thecate](#)

[amoeba](#)  
[Paulinella](#)  
covered  
by  
[Rutgers](#)  
[Research](#)

2022      Article by [Rutgers Research](#) on our short film [The Coral Holobiont Response to Climate Change](#) which won Best Trailer in the Kiez Berlin Film Festival.