

# Gregory Cary: curriculum vitae

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The Jackson Laboratory  
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Bar Harbor, ME 04609

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## **EDUCATION:**

- Ph.D. **University of Washington** (*December 2013*)  
Molecular & Cellular Biology Program
- A.B. **Colby College** (*May 2004*)  
*Major:* Biology; *Minor:* Chemistry

## **RESEARCH EXPERIENCE:**

- 2020 - present **Bioinformatics Analyst**, The Jackson Laboratory (JAX)  
*Advisor:* Gregory Carter, Ph.D.  
*Topic:* Supporting development of new therapeutic targets for Alzheimer's disease (Open-AD)
- 2017 - 2020 **Research Scientist**, Carnegie Mellon University (CMU)  
*Advisor:* Veronica Hinman, Ph.D.  
*Topic:* Echinoderm cis-regulatory sequence evolution and larval regeneration transcriptomics
- 2014 - 2016 **Postdoctoral Associate**, Carnegie Mellon University (CMU)  
*Advisor:* Veronica Hinman, Ph.D.  
*Topic:* Echinoderm cis-regulatory sequence evolution and larval regeneration transcriptomics
- 2009 - 2013 **Doctoral Student**, Institute for Systems Biology (ISB)  
*Advisor:* Aimee Dudley, Ph.D.  
*Topic:* Proteomic and transcriptomic assessment of post-transcriptional regulation in yeast
- 2006 - 2009 **Doctoral Student**, University of Washington (UW)  
*Advisor:* Albert La Spada, M.D., Ph.D.  
*Topic:* Polyglutamine neurodegeneration via transcriptional dysregulation
- 2004 - 2006 **Laboratory Technician**, Massachusetts General Hospital  
*Advisor:* Robert Brown Jr., M.D., D.Phil.  
*Topic:* Pre-clinical therapeutic trials in an ALS mouse model
- 2001 - 2004 **Undergraduate Research**, Colby College  
*Advisor:* Andrea Tilden, Ph.D.  
*Topic:* Effects of melatonin on the crustacean nervous system

## **TEACHING EXPERIENCE:**

- 2018, Fall **Special Faculty**, CMU - Genetics (03-220) (*50% of course*)  
**Lecturer**, CMU - Evolution of Regulatory Genomics (03-326): Genomics analyses (*2 lectures*)
- 2016, Spring **Adjunct Professor**, Chatham University - Biochemistry II (Bio439/539/Chem339)  
**Lecturer**, CMU - Evolution (03-125): Population Genetics (*7 lectures*)
- 2014, Fall **Lecturer**, CMU - Evolution (03-125): EvoDevo (*2 lectures*)
- 2008, Winter **Teaching Assistant**, UW - Advanced Cell Biology Lab (BIOL 402)  
2007, Fall **Teaching Assistant**, UW - Introductory Biology Lab (BIOL 200)
- 2004, Spring **Teaching Assistant**, Colby - Neurobiology Laboratory (BI274)  
**Chemistry Tutor**, Colby - General & Organic Chemistry
- 2009, 2014-18 **Undergraduate Research Mentor**, CMU and ISB - 15 students total

**PUBLICATIONS:** (<sup>[#]</sup> mentored undergraduate co-author; <sup>[†]</sup> mentored masters student co-author)

Analysis of sea star larval regeneration reveals conserved processes of whole-body regeneration across the metazoa  
**BMC Biol.** 2019. 17(1):16. DOI:[10.1186/s12915-019-0633-9](https://doi.org/10.1186/s12915-019-0633-9)  
Cary GA, Wolff A, Zueva O, Pattinato J<sup>[#]</sup>, Hinman VF

Genomic Resources for the Study of Echinoderm Development and Evolution  
**Methods Cell Biol.** 2019. 151:65-88 DOI:[10.1016/bs.mcb.2018.11.019](https://doi.org/10.1016/bs.mcb.2018.11.019)  
Cary GA, Cameron RA, Hinman VF

EchinoBase: Tools for Echinoderm Genome Analyses  
**Methods Mol Biol.** 2018. 1757:349-369. DOI:[10.1007/978-1-4939-7737-6\\_12](https://doi.org/10.1007/978-1-4939-7737-6_12)  
Cary GA, Cameron RA, Hinman VF

Echinoderm development and evolution in the post-genomic era  
**Dev Biol.** 2017. 427(2):203-211. DOI:[10.1016/j.ydbio.2017.02.003](https://doi.org/10.1016/j.ydbio.2017.02.003)  
Cary GA, Hinman VF

Genome-wide use of high and low affinity Tbrain transcription factor binding sites during echinoderm development  
**PNAS.** 2017. 114(23):5854-5861. DOI:[10.1073/pnas.1610611114](https://doi.org/10.1073/pnas.1610611114)  
Cary GA, Cheatle-Jarvela AM, Francolini RD<sup>[†]</sup>, Hinman VF

The evolution of gene regulation.  
**eLife.** 2017. 6. pii: e27291. DOI:[10.7554/eLife.27291](https://doi.org/10.7554/eLife.27291)  
Hinman V, Cary G

Proteomic analysis of Dhh1 complexes reveals a role for Hsp40 chaperone Ydj1 in yeast P-body assembly  
**G3 (Bethesda).** 2015. 5(11):2497-511. DOI:[10.1534/g3.115.021444](https://doi.org/10.1534/g3.115.021444)  
Cary GA, Vinh DBN, May P, Kuestner R, Dudley AM

Unidirectional P-body transport during the yeast cell cycle.  
**PLoS One.** 2014. 9(6):e99428. DOI:[10.1371/journal.pone.0099428](https://doi.org/10.1371/journal.pone.0099428)  
Garmendia-Torres C, Skupin A, Michael SA, Ruusuvuori P, Kuwada NJ,  
Falconet D, Cary GA, Hansen C, Wiggins PA, Dudley AM.

Identification and characterization of a drug sensitive strain enables puromycin-based translational assays in *S. cerevisiae*  
**Yeast.** 2014. 31(5):167-78. DOI:[10.1002/yea.3007](https://doi.org/10.1002/yea.3007)  
Cary GA, Yoon SH, Torres CG, Wang K<sup>[#]</sup>, Hays M, Ludlow C, Goodlett DR, Dudley AM.

Melatonin: neuritogenesis and neuroprotective effects in crustacean x-organ cells.  
**Comp Biochem Physiol A Mol Integr Physiol.** 2012. 161(4):355-60. DOI:[10.1016/j.cbpa.2011.12.005](https://doi.org/10.1016/j.cbpa.2011.12.005)  
Cary GA, Cuttler AS, Duda KA, Kusema ET, Myers JA, Tilden AR

Androgen receptor function in motor neuron survival and degeneration. (Review)  
**Phys Med Rehabil Clin N Am.** 2008. 19(3):479-94. DOI:[10.1016/j.pmr.2008.03.002](https://doi.org/10.1016/j.pmr.2008.03.002)  
Cary GA, La Spada AR

**DATASETS:**

Genome assembly of *Strongylocentrotus purpuratus* (GCA\_000002235.4)  
In: **Genbank** [Internet]. Bethesda, MD: NCBI Assembly; 2019 September.  
NCBI:Assembly: GCA\_000002235.4.

Regeneration time course of a larval sea star, *Patiria miniata*  
In: **Gene Expression Omnibus** [Internet]. Bethesda, MD: NCBI GEO; 2018 December.  
NCBI:GEO: GSE97230.

Genome-wide use of high and low affinity Tbrain transcription factor binding sites during echinoderm development

**In: Gene Expression Omnibus** [Internet]. Bethesda, MD: NCBI GEO; 2016 December.

**NCBI:GEO: GSE89865. [super-series]**

**NCBI:GEO: GSE89862. [ChIP-seq]**

**NCBI:GEO: GSE89863. [RNA-seq]**

Proteomic and transcriptomic analyses of Dhh1 complexes reveals mitochondrial RNP complex association with yeast P-bodies.

**In: Gene Expression Omnibus** [Internet]. Bethesda, MD: NCBI GEO; 2015 December.

**NCBI:GEO: GSE65989.**

## **CONFERENCE & MEETING PRESENTATIONS:**

**Talk:** Enhancing EchinoBase to Support Echinoderm Genomics and Regulatory Analyses  
2018 Developmental Biology of the Sea Urchin meeting XXV - Woods Hole, MA

**Talk:** Using Echinoderms for Youth Outreach and Undergraduate Genomic Education  
2018 Developmental Biology of the Sea Urchin meeting XXV - Woods Hole, MA

**Poster:** Regulatory Evolution of Cell Type Specification in Echinoderms  
2017 Society for Molecular Biology and Evolution - Austin, TX

**Talk:** Evolution of transcription factor binding site preference and contribution to GRN rewiring  
2017 Developmental Biology of the Sea Urchin meeting XXIV - Woods Hole, MA

**Talk:** Conserved features of metazoan whole-body regeneration  
2016 Elizabeth Jones Annual Retreat - Hidden Valley, PA

**Poster:** Conserved transcriptional changes during metazoan whole-body regeneration  
2016 Society for Developmental Biology meeting - Boston, MA

**Talk:** RNA-seq profiling of larval echinoderms highlights conserved features of metazoan regeneration  
2015 Developmental Biology of the Sea Urchin meeting XXIII - Woods Hole, MA

**Poster:** RNA-seq profiling of larval echinoderms highlights conserved features of metazoan regeneration  
2015 Comparative Biology of Tissue Repair, Regeneration, & Aging Symposium - Bar Harbor, ME

**Poster:** Evolution of Gene Regulatory Networks for Neurogenesis in Echinoderms  
2014 Systems Developmental Biology Symposium - Pittsburgh, PA

**Poster:** Systems-level analyses of yeast RNA granules as a model for function in higher eukaryotes  
2013 Systems Biology Symposium - Seattle, WA

**Poster:** Systems biology approaches for dynamic post-transcriptional regulatory processes  
2011 Eukaryotic mRNA Processing Meeting - Cold Spring Harbor, NY

**Poster:** TAPAS: a new technique for translation state analysis  
2010 Centers for Systems Biology Meeting - Seattle, WA

## **SERVICE ACTIVITIES:**

**Scientific Advisory Board:** Echinobase.org web information system 2020-present

**Software, Database, and Systems Manager:** Echinobase.org web information system 2016-2019

**Judge,** Undergraduate Research Symposium, CMU 2014-2019

**Public Outreach,** Hinman Lab/CMU Gelfand Center, Ocean Life (Grades K-2) 2018, Spring

**Public Outreach,** Hinman Lab/CMU Gelfand Center, Diversity in the Ocean (Grades 7-9) 2018, Winter

**Public Outreach,** Tour Your Future careers in STEM, CanTEEN Career Exploration 2015, '16 Fall

**Graduate Student Committee,** Institute for Systems Biology 2010-2013

**Panelist,** STEM Education Panel, Institute for Systems Biology Retreat 2011, Fall

**Public Outreach,** Life Science Research Weekend, NWABR 2010, Fall

**Peer Review:** BMC Genomics, Communications Biology, Nature Communications, Mechanisms of Development, Development Genes and Evolution, Archives of Microbiology

**WORKSHOPS ATTENDED:**

<b>NSF Grants Conference</b> , National Science Foundation; Pittsburgh PA	<i>2016, Fall</i>
<b>XSEDE HPC Workshop: Introduction to Bridges</b> , Pittsburgh Supercomputing Center	<i>2016, Spring</i>
<b>Proteomics Informatics Workshop</b> , Institute for Systems Biology	<i>2009, Spring</i>
<b>TA Conference on Teaching and Learning</b> , UW Center Instructional Development and Research	<i>2007, Fall</i>

**AWARDS AND HONORS:**

Graduated <i>summa cum laude</i> and with Honors in Biology, Colby College	<i>2004</i>
BRIN-ME Undergraduate Research Fellowship	<i>2003</i>
Colby College Presidential Scholarship	<i>2000-2004</i>

**SOCIETY AFFILIATIONS:**

Society for Molecular Biology and Evolution  
Society for Developmental Biology  
American Association for the Advancement of Science