Tenure Promotion

Ecology & Evolutionary Biology

Dr. Brian O'Meara

Fall 2014

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A. SUMMARY

SUMMARY SHEET

Recommendations for Promotion and/or Tenure

Name of faculty member: Brian (O'Meara			
Present rank: Assistant Profess	or			
Department: Ecology and Evolution	onary Biology Highes	t degree earned: _	Ph.D.	
Original UTK rank: Assist. P	rof.	Subsequent pro	omotions (year, ran	k):
UTK RECORD				
Date of original UTK appointment	t as a full-time probation	onary faculty mem	ber: 1 August	t 2009
Years of full-time teaching experie	ence at instructor rank	or above before U	TK probationary pe	riod: 0
Years of full-time teaching at UTk	X, as of the May 31 st pr	ior to the review:	5	
Total years of teaching: 5 Lates	t year for tenure review	as stipulated in a	ppointment letter: 2	2014-2015
RECOMMENDATIONS				
DEPARTMENTAL FACULTY Date of departmental discussion: Result of discussion: For: Recuse (attach explanation for condition of the condition of the candidate of	3 December 20 9 Against:	0 Abst N/A [X] No ttach) [2 COR (where appro (Provide letter) al [] Do n	X] No ppriate) ot recommend appr	
COLLEGE COMMITTEE For: Against: Recuse (attach explanation for cor A copy of the report of the departn report disagrees in any substantia listing of the vote to indicate as ful	nflict of interest): nental and college com I way with the departm	mittees must also ental recommendo	ation, this report mu	
DEAN	[] Approve	[]]	Disapprove (Provide	e letter)
CHIEF ACADEMIC OFFICER	[] Approve	[]]	Disapprove (<i>Provide</i>	e letter)
CHANCELLOR'S RECOMME	NDATION TO THE		Disapprove (<i>Provide</i>	e letter)

EDUCATIONAL HISTORY AND EMPLOYMENT HISTORY

Candidate Name: Brian Christopher O'Meara

Educational History

Institution	Program or Degree	Dates in Program	Degree
University of California, Davis	Ph.D.	2002-2008	Ph.D.
Harvard University	B.A. with magna	1997-2001	B.A.
	cum laude and highe	et honore in higlagy	

cum laude and highest honors in biology

Employment History

Ranks Held	<u>Institution</u>	<u>Department</u>	Effective Date of Rank
Assistant Professor	University of Tennessee	Ecology and Evolution	2009- present
Postdoc	NESCent		2008 - 2009

Department of Ecology and Evolutionary Biology

STATEMENT OF RESPONSIBILITIES

Dr. Brian O'Meara

I. Teaching

To receive tenure, a member of the faculty of the Department of Ecology and Evolutionary Biology (EEB) is expected to develop excellence in teaching at either or both of the undergraduate and graduate levels. It is expected that the teaching load of an individual faculty member be reviewed periodically in order to adjust for different levels of research and service activities. There is no set teaching load.

Dr. O'Meara initiated 400-level course in Macroevolution in his first year and has offered it every year since. It has grown from eight students to approximately thirty, including graduate students from the Geology and Ecology & Evolutionary Biology departments as well as undergraduate students majoring in Chemistry, Geology, Ecology & Evolutionary Biology, and Animal Science. This class features a mixture of lecture, clicker questions, discussions, student presentations, and use of statistical software for simulations. He has also taught in EEB511, the team-taught core course in evolution, every fall (and is one of only three faculty in our department to have done so). In Fall 2013 he became the instructor of record for this course. He has also taught Biology 130, Biodiversity, in Spring 2012 (206 students) and in Spring 2014, and has been involved in discussions regarding restructuring this course to meet core learning objectives of Biology. Dr. O'Meara has been an instructor in EEB607 twice, EEB503 twice, EEB504, and EEB511 twice. He has also guest lectured in courses in Sweden, Austria, Louisiana, North Carolina, Tennessee, and Switzerland (in addition to research talks).

II. Research

A junior member of the faculty of EEB is expected to develop a significant, externally funded program of original research in his or her area of expertise. Regular publication is expected, with at least some papers being in high-quality journals.

Dr. O'Meara is conducting research about macroevolutionary processes leading to current diversity. Much of his research involves the development and testing of phylogenetic methods. He has received grants of \$132,345, \$138,590, and \$98,252 (sole recipient) from the iPlant Collaborative (NSF-sponsored), a grant of \$141,143 from NSF as Co-PI (\$28,228 for his use), a grant of \$320,000 from NSF as PI (with additional funds going to Co-PI Bryan Carstens), a grant of \$520,000 from NSF as PI (the grant total includes funding for Co-PI Gilchrist), plus various smaller awards, generally to researchers he mentors, from places such as Google Summer of Code and the Encyclopedia of Life. He has published numerous peer-reviewed papers in high impact journals (*Nature*, *PNAS*), including invited syntheses of the field in journals such as the *Annual Review of Ecology, Evolution, and Systematics*.

III. Service

A junior member of the faculty of EEB is expected to develop a record of service to the department and the profession.

Locally, Dr. O'Meara serves on the web committee as sole faculty representative as well as the graduate admissions committee and the department head search committee; in the past he has also served on the undergraduate education committee. He served as a member of the Dean's Advisory Council for three years. He organized a multi-department speaker series on women in science using Haines-Morris funds as well as wrote a successful proposal to acquire a cart of laptops for teaching in the EEB department. Dr. O'Meara is faculty advisor for Darwin Day Tennessee. Dr. O'Meara has mentored ten postdocs while at Tennesee, including approximately 19% of all NIMBioS postdocs.

Internationally, he is an elected member of the Society of Systematic Biologists Council, a member of the Phylotastic Hackathon leadership team, organizer of lightning talks for the Evolution 2013 meeting in Snowbird, Utah (1400 participants from three international societies), co-organizer of the Evolution 2014 meeting in Raleigh, NC (nearly 2000 participants from three international societies), member of the Steering Committee for the iEvoBio meetings, and an associate editor for *Methods in Ecology and Evolution*. He has reviewed for journals and organizations such as NSF, *Science*, *Evolution*, *Systematic Biology*, and many more. Dr. O'Meara is the Society of Systematic Biologists' liaison to three-society initiative for removing barriers to women in science and pursues other work on removing barriers for women in science). He has also taught and organized numerous workshops in phylogenetics and computing, locally, nationally, and internationally.

B. TEACHING ABILITY AND EFFECTIVENESS

STATEMENT OF TEACHING PHILOSOPHY AND DESCRIPTION OF ITS IMPLEMENTATION

My overall teaching goal is to create students who have the skills of creative, skeptical scientists, curious about evolutionary processes, while being grounded with deep knowledge of biological patterns and mechanisms. This is accomplished through teaching a mixture of large introductory lecture courses, graduate seminars and courses, and most centrally, a mixed upper level undergraduate/introductory graduate student course in macroevolution.

My Macroevolution course (EEB464) was started in my first year here and has been growing since. The class explicitly targets four Biology degree-level learning outcomes ("Evolution: Populations of organisms and their cellular components have changed over time through both selective and non- selective evolutionary processes", "Formulate empirically-testable hypotheses", "Interpret visual representations (figures and diagrams)", and "Evaluate data and come to a conclusion (with evidence) (formulate an argument)"). The class is a mixture of lecture, small group discussions, class discussions, and investigations, both computational and empirical. Technology (laptops, clickers, homemade interactive white board, embedded videos) is used as appropriate to engage students and assess progress, but students also get hands on experience looking at fossils or other biological specimens. Lectures are recorded for students to review later if needed. For class discussions, I often will have them break into small groups to talk about an issue and then come together for an entire class discussion, so that students may try out their ideas on one or two peers before voicing them to the entire class. I also encourage class discussions to feature dialog with each other, rather than just response to me. Over the course of a semester the class covers various topics in macroevolution such as differential diversification and the history of life on earth. I emphasize how we know about these things, how to perform experimental tests of these ideas, and current work on the topics, so students see science as an ongoing process of discovery rather than a static set of facts. Students are assessed through clicker questions (some reviewing past topics, some based on a list of key taxa students are assigned to learn, some just to have students commit to hypotheses about data presented in class), essay-based tests, a report on a topic including what work still has to be done on it, and a pair presentation covering a graduate-level research proposal. In addition to positive reviews and growing numbers, one measure that shows the class's impact is the number of graduate students from the EEB and Geology departments who are encouraged by their committees to take it. Committees justifiably want to protect students from taking courses at the cost of their research, but several faculty feel that the content in this course is worth the time it takes for students to take it.

I do extensive teaching at the graduate level. Some of this is for classes that are reading groups, while another is for a shared multiple lab discussion group. This started as simply a joint meeting of four small labs (for example, my lab was originally just me) but is now an active discussion group with students from at least seven lab groups enrolled. It covers a mixture of work in progress as well as recent relevant papers to students' research. My goal in all these classes is to encourage students to focus on the questions being addressed by their work or others: it is easy for students to adopt popular methods without considering whether those are the best way to address their questions. I have also been heavily involved in our Evolution core course, being one of only three faculty to teach in the Evolution or Ecology core each of the past six years. I have been working in that course to have students focus on understanding the questions that can

be addressed and why someone might want to rather than focusing on the minutia of current methods. I try to teach students to see the connections across methods (for example, all the methods that use a discrete state transition matrix, even though some are for inferring trees while others are for investigating biogeographic history). We have also worked to make the core an early way for students to get feedback on their projects before they even set up their committees, which may have some role in EEB's remarkable success at getting NSF graduate research fellowships or honorable mentions (one third of all UT Knoxville students receiving such honors had been through the core course; two additional EEB students won honors before taking the core).

In 2012 and 2014 I taught the large introductory biology class Bio130, Biodiversity. I use readings, clicker questions, MasteringBiology assignments, and parts of lectures to teach basic material, and the remaining parts of lectures to having students learn to think about the material at higher levels of understanding. In addition to anonymous feedback during the class (see below), I embedded observers in the class to see how students were taking notes and engaging with the lectures. However, I think I was only partially successful at achieving my goals in this class. The presentation style I chose to use, with slides with primarily figures, images, and videos accompanying a lecture rather than a more bullet point-heavy presentation was a style many students had trouble with, especially in taking notes (one embedded observer found that the only note a student took from a hybridization lecture was the word "labradoodle"). As the course went on, I modified my slides (which were made available to students as pdfs and as recorded lectures) to include "speaker's notes" describing what was said and the point of each slide. I also assigned more homework, which students described as helping their understanding of the material. My clarity of speech was also difficult for some students and is clearly something to continue to improve. Successes in the class included transmitting enthusiasm for the subject (based on the student evaluations, as well as anecdotes such as students asking to bring a boyfriend or relative to observe the class) as well as creating an environment where students felt comfortable asking and answering questions: i.e., in just one 50 minute lecture, approximately 20 of 200 students volunteered in whole-class discussion. Students rated my enthusiasm high (4.25/5 average). Overall, I think I targeted the class at too high a level of understanding without providing enough basic framework for students. I will be incorporating these lessons when teaching the course again this spring through changes like including more structuring text in slides to make sure students recognize key points. I have also been involved in the ongoing discussion regarding class structure and content organized by Beth Schussler to align with the overall Biology framework.

Due to my expertise in the field I have frequently been asked to teach in short workshops (this is in addition to research talks). I often give these virtually to prevent disruption of regular duties as well as the expense and hassle of travel. I have taught in Sweden, Austria, Louisiana, North Carolina, and Switzerland. I also organized and taught at a workshop at NIMBioS on computational resources for phylogenetics. This course included students from around the world and had a fairly even gender balance.

I have certain approaches I try to extend across all my classes. One is real time assessment of my teaching: I give students a link to an anonymous form (http://www.brianomeara.info/feedback) they can use to give feedback on any part of the class at any time. This allows me to improve the class for the students in it rather than just waiting for SAIS reports at the end. I also try to use all the time students are in the room to promote learning, rather than just the assigned class times. For example, some students may show up half an hour early for some classes, and I typically have an educational video relevant to the day's focal topic running they can learn from while waiting. One consistent theme in student reviews is my enthusiasm for the subject matter. Based on student reactions, my classes are also seen as a safe place to ask questions about confusing material.

Mentoring of graduate students and postdocs is also an important part of my teaching. I have had ten postdocs come through my lab: some from internal funds (startup or NSF) and others as comentored NIMBioS postdocs. I also advise three graduate students and co-advise a fourth. In addition to my own students, I have served on the committees of eleven other graduate students in three graduate programs at UTK and one at Vanderbilt. With all of them, my goal has been to help the student or postdoc achieve her or his professional goals. This may require more teaching experience for some and more work on programming skills for others. I try to create a supportive environment where people are encouraged to propose new ideas but also know that they will receive constructive, honest feedback.

SUMMARY OF TEACHING ASSIGNMENTS FOR REVIEW PERIOD

Semester	Dept. Course #	Course Title	Credit Hours	14-day enrlmnt	% Respon	Lec (L)/ Lab (B)	# GTAs assisting	Honors Y/N
Fall 2009	EEB511*	Core	4	7	13	L	0	N
Spring 2010	EEB409	Perspectives	3	13	100	L	0	N
Spring 2010	EEB607	Speciation	1	13	100	L	0	N
Fall 2010	EEB511*	Core	4	8	13	L	0	N
Spring 2011	EEB503	Seminar	1	35	100	L	0	N
Spring 2011	EEB607	Speciation	1	9	100	L	0	N
Fall 2011	EEB464	Macroevolution	3	24	100	L	0	N
Fall 2011	EEB504*	HOFF	1	7	25	L	0	N
Fall 2011	EEB503	Seminar	1	44	100	L	0	N

Fall 2011	EEB511*	Core	4	12	13	L	0	N
Spring 2012	Bio130	Biodiversity	3	206	100	L	1	N
Fall 2012	EEB464	Macroevolution	3	22	100	L	0	N
Fall 2012	EEB511*	Core	4	14	25			
Fall 2012	EEB504*	HOFF	1	5	25	L	0	N
Spring 2013	EEB607*	HOFF	1	8	25	L	0	N
Fall 2013	EEB464	Macroevolution	3	30	100	L	0	Y
Fall 2013	EEB504*	HOFF	1	11	25	L	0	N
Fall 2013	EEB511*	Core	4	19	50	L	0	N
Spring 2014	EEB602	Phyloseminar	1	24	100	L	0	N
Spring 2014	EEB607	HOFF	1	6	25	L	0	N
Spring 2014	Bio130	Biodiversity	4	93	100	L	0	N
Fall 2014	EEB464	Macroevolution	3	28	100	L	0	Y
Fall 2014	EEB504	HOFF	1	9	33	L	0	N
Fall 2014	EEB511	Core	4	12	40	L	0	N
Fall 2014	EEB607	Speciation	1	15	100	L	0	N

^{*}My responsibility for the Core course varies from 13% to 50%. My involvement for the HOFF course varies from 25% to 33%.

SAIS REPORTS

Sem/ Year	Course/ Hrs	No. of Students	No. of Responses			Instructor's Contribution	.,
Spring 2010	EEB409/3	13	11	3.55	3.73	4.09	3.55
Spring 2010	EEB607/1	9	6	3.67	3.83	4.00	3.83

Fall 2011	EEB464/3	24	17	4.24	4.18	4.35	4.24
Fall 2011	EEB503/1	44	18	4.00	3.88	4.25	4.25
Spring 2012	Bio130/3	206	102	3.05	3.14	3.04	2.67
Fall 2012	EEB464/3	22	14	4.36	4.29	4.64	4.36
Spring 2013	EEB607/1*	5	5	4.80	4.80	5.00	5.00
Fall 2013	EEB511/4*	19	11	2.64	2.91	3.55	3.27
Fall 2013	EEB464/3	30	15	4.00	4.20	4.40	4.20
Spring 2014	Bio130/3	93	28	3.04	3.14	3.43	3.07
Spring 2014	EEB602/1	24	8	4.38	4.50	4.88	4.75

Range: 0-5; 5 = excellent, 0 = very poor

PEER REVIEW OF TEACHING

Sandy Echternacht has reviewed my teaching; Randy Small has also attended one of my lectures for Bio130. In Spring 2014 Mike Gilchrist, Ed Schilling, and Sally Horn evaluated my teaching.

SUMMARY OF STUDENT COMMENTS

To be supplied by Dept. Head.

GRADUATE SUPERVISION

Summary of Graduate Dissertation (Doctoral) Supervision

^{*}Courses with asterisks are team-taught.

Katie Massana	Started Fall 2012	Statistical biogeography
Jen Bosco (advisor	Started Fall 2012	Evolution of spider
Riechert; I am co-advisor)		behavioral traits
Orlando Schwery	Started Fall 2014	Phylogenetic methods

Summary of Graduate Dissertation (Masters) Supervision

Name	Graduation Year	Dissertation Title	Placement
Sam Borstein	Started Fall 2013	Cichlid evolution	

GRADUATE COMMITTEES

Name	Dept.	Degree	Project Title	Date Completed
Phillip Hollingsworth	EEB	PhD	Fish evolution	
Ivan Juric	EEB	PhD	Population evolution	
Leonidas Salichos	Vanderbilt	PhD	On the identification of factors influencing phylogenetic accuracy	
Will Atwood	Geology	PhD	Examining Pentremitidae: Species Discrimination, Systematics, and Ontogeny	
Mauricio González Forero	EEB	PhD	Major evolutionary transitions mediated by manipulation	
Sharon Clemmensen	EEB	PhD	Fish evolution	
Will Howell	EEB	PhD	Protein evolution	
Max Rupp	EEB	Masters	Fish evolution and behavior	
Troy Fadiga	Geology	PhD	Echinoderm evolution	
Geetha Saarunya S	GST	PhD	Function and evolution of permafrost bacteria	
Jordan Utley	GST	PhD	Rapid detection of pathogens	
Marisol Sanchez	EEB	PhD		

POSTDOCTORAL MENTEES

Name	Funding
Hugo Alamillo	Startup
Barb Banbury	iPlant
Jeremy Beaulieu	NIMBioS
JJ Chai	NIMBioS
Nathan Jackson	NSF
Tony Jhwueng	NIMBioS
Sandy Kawano	NIMBioS
Michelle Lawing	NIMBioS
Ryan Martin	NIMBioS
Nick Matzke	NIMBioS

C. RESEARCH, SCHOLARSHIP, CREATIVE ACTIVITY

CANDIDATE'S STATEMENT

My research addresses key questions in macroevolution to understand the processes leading to present diversity and disparity. I do this through construction and extensive testing of new approaches and the implementation of these in user-friendly software, as well as by examination of particular biological examples. These papers aim to be major works that move the field rather than mere examples of things we already know. One focus in my work has been dealing with heterogeneity of biological processes. The factors affecting oak tree evolution probably differ from those affecting water lily evolution, but nearly all methods apply the same set of parameters to all taxa being investigated. My work on dealing with different evolutionary rates (O'Meara et al. 2006) has been cited 241 times and has been used for things as varied as fish jaw evolution and flower size. This work has continued to deal with heterogeneity of multiple parameters for a single trait on a tree (Beaulieu et al. 2012). Work on discrete characters (O'Meara 2007) has also been extended to deal with hidden rates (Beaulieu et al. 2013). In the lab we have also created a general forward in time simulation model that was featured in an invited Evolution symposium in 2012. I have also coauthored other publications using or testing various phylogenetic methods (Farrell et al. 2001; Driskell et al. 2004; Collar et al. 2009; Smith and O'Meara 2009; Collar et al. 2010; Abercrombie et al. 2011; Hulsey et al. 2013; Soltis et al. 2013; Jhwueng et al. 2014; Williams et al. 2014).

I have also worked on species delimitation (O'Meara 2010). An NSF grant based on using these techniques, with others, on fungal datasets, with PI Karen Hughes and me as one of the CoPIs, was recently funded. I have continued to develop the method, which was presented at a different Evolution symposium in 2013. I was also PI on a preproposal and proposal based on this work which resulted in a grant of \$340,000 for UTK as well as additional funds for my CoPI Bryan Carstens. This is funding another postdoc in my lab; we are currently finishing simulations for a manuscript we will submitting to *PNAS*.

Another aspect of research effort has been on extending phylogenetic models. From collaborations with JJ Chai, a NIMBioS postdoc, and Mike Gilchrist, a faculty member in EEB at UTK, I have helped to create codon and amino acid models which use explicit parameters about optimal amino acids, strength of selection, and amino acid similarities to create transition

matrices for phylogenetic inference with realistic features such as different rates for different optimal amino acids and unequal gain and loss rates for amino acids or codons (it is more likely to evolve towards a codon representing the optimal amino acid than away from it, for example). We have recently been awarded \$520K from the NSF to continue this work (me as PI, Dr. Gilchrist as Co-PI) and are readying a publication on the approaches. Our source code for the model is an R package available on github.

Another creative activity has been making sure that other biologists can do analyses. This has resulted in papers on re-usability of trees (Stoltzfus et al. 2012), hackathons to create new tools (Stoltzfus et al. 2013), and cyberinfrastructure for phylogenetics (Goff et al. 2011), for which my lab received a total of \$369,187 from three sets of funding. We have also received \$50,000 from the Encylopedia of Life as a Rubenstein fellowship to Barb Banbury with summer funding for me as well for an R interface to the Encyclopedia of Life (Banbury and O'Meara 2014). Work has also involved developing fast algorithms for analyses (Smith and O'Meara 2012) and new implementations of my software (Stack et al. 2011) (funded by Google Summer of Code).

I have also been involved in reviews, often invited, of comparative methods or phylogenetics, including a review of methods for the *Annual Review of Ecology, Evolution, and Systematics* (O'Meara 2012) (which was the most downloaded paper in the journal), a chapter on phylogenetic reconstruction in the *Encylopedia of Theoretical Ecology* (Hastings and Gross 2012), a review of phylogenetic methods in R (O'Meara 2014), and book chapters (Beaulieu and O'Meara 2014; O'Meara and Beaulieu 2014) as well as other invited works in prep.

There is other work in the pipeline that is not listed here. For example, I am lead author on a manuscript on floral evolution that pioneers use of multiple characters simultaneously to look at their joint effect on diversification and transition and show the importance of non-equilibrium processes in structuring life; this was reviewed and rejected for *PNAS* and is being revised for a mid-tier journal. I am revising a manuscript for *Systematic Biology* on extending comparative methods for a network. By the time this tenure packet is evaluated (November 2014), I expect to have a manuscript about settlement bias of plants on Hawaii and a manuscript about a new general diversification model also submitted. A paper questioning the widespread belief that

extinction cannot be estimated from neontological phylogenies is currently being revised *Evolution*.

RESEARCH AND SCHOLARY PUBLICATIONS

- My work is highly cited, with 1138 citations overall, 925 since 2009 alone.
- Abercrombie, J. M., B. C. O'Meara, A. R. Moffatt, and J. H. Williams. 2011. Developmental evolution of flowering plant pollen tube cell walls: callose synthase (CalS) gene expression patterns. EvoDevo 2:14.
- Banbury, B. L. and B. C. O'Meara. 2014. Reol: R interface to the Encyclopedia of Life. Ecology and Evolution.
- Beaulieu, J. M., D.-C. Jhwueng, C. Boettiger, and B. C. O'Meara. 2012. Modeling Stabilizing Selection: Expanding the Ornstein-Uhlenbeck Model of Adaptive Evolution. Evolution 66:2369-2383.
- Beaulieu, J. M. and B. C. O'Meara. 2014. Hidden Markov Models for Studying the Evolution of Binary Morphological Characters. Pp. 395-408 in L. Z. Garamszegi, ed. Modern Phylogenetic Comparative Methods and Their Application in Evolutionary Biology. Springer-Verlag, Berlin.
- Beaulieu, J. M., B. C. O'Meara, and M. J. Donoghue. 2013. Identifying hidden rate changes in the evolution of a binary morphological character: the evolution of plant habit in campanulid angiosperms. Systematic biology.
- Collar, D. C., B. C. O'Meara, P. C. Wainwright, and T. J. Near. 2009. Piscivory Limits

 Diversification of Feeding Morphology in Centrarchid Fishes. Evolution 63:1557-1573.

- Collar, D. C., J. A. Schulte, B. C. O'Meara, and J. B. Losos. 2010. Habitat use affects morphological diversification in dragon lizards. Journal of Evolutionary Biology 23:1033-1049.
- Driskell, A. C., C. Ane, J. G. Burleigh, M. M. McMahon, B. C. O'Meara, and M. J. Sanderson. 2004. Prospects for building the tree of life from large sequence databases. Science 306:1172-1174.
- Farrell, B. D., A. S. Sequeira, B. C. O'Meara, B. B. Normark, J. H. Chung, and B. H. Jordal. 2001. The evolution of agriculture in beetles (Curculionidae: Scolytinae and Platypodinae). Evolution 55:2011-2027.
- Goff, S. A., M. Vaughn, S. McKay, E. Lyons, A. E. Stapleton, D. Gessler, N. Matasci, L. Wang, M. Hanlon, A. Lenards, A. Muir, N. Merchant, S. Lowry, S. Mock, M. Helmke, A. Kubach, M. Narro, N. Hopkins, D. Micklos, U. Hilgert, M. Gonzales, C. Jordan, E. Skidmore, R. Dooley, J. Cazes, R. McLay, Z. Lu, S. Pasternak, L. Koesterke, W. H. Piel, R. Grene, C. Noutsos, K. Gendler, X. Feng, C. Tang, M. Lent, S.-J. Kim, K. Kvilekval, B. S. Manjunath, V. Tannen, A. Stamatakis, M. Sanderson, S. M. Welch, K. A. Cranston, P. Soltis, D. Soltis, B. O'Meara, C. Ane, T. Brutnell, D. J. Kleibenstein, J. W. White, J. Leebens-Mack, M. J. Donoghue, E. P. Spalding, T. J. Vision, C. R. Myers, D. Lowenthal, B. J. Enquist, B. Boyle, A. Akoglu, G. Andrews, S. Ram, D. Ware, L. Stein, and D. Stanzione. 2011. The iPlant Collaborative: Cyberinfrastructure for Plant Biology. Frontiers in plant science 2:34-34.
- Hastings, A. and L. J. Gross. 2012. Encyclopedia of theoretical ecology. University of California Pr.
- Hulsey, D. C., B. P. Keck, H. Alamillo, and B. C. O'Meara. 2013. Mitochondrial genome primers for Lake Malawi cichlids. Molecular ecology resources 13:347-353.

- Jhwueng, D.-C., S. Huzurbazar, B. C. O'Meara, and L. Liu. 2014. Investigating the performance of AIC in selecting phylogenetic models. Statistical applications in genetics and molecular biology.
- O'Meara, B. C. 2007. Estimating Different Rates of Gene Loss on a Tree. Genetics 177:1415-1416.
- O'Meara, B. C. 2010. New Heuristic Methods for Joint Species Delimitation and Species Tree Inference. Systematic Biology 59:59-73.
- O'Meara, B. C. 2012. Evolutionary Inferences from Phylogenies: A Review of Methods. Annual Review of Ecology, Evolution, and Systematics 43.
- O'Meara, B. C. 2014. CRAN Task View: Phylogenetics. Version 2014-07-17, URL http://cran.r-project.org/web/views/Phylogenetics.html.
- O'Meara, B. C., C. Ane, M. J. Sanderson, and P. C. Wainwright. 2006. Testing for different rates of continuous trait evolution using likelihood. Evolution 60:922-933.
- O'Meara, B. C. and J. M. Beaulieu. 2014. Modelling Stabilizing Selection: The Attraction of Ornstein–Uhlenbeck Models. Pp. 381-393 in L. Z. Garamszegi, ed. Modern Phylogenetic Comparative Methods and Their Application in Evolutionary Biology. Springer-Verlag, Berlin.
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- Stoltzfus, A., B. O'Meara, J. Whitacre, R. Mounce, E. L. Gillespie, S. Kumar, D. F. Rosauer, and R. A. Vos. 2012. Sharing and re-use of phylogenetic trees (and associated data) to facilitate synthesis. BMC research notes 5:574-574.
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PROJECTS, GRANTS, COMMISSIONS, AND CONTRACTS

Since starting at UTK a bit over five years ago, I have turned in 28 grants through Tera-PAMS; this includes 21 I have submitted through NSF Fastlane (this includes 16 full proposals and 5 preproposals). Of full NSF proposals which have had decisions (several are pending), I have a 23% success rate; for preproposals, I have a 60% success rate. While at UTK I have been

awarded two NSF grants as a PI, one as a Co-PI, plus \$369,187 in funding from the NSFsponsored iPlant Collaborative as a working group lead, totaling \$1.4M in NSFderived funds as an assistant professor. NSF funding rates are between five and ten percent in the programs I apply to, so this represents an unexpected string of successes. As funding amounts vary by discipline, for the convenience of the Promotion and Tenure Committee and other evaluators I have attempted to construct a baseline, using appropriate aspirational peer departments for our Ecology and

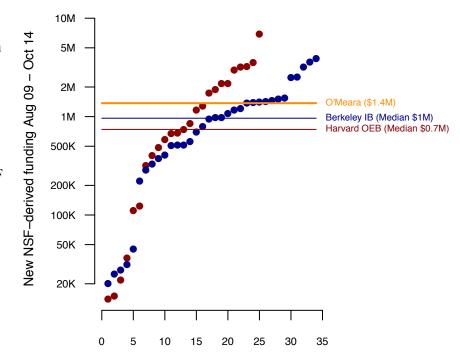


Figure C-1: Funding at appropriate aspirational peer departments for UTK EEB. This uses public data from NSF (acquired Oct. 7, 2014). It includes all faculty listed on peer departments' webpages who have received NSF funding in the time window; faculty with no returned funding were excluded from calculation of the median or plotting. Note that this plot does not include funding from NIH, EPA, foundations, or other sources, though in ecology and evolutionary biology departments non-NSF sources are relatively infrequent. For faculty who have entered into either peer department during the focal time period, the totals include all their funding, even that awarded when they were at different institutions.

Evolution department (see Fig. C-1). Compared to faculty (at all levels, not just assistant professors) in Harvard's Organismic and Evolutionary Biology department and UC Berkeley's Integrative Biology department, my new NSF-derived funding is well above the median for that of faculty in either department for the relevant time period (85% and 42% higher, respectively). There are various caveats to that figure (for example, it does not include non-NSF funding; it includes full faculty but also some emeritus faculty (though only if they got NSF funding in the interval)), but it suggests at the least that by the standards of my field I have been fairly successful at acquiring funding. I have also been responsible for mentoring other proposals to Encyclopedia of Life (funded) and Google Summer of Code (funded). Funds for working groups at NIMBioS and NESCent (five while an assistant professor), funds for workshops or hackathons not part of research grants, and large group proposals in which I am not at least a Co-PI (such as a recent grant to NSF's ADVANCE program) are not included in that total or below.

External Funding

Completed

2013: Encyclopedia or Life: "Reol, an R interface to Encyclopedia of Life data." PI Barb Banbury, Subcontractor Brian O'Meara. **\$50,000.**

2010-2012: NSF/iPlant: "iPlant-iPToL" PI Mike Sanderson, Co-PIs Michael Donoghue, Pam Soltis, Doug Soltis, Val Tannen, Alexis Stamatakis, Todd Vision. I ran the trait evolution working group, which included two months of summer salary for me and funding for a postdoc for this year and next year. Note that I was not part of the initial proposal (thus not an official PI/co-PI), but was invited to become part of it once it was funded and I function as a co-PI. Funds from NSF via iPlant via University of Arizona. \$132,345 to UTK in year 1, \$138,590 in year 2 for \$270,935 in total.

2012-2013: NSF/iPlant: "rPlant" funding for a postdoc and summer salary based on a white paper I submitted. \$98,252

2012: Google: "Optimizing R code for approximate Bayesian computing" Google Summer of Code 2012. **\$5000** by Google to pay for U of Nebraska graduate student Daniel Gates to work on speeding up code I wrote. Note that my postdoc, Barb Banbury, and a colleague at U. of Arizona, Derrick Zwickl, handled the mentoring. Funds went directly to the student, not through UTK.

2010: Google: "Ancestral State Reconstruction in R" Google Summer of Code 2010. **\$5000** by Google to pay for Penn State graduate student Conrad Stack to work on putting my program Brownie into the R programming language. I was primary mentor, Luke Harmon was co-mentor. Funds went directly to the student, not through UTK.

Funded and In Progress

2014: NSF: "Population Genetics-based Codon Models." PI Brian O'Meara, Co-PI Mike Gilchrist. Amount for UTK: **\$520,000**. Funding years 2014-2016.

2013: NSF: "Phylogeographic inference using approximated likelihoods." PI Brian O'Meara, Co-PI Bryan Carstens. Amount for UTK: **\$340,000**. Funding years 2013-2015.

2012: NSF: "Historical naming traditions and cryptic speciation bias biodiversity estimates in transatlantic agaric fungi." PI: Ronald Petersen. Co-PIs Brian O'Meara and Karen Hughes. Amount requested for UTK: **\$141,143** [though note that the bulk of this goes to Petersen and Hughes, with just summer salary going to me]. Funding years 2012-2014

Under review

(Note that this section still needs to be updated)

Successful Pre-proposals

Some NSF programs now require a preproposal to be approved before a full proposal is invited; invitation rates are approximately 25%.

2013: NSF: "Preproposal: Population Genetics-based Codon Models". PI Brian O'Meara, Co-PI Mike Gilchrist, Co-PI Russell Zaretzki.

2011: NSF: "Preproposal: Phylogeny and diversification of Lake Malawi Cichlid Flock". PI Darrin Hulsey, Co-PI Brian O'Meara.

2011: NSF: "Preproprosal: Phylogeographic inference using approximated likelihoods". PI Brian O'Meara, Co-PI Bryan Carstens.

Unsuccessful grant proposals

2013: Templeton Foundation: Evolution of Play. PI Gordon Burghardt, Co-PI Brian O'Meara. Declined. Budget: \$199,912.

2012: NSF: "Phylogeny and Diversification of Lake Malawi Cichlid Flock". PI Darrin Hulsey, Co-PI Brian O'Meara. Declined. Budget: \$626,470.

2012: Templeton Foundation: Evolution of Play. PI Gordon Burghardt, Co-PI Brian O'Meara. Declined. Budget: \$263,763.

2011: NSF: "Collaborative Research: Phylogeographic Inference using approximated likelihoods" PI Brian O'Meara, Co-PI Bryan Carstens. Declined. Amount requested for UTK: \$329,790

2012: NSF: "Collaborative Research: Assembly the Tree of Life: Interactive Heuristics in Tree Search (inHits)". PI Michael Sanderson, Co-PI Michelle McMahon, Co-PI Derrick Zwickl, Co-PI Brian O'Meara, Co-PI Oliver Eulenstein, Co-PI David Fernandez-Baca, Co-PI Stephen Smith. Declined. Amount requested for UTK: \$220,115 (approx. \$2M in funding for entire grant)

2011: NSF: "Making comparative methods as easy as ABC" PI: Brian O'Meara. Declined. Amount requested for UTK: \$301,016

2011: NSF: "Preproposal: Participation in AVAToL ideas lab" PI Brian O'Meara. Declined.

2011: NSF: "Dimensions: Collaborative Research: How do Plant-Insect Interactions Generate Biodiversity? -A Case Study of Euphorb Vines (Euphorbiaceae: Plukenetieae) and Their Insect Associates" proposal to NSF Dimensions of Biodiversity. Modification and resubmission of above grant. PI Scott Armbruster, Co-PI Bruce Baldwin. Senior collaborators: André Freitas, Niklas Wahlberg, Silvana Buzato, Narcísio Costa Bigio, Ricardo Secco, Lynn Gillespie, Christophe Pélabon, Thomas Hansen, Hans Stenoien, Terry Griswold, Bryan Danforth, Brian O'Meara. Declined. Amount requested for UTK: \$298,826

2010: NSF "SI2-SSE: Phylogenetic needs discovery environment". PI: Brian O'Meara. Declined. Amount requested for UTK: \$454,353

2010: NSF: "Phylogeny and Diversification of Lake Malawi Cichlid Flock" PI Darrin Hulsey, Co-PI Brian O'Meara. Declined. Amount requested for UTK: \$527,895

2010: NSF: "Making comparative methods as easy as ABC" PI: Brian O'Meara. Declined. Amount requested: \$275,288

2010: NSF: "Dimensions: Collaborative Research: How do Plant-Insect Interactions Generate Biodiversity? -A Case Study of Euphorb Vines (Euphorbiaceae: Plukenetieae) and Their Insect Associates" proposal to NSF Dimensions of Biodiversity. PI Scott Armbruster, Co-PI Bruce Baldwin. Senior collaborators: André Freitas, Niklas Wahlberg, Silvana Buzato, Narcísio Costa Bigio, Ricardo Secco, Lynn Gillespie, Christophe Pélabon, Thomas Hansen, Hans Stenoien, Terry Griswold, Bryan Danforth, Brian O'Meara. Declined. Amount requested for UTK: \$205,836

OTHER EVIDENCE OF RESEARCH OR SCHOLARSHIP

I am the maintainer for the CRAN Phylogenetics Task View. This is an overview of the implementations of phylogenetic methods in the popular programming language R. According to statistics compiled by Bob Muenchen, R is used in over 14,000 scientific papers per year and its help list has over 13,000 subscribers. I am one of only 31 maintainers of task views for all these users.

While many people who develop new techniques merely publish them, one focus of my research is making well-tested software implementations available. This software is all open source, allowing others to improve upon it. Some of this software includes:

Lampyr: A web app for teaching about biodiversity. http://lampyr.org

corHMM: R package for dealing with discrete trait evolution. http://cran.r-

project.org/web/packages/corHMM/index.html

OUwie: R package for dealing with continuous trait evolution. http://cran.r-project.org/web/packages/OUwie/

Brownie: C++ software for species delimitation and trait evolution.

http://brianomeara.info/brownie

DateLife: Web service and underlying software for estimating divergence dates for taxa. http://datelife.org/

Reol: R package for extracting information from the Encyclopedia of Life. http://reolblog.wordpress.com/

rPlant: R package to interact with NSF-sponsored high performance computing. http://cran.r-project.org/web/packages/rPlant/index.html

phrapl: R software for testing phylogeographic hypotheses. https://r-forge.r-project.org/projects/phrapl/

PUBLIC TALKS

Please also see the discussion of workshops taught for other talks; this section only includes scientific talks about my research, not teaching at other venues.

- June 24, 2014: "Non-equilibrium dynamics lead to long-term persistence of ancestral floral forms in ancestral floral forms in modern angiosperms" talk at Evolution 2014 meeting; coauthors were S Smith, W SArmbruster, L Harder, C Hardy, L Hileman, L Hufford, A Litt, S Magallon, S Smith, P Stevens, C Fenster, P Diggle.
- June 24, 2014: "Phylogeographic model selection using approximated likelihoods" talk at Evolution 2014 meeting; lead author and speaker was Nathan Jackson, other authors were A. Garcia, B. Carstens, and B. O'Meara.
- June 23, 2014: "Non-null effects of a null range: Exploring parameter estimation in the dispersal-extinction-cladogenesis model" talk at Evolution 2014; lead author and speaker was Katie Massana (grad student), coauthors were J. Beaulieu, B. O'Meara, and N. Matzke.
- June 23, 2014: "Estimating how contemporary taxa will evolve in the future to understand how island communities were assembled in the past" talk at Evolution 2014; lead author and speaker was Jeremy Beaulieu, coauthor was Brian O'Meara.
- May 13, 2014: "Dealing with the heterogeneity of life for comparative methods" Smithsonian Phylopizza (attended by biologists throughout the DC area, including NSF; this was their best-attended seminar to date)
- June 22, 2013: Invited symposium talk at Evolution on "Simulated likelihood for species delimitation and phylogeography"
- Aug 17, 2012: Invited talk on "Making comparative methods as easy as ABC" to the U. of Georgia Institute of Bioinformatics
- July 9, 2012: Talk at iEvoBio on "DateLife: When lineages meet"
- July 7, 2012: Invited symposium talk at Evolution on "Approximate Bayesian computation for continuous characters"
- April 17, 2012: Invited talk on "Making comparative methods as easy as ABC" at the Smithsonian.
- April 16, 2012: Invited talk on "Making comparative methods as easy as ABC" as one of three speakers for U. of Maryland's "Organismal Biology Day"
- Oct. 3, 2011: Invited departmental talk on "Making comparative methods as easy as ABC" at U. of Florida

Sept. 15, 2011: Invited departmental talk on "Making comparative methods as easy as ABC" at U. of Alabama

May 1, 2011: Invited talk on "Phylogenetics" to Rationalists of East Tennessee

March 30, 2011: Invited talk on "Making comparative methods as easy as ABC" at phyloseminar.org (meeting hosted online)

Nov. 8, 2010: Invited talk on "Phylogenetics" at iPlant site visit by NSF

May 24, 2010: Invited talk on "Phylogenetics" at iPlant community outreach meeting in Las Vegas.

April 26, 2010: "Species delimitation": Invited departmental talk at Louisiana State University.

November 3, 2009: Invited talk on "Species delimitation" at National Institute for Biological and Mathematical Synthesis

Oct. 14, 2009: Invited talk on my history in Science at National Institute for Biological and Mathematical Synthesis

D. INSTITUTIONAL, DISCIPLINARY, AND/OR PROFESSIONAL SERVICE

CANDIDATE'S STATEMENT

I contribute to service within the department, throughout the University, and in the broader academic community. Within the department I serve as the head of the web committee, and for years maintained the department website personally, including 76 blog posts with department news. I was the point contact for the recent move of the department web site to new templates by the UT Communications group, and I continue to oversee our website, including our extension into social media. While doing this service, I have also served on the undergraduate curriculum committee and, currently, on the graduate admissions committee. I am also an active participant in departmental questions such as the search for the next head or ways to improve our core course.

In the College and University, I contribute through serving on the Dean of Arts and Sciences' Advisory Committee, both by giving feedback on various questions the Dean poses as well as, when appropriate, communicating ideas from the Dean to our department's faculty. I also am the faculty advisor for Darwin Day Tennessee, a long-running student-organized event that successfully educates students and members of the local community about evolution and related topics. Last year we invited Nobel Prize winner Camille Parmesan to campus to talk about evolution and climate change, while also educating local teachers through a workshop. I organized a cross-department discussion and speaker series about women in science issues, funded via an internal Haines-Morris grant.

I contribute extensively to service in the broader scientific community. I am an elected member of the Council for the Society of Systematic Biologists, publisher of one of the most highly cited journals in evolutionary biology. I am a member of the Phylotastic hackathon leadership team, a group that arranges funding for developers to meet to work together on coding solutions to problems in evolutionary biology. I have been in two invited symposia at our field's major international meetings, the Evolution meetings. I initiated and organized the lightning talks at the most recent Evolution meeting and have been asked to do it at the next Evolution meeting. I have recently been asked to join the leadership team of the iEvoBio meeting, a meeting that overlaps with the Evolution meeting and which focuses on bioinformatics research for evolutionary biology. I review for NSF, Science, and many other journals. I also have a strong online media presence as a scientist, with 52 YouTube videos of my talks and lectures (watched a total of 3,282 minutes by the public, and available at https://www.youtube.com/user/bcomeara), 459 followers on Twitter (including faculty from around the world, and available at https://twitter.com/omearabrian), and prominent placement of my personal website in search results (for example, for a widely used statistical approach, the Akaike Information Criterion (AIC), my personal website is the second listing on Google and third listing on Bing; my website overall received 3,200 hits last month alone).

INSTITUTIONAL SERVICE

Record of committee work at department, college, and university levels

Served on strategic planning committee

Served on seminar planning committee

Served on website committee: created department blog, updated website.

Served on undergraduate curriculum committee

Served on graduate admissions committee

Wrote and received grant for laptop teaching cart for EEB

Dean's Advisory Committee

DISCIPLINARY SERVICE

Record of membership and active participation in professional societies Society of Systematic Biologists, elected member of Council

Evaluation of peer research-

I have been a reviewer for Science, NSF, Systematic Biology, Bioinformatics, Evolution, Nucleic Acids Research, Ecology, and iEvoBio, among others.

E. APPENDICES

All comments from classes with electronic SAIS returns (classes 2012 or later), and at least 5 responses (which excludes three HOFF classes only) are listed below.

Key to color scheme:

Bio130: Biodiversity (2014 and 2012)

EEB602: Phyloseminar (2014)

HOFF (2013)

EEB464: Macroevolution (2013 and 2012)

EEB511: CORE Evolution (2013). Note that this was team taught: the SAIS scores reflect the average for the team, while the comments have been pruned to only include those referring to my section.

Year	Semester	Course	Question	Response
2014 Spring Bio130	Spring	Bio130	1: Intellectually	yes
		stimulating	Yes. The information that was presented offered new insight on topics that I learned in high school	
			It did stimulate my mind. It made me think about all living things as a whole, and everything be affected by greater causes (like evolution, genetics, etc.).	
				yes. yes.
			Yes and Yes	
		This class was intellectually stimulating, but was more fact-based, rather than concept-based.		
			It was a very intriguing and complicated subject, great teacher though	
			This class was intellectually stimulating and it required that I applied what I learned.	
		Yes, excellent examples, everything was explained really well and relative to contemporary Biology. Enthusiasm was good, and in general had a helpful attitude.		
			Not too much. it was pretty basic concepts, but sometimes I was asked to think outside the box	
			It was intellectually stimulating, yes. I enjoyed learning more about how living things came to be and how they continue to develop.	
			yes, I especially like when he opens the class up for discussion.	
				Biology does not really interest me that much. Biology in humans is somewhat interesting but definitely the biology of plants.
				it was but it was still some what difficult to understand.
		2: Greatest contribution to learning	content	
			The instructor was very animated and took time to explain key concepts within the course material. The online questions were also beneficial to fully understanding the current topics.	

Reading the book contributed most to my learning. lecture Conversing with the teacher Studying for tests. Dr. O'Meara showed us videos before class and used some as examples of concepts which I found really helpful. They are both interesting and informative. The teacher is phenomenal Studying and going to lab. Tying what we were learning to what actual Biologists do today (but I just enjoy that kind of thing). Also, that review questions before exams really helped so I knew what I didn't know. The open dialogue we had in every lecture The entire course rounded out and brushed me up on my biology. The study guide was probably the most helpful thing even though about 3/4 of the questions were not on the exam. 3: Detracted what we learned did not show on the slides from learning Too many power points Sometimes the professor talked too fast, but when addressed, this problem was fixed. The lectures themselves seemed disorganized and confusing. I had trouble following along during class. speed The labs The speed with which the instructor spoke. Some parts of lecture felt boring and unnecessary. I would rather hear about examples or more information than take ten minutes to "discuss" basic concepts with people in the class. Not being able to understand what Dr. O'Meara was saying. Sometimes, the lectures would go a little too specific and I would have trouble paying attention. I think a little more general and maybe more rapid movement through the material might have worked better for me. Sometimes the slides were really boring Other students not paying attention or talking while the professor is teaching. nothing The professor mumbled a lot and talked really fast so it made it hard to understand him. He would also cough a lot into the microphone. 4: Suggestions studyguide Examinations should be over less chapters. Then exams are also not as general as they should be when covering a mass amount of information. Have more clear, concise lectures. Work on speech clarity

Make the class and lab separate

There is too much information on the slides. Almost like they contain as much information as the book. It's almost impossible to follow along.

Get rid of normal curve. It does not make sense - logically or morally - and is especially frustrating for trying to figure out grades.

A different microphone or speaking slower.

Maybe a little more warning before the exams, and a little more help preparing, or emphasis for things to look at (like memorizing a plant phylogeny). I studied a lot and they were still really difficult. Also, as I said ^^ maybe lectures that don't go into detail about such specific topics.

less words on the slides. and slowing down wayyyy more.

Excellent class overall!

I wish that the week could consist of going over the chapters and then on Friday we would have an open discussion over all the chapters we read. I understand that this might be an unreasonable suggestion due to scheduling.

Talk slower

I think that you should have more clicker questions to encourage students to come to class and the powerpoints that you upload maybe leave some parts blank to also encourage attendance and insure that they are following along.

I feel like most of the exam is not talked about in class. We covered so much information and then had to learn and retain all that information. It was too much for this class. I think the amount of information should be cut down so that it can be further explained and better learned.

2014 Spring

EEB602

1: Intellectually stimulating

Yes, this class was effective in getting me familiar and thinking about different methods in phylogenetics that I haven't used before.

I was able to learn more about how systematics was done especially in computer based techniques. It really opened my eyes to areas in which I need to gain further understanding and ability.

Yes, some of the topics were new and I learned a lot

Yes, the course was intellectually stimulating. It made me think of different ways we could use phylogenetics to answers questions of scientific interest. The class was awesome and I wish it would continue. The instructor is very knowledgeable and helpful with clarifying doubts and ambiguities from the publications.

This course exposed me to a world of

paleogeographic/biogeographic/phylomimetic/phylogenetic models that I never would have been exposed to had I not taken this course. Drs. O'Meara, [and two postdocs, names redacted] displayed extraordinary knowledge of the subject matter and this instilled much confidence in those taking the course of the quality of the course being taken. The topics discussed were challenging to me, as I am not a phylogeneticist by training. However, I am happy to have taken the course and believe that I can use what I have learned to improve upon my own research and this makes me happy.

2: Greatest contribution to

Enjoyed the discussions, to be sure. I learned a lot both from listening and asking questions.

learning

The fact that there were people from several disciplines and levels of understanding all asking questions from different viewpoints. Also, toward the end of the semester we looked at some papers that were applying the methods to different disciplines, so i think that everyone got quite a bit out of this course for relatively little time commitment.

The dicussions clarified most of the readings. A couple of times the readings and the discussion topics were beyond any of my prior experience, but It was still worth sitting on the periphery of those discussions and eventually receive a modicum of comprehension.

The explanations given during the class, reading amazing papers

Reading through the publications and discussing them afterward.

Multiple times, great detail was given to discussing the mechanism of a tree-building algorithm or evolutionary model and this made using the algorithm or model in context of the research article much clearer.

3: Detracted from learning

not having enough time to read the papers. often sent out too late to do more than a thorough skim, as opposed to reading them well.

Nothing

while extemporaneous discussions about various aspects of the papers produces a good discussion, I would have occasionally liked a more structured discussion with some topics from the papers to discussed layed out beforehand, and occasionally some better direction through the discussions.

None

There were participants in the class that at times detracted from my learning with random interruptions, or criticism that was not constructive.

At various times during each discussion, the discussion tended to stray from the main points at hand and would be longer than I would have liked. Certain aspects of the underpinnings of a tree-building algorithm or evolutionary model were discussed in some detail, but at other times this was lacking and detracted from take-home message to be had from the article (e.g., is this appropriate for my research, and if not, how could I adapt it to be). Dr. O'Meara is brilliant but sometimes speaks very swiftly. Quick replies made it hard to understand what he had said. He has answered the question asked, but it was so quick that many times I missed what I would have liked to have heard because it was probably worth its weight in gold.

4: Suggestions

A little more structure would have been helpful, set learning goals for the week or so.

Nothing that I can think of. Offer it again!

Make sure all students contribute to discussions. I felt a core group of students contributed to discussions, but there were a number of students who may have spoken only once in the semester and a small few who never contributed.

It is good the way it is, continue organizing this type of seminars, they are very helpful

				I would suggest to send out the publication or publications of interest at least 3-4 days beforehand. It was hard to find time to read everything and understand it when the reading material was sent 1-2 days before. Also, maybe allow people enrolled in the class to suggest papers (they need not be selected, but nice if this option is available). It would be good to have objectives that we should get from each paper (not necessary though). Make an outline of the key points that need to be discussed for the paper, get through them, and then if there needs to be extra discussion let people have a free for all. It seemed a bit chaotic and disorganized as times because of the lack of progression in the discussion.
2013	Spring	EEB607	1: Intellectually stimulating	Yes-there is a wide range of content we discuss, and a diverse group of participants who bring up details more specific to their own background. this makes the discussions very stimulating typically.
				Yes-Yes, this discussion group/lab meeting was stimulating, and challenged me to think about things "outside the box," so to speak. We read papers and discussed their strengths and weaknesses which helped better myself as a scientist. Brian always tries and succeeds in making the discussions intellectually stimulating.
				Yes-It is always nice to see what other people are doing
			2: Greatest contribution to learning	content and format (ie discussions rather than presentations)
				Discussing research project ideas and picking apart papers really helped me learn more about current methods and interesting hypotheses in science.
				I was able to bounce some ideas off peers for my research
			3: Detracted	NA
			from learning	Nothing in particular detracted from my learning.
				The class time was not great. By 5 pm everyone is slowing down and wants to be doing something else. Also, I got sick of trees. The bifurcating kind.
			4: Suggestions	keep it open as a class section and continue to allow anyone to show up. i think this helps bring in more students than just HOFF lab members.
				-Sending out papers (if a person is presenting one) earlier than 1-2 days aheadHaving discussion questions to keep the discussion going.
				Switch the timing. Early is better than late.
2013	Fall	EEB464	1: Intellectually stimulating	It did! I enjoyed your class very much. Your lectures are engaging and the material is good-I like your use of journal articles to teach real examples and applications. I love how much effort you put into it!
				Yes, Lots of good examples and opportunities for critical thinking and discussion
				Yeah, I actually learned quite a bit in this class, it just sucked a lot of the time some of the material was a bit new and I had not much to say most of the time or it was discussing data types not previously encountered and I did not always know how to interpret the data.
				Yes, although much of what was in the course was stuff that I had been taught before.
				There were concepts approached in ways that I had not thought about before. I enjoyed the class discussions.

It was a great class.

Yes, I had to switch gears from a case by case kind of study in Animal Science to a much broader, more encompassing thought process. It was definitely different from what I am used to.

Yes, this was my first evolution course at UT, and I felt it addressed topics relevant in the field today. It was difficult for me to think in terms of evolution, as I primarily study ecology.

Yes, great powerpoints and discussion.

The course did present many new topics and the way it was run required you to come up with many of your own ideas.

Yes it was and yes it did.

2: Greatest contribution to learning

The lectures were helpful, but I feel when we discussed the material I learned much more.

The lecture part of the class without the frequent student contributions.

Your enthusiasm really makes this class easy to learn! you always sound so excited about the topic, so it in turn makes me want to learn about it too!

I really liked the presentations where we voted on the topic that we wanted to learn about. I liked the one on building models, because it made you think of different ways of testing hypotheses.

Class discussions

The lecture format was great. Dr. O'Meara was a great professor

Listening to the other students helped tremendously since I had very little background in EEB.

Examples, running through simulations, learning macroevolutionary study techniques

How well the teacher broke down the material.

3: Detracted from learning

I feel like there were a few irrelevant topics that made me kind of bored and not really wishing to pay attention. I think Darwin and the other people we have discussed should have had there material discussed not so much their lives.

I don't think as much time should be devoted to the students voicing ideas in each class period. A separate class period, or one per week might be beneficial to have as an open discussion lecture in which the students can share thoughts, ideas, and ask questions. It seems like a lot of each period was filled with quietly asked questions that sometimes had no answer other than acknowledgement of the response being correct, or periods of silence.

Too many questions trying to get student involvement. A lot of time can get wasted waiting for students to answer...I like letting students participate a lot, but I think there can be too much. I know this is tough, but sometimes your enunciation is difficult to understand. You speak very quickly, which is fine, but the words can often run together

Lack of powerpoints to review after class/lag time in uploading of presentations. Open ended assignment directions, so I didn't know how much or how little was expected of me. The in-class discussions were interesting, but not always necessary. Often we don't respond to questions because it seems silly to state the obvious.

Not having specific background reading. I did enjoy researching certain things on my own, but some concepts would have been better with background reading.

Maybe the amount of participation involved.

None that I can think of.

Occasionally, Dr. O'Meara was difficult to understand because he spoke quickly and sometimes without articulating fully.

Group presentation preparation period was a bit rushed (we were assigned partners less than 2 weeks before presentation)

I could almost NEVER understand the professor. He speaks very fast and mumbles. He does recored the lectures and put them on line which is nice of him but often during the lecture he walks away from his computer so you cannot hear anything he his saying. Also a lot of the class involves discussion where he will as a general question and get a response from students. This is a good way to stimulate thinking but I very often could not hear other students responses and Dr. O'Meara's response would be something vague like yes or why? but I could get nothing out of it because if I understood his original question I likely didn't understand the other students response. Also questions were often posed that were not clearly answered. I understand that it is a discussion class but if no one give a good answer it would be beneficial to have the professor provide useful information. Finally, the requirements for most assignments were very vague and hard to judge exactly what needed to be done and grading of assignments was very slow so it was hard to judge how you were doing in class and if there was anything you needed to improve upon.

No aspect distracted me.

4: Suggestions

Maybe a readings folder for further readings for the students relevant to the class topic should they want to learn more.

A peer review for the papers might be useful so students can develop their ideas better. Maybe some short essay homework assignments to reinforce main ideas in macroevolution in some digital form or online discussion group via Blackboard (these are frequently suggested as a method to increase participation for many classes, but rarely used as a graded part which might encourage its utility).

I'd like to suggest making the class a bit more challenging. I enjoyed it- but the assignments given were very minimal. I was really looking forward to coding in class like we talked about, or getting really into the topic reviewbut there wasn't enough space. I had a hard time getting motivated to spend so much time researching new things for a 1,000 word review. I know there are time-limits to your grading, but this is where I'd suggest to the university that for a class this size, you deserve a TA to help with grading! Also, I was a bit disappointed over the clickers-a lot of us had to purchase them for this class only and it's near impossible to get one used, so I had to pay 50 at the bookstore to use it a very limited number of times. I'd suggest either making their use mandatory and much more frequent, or stick to paper quizzes.

Instead of skimming the surface of a lot of topics, go into topics in more detail. Actually building a model in R probably made lots of people confused if they were not familiar with it. I would not spend class time writing code, but instead focus on conceptually how you would build one (on paper). The parameters of the model you created and what it was doing was not obvious.

Add assigned reading for difficult topics and do more with R (and get everyone involved like was planned). Also, maybe have a graded topic earlier in the semester... I was stressed out until I got my midterm back! Clickers are the devil.

More lecturing less group work.

I think it would be beneficial if the student answers were recorded in the posted lectures. It would certainly help when reviewing them if we could hear both halves of the conversation.

At the beginning of the course, detail how macroevolution is studied, giving concrete examples: showing packages, softwares, and using real data. I struggled to understand HOW to address macroevolutionary questions because the methods were always unclear to me.

More groupwork

Please speak more clearly. This is by far my biggest and most important recommendation. Also providing more assignments or spreading out the current assignments so they are not all in the second half of the semester would be helpful in the learning process. And providing more and quicker feedback would be very helpful.

Keep doing what you're doing.

2013 Fall

EEB511

1: Intellectually stimulating

This class was intellectually stimulating. [comments about other instructor(s) redacted] Brian's section of the class has been really great. I am getting something from it, and it actually seems interesting to me (a surprise).

Yes, it was very stimulating as it offered relevant knowledge for modern research in evolution and phylogenetics.

[comments about other instructor(s) redacted] Phylo: Intellectually stimulating in that concepts were applied to real scenarios, but I wouldn't say it stretched my thinking per-say (maybe because I already had a background in phylogenetics)

It was, Both the Population Genetics and Phylogenetics was new material to me.

Yes. [comments about other instructor(s) redacted] Brian's section seemed more organized, clear, and challenging. [comments about other instructor(s) redacted]

No

Yes

Yes and no. Brian O'Meara and [postdoc assisting this section] did an excellent job at this. The material they taught was intellectually challenging yet had very clear goals, applications and usefulness. [comments about other instructor(s) redacted]

2: Greatest contribution to

Brian's section, where we did more "real life" examples.

Lectures and quizzes

learning

[comments about other instructor(s) redacted] Phylo: Examples used in lecture. Thought questions given during lecture to evaluate understanding before moving on to other material. Clear learning objectives and immediate feedback about quiz answers.

[comments about other instructor(s) redacted] The powerpoint slides used by Brian were intuitive and a great learning aide. Having a group of people to look over my NSF GRFP was a great component of this class. All instructors were very helpful in one-on-one meetings with me. They were all willing to schedule times outside of normal office hours. Brian in particular made it very easy to come and talk to him by posting his weekly schedule online.

There were two professors who co-taught this class, each with their specialty subject: [other faculty member] and O'Meara (phylogenetics). I put fair on all the evaluations because I would grade these two teachers very differently. For O'Meara's portion I would put excellent or very good on everything. He is organized and prepared for each lecture and clearly communicates the learning objectives and requirements. Always willing to engage students, O'Meara offered unique methods to encouraging class participation and was willing to assist our learning in any way he could. He also encouraged feedback and discussion. The subject is a difficult one, but O'Meara communicated complex topics in a very understandable way. In short, he is a great teacher with many great qualities (organized, attentive, intelligent but relatable).

The phylogenetics portion of this class was very informative and useful. [comments about other instructor(s) redacted] Just listening to and reading the slides or board from both instructors was by far the best part of CORE. Both professors were very knowledgeable and where able to explain the topics in a clear manner.

Reading, discussion in class

The in-class activities with Brian and Jeremy, i.e. using R, BEAST, RAxML etc. programs, practicing with interpreting trees, question and answer...

3: Detracted from learning

[comments about other instructor(s) redacted]

[comments about other instructor(s) redacted]

[comments about other instructor(s) redacted]

[comments about other instructor(s) redacted] Phylo: The mini project did not really contribute much to my learning (too short of a time frame to get what you wanted us to get out of it). Not that it detracted from my learning per-say, but it stole valuable lecture time away from learning more-loading programs and running some of the R code (took to long for everyone to get set up)- maybe have everyone load programs before class then we can jump right into using them during class time.

[comments about other instructor(s) redacted]

				[comments about other instructor(s) redacted] If this course is designed to get students "up to speed" on current research directions of evolution, I'd say Brian/[assisting postdoc] did a great job, [comments about other instructor(s) redacted] I feel like I could sit down with someone now and tell them WHY phylogenetics is important (Brian/[assisting postdoc]'s section) and how to work with it in various applications [comments about other instructor(s) redacted]
			4: Suggestions	[comments about other instructor(s) redacted]
				I really enjoyed thinking about a phylogenetics project that was presented as an exercise near the end of the course. Going a few steps further into this would be great.
				Mentioned above.
				[comments about other instructor(s) redacted]
				[comments about other instructor(s) redacted]
				[comments about other instructor(s) redacted]
				[comments about other instructor(s) redacted] I REALLY enjoyed the "project" that Brian has had us work on (i.e. ask a phylogenetics research question, work as a group to solve the question using the programs/resources we learned in class). It would be cool if we could have a little more time/incentive (graded?) to complete this project and write up results. That is also a potential publicationif a group is really interested in their topic they could potentially pursue publishing a manuscript on the subject, much like courses such as Field Ecology where students routinely get a publication out of a paper in the course. [comments about other instructor(s) redacted] The BIG PICTURE [] was definitely there for Phylogenetics however.
2012	Spring	Bio130		Yes-It made me do a lot more critical thinking than any other class.
2012			stimulating	Yes-I like biology
2012				1 to 1 mile elelegy
2012				No-Because I was force to take this class by the university
2012				
2012				No-Because I was force to take this class by the university
2012				No-Because I was force to take this class by the university Yes-Biology is very complex and interesting, but very hard. Yes-I am a Biology major, so I enjoyed the class content. I also liked how Mr. O'Meara used examples and showed us via tactile methods. I think he
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Yes-It was almost a little too vast.

Yes-I learned about several things that I haven't heard before.

Yes-It got one thinking about species extinction and all the tpyes of science that still is going on and is recent, also relates to us.

Yes-yes it helped to consider biological aspects that naturally occur and the relatedness to every day life.

Yes-It made me aware of the diversity of the world i live in.

No-It wasn't a course that I was very interested in taking.

Yes-The class itself was directed towards a wide variety of biological aspects, and did broaden my horizons.

Yes-it presented me with some interesting concepts, but the instructor's bias towards his particular field was evident throughout the entire course-as well as his expectation that we have as good a grasp on that field as he.

Yes-It was presented in an interesting way.

Yes-The text used applied knowledge so you could not read your book and notes and get an A.

Yes-This class was a very intellectually stimulating class because it was clear the subject is very important to the professor. It was great to see parts of the world that most do not even know are out there.

Yes-I had taken Bio 101 last semester, and in comparison, this class requires much more thinking.

Yes-It taught me something that I didn't know. Especially about fungi and bacteria.

No-I did not enjoy the topic so it did not stretch my thinking.

Yes-Alot of interesting concepts, topics and examples

Yes-We were required to take clicker quizzes on a regular basis.

Yes-Mr. O'Meara has a way of engaging his students while talking about Biology that forces you to think of the larger picture. Instead of just presenting facts he was able to focus on what the content meant on a larger scale.

Yes-Fantastic teacher. Definitely inspiring

Yes-Yes I loved the content

No-While the amount of information was great -expecting us to assume certain details with other examples in class, the information was not difficult. The concepts were overly simple. The difficult aspect was applying the information. I can know all of the characteristics of a protist but unless I have background knowledge of organisms, I'm not going to be able to identify them as a protist unless they were specifically "emphasized" in class. It was hard to study for this class because you had to know everything; it's not like you can choose the important information. If you don't know a detail, despite knowing the concept, you miss the test question.

No-Very Basic, not what I was expecting

Yes-Yes, it brought up many ideas that I feel are very important for a college student to know.

Yes-Yes, as far as biodiversity it stretches your thinking deeply into biology. It covers all aspects of biology to get you ready for future classes. Yes-I loved the biodiversity examples. Opened my eyes to some things I 2: Greatest Lab, reading, and the very few assignments we had. contribution to cellular learning The videos The instructors enthusiasm and the short web assigns that were clearly announced on black board. The application and expansion of points in lecture lectures Teaching myself. The pictures and videos on the powerpoint slides. Taking notes Being interested in what you were teaching. The lectures assisted a little, the reading assignments did not help at all. How interesting it was and how he presented it. It made Biology fun rather than just feeding us facts. The lectures where we were able to take notes rather than the lectures that had slides with pictures. The comfort level of the class, meaning I could ask questions and not feel awkward. Taking good notes Slides with more pictures than words Dr.O'Meara was always excited for class and made coming to a large lecture actually not terrible. The videos gave good illustrations. Powerpoint and lecture were good. Clicker questions and readings. the comparisons to noticle things in life. Sitting in class. All of the examples that were used. The use of Mastering Biology as reviews before exams. the text book The slides were a big help also posting them online. The lectures were the most helpful. the highlighted text in the book I really liked the use of graphs and phylogenetic trees to connect ideas The clicker questions were a fun way to test my knowledge. Nothing really, maybe the videos he showed in the class. **Powerpoints** Study guides Many of the lectures were informative

The Lampyr assignments

The instructors enthusiasm made what could have been a boring lecture into a truly interesting course.

The instructor's teaching methods.

Lectures and readings

Every time a point was made in a form other than an example. While teaching through examples might seem more interesting, it's not good for learning. Make a point, than make an example, then make the point again.

The lectures

Dr O'Meara was a HUGE contributor to the class. His interest and enthusiasm really made me want to learn more about biology and biological processes.

The homework assignments was a big part of helping me learn the material.

Videos and examples

3: Detracted from learning

The lectures.

none

The sleeping girl in front of me

I forgot my clicker several times from class and could not get credit for attending class.

Just some tricky test questions

test questions were abstract

Unclear what content to focus on for exams.

The graphs. Some of the graphs didn't make sense or I couldn't see how they related to the subject matter.

The lack of information on his slides made it hard to pull our own notes out of thin air.

Not enough information in slides

The inconsistency of assignment vs test. At this point i find it useless even reading or studying because none of the work i put in will help me on the tests. I'm not even quite sure what i am supposed to be learning now. The requirements to pass his test are not designed for a Freshman but rather a Junior or Senior who is able to use the material given in a laboratory environment.

I wish there were less graphs or for him to tell us how to read the graphs before explaining them.

The pictures on the slides

lecture videos the entire time were no help.

None

Lack of explanation for some things, gets off topic

Dr. Omeara's speech was sometimes unintelligible

There was a lot of material that was really broad rather than going indepth with some things, but that is probably the way with most 100 level science classes

He rarely wrote any notes on the slides, and that made it difficult to take notes and to go back and study his slides.

Couldn't hear you most of the time because you get over entusiastic sometimes.

my computer, and failure for instructor to narrow material and clarify concepts.

The way the notes were being presented.

Professor was often difficult to understand and follow.

The inability to know exactly what to take notes on

Too much detail on individual species.

the lectures were interesting but the instructor often would get excited and become almost impossible to understand

Dr. Omera gets really excited about what he is talking about and then speeds up and becomes difficult to understand.

I was not interested in the topics.

I would recommend not using as many videos unless some sort of subtitles were added, maybe note titles. People who talk during class can be distracting at those times.

Bad sound system

The powerpoints can tend to get boring after so many minutes, but the videos you include help to break it up. However, from these videos, it is sometimes difficult what we were supposed to take away.

nothing

He did. I can tell he is excited about teaching, but he talks and about towards the end of what he is saying you cant understand. He also asked clicker questions and test questions that were tricky. They were worded so weird sometimes you didn't even know what he was asking for. He also asked things that were not in his lectures or book on the tests. He is a good person and you can tell he cares, but he is not a good teacher. He does not relay what he is trying to say or get across to the students well. I don't think he should be teaching about viruses and how they work to a 130 class. The information on a freshman level is too hard for them. I think that when a freshman teacher has to curve every test he has and his overall final grades to make students pass, that says a lot about what kind of teacher he is.

The lectures were extremely boring

Many other lectures were very boring

none

Large class

Had a hard time hearing sometimes

Table weren't very helpful because you have to study them to draw out information and there is not enough time for it to be used effectively in a lecture. This is only for tables, not for graphs.

Having multiple videos on a single slide

Some students were not interested in the class and their distraction took away from my learning.

In some of the lectures O'Meara used other things that had nothing to do with biology to relate it to a biological term or experiment. But that kinda threw me off.

Sometimes hard to parse what was going on on the slides... More text would have helped.

4: Suggestions

Put more text on the slides, assign more homework (since you made us spend a good amount of money on the freaking masteringbiology), Make the tests more straight forward, speak clearer, and since your tests are so full of critical thinking make the students do more throughout the course on slides and assignments or make your tests easier.

Just be more clear in the lectures. And put words into the slides

Study guidex

more forward questions

Poor quality power points and makes it very unclear what content would be on the exams. Tests barely reflected any power points, honestly have no idea where the questions came from. Would have assignments like "skim chapters 2-18" which is beyond ridiculous. Atleast invest in a studyguide or make it clear what were supposed to study to prepare ourselves. Didn't really learn much, stressed more than anything

Connect the topics better (relate them to each other).

Make the tests easier rather than manipulating the grading scale. The objective of the course is to teach us, not make us fail but change our failing grades to Bs

I didn't know what to expect from the test; study guides help. Better questions.

There are so many, he is a wonderful and an enthusiastic teacher but I learned nothing in his course. We would go over power points during class as we take notes, but reviewing these power points for the test did not help because they were simply repeating images on slides. If you are not a good note-taker, you were simply left behind. I also missed one test because of odd circumstance and was only allowed half the points. I could not change nor predict these circumstances but it did not change the outcome. His test are way too difficult and are nothing that we went over, it is like coming into a test blindly, knowing that everything you learned was pointless. It was a very difficult class and do not wish to retake it. I reconsider my major because of this class, although i know other biology classes could have resulted in better outcomes.

Label your slides and point the key aspects that you wish for us to obtain. Some times he gets so into what he is saying, I'm not sure how to take away from it. I loved this class, but had a hard time following with the slides.

More Notes on slides.

having notes to take instead of watching pointless videos that did NOT help with the test.

Stay on topic with slides

Get some different labs.

Be sure to wear the microphone when teaching.

Put more material on the slides and be more specific on what we are tested on.

Better powerpoints. Speak slower and calmly, use MasteringBio more

more focus on specific parts of the course

More written notes. It was hard to pass the test.

More elaborate powerpoints with things other than just pictures or videos for note taking.

Make clear, maybe in text on the slides, exactly what the students need to know or take notes on. I think the reason the majority of the class struggled to do well on the class was because they had no idea what to study. Also try to relate the questions on the test to the lectures better.

Be more clear on the lampyr project, i had no idea we were turning the three pieces at separate times, I ended up turning them in all at once which may have faltered my grade some because i rushed.

the instructor needs to understand it is a freshman level course and as such student's will not have a large background knowledge of taxonomy- which he seemingly expects everyone to have.

Slow your speech down a little bit.

Lampyr mini-essays did not seem necessary.

The professor mumbles a tad, improving that would make the course better.

more online homework cause we only had like 3 of them and we had to pay like 60 dollars for the code

The only complaints I have is in the organization of the content. It could have been better explained in lecture the organization of the organisms, sometimes I felt lost even while paying attention the entire time. It would have also been helpful to have a few more key words written so that we know how to spell things in our notes, especially unfamiliar words. This got better as the semester went on though.

The Lampyr assignments were not as spread out as I would have liked for them to be. We had ample amount of time before the first one was due and after it was due to start the second one, but the third one was due just a week after the second, during the last week of school. Spreading out the assignments would help students in the busy times of the semester.

Talk a little slower so that people can understand what you are saying. Put more words on the slide to explain some of the pictures.

a new instructor or at least some teaching classes on how to relay the information correctly, because obviously he is not.

Actually putting words on his slide show instead of just pictures.

More focus on making sure everyone understands each concept, and the tests actually then relating to those concepts instead of all the technics and theoretical questions that I've seen on the tests

None. Mr. O'Omeara did a fine job of keeping things interesting. His excellent use of multimedia was impressive to say the least.

Break it up a bit more?

Instructor to speak louder

				While teaching, make a point, tell an example or examples, and make the point again. In this way, we can understand points and know when you are moving to the next one. Moving flawlessly from example to example does not give your points enough distinction to actually be retained.
				Be clear about how the grading works
				I suggest that O'Meara talk more slowly and use more slides with words on it and also make test questions that make sense. instead of tricky ones.
				Overall, it was very good. I would suggest the Lampyr assignments be due earlier in the semester, because I had 800 other things due when they were assigned. I understand that is sometimes unavoidable though.
2012	Fall	EEB464		Yes-I actively learned something everyday
			stimulating	Yes-The instructor forced me to view things differently than I have in the past. I really learned a lot
				Yes-I found this class wonderfully stimulating and I really appreciated that the Prof. O'Meara would stop in the middle of his lecture to answer any and all questions without it bothering him.
				Yes-I didn't know ,inch about macro evolution before this class so I definitely learned a lot and discovered evidence for macro evolution
				Yes-The instructor presented good examples, and challenged us to think about explanations on our own first.
				Yes-The use of peer reviewed papers as examples for each topic in conjunction with Brian's socratic method of teaching really allowed us to discuss the ideas and develop a fuller understanding.
				Yes-Yes, the instructor did a good job of creating an atmosphere encouraging student response and discussion. The course made you think deeper about macroevolutionary concepts.
				Yes-Topics covered were always interesting and multiple approaches to topics were often brought up
				Yes-I enjoyed Dr. O'Meara as an instructor. Several scientific studies were presented and I learned a lot about general concepts in macroevolution.
			2: Greatest contribution to learning	The powerpoints were great and posting them really helped
				The instructor's knowledge and enthusiasm
				Prof. O'Meara contributed the most. I would think that for some students the class material would be a little dry, but generally the class was very engaged.
				Class lectures were the most helpful sources of information, but the assignments also gave us a chance to see what was in the literature about macro Evolution.
				The lecture material
				Dr. O'Meara was very technological savvy and posed questions for students to answer and even let us choose lectures we wanted to learn about.
				The coverage of ideas was nearly exhaustive within the context of Macroevolution and filled gaps in my knowledge of which I was unaware prior to this course.
				in class discussion

Class time was well organized and instructor definitely made an effort to provoke critical thinking from students

3: Detracted from learning

Rushed assignments. The partner presentation wasn't given with enough time, so it was more that we had to get this done rather than learn.

All the pretty girls, and me wanting to play super smash bros. and zelda on the instructor's wii.

I have no comments to make here.

It was hard to gauge what the instructor wanted from us studs to o. Our assignments because we didn't get grades back in time for the subsequent assignments. It would be nice to know his grading style and what kind of results he expected from us.

Assignments were not very well explained, and none of them were graded throughout the semester (still haven't been) so I have no idea what was expected of me, or how I'm doing in the class overall.

The seating in the room was not ideal, it seemed crowded despite the small class size.

slow response time on grades. it was hard to know how well you were doing in the course.

Pace was a little fast at times, but other than that the course as a whole was excellent

4: Suggestions

Grade the assignments faster. Its much less stressful to know where you stand within the class

maybe eliminate the group presentation, and make it a solo 5 minute presentation. It's hard to meet up outside of class when you work 45 hours a week

The only aspect of the class that I felt that I performed poorly in were the quizzes. That is almost entirely my fault, but the expectations for outside of the class learning were somewhat poorly defined. 1 quiz a month desensitizes you to the need to keep up with the taxa you're supposed to study before class.

Quicker in returning grades

More feedback on graded assignments!!

Recieving grades back in a more timely manner. Group presentations were difficult, grad students do much more research than we were able to in a weeks time.

Maybe some assigned reading or topical links that would prepare us to contribute more to the discussion. Even a list of jargon terms to know prior to each class which would allow discussion without having to waste time on defining ideas.

Faster returns on grades

Perhaps spreading assignments out across the semester would be helpful. The last few weeks were disproportionately heavy on the work load