Proposal Strategy/Structure II Significance

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Proposal Structure: Significance

Page Limits

Section of Application	Section of Application	
Introduction to Revision or Resubmission Applications	1 page	
Introduction to Revision or Resubmission Applications	1 page	
For each project and core of multi-component applications		
Specific Aims	1 page	
Research Strategy (Item 5.5.3 of Research Plan)	6 pages	
For Activity Codes R03, R13/U13, R21, R36, R41, R43, Fellowships (F), SC2, SC3		
Research Strategy (Item 5.5.3 of Research Plan)	12 pages	
For Activity Codes R01, single project U01, R10, R15, R18, U18, R21/R33, R24, R33, R34, U34, R42, R44, DP3, G08, G11, UH2, UH3, SC1, X01		
Research Strategy (Item 5.5.3 of Research Plan)		
For all other Activity Codes, including Cs, Ps, Ss, Ts, Us, etc.	follow FOA instructions *	
Biosketch (per person)		
For all Activity Codes except DP1 and DP2	4 pages	
Biosketch (per person)		
For DP1 and DP2	2 pages	
Appendix **	No page limits, but content limitations.	
	See relevant section of instructions and FOA	

Templates

• Mechanics of document preparation (L02-mechanics.pdf 🗟 🗹) and the LaTeX proposal template (proposal-latex-template.zip 🗹)

			% -*-latex-*- % Document name: proposal.tex, a template for the PhD Proposal
Fillable Individual PHS 398 Forms (These forms are to be used only with paper submissions using the PH samples provided below in an SF424 (R&R) application. These are filla cause an error in the electronic submission of an SF424 (R&R) applicat application page for appropriate formats to be used for electronic subr	ble PDF forms tion. See the <mark>SF</mark>	which will	z ubcument name: proposal.tex, a template for the PhD Proposal Z Authors: Rob MacLeod Z Last update: Mon Jan 30 07:10:25 2012 by Rob Macleod Z - created Z
Form Page 1: Face Page	MS Word (88 KB)	PDF (310 KB)	<pre>XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX</pre>
Form Page 1-continued: Additional form for use only if Multiple PD/PIs are proposed. Do not include if submitting a single-PD/PI application.	MS Word (76 KB)	PDF (254 KB)	Nusepackage{nih-proposal} XXXXXXXXXXXXXXXXXXXX New commands
Form Page 2: Summary, Relevance, Project/Performance Sites, Senior/Key Personnel, Other Significant Contributors, and Human Embryonic Stem Cells	MS Word (117 KB)	PDF (369 KB)	<pre>% Change the footnotes from numbers to a series of symbols %Prenewcommand(\thefootnote}\\fraymbol(footnote}) /newcommand(\tell)(ken et al.)</pre>
Project/Performance Site Format Page - use only if additional space is needed.	MS Word (92 KB)	PDF (269 KB)	\newcommand{\etc\f{\em_stc.}}
Form Page 3: Research Grant Table of Contents	MS Word (79 KB)	PDF (701 KB)	<pre>\newcommand(\eg){(\em e.g.,}) \newcommand(\infty){\em i.e.,}} \newcommand(\ep){(\ransebox(0.5ex}{\tinu++})}</pre>
Form Page 4: Detailed Budget for Initial Budget Period	MS Word (89 KB)	PDF (309 KB)	\newcommand{\degrees}{{\$^(\ <mark>c</mark> irc)\$}} \newcommand{\splitline}{\begin{center}\rule{\columnwidth}{,7mm}center
Form Page 5: Budget for Entire Proposed Project Period	MS Word (86 KB)	PDF (573 KB)	\newcommand{\nuv}{\${\rm \mu V}\$} \newcommand{\ohm}{\$\Omega\$}
Biographical Sketch Format Page	MS Word (38 KB)	PDF (599 KB)	\newcommand{\sft}{\${\rm ft^2}\$}
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Checklist Form Page	MS Word (84 KB)	PDF (506 KB)	
Continuation Format Page	MS Word (36 KB)	PDF (202 KB)	<pre>XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX</pre>
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Who is the Audience?







Don't assume too much!!

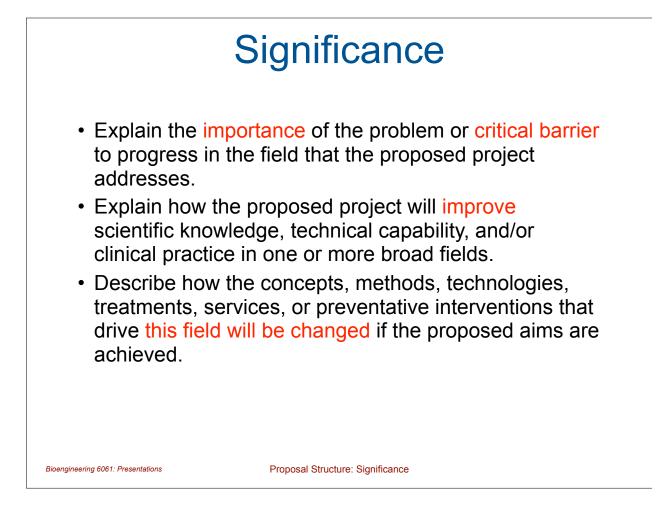
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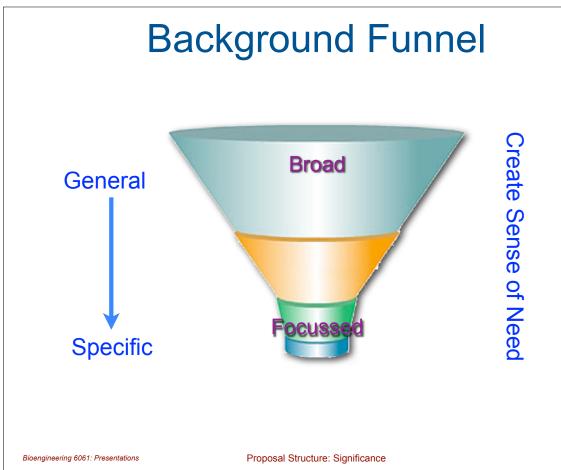
NIH Research Strategy Structure

- Significance
- Innovation
- Approach
- Investigator(s)
- Environment
- Impact

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Background Tips

- Describe what is known, but only what the reader needs to know to appreciate your proposal.
- Create need and also rationale for your research.
- Be objective, do not criticize others.
- Develop clear structure, with subsections.
- Use literature extensively.
- Background funneling to unknown.
- Be direct and use signaling words: "unknown", "problem", "impediment", "challenge".
- · Link the need to the proposed research

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Proposal Structure: Significance

Background funneling to unknown

- "Global warming is arguably one of the most pressing concerns of our time. It has been linked to the rapidity of observed climate change--the fact the the Earth's temperature rose by approximately 0.7C over the last century (the most dramatic increase documented in historic times)[1-3] and the attendant threat posed by melting polar icecaps, rising seal levels, and potentially, more severe weather patterns. "
- "We do not know yet what proportion of this global warming is due to human activity and what is due to natural variations. More important, we lack an effective model to predict precisely by how much the temperature will rise as a consequence of the increase of levels of CO2 and other greenhouse gases in the atmosphere of the earth."

Link Problem to the Research

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Another Example

"These treatments are only moderately effective and are often accompanied by severe side effects and viral resistance. Thus, there remains a need for new therapies for this serious disease. We propose to use a novel class of experiments to test...."

Possible Elements of Significance

- Claim timeliness
- Propose practical solution to a problem
- · Identify a large population affected
- Address a Gap in knowledge
- Outline implications over a wide range of practical problems
- Seek improvement to instrumentation
- Open new research directions
- · Improve quality of life
- Bridge from theoretical to practical knowledge

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Proposal Structure: Significance

Useful Signal Words/Phrases

Background

– "X is"

- "X affects"

- Unknown/Need
 - "...is unknown"
 - "...is unclear"
 - "...has not been determined"
- Objectives/Aims
 - "We propose to..."
 - "Our objective is..."
 - "We will examine the hypothesis that..."
- Significance/impact
 - "...may result in..."
 - "...will contribute to ... "
 - "...may be used to..."

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Image Analysis of C. Elegans Proposal Score: 10, Percentile: 2

Carolyn Wahlby, PhD, Computational Biologist, PI, Imaging Platform, Broad Institute of Harvard & MIT, Cambridge, MA, USA Associate Professor in Quantitative Microscopy, Centre for Image Analysis, Uppsala University, Sweden

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The C. Elegans Imaging Example: The Background and Need

• "The NIH is committed to translating basic biomedical research into clinical practice and thereby impacting global human health¹, and Francis Collins identifies high-throughput technology as one of five areas of focus for the NIH's research agenda². For many diseases, researchers have identified successful novel therapeutics or research probes by applying technical advances in automation to high-throughput screening (HTS) using either biochemical or cell-based assays^{3–6}. Researchers are using genetic perturbations such as RNA interference or gene overexpression in cell-based HTS assays to identify genetic regulators of disease processes as potential drug targets^{7–9}. However, the molecular mechanisms of many diseases that deeply impact human health worldwide are not well-understood and thus cannot yet be reduced to biochemical or cell-based assays."

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The C. Elegans Imaging Example: Tie to Pl's Recent Research

 "Ideally, researchers could approach disease from a phenotypic direction, in addition to the traditional molecular approach, by searching for chemical or genetic regulators of disease processes in whole model organisms rather than isolated cells or proteins. Moving HTS towards more intact, physiological systems also improves the likelihood that the findings from such experiments accurately translate into the context of the human body (e.g., in terms of toxicity and bioavailability), simplifying the path to clinical trials and reducing the failure of potential therapeutics at later stages of testing. In fact, for some diseases, a whole organism screen may actually be necessary to break new therapeutic ground; in the search for novel therapeutics for infectious agents, for example, it is widely speculated that the traditional approach of screening for chemicals that directly kill bacteria in vitro has been largely exhausted¹⁰. Our work recently identified six novel classes of chemicals that cure model organisms from infection by the important human pathogen E. faecalis through mechanisms distinct from directly killing the bacterium itself¹¹. Anti-infectives with new mechanisms of action are urgently needed to combat widespread antibiotic resistance in pathogens.' Bioengineering 6061: Presentations

The C. Elegans Imaging Example: Justify Basic Methodology

 "Enabling HTS in whole organisms is therefore recognized as a high priority (NIH PAR-08-024)^{12,13}. *C. elegans* is a natural choice. Manually-analyzed RNAi and chemical screens are well-proven in this organism, with dozens completed^{14–16}. Many existing assays can be adapted to HTS; instrumentation exists to handle and culture *C. elegans* in HTS-compatible multi-well. Its organ systems have high physiologic similarity and genetic conservation with humans^{17,18}. *C. elegans* is particularly suited to assays involving visual phenotypes: physiologic abnormalities and fluorescent markers are easily observed because the worm is mostly transparent. The worms follow a stereotypic development pattern that yields identicallyappearing adults^{19,20}, such that deviations from wild-type are more readily apparent."

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Proposal Structure: Significance

C. Elegans Example: Establish the Problem, Point to a Solution

 "The bottleneck that remains for tackling important human health problems using C. elegans HTS is image analysis (NIH PA-07-320)^{21,22}. It has been recently stated, "Currently, one of the biggest technical limitations for large-scale RNAi-based screens in *C. elegans* is the lack of efficient highthroughput methods to quantitate lethality, growth rates, and other morphological phenotypes"²³. Our proposal to develop image analysis algorithms to identify regulators of infection and metabolism in highthroughput C. elegans assays would bring image-based HTS to whole organisms, and have the following impact."

C. Elegans Example: Impact linked to Aims

• "Identifying novel modulators of infection by the NIH priority pathogen Microsporidia (Aim 1). Microsporidia are emerging human pathogens whose infection mechanisms are almost completely unknown. Further, they inflict agricultural damage and are on the EPA list of waterborne microbial contaminants of concern^{24,25}. Identifying anti-microsporidian therapeutics is a special challenge because they are eukaryotes. Moreover, they are obligate intracellular pathogens so they are not amenable to traditional antibiotic screens; screening for drugs to kill them requires the presence of a validated, infectible host whose immune system is homologous to mammals, such as *C. elegans*^{26,27}. This screen could identify not only useful chemical research probes and compounds that kill these pathogens outright, but also those that block microbial virulence, are modified by the host for full efficacy (prodrugs), or enhance host immunity."

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Proposal Structure: Significance

C. Elegans Example: Impact linked to Aims

"Identifying novel regulators of fat metabolism (Aim 2). Disregulation of metabolism results in many common and expensive chronic health conditions; diabetes alone affects 24 million Americans²⁸. Energy centers must receive and integrate nutritional information from multiple peripheral signals across multiple tissues and cell types to elicit appropriate behavioral and metabolic responses; screening in a whole organism is important. In particular, screening with a strain of *C. elegans* with an RNAi-sensitive nervous system will likely reveal novel energy regulators of therapeutic and research value."

Link back to human health.

C. Elegans Example: Impact linked to Aims

 "Identifying novel regulators of infection by the pathogen Staphylococcus aureus (Aim 3). S. aureus is life-threatening for immunecompromised patients. Recently, antibiotic-resistant MRSA strains have created an urgent need for therapeutics with a new mechanism of action²⁹. We will identify genetic regulators of the C. elegans host's response to infection by S. aureus³⁰. These will lead to potential drug targets useful for boosting humans' innate immunity."

States significance to human health

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Proposal Structure: Significance

C. Elegans Example: Explicit Statement of Significance

- Enabling the automated analysis of a wide variety of C. elegans screens. Because C. elegans has proven to be an excellent model for many human organs and processes, the impact of algorithms for automated scoring for currently intractable *C. elegans* image-based screens on our understanding and treatment of a variety of human diseases will be substantial. Adding novel C. elegans algorithms to existing open-source software will create a flexible toolbox that can be applied to other types of assays (including alternative formats such as microfluidics chambers; see Yanik support letter) with minimal modification:
- Aim 1: The algorithms developed for Aim 1 will enable scoring viability and other body morphology
 assays probing a number of biological processes. Our collaborators plan several RNAi and chemical
 screens using live/dead assays to identify modulators of many other clinically relevant pathogens (see
 Ausubel and Mylonakis support letters).
- Aim 2: The algorithms developed for the fat metabolism assay can also be used to quantify the levels of any stain within worms, to measure protein expression levels, the degree of staining by fluorescent dyes or antibodies, and promoter activity in reporter assays probing a wide range of biological processes.
- Aim 3: Where localization patterns are of interest, **the algorithms developed** for the gene expression pattern assay will often be directly applicable, especially given the proposed machine learning capabilities.
- Many benefits come from the automation of image analysis for such screens: (a) increased throughput so as to enable genome-scale RNAi and large-scale chemical screens in whole animals; (b) quantitative results amenable to data mining31–33; (c) increased objectivity and consistency; and (d) increased sensitivity to subtle phenotypes, which often can not be scored reliably by eye. The requisite automation of sample preparation and image acquisition has the welcome side effect of improving consistency and providing a permanent record of the experiment.

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C. Elegans Example: Explicit Statement of Significance

"Creating open-source software for the C. elegans community. C. elegans is used for studying complex multicellular biological processes by more than 11,000 researchers in 750 laboratories worldwide (http:// www.WormBase.org, January 2010), and the close-knit community rapidly shares methods^{17,18,34,35}. Based on our experience developing the CellProfiler software system (see Preliminary studies), packaging automated image analysis algorithms in user-friendly software encourages their use by the broader research community. Although we developed CellProfiler solely for high-throughput screening, 70% of studies citing it actually used it to quantify low-throughput assays (fewer than 100 samples). In this proposal we focus on developing algorithms that are robust and efficient for large-scale experiments, but we anticipate they will become an everyday tool for many researchers in the *C. elegans* community, a good investment since many of these are funded by the NIH."

Identify need, solution, impact

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Proposal Structure: Significance

The C. Elegans Example: Explicit Statement of Significance

 "Thus, in addition to the discovery of potential drugs and drug targets related to metabolism and infection, which could significantly impact the global burden of human disease, our aims will yield open-source software for automated, accurate, quantitative scoring for a wide range of *C. elegans* image-based assays that are currently intractable. The impact will be multiplied by *C. elegans* laboratories worldwide using the resulting software to study a wide variety of pathways relevant to basic biological research and human disease, in both low-throughput and highthroughput experiments"

Summary paragraph

The C. Elegans Example: Innovation

- "In response to the strong demand for C. elegans screening, we propose to build on our technological innovations in sample preparation and imaging and our computational innovations for cells and brains to now create a novel technology for C. elegans. Our proposed work to develop novel algorithms for identifying and characterizing worms in microscopy images will bridge the final gap, for the first time enabling widespread identification of genetic and chemical regulators of human biological processes and diseases via wholeorganism screening."
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Describes Innovation and shows impact on the field.

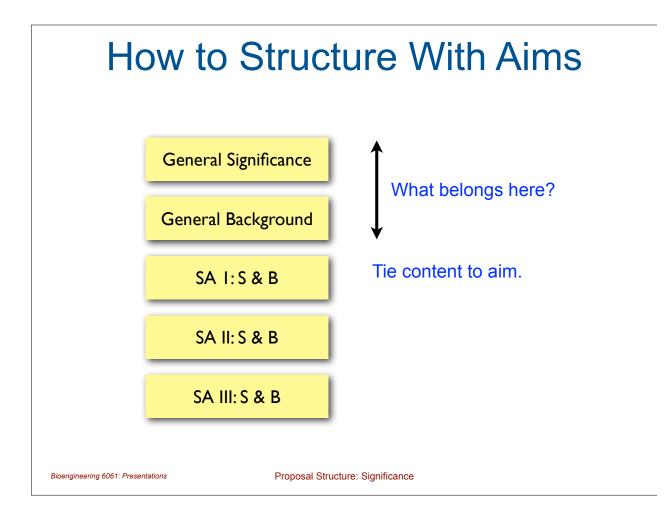
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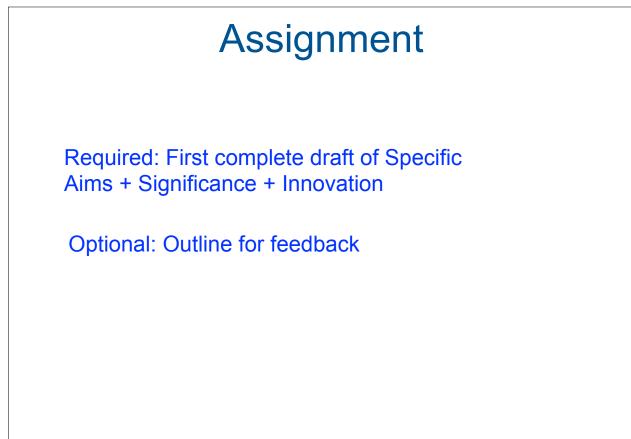
Proposal Structure: Significance

The C. Elegans Example: Innovation

 "Several algorithmic innovations are necessary in order to quantify a variety of C. elegans phenotypes and attain the robustness required for routine high-throughput screening. We propose a novel, simplified represen- tation for worm shapes that lends itself to a probabilistic interpretation. This allows us to adapt shape models to identification of worms in a high-throughput context, and leads to a novel algorithm for detangling worms by morphology-guided graph search. We will also build upon methods from our work in deformation analysis54 and per-cell classification of cellular phenotypes by machine learning55 to quantify phenotypic variation and fluorescence localization in individual worms."

Highlights novel approach





Information



Scientific Writing and Communication



Rob's Grant Information Page

A list of granting sources and links to grant applications. The choices reflect my biomedical bias and is in no way comprehensive.

Granting Agency Policy and Program Information

Other good grants sites

- University of Utah Health Science Resources for Basic Scientists including Research grant information College of Engineering grant information

NIH General Information

http://www.sci.utah.edu/~macleod/grants/

Grant Writing Tips

Some of these are specific to grants, others simply useful for any writing project.

- Rob's Writing page

- Rob's Latex Page.

 NIH Insider Guide, A set of tips from former NIH study section chairs.

 Proposal Writing: The Business of Science (pdf) by Wendy Sanders. Great advice for any grant writer.

 NIH Gludlines (also for review)

 Grant writing tips from NIH

 General reviewer guidlines

 Reviewer guidance on the shortened applications

 FAQ for review of short grants.

 RO1 Grant writing tips

 AHA Grant Writing Tips

 Proposal Writer's Guide by Don Thackrey

 The SCI text markup scheme

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