Electronic Research Administration (ERA) Forum Webinar Series



Research.gov Updates, SciENcv: Overview of Capabilities, and New Award Notice - PI Transfer

> National Science Foundation November 5, 2020

Welcome and Webinar Administrative Logistics

- Attendees will be in listen-only mode during the webinar.
- View real-time captions during the webinar at URL: <u>https://www.captionedtext.com/client/event.aspx?EventID=4569</u>
 252&Customer
 - Enter confirmation ID # 4569252
- Presentation, Agenda, and Webinar recording will be available on the NSF ERA Forum Website https://www.nsf.gov/bfa/dias/policy/era forum.jsp.
- For Zoom Support, call +1-833-966-6468 (+1-833-Zoom-Gov) or email support@zoom.us.

Agenda

- Welcome, Forum Webinar Logistics, and How to Engage with the Forum
- The ERA Forum
- Speakers
- Forum Webinar Discussion Topics:
 - Research.gov Updates
 - Migrating Proposal Preparation from FastLane to Research.gov
 - System Enhancements
 - New Proposal Preparation Demo Site
 - Proposal Preparation Demo
 - SciENcv Overview of Capabilities
 - New Award Notice PI Transfer
- Questions and Answers
- Next Steps and ERA Forum Resources

How to Engage with the Forum

Forum Participation

- Questions can be sent to nsferaforum@nsf.gov during the webinar
 - Responses will be provided in a Frequently Asked Questions (FAQs) document and posted to the ERA Forum webpage
- Post-webinar Survey
- Future Forum Topics
- Forum Listserv and Email
 - Subscribe: <u>NSF-ERA-FORUM-subscribe-request@listserv.nsf.gov</u>
 - Email us: nsferaforum@nsf.gov
- General Information



The ERA Forum

Background and Purpose

 Provides a new approach to regularly engaging the research community, which includes collecting individual opinions and perspectives on ERA activities, gathering topics of interest for future Forum Webinars, and soliciting volunteers for usability studies.

Past Forum Webinars:

- May 2020:
 - Key updates included the availability of both NSF-approved formats for the Biographical Sketch and Current and Pending Support sections of National Science Foundation (NSF) proposals that fall under the revised <u>Proposal & Award Policies & Procedures Guide</u> (PAPPG) (NSF 20-1); the availability of Separately Submitted Collaborative Proposals from Multiple Organizations on Research.gov.

Speakers



Maria	Jean	Sonya	Sherri Bailey	Jeff
Koszalka	Feldman	Arora		Vieceli
Staff Associate, Research.gov, Division of Institution and Award Support	Head, Policy Office, Division of Institution and Award Support	IT Project Manger, Division of Information Systems	Product Manager, National Institute of Health	Head, Systems Office, Division of Institution and Award Support

Research.gov Updates: Migrating Proposal Preparation from FastLane to Research.gov

Research.gov – Important Notice No. 147

- NSF is taking proactive steps to incrementally move the preparation and submission of all proposals from FastLane to Research.gov by 2022
- Some funding opportunities will require the use of Research.gov

Important Notice No. 147

Research.gov Implementation Update

Important Notice to Presidents of Universities and Colleges and Heads of Other National Science Foundation Grantee Organizations

September 22, 2020

The National Science Foundation (NSF) has been at the forefront in the development of Federal agency electronic systems designed to prepare and submit proposals for Federal financial assistance. From the introduction of FastLane in 1994, to the incremental development of Research.gov as its eventual replacement, NSF has led the way with modern, agile systems tailored to meet the needs of the research community.

While NSF's FastLane system has been a resounding success story, it is now an aging, antiquated system that has become increasingly expensive to maintain and even harder to improve. Over the past few years, NSF has partnered with and received valuable input from the research community, resulting in the development of a modern, flexible Research gov system that reduces administrative burden to meet the current and future needs of researchers, administrators and organizations. As a result, NSF has successfully migrated important research functions from FastLane to Research gov including the preparation and submission of annual and final project and outcomes reports, most notifications and requests and award payments. NSF is now taking proactive steps to incrementally move the preparation and submission of all proposals from FastLane to Research.gov with a tentative target date for completion by 2022.

In support of this effort, in the coming weeks and months, NSF will begin making changes to specific funding opportunities to require the use of Research gov for the preparation and submission of proposals to NSF. The Directorate for Biological Sciences (BIO) will soon require the use of Research gov for the preparation and submission of proposals in response to its core programs that do not have deadline dates. NSF funding opportunities will clearly specify whether submission via Research.gov is available or required.

To ensure that researchers and administrators are prepared for these changes, NSF is developing additional training materials to meet the needs of the community. This includes video tutorials, Frequently Asked Questions, step-by-step guides and a demonstration site. Current training materials are available on the About Research gov site.

NSF encourages the community to become familiar with Research gov and to begin using it for the preparation and submission of proposals, as well as to provide NSF with valuable feedback. For additional information, FAQs, opportunities for training and to provide feedback, please visit Research.gov.

> Dr. Sethuraman Panchanathan Director

¹During this time, NSF will continue to permit proposals to be prepared and submitted via Grants.gov.

Research.gov – Directorate for Biological Sciences

- Core BIO program solicitations without deadline dates will require submission via Research.gov
- 90-day grace period before solicitations are removed from FastLane
- Visit the BIO website for details

NSF 20-129

Dear Colleague Letter: Implementation of Full Proposal Submission through Research.gov for "No Deadline" Program Solicitations in the Directorate for Biological Sciences

September 29, 2020

Dear Colleagues:

The Directorate for Biological Sciences (BIO) is notifying members of the research communities of an important change to the program solicitations noted below, effective in calendar year 2020.

Innovating and migrating proposal preparation and submission capabilities from FastLane to Research.gov is part of the ongoing NSF IT modernization effort and is intended to provide the next generation of grants management capabilities for the research community, as described in Important Notice No. 147. The Research.gov Proposal Submission System improves the user experience while also reducing administrative burden through an intuitive interface and is built on flexible system architecture with the ability to meet both users' changing needs and emerging government requirements.

In accordance with the modernization effort, BIO will implement a requirement for submission of full proposals via Research.gov or Grants.gov for the program solicitations listed below in the Division of Environmental Biology (DEB), the Division of Integrative Organismal Systems (IOS), the Division of Molecular and Cellular Biosciences (MCB), and in the Research Resources Cluster of the Division of Biological Infrastructure (DBI). These represent the set of BIO solicitations that accept proposal submission at any time (i.e., have no deadlines). This is the first phase in an eventual shift to require all proposals be submitted via Research.gov or Grants.gov.

By using a phased approach and starting with the no-deadlines solicitations, BIO hopes to provide researchers with sufficient time to adjust to submissions through Research.gov and the sunsetting of FastLane. We do not anticipate that the change to Research.gov will have significant impacts on the submission process, and this migration will not affect the merit review process in any way. Additional technical support is available for those who are new to Research.gov and to those who encounter any issues with proposal preparation and submission.

Revised solicitations for the below impacted programs will be published in the coming weeks. After a 90-day grace period, during which submissions will still be allowed through FastLane, the new solicitations will no longer be available in FastLane, and any new proposals must be submitted through Research.gov or Grants.gov.

Research.gov Updates: System Enhancements

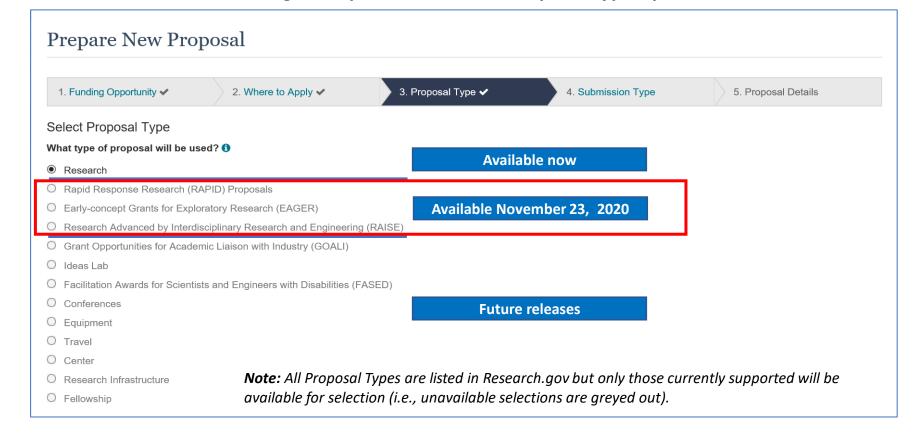
Font Warnings – Changes Effective November 23

- The compliance checks and associated compliance warnings for font type and font size will be removed in Research.gov to align with FastLane
- PAPPG (NSF 20-1) <u>Chapter II.B.2.a.</u> proposal font requirements must be followed
- Proposals may be Returned Without Review if the font type or font size are not compliant with the PAPPG
- NSF is making these changes per feedback from the research community

Additional Proposal Types – Available November 23

- Rapid Response
 Research (RAPID),
 EArly-concept Grants for
 Exploratory Research
 (EAGER), and Research
 Advanced by
 Interdisciplinary Science
 and Engineering (RAISE)
 proposal types will be
 available in
 Research.gov
- New associated
 <u>automated compliance</u>
 <u>checks</u> will also be
 added

Research.gov Proposal Screenshot: Proposal Type Options



Research.gov Proposals: Capabilities Overview

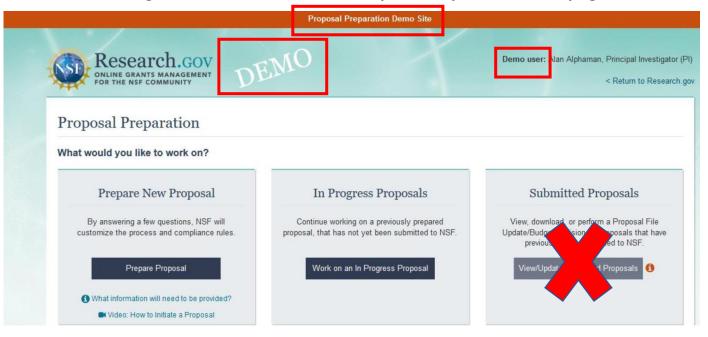
Current Capabilities	Upcoming Capabilities	Future Capabilities
Proposal Types ✓ Research: Single Submissions from One Organization (April 2018) ✓ Research: Single Submission Collaborative Proposals with Subawards (June 2019) ✓ Research: Separately Submitted Collaborative Proposals from Multiple Organizations (March 2020) Training ✓ Research.gov Proposal Preparation Demo Site (October 2020)	 Proposal Types Available November 23, 2020: Rapid Response Research (RAPID) EArly-concept Grants for Exploratory Research (EAGER) Research Advanced by Interdisciplinary Science and Engineering (RAISE) Proposal Features Withdrawal of Single and Collaborative Submissions 	 Proposal Types Conference Equipment Travel Grant Opportunities for Academic Liaison with Industry (GOALI) Ideas Lab Facilitation Awards for Scientists and Engineers with Disabilities (FASED) Center Research Infrastructure Fellowship
Note: Since July 2020, Research.gov is able to support all Research proposals that conform to standard PAPPG requirements for formatting and document content.		 SBIR and STTR Phases I/II Submission Types Letter of Intent (LOI) Preliminary Proposal Full Proposal related to a Preliminary Proposal Renewal Proposal Accomplishment-Based Renewal Proposal

Research.gov Updates: New Proposal Preparation Demo Site

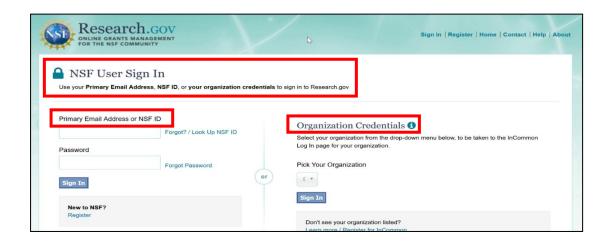
Research.gov Proposal Preparation Demo Site

- Now available for use by the research community and NSF staff
- All users can perform the proposal <u>preparation</u> functions that can be executed by a Principal Investigator (PI) in the actual Research.gov Proposal Submission System. Proposal submission capability is disabled
- All users must sign in to Research.gov to access the demo site, and each user will have the role of PI for demo site purposes. No other user roles (e.g., Sponsored Project Officer or Authorized Organizational Representative) are available or needed

Research.gov Demo Site Screenshot: Proposal Preparation Homepage



Research.gov Sign-in Options





- As of September 28, the Research.gov Sign-in page includes three sign-in options:
 - Primary email address + password (new option)
 - NSF ID + password (existing option)
 - Organization-issued Credentials (InCommon Federation participating organizations only) (existing option)
- Users can enter either their primary email address or their NSF ID for NSF account password recovery

Research.gov Proposal Demo

Research.gov Proposal Demo

- Proposal Creation Wizard
- Main Proposal Page
- Document Upload and Compliance Error/Warning Messages
- Separately Submitted Collaborative Proposal
- Senior Personnel Documents
- Single Copy Documents
- Budget
- Proposal File Update (PFU)
- Budget Revision



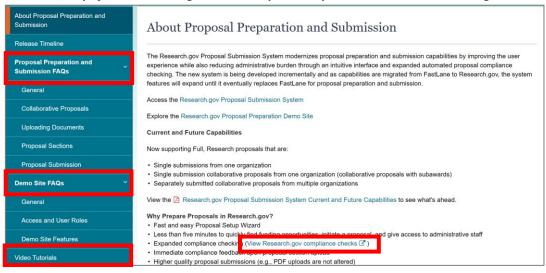
Research.gov Proposal Training Resources

- Resources available when preparing proposals on both the actual Research.gov site and the demo site:
 - Information tool tips
 - Links to the PAPPG
 - Links to FAQs
 - Links to video tutorials
- Resources on Research.gov <u>About</u>
 <u>Proposal and Submission</u> page:
 - FAQs by topic
 - Links to video tutorials
 - Link to <u>Research.gov proposal</u> automated compliance checklist





Excerpt from Research.gov About Proposal Preparation and Submission Page



Information Resources for the Research Community

- NSF System Updates listserv to receive the latest Research.gov system enhancement information and FastLane news. Sign up by sending a blank email to: system updates-subscriberequest@listserv.nsf.gov
- Automated Compliance Checking of NSF Proposals webpage
- NSF <u>Biographical Sketch</u> and <u>Current and Pending Support</u> websites with FAQs, as well as links to SciENcv video tutorials and reference resources with step-by-step instructions and screenshots
- Research.gov About Proposal Preparation and Submission webpage with new and updated FAQs organized by topic and links to video tutorials (new information is being added on an ongoing basis, so please check back frequently....)
- Research.gov About Account Management webpage
- NSF Electronic Research Administration (ERA) Forum webpage
- NSF ERA Forum listserv to receive updates on ERA Forum activities. Sign up by sending a blank email to: NSF-ERA-FORUM-subscribe-request@listserv.nsf.gov

SciENcv Overview

Sherri Holland Bailey, MLIS



SciENcv = Science Experts Network Curriculum Vitae

Use your data from multiple systems to support funding applications, reporting, and collaboration with less burden and complexity

Goals

Reduce burden of applying for federal funds and maintaining federal profiles

Leverage and extend data to/from external systems

Track impact of federal investments in science and scientist careers through scientist-curated data

Current Features

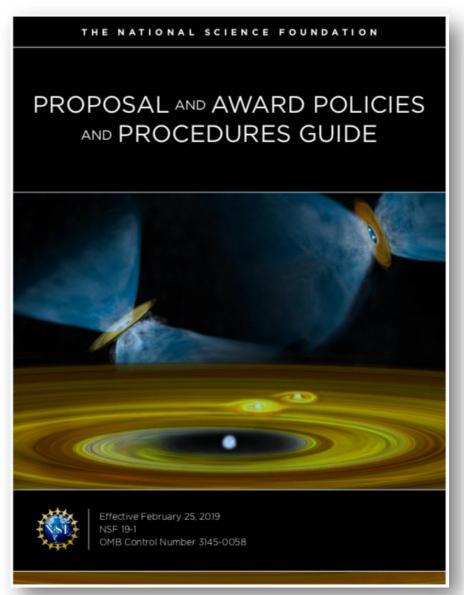
NIH biosketches, NSF biosketch, NSF Current and Pending Support, Dept. of Ed. IES biosketch Integration with ORCID, Fastlane, PubMed/PMC and eRA Commons Bulk upload of citations from reference manager software



NIH SF424 Application guide, 245 pages

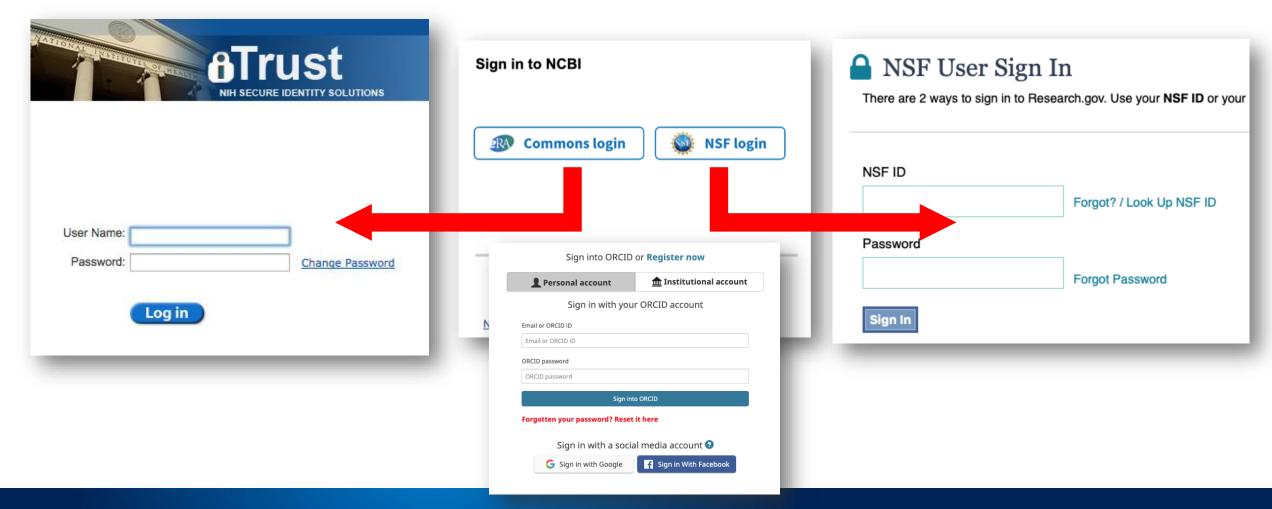
GENERAL APPLICATION GUIDE FOR NIH AND OTHER PHS AGENCIES SF424 (R&R) - Forms Version D Released: March 25, 2016 Revised: June 10, 2016 G.100 How to use the Application G.100 - How to Use the Application Instructions Use these application instructions to fill out the forms that are posted in your funding opportunity announcement. Tour the application guide Quick Links - Step 1. Become familiar with the application process Step 2. Select the correct application instructions. - Step 3. Choose an Application Guide format. · Step 4. Complete the appropriate forms. Step 5. Stay informed of policy changes and updates. Step 1. Become familiar with the application submission process Understanding the application process is critical to successfully submitting your application. Use the Application Process section of these instructions to learn the importance of completing required registrations before submission, how to submit and track your application, where to find page limits and formatting requirements, and more. Step 2. Use these instructions together with the forms and information found in the funding opportunity announcement The funding opportunity announcement will include the forms needed for your application submission Remember that the funding opportunity announcement instructions always supersede these application instructions. Step 3. Choose which application instruction format you prefer to use.

NSF Proposal and Award Policies and Procedures Guide, 192 pages

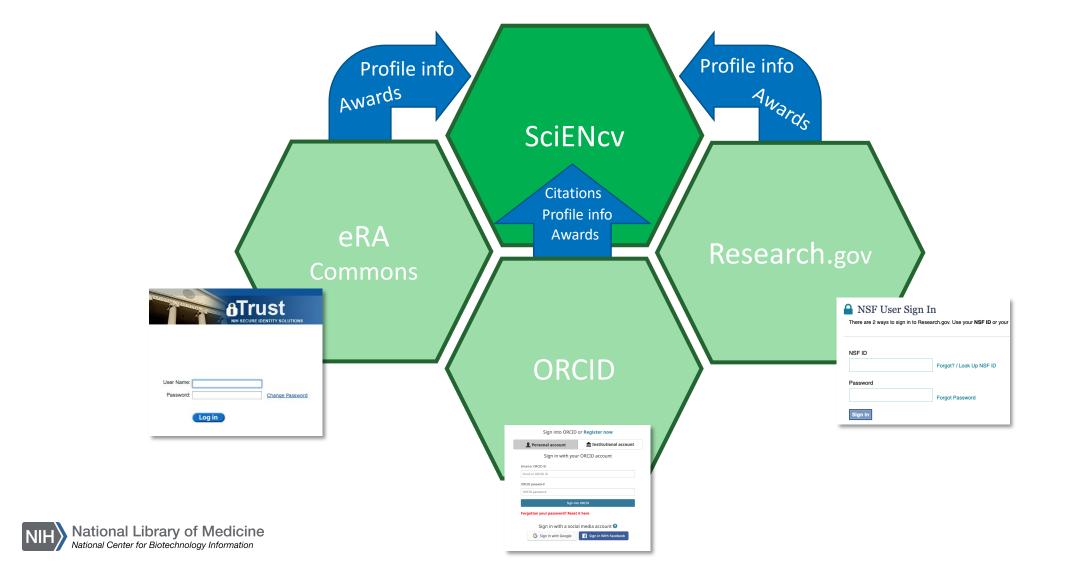




Link to external accounts

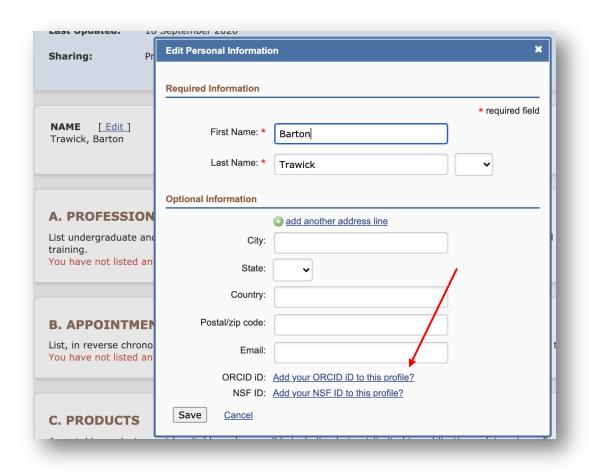


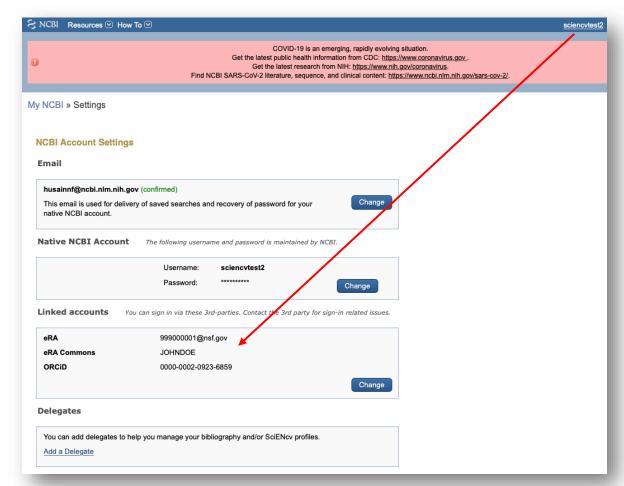
Linking external accounts





Link to external accounts



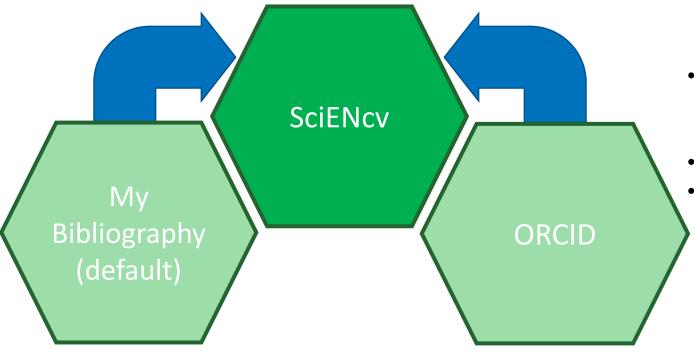


Using citations in SciENcv

Pulls from PubMed

 Tracks NIH Public Access Compliance

- Populates NIH RPPR
- Supports RIS upload
- Can create manual citations



- Pulls from multiple sources (CrossRef, Scopus, etc.)
- Supports BibTex upload
- Can create manual citations





C. PRODUCTS

Acceptable products must be citable and accessible including but not limited to publications, data sets, software, patents, and copyrights. Unacceptable products are unpublished documents not yet submitted for publication, invited lectures, and additional lists of products

citation information including (where applicable and practicable) names of all authors, date of publication or relea journal or book, volume, issue, pages, website and Uniform Resource Locator (URL) or other Persistent Identifier.

PRODUCTS MOST CLOSELY RELATED TO THE PROPOSED PROJECT [Select citations]

- 1. Edelsbrunner H, Guibas L, Hershberger J, Seidel R, Sharir M, Snoeyink J, Welzl E. Implicitly representing Computational Geometry, 1989 October 1; 4(5):433-466. Available from: http://link.springer.com/10.100
- 2. Scott RA, Lagou V, Welch RP, Wheeler E, Montasser ME, Luan J, Mägi R, Strawbridge RJ, Rehnberg E, Gust Lecoeur C, Shungin D, Sanna S, Sidore C, Johnson PC, Jukema JW, Johnson T, Mahajan A, Verweij N, Tho Sennblad B, Gieger C, Salo P, Perola M, Timpson NJ, Evans DM, Pourcain BS, Wu Y, Andrews JS, Hui J, Bie O'Connell JR, Stirrups K, Vitart V, Hayward C, Esko T, Mihailov E, Fraser RM, Fall T, Voight BF, Raychaudhu Sibl Robertson N, Rybin D, Liu CT, Beckmann JS, Willems SM, Chines PS, Jackson AU, Kang HM, Stringham HW P, Müller-Nurasyid M, Franco-Cereceda A, Folkersen L, Marullo L, Jansen H, Oldehinkel AJ, Bruinenberg M, U D M S, Varga TV, Hallmans G, Oksa H, Antonella M, Nagaraja R, Trompet S, Ford I, Bakker SJ, Kong A, Kumari Antti J, Mangino M, Small K, Miljkovic I, Liu Y, Atalay M, Kiess W, James AL, Rivadeneira F, Uitterlinden AG Campbell S, Polasek O, Bonnycastle LL, Hercberg S, Dimitriou M, Bolton JL, Fowkes GR, Kovacs P, Lindstre Rathmann W, Grallert H, Maerz W, Kleber ME, Boehm BO, Peters A, Pramstaller PP, Province MA, Borecki I Farrall M, Stumvoll M, Ferrucci L, Waterworth DM, Bergman RN, Collins FS, Tuomilehto J, Watanabe RM, d Psaty BM, Vollenweider P, Wilson JF, Wright AF, Hovingh GK, Metspalu A, Uusitupa M, Magnusson PK, Kyvil Meneton P, Lind L, Boehnke M, Shuldiner AR, van Duijn CM, Morris AD, Toenjes A, Peyser PA, Beilby JP, Kö Schwarz PE, Lakka TA, Rauramaa R, Adair LS, Smith GD, Spector TD, Illig T, de Faire U, Hamsten A, Gudn Kiukaanniemi SM, Saaristo TE, Boomsma DI, Stefansson K, van der Harst P, Dupuis J, Pedersen NL, Satta Mohlke KL, Balkau B, Froquel P, Pouta A, Jarvelin MR, Wareham NJ, Bouatia-Naji N, McCarthy MI, Franks C, Ingelsson E, Prokopenko I, Barroso I. Large-scale association analyses identify new loci influencing glye biological pathways. Nat Genet. 2012 Sep;44(9):991-1005. PubMed PMID: 22885924; NIHMSID: UKMS49

Berndt SI, Gustafsson S, Mägi R, Ganna A, Wheeler E, Feitosa MF, Justice AE, Monda KL, Croteau-Chonka

G, Yang J, Albrecht E, Amin N, Bragg-Gresham JL, Cadby G, den Heijer M, Eklund N, Fischer K, Goel A, Ho Johnson T, Kanoni S, Kleber ME, König IR, Kristiansson K, Kutalik Z, Lamina C, Lecoeur C, Li G, Mangino I M, Ngwa JS, Nolte IM, Paternoster L, Pechlivanis S, Perola M, Peters MJ, Preuss M, Rose LM, Shi J, Shungi Hall P, Hartikainen AL, Hayward C, Heard-Costa NL, Heath AC, Hebebrand J, Homuth G, Hu FB, Hunt SE, Jula A, Kähönen M, Kathiresan S, Kee F, Khaw KT, Kivimäki M, Koenig W, Kraja AT, Kumari M, Kuulasmaa Launer LJ, Lind L, Lindström J, Liu J, Liuzzi A, Lokki ML, Lorentzon M, Madden PA, Magnusson PK, Manunt Medland SE, Mihailov E, Milani L, Montgomery GW, Mooser V, Mühleisen TW, Munroe PB, Musk AW, Narisu BA, Palmer CN, Palotie A, Peden JF, Pedersen N, Peters A, Polasek O, Pouta A, Pramstaller PP, Prokopenko A, Rivadeneira F, Rudan I, Saaristo TE, Sambrook JG, Sanders AR, Sanna S, Saramies J, Schipf S, Schreiber S, Schunkert H, Shin SY, Signorini S, Sinisalo J, Skrobek B, Soranzo N, Stančáková A, Stark K, Stephens JC, Stirrups K, Stolk RP, Stumvoll M, Swift AJ, Theodoraki EV, Thorand B, Tregouet DA, Tremoli E, Van der Klauw MM, van Meurs JB, Vermeulen SH, Viikari J, Virtamo J, Vitart V, Waeber G, Wang Z, Widén E, Wild SH, Willemsen G, Winkelmann BR, Witteman JC, Wolffenbuttel BH, Wong A, Wright AF, Zillikens MC, Amouyel P, Boehm BO, Boerwinkle E, Boomsma DI, Caulfield MJ, Chanock SJ, Cupples LA, Cusi D, Dedoussis GV, Erdmann J, Eriksson JG, Franks PW, Froguel P, Gieger C, Gyllensten U, Hamsten A, Harris TB, Hengstenberg C, Hicks AA, Hingorani A Hinney A, Hofman A, Hovingh KG, Hyeem K, Illig T, Jaryelin MR, Jöckel KH, Keinanen-Kiukaanniemi SM, Kiemeney LA, Kuh D, Laakso M, Lehtimäki T, Levinson DF, Martin NG, Metspalu A, Morris AD, Nieminen MS, Njølstad I, Ohlsson C, Oldehinkel AJ, Ouwehand WH, Palmer LJ, Penninx B, Power C, Province MA, Psaty BM, Qi L, Rauramaa R, Ridker PM, Ripatti S, Salomaa V, Samani NJ, Snieder H, Sørensen TI, Spector TD, Stefansson K, Tönjes A, Tuomilehto J, Uitterlinden AG, Uusitupa M, van der Harst P, Vollenweider P, Wallaschofski H, Wareham NJ, Watkins H, Wichmann HE, Wilson JF, Abecasis GR, Assimes TL, Barroso I, Boehnke M, Borecki IB, Deloukas P, Fox CS, Frayling T, Groop LC, Haritunian T, Heid IM, Hunter D, Kaplan RC, Karpe F, Moffatt MF, Mohlke KL, O'Connell JR, Pawitan Y, Schadt EE, Schlessinger D, Steinthorsdottir V, Strachan DP, Thorsteinsdottir U, van Duijn CM, Visscher PM, Di Blasio AM, Hirschhorn

NSF Policy requires that each biographical sketch be no longer than two (2) pages unless otherwise specified in a

It is the responsibility of the submitter to ensure compliance with the two (2) page maximum.

Close

J, Seidel R, Sharir M, Snoeyink J, Welzl E. Implicitly representing arrangem Berndt St, Gustafsson S, Magi R, Ganna A, Wheeler E, Feltosa MF, Justice AE, Monda KL, Croteau-Chonka Dackson AU, Luan J, Randad IJC, Vedantam S, Willer CJ, Winkler TW, Wood AR, Workalemahu T, Hu J, Lee'r 1; 4(5):433-466. Available from: http://link.springer.com/10.1007/BF021

A, Trip MD, Tyrer J, Van Vliet-Ostaptchouk JV, Vandenput L, Waite LL, Zhao JH, Absher D, Asselbergs FW, E, Bellby J, Bonnycastle LL, Brambilla P, Bruinenberg M, Campbell H, Chasman DI, Chines PS, Collins FS, Col Dimitriou M, Edkins S, Estrada K, Evans DM, Farrall M, Ferrario MM, Ferrières J, Franke L, Frau F, Gejman C, Johnson PC, Jukema JW, Johnson T, Mahajan A, Verweij N, Thorleifsson Timpson NJ, Evans DM, Pourcain BS, Wu Y, Andrews JS, Hui J, Bielak LF, Z vard C, Esko T, Mihailov E, Fraser RM, Fall T, Voight BF, Raychaudhuri S, Che

JN, Lindgren CM, Morris AP, Meyre D, Scherag A, McCarthy MI, Speliotes EK, North KE, Loos RJ, Ingelsson E. Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. Nat Genet. 2013 May; 45(5):501-12. PubMed PMID: 23563607; NIHMSID:

solicitation.



NAME: test2, sciency NSF ID: 999000001@nsf.gov ORCID: 0000-0002-0923-6859 POSITION TITLE & INSTITUTION: (a) PROFESSIONAL PREPARATION INSTITUTION LOCATION

(b) APPOINTMENTS

(c) PRODUCTS

Products Most Closely Related to the Proposed Project

- . Edelsbrunner H, Guibas L, Hershberger J, Seidel R, Sharir M, Snoeyink J, Welzl E. Implicitly representing arrangements of lines or segments. Discrete & Computational Geometry 1989 October 1; 4(5):433-466. Available from: http://link.springer.com/10.1007/BF02187742 DOI: 10.1007/BF02187742
- 2. Scott RA, Lagou V, Welch RP, Wheeler E, Montasser ME, Luan J, Mägi R, Strawbridge RJ, Rehnberg E, Gustafsson S, Kanoni S, Rasmussen-Torvik LJ, Yengo L, Lecoeur C, Shungin D, Sanna S, Sidore C, Johnson PC, Jukema JW, Johnson T, Mahajan A, Verweij N, Thorleifsson G, Hottenga JJ, Shah S, Smith AV, Sennblad B, Gieger C, Salo P, Perola M, Timpson NJ, Evans DM, Pourcain BS, Wu Y, Andrews JS, Hui J, Bielak LF, Zhao W, Horikoshi M, Navarro P, Isaacs A. O'Connell JR. Stirrups K. Vitart V. Hayward C. Esko T. Mihailoy E. Fraser RM. Fall T, Voight BF, Ravchaudhuri S, Chen H, Lindgren CM, Morris AP, Ravner NW, Robertson N, Rybin D, Liu CT, Beckmann JS, Willems SM, Chines PS, Jackson AU, Kang HM, Stringham HM, Song K, Tanaka T, Peden JF, Goel A, Hicks AA, An P, Müller-Nurasvid M, Franco-Cereceda A, Folkersen L, Marullo L, Jansen H, Oldehinkel AJ, Bruinenberg M, Pankow JS, North KE, Forouhi NG, Loos RJ, Edkins S, Varga TV, Hallmans G, Oksa H, Antonella M, Nagaraja R, Trompet S, Ford I, Bakker SJ, Kong A, Kumari M, Gigante B, Herder C, Munroe PB, Caulfield M, Antti J, Mangino M, Small K, Milikovic I, Liu Y, Atalay M, Kiess W, James AL, Rivadeneira F, Uitterlinden AG, Palmer CN, Doney AS, Willemsen G, Smit JH, Campbell S, Polasek O, Bonnycastle LL, Hercberg S, Dimitriou M, Bolton JL, Fowkes GR, Kovacs P, Lindström J. Zemunik T. Bandinelli S. Wild SH. Basart HV. Rathmann W. Grallert H. Maerz W, Kleber ME, Boehm BO, Peters A, Pramstaller PP, Province MA, Borecki IB, Hastie ND, Rudan I, Campbell H, Watkins H, Farrall M, Stumvoll M, Ferrucci L, Waterworth DM, Bergman RN, Collins FS, Tuomilehto J, Watanabe RM, de Geus EJ, Penninx BW, Hofman A, Oostra BA, Psaty BM, Vollenweider P, Wilson JF, Wright AF, Hovingh GK, Metspalu A, Uusitupa M, Magnusson PK, Kyvik KO, Kaprio J, Price JF, Dedoussis GV, Deloukas P, Meneton P, Lind L, Boehnke M, Shuldiner AR, van Duijn CM, Morris AD, Toenjes A, Peyser PA, Beilby JP, Körner A, Kuusisto J, Laakso M, Bornstein SR, Schwarz PE, Lakka TA, Rauramaa R, Adair LS, Smith GD, Spector TD, Illig T, de Faire U, Hamsten A, Gudnason V, Kivimaki M. Hingorani A. Keinanen-Kiukaanniemi SM, Saaristo TE, Boomsma DI, Stefansson

BS-1 of 3

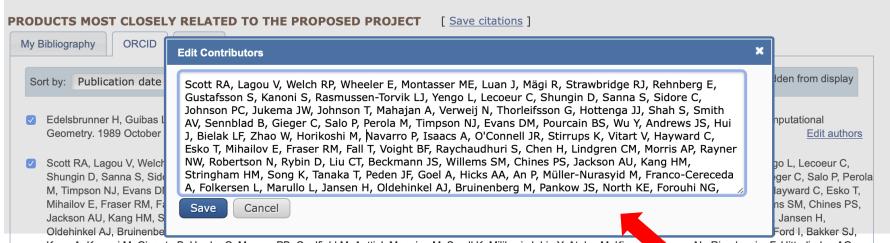
NIHMS536922; PubMed Central PMCID: PMC3973018.

PRODUCTS MOST CLEELY RELATED TO THE PROPOSED PROJECT [Save citations]

My Bibliography Edited unchecked entries are hidden from display Sort by: Publication date \$ Select: None 3 item(s) selected Refresh the list from ORCID Edelsbrunner H, Guibas L, Hershberger J, Seidel R, Sharir M, Snoeyink J, Welzl E. Implicitly representing arrangements of lines or segments. Discrete & Computational Geometry. 1989 October 1; 4(5):433-466. Available from: http://link.springer.com/10.1007/BF02187742 DOI: 10.1007/BF02187742 Edit authors Scott RA, Lagou V, Welch RP, Wheeler E, Montasser ME, Luan J, Mägi R, Strawbridge RJ, Rehnberg E, Gustafsson S, Kanoni S, Rasmussen-Torvik LJ, Yengo L, Lecoeur C, Shungin D, Sanna S, Sidore C, Johnson PC, Jukema JW, Johnson T, Mahajan A, Verweij N, Thorleifsson G, Hottenga JJ, Shah S, Smith AV, Sennblad B, Gieger C, Salo P, Perola M, Timpson NJ, Evans DM, Pourcain BS, Wu Y, Andrews JS, Hui J, Bielak LF, Zhao W, Horikoshi M, Navarro P, Isaacs A, O'Connell JR, Stirrups K, Vitart V, Havward C, Esko T, Mihailov E, Fraser RM, Fall T, Voight BF, Raychaudhuri S, Chen H, Lindgren CM, Morris AP, Rayner NW, Robertson N, Rybin D, Liu CT, Beckmann JS, Willems SM, Chines PS, Jackson AU, Kang HM, Stringham HM, Song K, Tanaka T, Peden JF, Goel A, Hicks AA, An P, Müller-Nurasyid M, Franco-Cereceda A, Folkersen L, Marullo L, Jansen H, Oldehinkel AJ, Bruinenberg M, Pankow JS, North KE, Forouhi NG, Loos RJ, Edkins S, Varga TV, Hallmans G, Oksa H, Antonella M, Nagaraja R, Trompet S, Ford I, Bakker SJ, Kong A, Kumari M, Gigante B, Herder C, Munroe PB, Caulfield M, Antti J, Mangino M, Small K, Milikovic I, Liu Y, Atalay M, Kiess W, James AL, Rivadeneira F, Uitterlinden AG, Palmer CN, Doney AS, Willemsen G, Smit JH, Campbell S, Polasek O, Bonnycastle LL, Hercberg S, Dimitriou M, Bolton JL, Fowkes GR, Kovacs P, Lindström J, Zemunik T, Bandinelli S, Wild SH, Basart HV, Rathmann W, Grallert H, Maerz W, Kleber ME, Boehm BO, Peters A, Pramstaller PP, Province MA, Borecki IB, Hastie ND, Rudan I, Campbell H. Watkins H, Farrall M, Stumvoll M, Ferrucci L, Waterworth DM, Bergman RN, Collins FS, Tuomilehto J, Watanabe RM, de Geus EJ, Penninx BW, Hofman A, Oostra BA, Psaty BM, Vollenweider P, Wilson JF, Wright AF, Hovingh GK, Metspalu A, Uusitupa M, Magnusson PK, Kyvik KO, Kaprio J, Price JF, Dedoussis GV, Deloukas P, Meneton P, Lind L, Boehnke M, Shuldiner AR, van Duijn CM, Morris AD, Toenjes A, Peyser PA, Beilby JP, Körner A, Kuusisto J, Laakso M, Bornstein SR, Schwarz PE, Lakka TA, Rauramaa R, Adair LS. Smith GD. Spector TD. Illig T. de Faire U. Hamsten A. Gudnason V. Kivimaki M. Hingorani A. Keinanen-Kiukaanniemi SM. Saaristo TE. Boomsma DI. Stefansson K. van der Harst P, Dupuis J, Pedersen NL, Sattar N, Harris TB, Cucca F, Ripatti S, Salomaa V, Mohlke KL, Balkau B, Froquel P, Pouta A, Jarvelin MR, Wareham NJ, Bouatia-Naji N, McCarthy MI, Franks PW, Meigs JB, Teslovich TM, Florez JC, Langenberg C, Ingelsson E, Prokopenko I, Barroso I, Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. Nat Genet. 2012 Sep;44(9):991-1005. PubMed PMID: 22885924; NIHMSID: UKMS49338; PubMed Central PMCID: PMC3433394. Berndt SI, Gustafsson S, Mägi R, Ganna A, Wheeler E, Feitosa MF, Justice AE, Monda KL, Croteau-Chonka DC, Day FR, Esko T, Fall T, Ferreira T, Gentilini D, Jackson AU, Luan J. Randall JC. Vedantam S. Willer CJ. Winkler TW. Wood AR. Workalemahu T. Hu YJ. Lee SH. Liang L. Lin DY. Min JL. Neale BM. Thorleifsson G. Yang J. Albrecht E. Amin N. Bragg-Gresham JL, Cadby G, den Heijer M, Eklund N, Fischer K, Goel A, Hottenga JJ, Huffman JE, Jarick I, Johansson Å, Johnson T, Kanoni S, Kleber ME, König IR, Kristiansson K, Kutalik Z, Lamina C, Lecoeur C, Li G, Mangino M, McArdle WL, Medina-Gomez C, Müller-Nurasvid M, Ngwa JS, Nolte IM, Paternoster L, Pechlivanis S, Perola M. Peters MJ, Preuss M, Rose LM, Shi J, Shungin D, Smith AV, Strawbridge RJ, Surakka I, Teumer A, Trip MD, Tyrer J, Van Vliet-Ostaptchouk JV, Vandenput L, Waite LL, Zhao JH, Absher D, Asselbergs FW, Atalay M, Attwood AP, Balmforth AJ, Basart H, Beilby J, Bonnycastle LL, Brambilla P, Bruinenberg M, Campbell H, Chasman DI, Chines PS, Collins FS. Connell JM. Cookson WO, de Faire U, de Vegt F, Dei M, Dimitriou M, Edkins S, Estrada K, Evans DM, Farrall M, Ferrières J, Franke L, Frau F, Geiman PV,

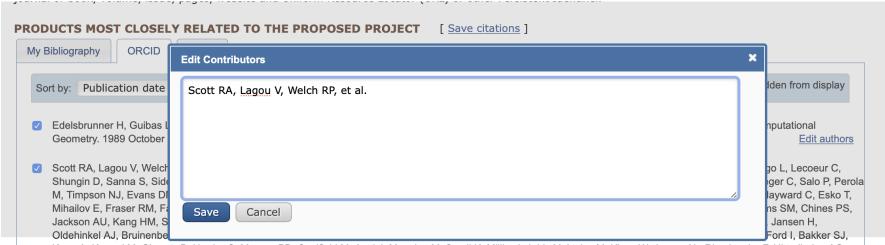
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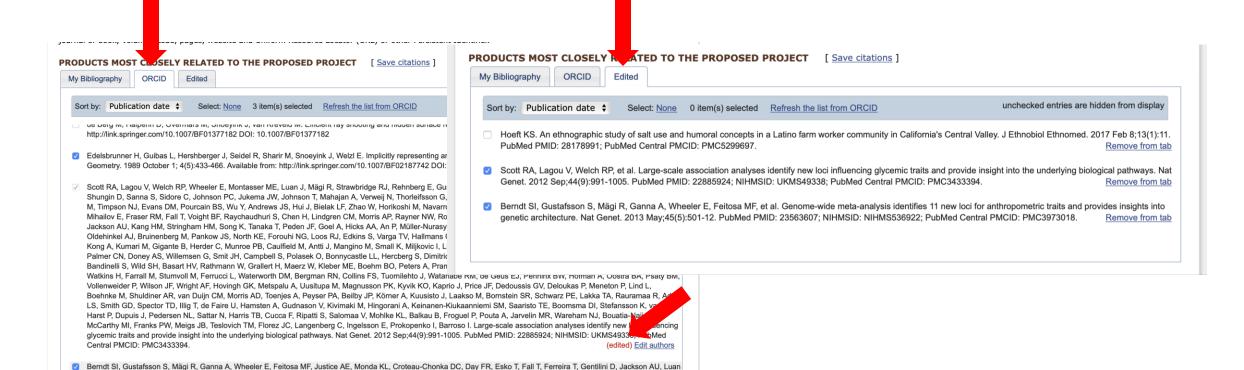
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Kong A, Kumari M, Gigante B, Herder C, Munroe PB, Caulfield M, Antti J, Mangino M, Small K, Miljkovic I, Liu Y, Atalay M, Kiess W, James AL, Rivadeneira F, Uitterlinden AG, Palmer CN, Doney AS, Willemsen G, Smit JH, Campbell S, Polasek O, Bonnycastle LL, Hercberg S, Dimitriou M, Bolton JL, Fowkes GR, Kovacs P, Lindström J, Zemunik T, Bandinelli S, Wild SH, Basart HV, Rathmann W, Grallert H, Maerz W, Kleber ME, Boehm BO, Peters A, Pramstaller PP, Province MA, Borecki IB, Hastie ND, Rudan I, Campbell H, Watkins H, Farrall M, Stumvoll M, Ferrucci L, Waterworth DM, Bergman RN, Collins FS, Tuomilehto J, Watanabe RM, de Geus EJ, Penninx BW, Hofman A, Oostra BA, Psaty BM, Vollenweider P, Wilson JF, Wright AF, Hovingh GK, Metspalu A, Uusitupa M, Magnusson PK, Kyvik KO, Kaprio J, Price JF, Dedoussis GV, Deloukas P, Meneton P, Lind L, Boehnke M, Shuldiner AR, van Duijn CM, Morris AD, Toenjes A, Peyser PA, Beilby JP, Körner A, Kuusisto J, Laakso M, Bornstein SR, Schwarz PE, Lakka TA, Rauramaa R, Adair LS, Smith GD, Spector TD, Illig T, de Faire U, Hamsten A, Gudnason V, Kivimaki M, Hingorani A, Keinanen-Kiukaanniemi SM, Saaristo TE, Boomsma DI, Stefansson K, van der Harst P, Dupuis J, Pedersen NL, Sattar N, Harris TB, Cucca F, Ripatti S, Salomaa V, Mohlke KL, Balkau B, Froguel P, Pouta A, Jarvelin MR, Wareham NJ, Bouatia-Naji N, McCarthy MI, Franks PW, Meigs JB, Teslovich TM, Florez JC, Langenberg C, Ingelsson E, Prokopenko I, Barroso I. Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. Nat Genet. 2012 Sep;44(9):991-1005. PubMed PMID: 22885924; NIHMSID: UKMS49338; PubMed Central PMCID: PMC3433394.







J, Randall JC, Vedantam S, Willer CJ, Winkler TW, Wood AR, Workalemahu T, Hu YJ, Lee SH, Liang L, Lin DY, Min JL, Neale BM, Thorleifsson G, Yang J, Albrecht E, Amin N, Bragg-Gresham JL, Cadby G, den Heijer M, Eklund N, Fischer K, Goel A, Hottenga JJ, Huffman JE, Jarick I, Johansson Á, Johnson T, Kanoni S, Kleber ME, König IR, Kristiansson K, Kutlik Z, Lamina C, Lecour C, Li G, Mangino M, McArdle WL, Medina-Gomez C, Müller-Nurasyid M, Ngwa JS, Nolte IM, Paternoster L, Pechlivanis S, Perola M, Peters MJ, Preuss M, Rose LM, Shi J, Shi J,

PRODUCTS MOST CLOSELY RELATED TO THE PROPOSED PROJECT [Select citations]

- 1. Edelsbrunner H, Guibas L, Hershberger J, Seidel R, Sharir M, Snoeyink J, Welzl E. Implicitly representing arrangements of lines or segments. Discrete & Computational Geometry. 1989 October 1; 4(5):433-466. Available from: http://link.springer.com/10.1007/BF02187742 DOI: 10.1007/BF02187742
- 2. Scott RA, Lagou V, Welch RP, et al. Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. Nat Genet. 2012 Sep;44(9):991-1005. PubMed PMID: 22885924; NIHMSID: UKMS49338; PubMed Central PMCID: PMC3433394.
- 3. Berndt SI, Gustafsson S, Mägi R, Ganna A, Wheeler E, Feitosa MF, Justice AE, Monda KL, Croteau-Chonka DC, Day FR, Esko T, Fall T, Ferreira T, Gentilini D,





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Profile name: Bart Trawick NIH Biosketch format [Edit]

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Last Updated: 30 November 2017

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NAME [Edit] Trawick, Barton

Houston, Houston, TX

ORCID ID eRA COMMONS ID

http://orcid.org/0000-0002-3635-3508 trawick

EDUCATION/TRAINING [Edit entries]

(Begin with baccalaureate or other initial professional education, such as nursing, include

INSTITUTION AND LOCATION	DEGREE
	(if applicable

Texas A&M University, College Station, TX BACHELOR OF /

University of Texas Health Science Center at MASTER OF SCII Houston, Houston, TX

University of Texas Health Science Center at DOCTOR OF PHILC NIH Biosketch

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Last Updated: 9 July 2019

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OMB-3145-0058

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NAME [Edit]

Trawick, Bart

http://orcid.org/0000-0002-3635-3508

A. PROFESSIONAL PREPARATION [Edit entries]

List undergraduate and graduate education and postdoctoral training. List the year the degree was received as well as inclusive dates of postdoctoral training.

INSTITUTION	LOCATION	MAJOR/AREA OF STUDY	DEGREE (if applicable)	YEAR
Texas A&M University	College Station, TX, US	Chemistry		1992
University of Texas Health Science Center at Houston	Houston, TX, US	Biochemistry	MASTER OF SCIENCE	1996
University of Texas Health Science Center at Houston	Houston, TX, US	Immunology	DOCTOR OF PHILOSOPHY	2000





SciENcv Biosketch Output

NIH Biosketch

NSF Biosketch

OMB No. 0925-0001 and 0925-0002 (Rev. 09/17 Approved Through 03/31/2020)

BIOGRAPHICAL

Provide the following information for the Senior/key pe

NAME: Trawick, Bart

eRA COMMONS USER NAME (credential, e.g., agency POSITION TITLE: Director, Customer Services Divsion

EDUCATION/TRAINING (Begin with baccalaureate or of nursing, include postdoctoral training and residency traini

INSTITUTION AND LOCATION

Texas A&M University, College Station, TX

University of Texas Health Science Center, Houston, TX University of Texas Health Science Center, Houston, TX

A. Personal Statement

I work to promote access to government-funded science

- Database resources of the National Center for Biot Jan 4;46(D1):D8-D13. PubMed PMID: 29140470;
- Database resources of the National Center for Biot Jan 4;44(D1):D7-19. PubMed PMID: 26615191; Pu
- Database resources of the National Center for Biot Jan;43(Database issue):D6-17. PubMed PMID: 2 PMC4383943.

B. Positions and Honors

Positions and Employment

2002 - 2005	Staff Scientist, Book melf project, Nationa Bethesda, MD
2005 - 2009	Project Supervisor, NIHMS, National Cer
	MD
2009 - 2017	Public Services Section, National Center
2017 -	Director, Customer Services Divsion, Na
	Bethesda, MD

Other Experience and Professional Membershi

Honors

11011010	
2004	NLM Special Achievement Award, Nation
2008	National Institutes of Health Director's Av
2009	NLM Special Achievement Award, Nation
2009	National Institutes of Health Merit Award,
2011	NLM Special Achievement Award, Nation

<citation type="journal" id="1065150"> <title>Large-scale association analyses identify new loci influencing glycemic 3508

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contributor type="author">7han W</contrib

NSF BIOGRAPHICAL SKETCH

ITUTION: Staff Scientist, National Institutes of Health

EPARATION

	LOCATION	MAJOR / AREA OF STUDY	DEGREE (if applicable)	YEAR YYYY
rsity	College Station, TX	Chemistry	BA	1992
th Science on	Houston, TX	Biochemistry	MS	1996
th Science on	Houston, TX	Immunology	PHD	2000

ntist, National Institutes of Health, National Library of Medicine,

elated to the Proposed Project

FS, Machado FP, Paploski IAD, Martins TF, Teider-Junior PI, Serpa ach RVW, Barros-Filho IR, Lipinski LC, P Dos Santos A, Labruna MB, serosurvey of anti-Rickettsia spp. antibodies in wild boars (Sus scrofa), of Brazil, PLoS Negl Trop Dis. 2019 May;13(5):e0007405. PubMed Med Central PMCID: PMC6542515.

RA, N. cy ZS, Trawick BW, Wang M, Tejpal N, Wang ME, Tian L, role of \$2.5 in the induction of regulatory T cells in transplantation oc. 2001 Nov Dec;33(7-8):3835-6. PubMed PMID: 11750634.

he National Cente, for Biotechnology Information. Nucleic Acids Res. -D17. PubMed PMID. 27899561; PubMed Central PMCID:

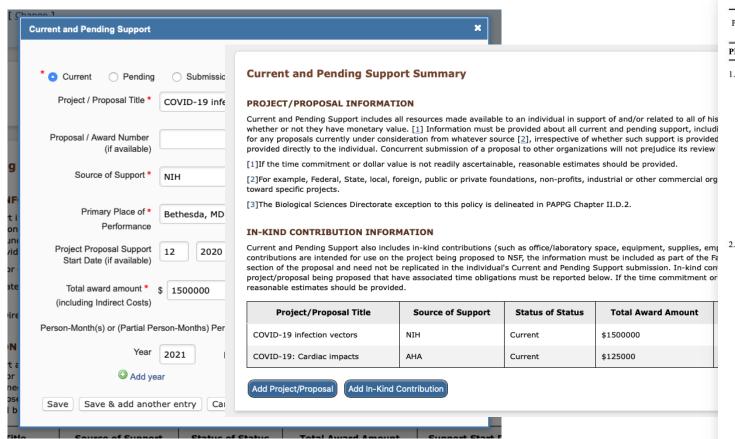
ts, Whether or Not Related to the Proposed Project

am GS, Guha R, Brimacombe K, Grossman A, Chung TDY, Weidner uld D, Dahlin JL, Devanaryan V, Foley TL, McGee J, Kahl SD, Kales ejcek B, Gal-Edd N, Glicksman M, Haas JV, Iversen PW, Hoeppner M, H, Trawick B, McVey J, Lemmon VP, Li Z, McManus O, Minor L, Pacifici R, Chin WW, Xia M, Xu X, Lal-Nag M, Hall MD, Michael S, Austin CP. Assay Guidance Manual: Quantitative Biology and nical Drug Discovery. Clin Transl Sci. 2018 Sep;11(5):461-470. PubMed Med Central PMCID: PMC6132369.

he National Center for Biotechnology Information, Nucleic Acids Res. D13. PubMed PMID: 29140470; PubMed Central PMCID:



SciENcv: NSF Current & Pending Support



NSF CURRENT AND PENDING SUPPORT

PI/co-PI/Senior Personnel: Trawick, Barton

PROJECT/PROPOSAL CURRENT SUPPORT

 Project/Proposal Title: COVID-19 infection vectors Proposal/Award Number (if available):

Source of Support: NIH

Primary Place of Performance: Bethesda, MD

Project/Proposal Support Start Date (if available): 2020/12

Project/Proposal Support End Date (if available): 2024/12

Total Award Amount (including Indirect Costs): \$1,500,000

Person-Month(s) (or Partial Person-Months) Per Year Committee

Year	Person-months per year committed
2021	12

2. Project/Proposal Title: COVID-19: Cardiac impacts

Proposal/Award Number (if available): 12345

Source of Support: AHA

Primary Place of Performance: Bowling Green, KY

Project/Proposal Support Start Date (if available):

Project/Proposal Support End Date (if available):

Total Award Amount (including Indirect Costs): \$125,000

Person-Month(s) (or Partial Person-Months) Per Year Committed

Year	Person-months per year committed	
2020	1	



PDF

CPS-1 of 1



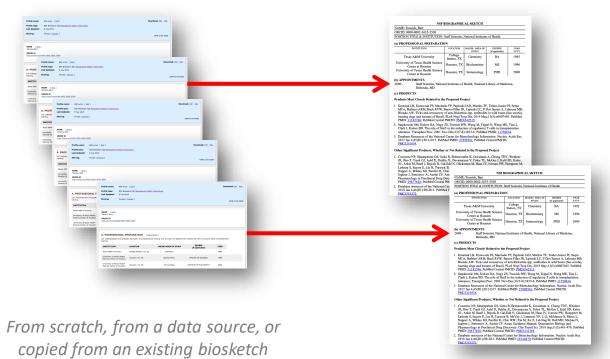


SciENcv Features

Create & store multiple biosketches plus C&PS forms

Output PDFs on demand

Have a delegate manage accounts







More information

- SciENcv is available at: http://www.ncbi.nlm.nih.gov/sciencv
- My NCBI documentation: http://www.ncbi.nlm.nih.gov/books/NBK3843/

- SciENcv video overview: http://youtu.be/PRWy-3GXhtU
- SciENcv for NSF Users: Biographical Sketches: https://youtu.be/nk7qlbele0k
- SciENcv for NSF Users: Current and Pending Support https://youtu.be/bhWQYvEEtJs

SciENcv questions/feedback: info@ncbi.nlm.nih.gov

NSF Login and usage questions: 1-800-381-1532 or rgov@nsf.gov

NSF Policy questions: policy@nsf.gov

New Award Notice PI Transfers

New Award Notices

- Multi-Year Project to Modernize NSF's Awards System
- Slowly Migrated New Functionality and Other Notices

- ✓ No Cost Extensions March 2018
- ✓ Other Administrative Actions June 2018
- ✓ Post-Award Funding Actions April 2019
- ✓ New and Renewal Awards January 2020
- ✓ Award Cancellations March 2020
- ✓ Award Reductions June 2020

New Award Notices – PI Transfers

- Same Standard, Consistent Layout
- No Longer a Letter
- Copy Sent to PI and co-PI(s)
- Clearly Identifies Information by Section

Scheduled for Release in late November 2020



Next Steps and ERA Forum Resources

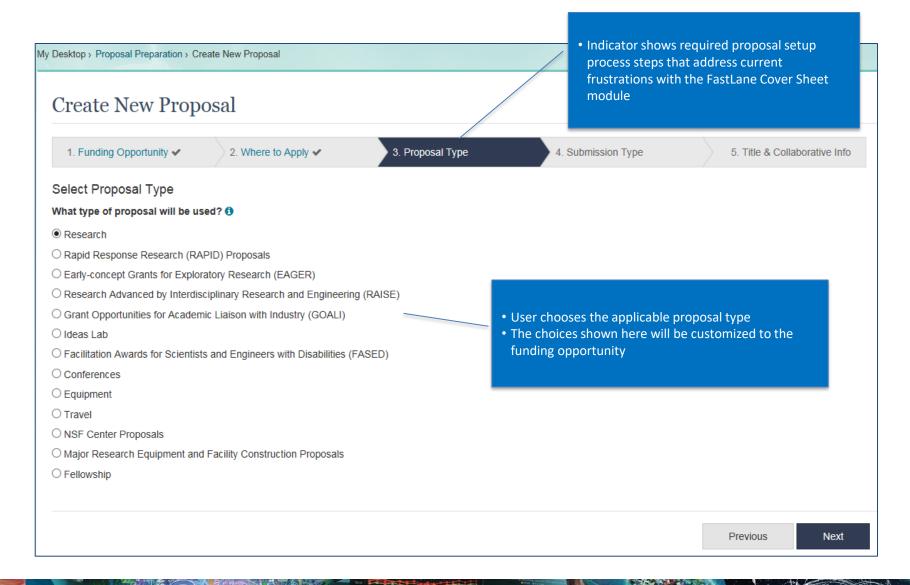
- ☐ Post-webinar Survey
 - https://www.surveymonkey.com/r/ERAForumSurveyNovember2020
- □ERA Forum Email
 - Questions pertaining to the presentation can be sent to nsferaforum@nsf.gov
 - If you are interested in participating in future application usability studies, please send an email with your contact information to nsferaforum@nsf.gov
- ☐ Subscribe to ERA Forum Listserv
 - Send an email to <u>Nsf-era-forum-subscribe-request@listserv.nsf.gov</u>
- ■NSF ERA Forum Website:
 - Agenda and presentation slides will be available soon on the ERA Forum website at: https://www.nsf.gov/bfa/dias/policy/era_forum.jsp.

Thank you for participating in today's ERA Forum Webinar

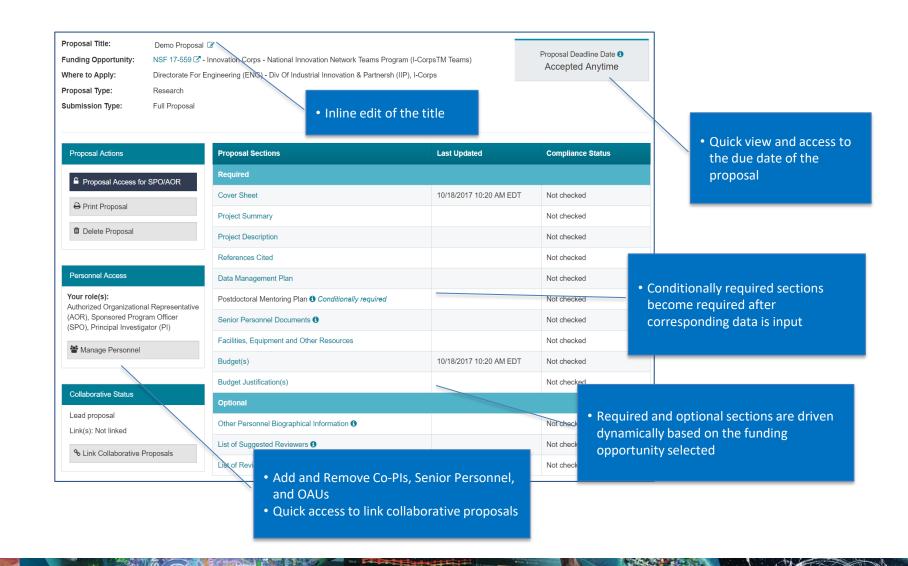
Appendix

Research.gov Demo Screenshots

Demo: Create New Proposal Wizard

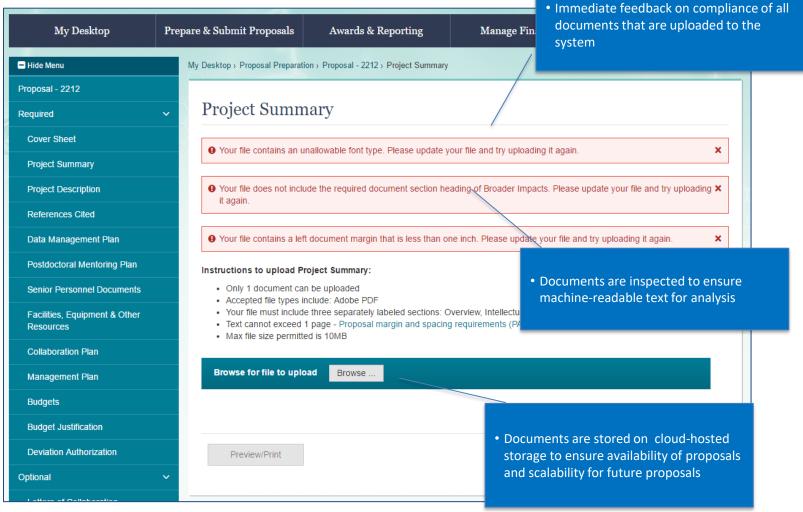


Demo: Main Proposal Page

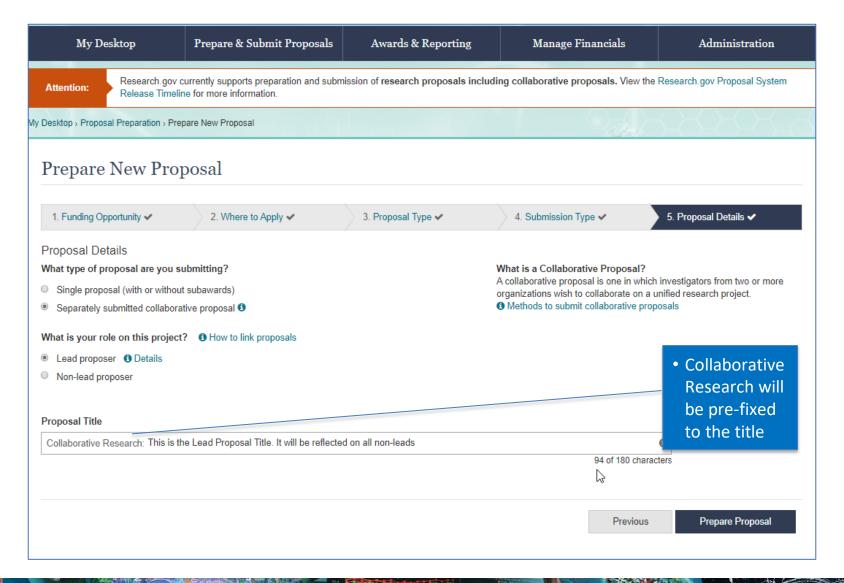


Demo: Document Upload and Compliance Error/Warning

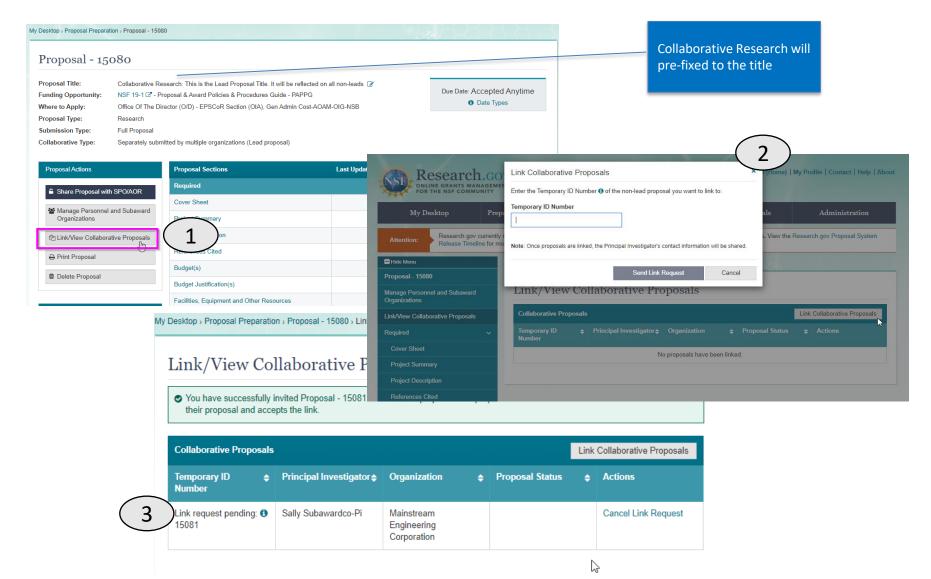
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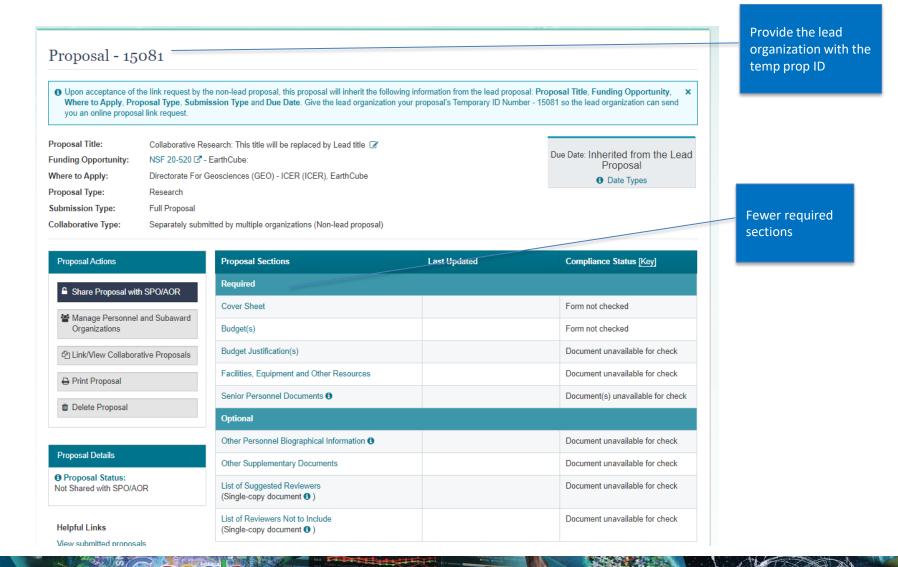
Demo: Separately Submitted Collaborative Proposals



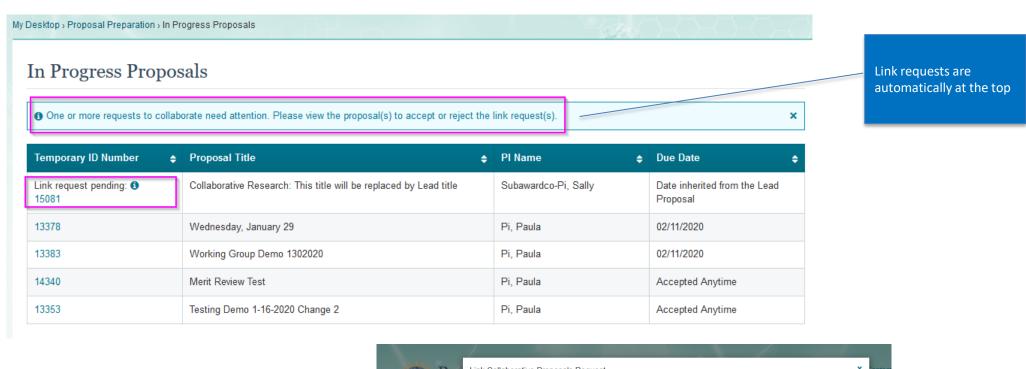
Demo: Separately Submitted Collaborative Proposals - Linking



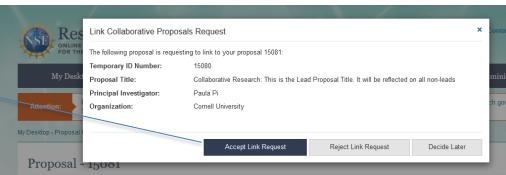
Demo: Separately Submitted Collaborative Proposals - Non-lead



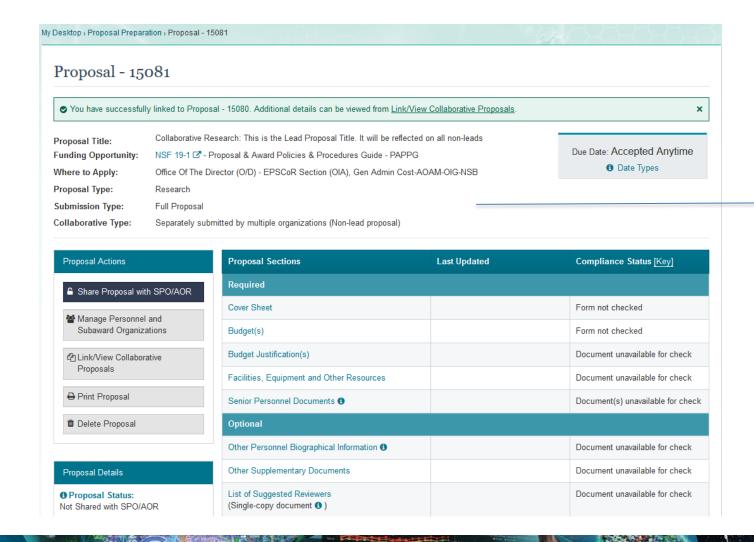
Demo: Separately Submitted Collaborative Proposals - Non-lead



Once link is accepted, non-lead cannot unlink. Must contact the lead

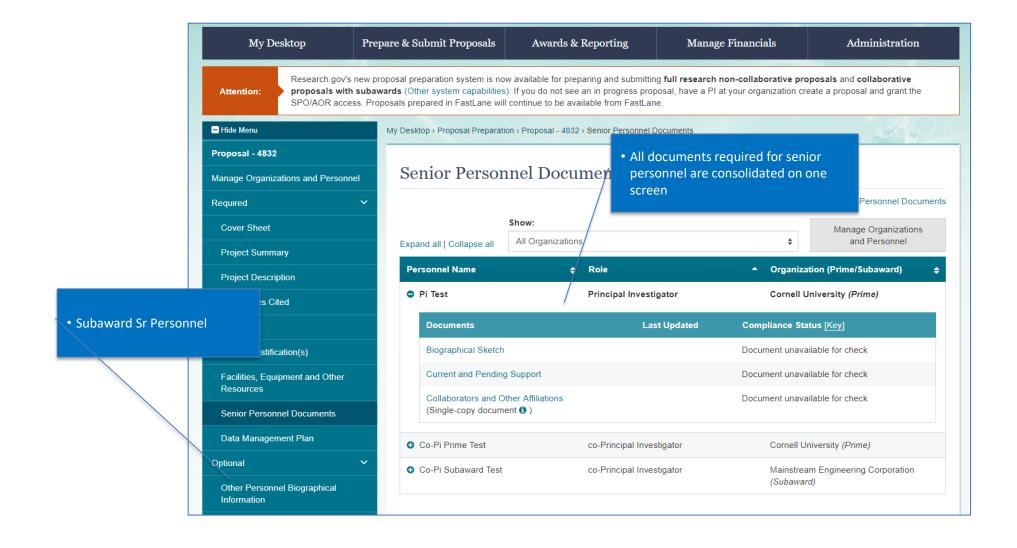


Demo: Separately Submitted Collaborative Proposals - Non-lead

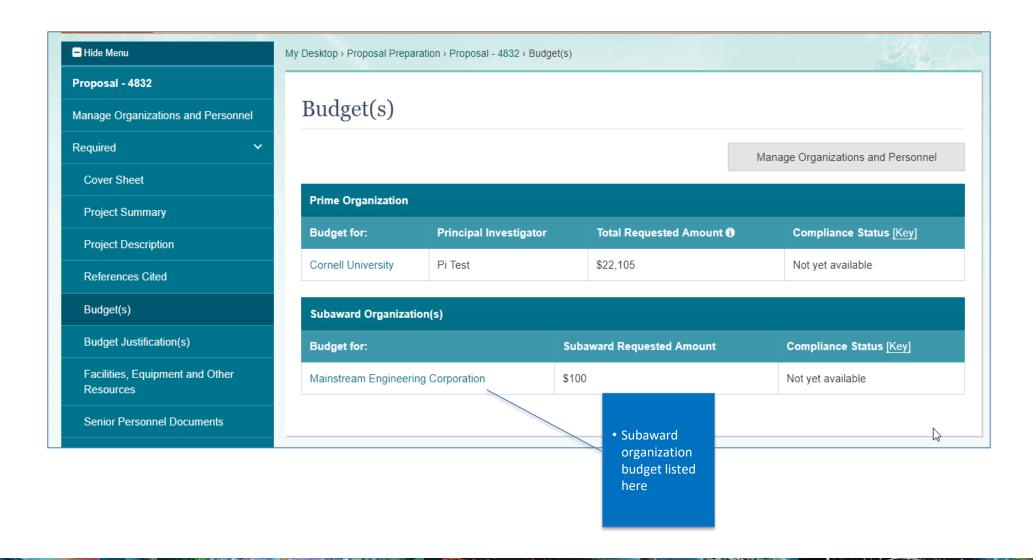


Once linked, non-lead will inherit Title, Funding Opportunity, Where to Apply (units of consideration), Proposal Type, Submission Type, and Due Date

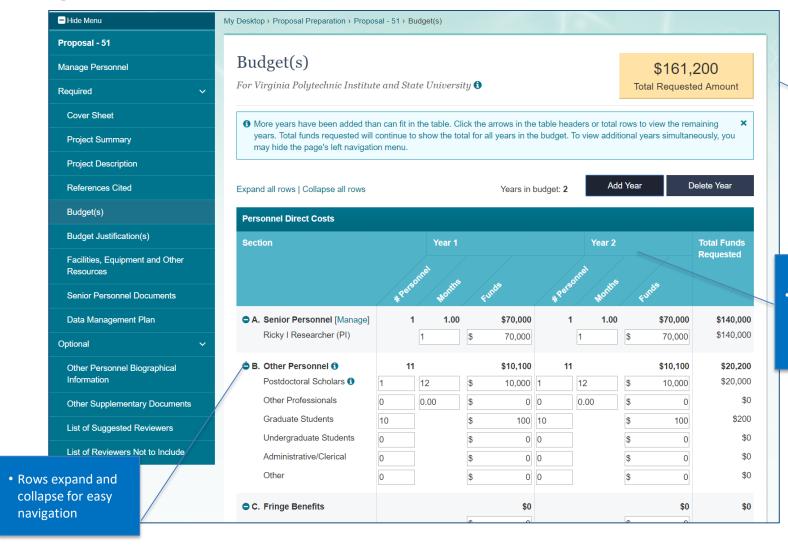
Demo: Senior Personnel Documents



Demo: Budget



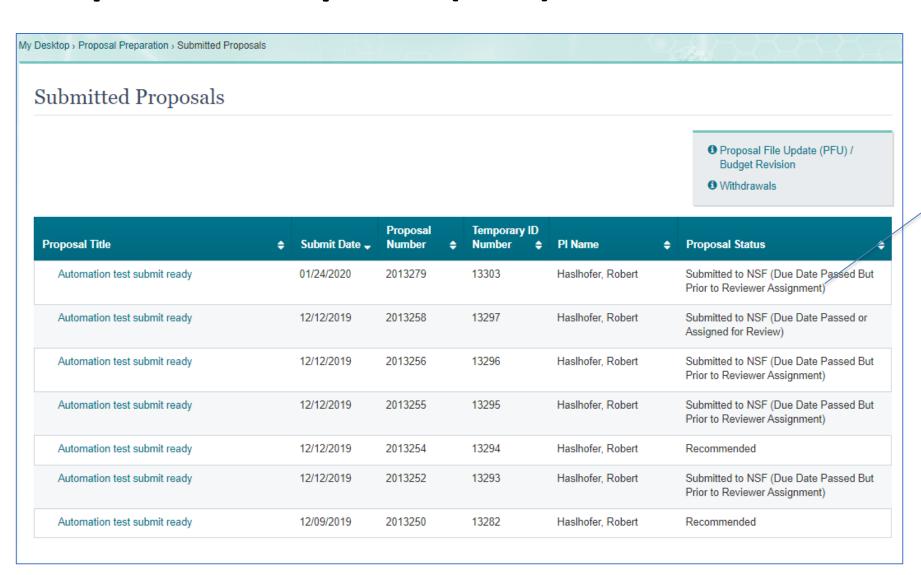
Demo: Budget



 All totals are dynamically summed and the total amount is placed on the **Cover Sheet**

 Multiple years displayed on one screen

Demo: Proposal File Update (PFU)



Prior to Review Assignment

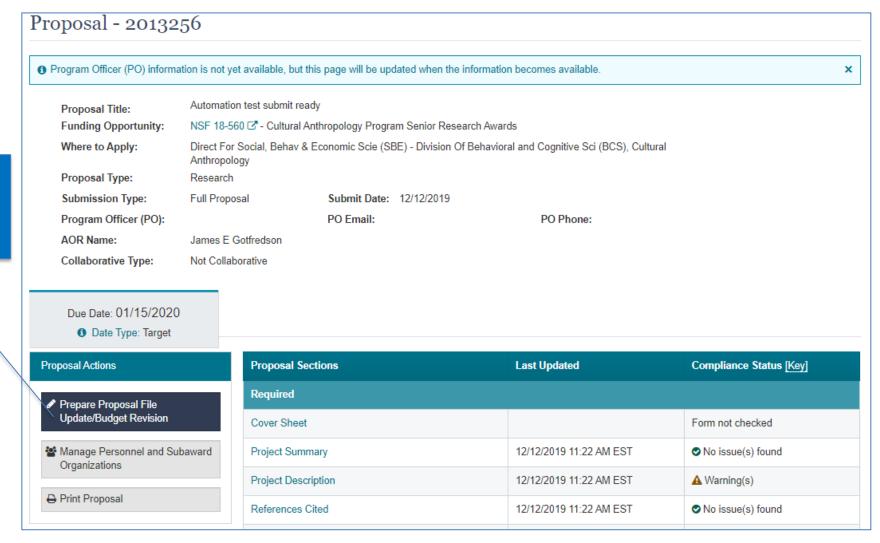
Demo: PFU

Prepare

Proposal

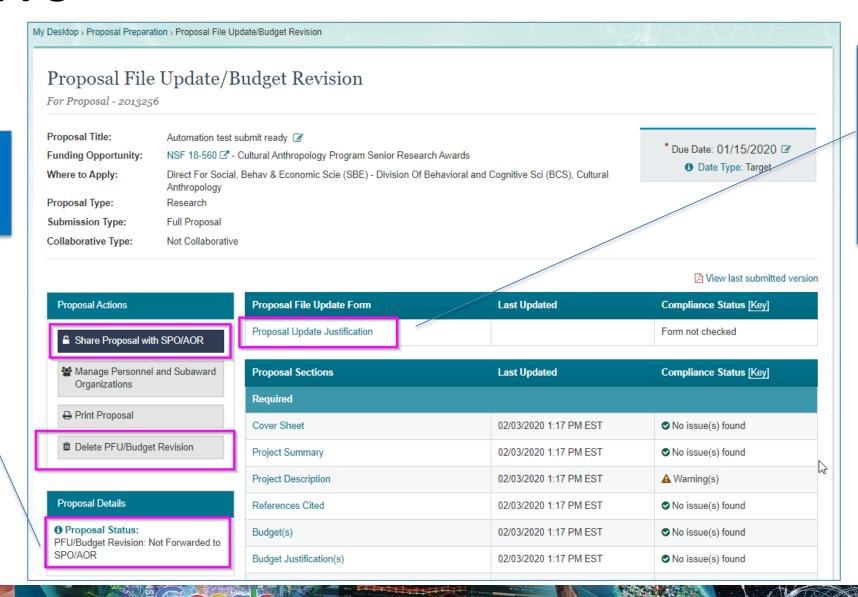
Revision

File/Budget



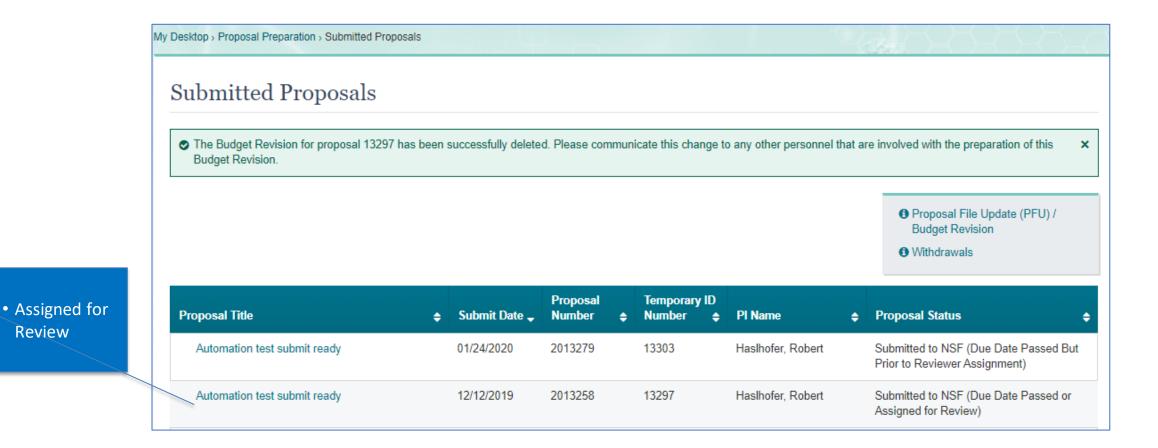
Demo: PFU

 Budget Revision Status



 Proposal Update Justification is required if there is a change in proposal content, not budget content

Demo: Budget Revision



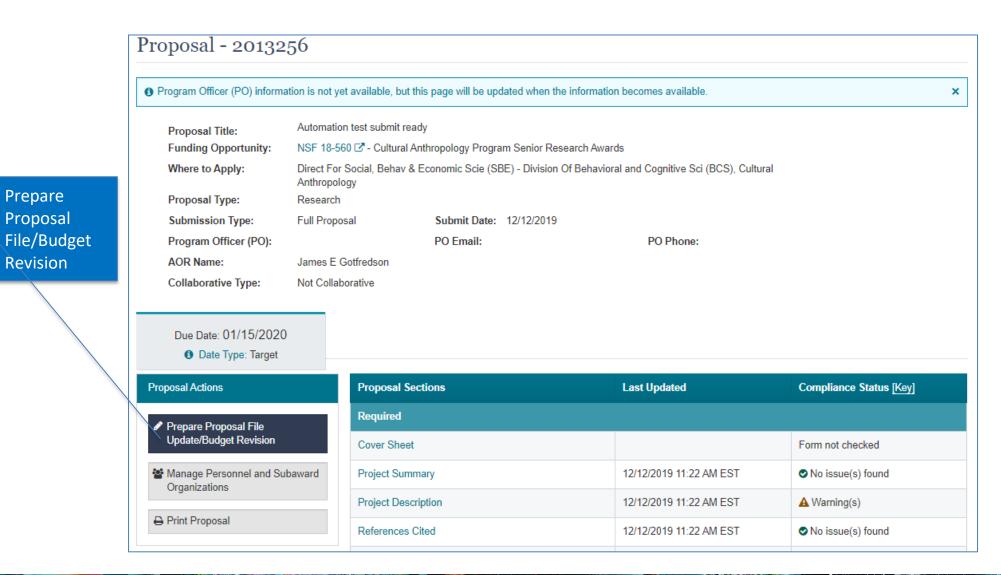
Review

Demo: Budget Revision

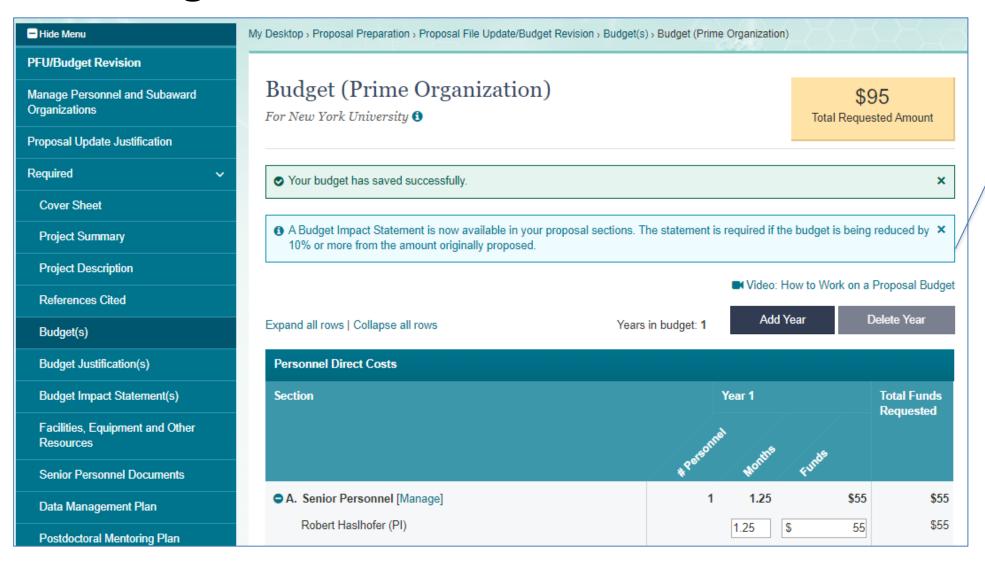
Prepare

Proposal

Revision



Demo: Budget Revision



 System will not enforce that the Budget Impact Statement exists if the budget is reduced by 10%