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SANTA BARBARA • SANTA CRUZ

THE CENTER FOR BIOMOLECULAR SCIENCE & ENGINEERING ENGINEERING BUILDING 2, ROOM 501 SANTA CRUZ, CALIFORNIA 95064

(831) 459-1477 (831) 459-1809 FAX

June 9, 2014

Dr. Simona Volpi, Program Director Genotype-Tissue Expression Project (GTEx) Division of Extramural Research 5635 Fishers Ln, Room 4107 Rockville, MD 20892

Dear Dr. Volpi,

In response to our recent communication regarding integration of GTEx data into the UCSC Genome Browser and the development of visualization for this, we provide here a proposal for a three year supplement to our Browser award (grant # 5U41HG002371-14).

We have also provided a detailed budget and justification, and a checklist with details on indirect costs (the university rate for this project is 53%, 53.5% and 54% successively in years 1-3 of the supplement).

We are requesting direct costs of \$710,655 and total costs of \$1,090,942.

Please let us know if you have any questions or need anything further for this proposal. We are extremely excited to be able to expand our browser to encompass the GTEx data.

Sincerely,

W. James Kent

Director, UCSC Genome Browser

Center for Biomolecular Science and Engineering

UC Santa Cruz

14. APPLICANT ORGANIZATION CERTIFICATION AND ACCEPTANCE: I certify that the statements herein are true, complete and accurate to the best of my knowledge, and accept the obligation to comply with Public Health Services terms and conditions if a grant is awarded as a result of this application. I am aware that any false, fictitious, or fraudulent statements or claims may subject me to criminal, civil, or administrative penalties.

lcrowder@ucsc.edu SIGNATURE OF OFFICIAL NAMED IN 13. (In ink. "Per" signature not acceptable.)

Tel: (831) 459-1378

DATE 6/9/14

FAX:

Tel: (831) 459-2778

dbeach@ucsc.edu

Name

Title

E-Mail:

Project Summary

In this supplement we propose one new specific aim – integrating the GTEx data into the UCSC Genome Browser. The NIH Common Fund's Genotype-Tissue Expression (GTEx) program measures genotype and gene expression in multiple tissues for many human donors. It has already released a substantial pilot data set at http://www.gtexportal.org, and is working on a larger production data set. We propose to integrate the data into the UCSC Genome Browser starting with the pilot phase data in the first year of this supplement, and extending to production data sets in later years. The integration will include at least two new tracks in the Genome Browser, one that shows the tissue-by-tissue expression data associated with each gene, and another that shows significant allele-specific differences in expression. We will also add new columns for the GTEx data to the Gene Sorter, and new panels to the existing gene details display. As the project progresses we anticipate adding a "phenotype associated significant gene expression differences" track as well.

Relevance

Gene RNA expression levels provide a detailed view of the inner state of a cell. Both genetically caused and other diseases are often associated with gene expression changes. RNA expression levels are helpful in understanding the normal as well as disease physiology of an organism. This work will thus help scientists develop better treatments for a broad range of diseases.

Supplement to 5U41HG002371-14, UCSC Genome Browser: Integrate GTEx data into the UCSC Genome Browser

In this supplement we propose to integrate the GTEx data into the UCSC Genome Browser for visualization and analysis. This will make the carefully collected, genotyped, and analyzed human RNA expression data from GTEx more accessible by the research community. It will also result in new display capabilities in the Genome Browser that may be useful for other data sets as well. We will use these capabilities to display allele-specific expression by SNPs that show significant eQTLs, as well as overall gene expression levels in a variety of tissues. The GTEx data will be displayed as annotation tracks aligned to the genome assembly, as part of the gene details page, and as columns in the Gene Sorter. We will update the GTEx data in the Genome Browser at least once yearly, and will synchronize new releases to match major data releases from GTEx when possible. UCSC will collaborate with the GTEx analysis team, integrating new analyses as well as new data, and advising on analysis methods when useful.

The pilot phase GTEx data is available now and includes RNA levels measured by a variety of methods on 45 tissues collected from 185 phenotypically normal donors, who were genotyped as well. Most of the tissues have samples from at least 20 different donors. This is enough to obtain very robust gene expression data compared to earlier projects, and in some cases to tie variations in gene expression to specific SNPs in a significant way as eQTLs. The GTEx project plans to extend these data, and to add data from donors with various phenotypes as well.

The first step in integrating the GTEx data into the browser will be to parse it, after which it will be formatted and indexed for fast display. Because the data are available and documented in reasonably simple formats at the GTEx portal, we anticipate that this step will take about 3 months. At the end of this step the pilot data will be made available in the UCSC Table Browser and on the Genome Browser download server.

Next we will develop a display for the expression information for a gene, similar to that shown in Figure 1. We will add this as a new section on the genes details page that displays when a user clicks on a gene in the Genome Browser. This step should take about 2 months.

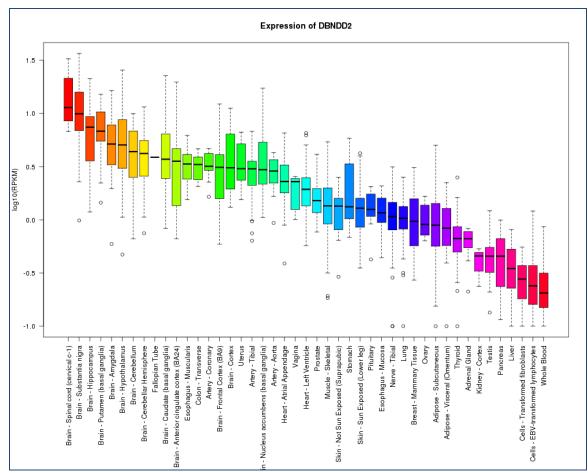
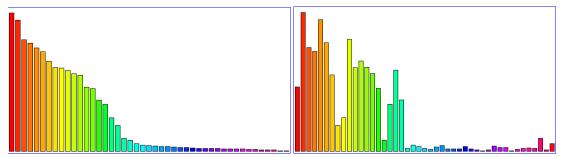


Figure 1. Gene expression Tukey boxplots for a single gene from the GTEx Portal. The order shown in these plots is always from highest expressed to lowest expressed tissue.

With this groundwork in place we will develop tracks for the Genome Browser. Since the browser displays data at many scales, from base level to complete chromosomes, the track graphics must work at many levels as well. For this reason the predominant view at the gene level will be a bar graph (Figure 2a), which is readable at smaller scales than the Tukey boxplots in Figure 1. Frequently users look at multiple genes simultaneously. Because of this we will keep a uniform, logarithmic-based scale for all genes, and we will order the tissues identically for all genes. The uniform ordering will be arranged such that related tissues are in close proximity (Figure 2b).



Figures 2a & 2b. Bar graphs sorted by gene expression (left (2a)) and by a canonical ordering of tissue types (right (2b)). Though some detail is lost, the bar graph is legible at a greater range of scales than the Tukey plot in Figure 1. The canonical ordering is preferred when expression patterns for multiple genes or multiple alleles of the same gene are shown together, since the two figures can be compared visually.

We will develop two Genome Browser tracks to represent the pilot data. One track will show an item for each gene, and will display a bar graph with a height of ~ 100 pixels at each gene for which there is GTEx data. The

second track will show an item for each SNP associated with an eQTL. It will display expression graphs for both alleles, one extending up and one extending down from a common axis (Figures 3 and 4). These new tracks will require 3-5 months to develop.

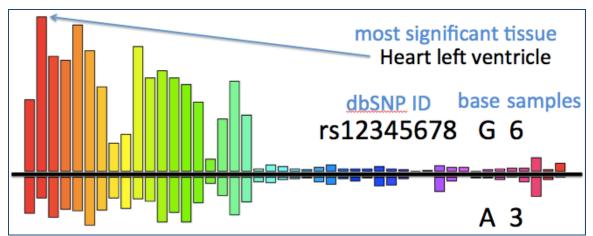


Figure 3. Mock-up of imagery and labeling for an eQTL SNP track item. Here the G allele would be associated with higher expression in general, and particularly higher on the biosample second from left.

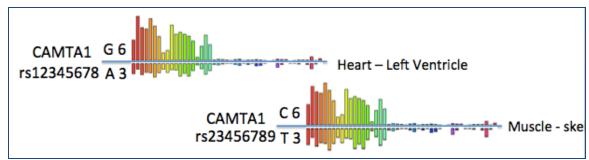


Figure 4. Mock-up showing how two nearby eQTL SNPs might appear in a track. Each item would include the gene name, SNP name, tissue where most significant difference is observed, allele frequencies at that SNP for that tissue, and median expression graph for each allele. It is possible that we could include an overall p-val for the QTL in the display at this level. As with all our data tracks, more detail about an item would be available by clicking on it. This would include links back to the GTEx portal where appropriate.

We will finish the first year of the supplement and the integration of the pilot phase GTEx data by adding the data to the UCSC Gene Sorter. The Gene Sorter display is useful for visualizing expression levels on multiple genes at once, and for selecting gene sets by expression, GO annotation, and many other criteria.

In subsequent years of this project we will update the UCSC databases with new data from GTEx, and as necessary, develop new tracks and other new displays as novel types of data become available. We will answer user questions about the GTEx data, and when possible enhance the GTEx displays and analysis according to user suggestions, and according to feedback from the GTEx scientists.

DETAILED BUDGET FOR INITIAL BUDGET PERIOD DIRECT COSTS ONLY

FROM 7/1/14 THROUGH 6/30/17

List PERSONNEL (Applicant organization only)
Use Cal, Acad, or Summer to Enter Months Devoted to Project
Enter Dollar Amounts Requested (omit cents) for Salary Requested and Fringe Benefits

CONSORTIUM/CONTRACTUAL COTAL DIRECT COSTS FO	COSTS	GET PE	RIOD	FAG	CILITIES AND	ADMINISTRATI			228,659
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CONSORTIUM/CONTRACTUAL (.				CT COSTS		
OTHER EXPENSES (Itemize by o	category)								
ALTERATIONS AND RENOVATION	ONS (Itemize by cate	gory)							
OUTPATIENT CARE COSTS									
INPATIENT CARE COSTS	gs and other pro	ject-reid	aleu ii a	ivei (4 ii	ips at \$1,0	boo per mp)			6,000
TRAVEL Travel to agency meetings and other project-related travel (4 trips at \$1,500 per trip)									
SUPPLIES (Itemize by category) Minor equipment (e.g. lap	otop workstation	i) - \$2,5	600; pro	ject-spe	ecific supp	lies, software	e - \$1,000		
EQUIPMENT (Itemize)									
CONSULTANT COSTS									
	SUBTOTALS				→	154,784	64,37	5	219,159
TBD	Grant Administratio	1							
Donna Karolchik	Genome Browser	1							
Ann Zweig	Eng & Proj Mngr	1						D	
TBD	Quality Assurance	4							
Kate Rosenbloom Pohl	Tech. Proj. Mngr	12							
Jim Kent	PD/PI	1							
		Mnths	Mnths	Mnths	SALARY	REQUESTED	BENEFITS		TOTAL

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BUDGET FOR ENTIRE PROPOSED PROJECT PERIOD DIRECT COSTS ONLY

BUDGET CATEGORY	INITIAL BUDGET	2nd ADDITIONAL	3rd ADDITIONAL	4th ADDITIONAL	5th ADDITIONAL
TOTALS	PERIOD (from Form Page 4)	YEAR OF SUPPORT REQUESTED	YEAR OF SUPPORT REQUESTED	YEAR OF SUPPORT REQUESTED	YEAR OF SUPPORT REQUESTED
PERSONNEL: Salary and fringe benefits. Applicant organization only.	219,159	228,923	239,073		
CONSULTANT COSTS	210,100	220,323	200,070		
EQUIPMENT					
SUPPLIES	3,500	1,000	1,000		
TRAVEL	6,000	6,000	6,000		
INPATIENT CARE COSTS					
OUTPATIENT CARE COSTS					
ALTERATIONS AND RENOVATIONS					
OTHER EXPENSES					
DIRECT CONSORTIUM/ CONTRACTUAL COSTS					
SUBTOTAL DIRECT COSTS (Sum = Item 8a, Face Page)					
F&A CONSORTIUM/ CONTRACTUAL COSTS					
TOTAL DIRECT COSTS	228,659	235,923	246,073		

JUSTIFICATION. Follow the budget justification instructions exactly. Use continuation pages as needed.

See continuation page

Personnel

Salaries are projected to increase 35 annually.

<u>PI W. James Kent (0.083 FTE; 1 month/year)</u> – Design and prototype of software tools. Scientific oversight. Communicate with GTEx project and with funding agency.

<u>Technical Project Manager Kate Rosenbloom Pohl (1.0 FTE; 12 months/year)</u> – Refine design and implement production software associated with tracks and other features. Communicate with GTEx project.

QA TBD (0.33FTE; 4 months/year) – Test software, provide usability feedback, and help move software from development to test to production servers.

<u>Engineering & Project Manager Ann Zweig (0.083 FTE; 1 month/year)</u> – Project manager for genome browser. Tracks and coordinates work and release on new features and new data sets.

<u>Genome Browser Manager Donna Karolchik (0.083 FTE; 1 month/year)</u> – Provides personnel management for programming staff, helping communication, managing performance, and assembling yearly performance reviews.

<u>Grant Administration TBD (0.083 FTE; 1 month/year)</u> — Provide oversight and coordination of proposals, reporting, financial management, resources and personnel.

Supplies

Funds are requested for a laptop workstation or other minor equipment in year 1 (\$2,500). An additional \$1,000 per year is requested each year for project-specific supplies and software.

Travel

Funds are requested for 1-2 travelers to GTEx meetings and workshops in the Washington DC area (4 trips at \$1,500 each).

4. DISCLOSURE PERMISSION STATEMENT: If this application does not result in an award, is the Government permitted to disclose the title of your proposed project, and the name, address, telephone number and e-mail address of the official signing for the applicant organization, to organizations that may be interested in contacting you for further information (e.g., possible collaborations, investment)?

OMB No. 0925-0001
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Checklist Form Page