

The DNA sequences of the human genome are now accessible by more than a hundred databases and searching tools on the World Wide Web. In this brief guide, however, I'll focus on just one (relatively) easy-to-use tool, the UC Santa Cruz (UCSC) Genome Browser:

Primary Resource:

The UCSC Genome Browser: http://genome.ucsc.edu/cgi-bin/hgGateway

The UCSC Browser can be used in a number of ways. In this guide, I'll focus on just one — using the Browser to locate and investigate genes of interest. As an example, let's suppose that you've been asked to locate and investigate a gene called **RS11**. First, open a web browser and type in the UCSC Genome Browser address:

http://genome.ucsc.edu/cgi-bin/hgGateway

A window will open, the top part of which looks something like this:

| Human Genor | ne Browser G | ateway | | | |
|--|--------------------|---------------------|------------------------------|----------------|--|
| The UCSC Genome Browser was created by the <u>Genome Bioinformatics Group of UC Santa Cruz</u> . Software Copyright (c) The Regents of the University of California. All rights reserved. | | | | | |
| clade | genome | assembly | position | image width | |
| Vertebrate 🛟 | Human 🛟 | May 2004 🛟 RS | 11 | 620 submit | |
| <u>Click here to reset</u> the browser user interface settings to their defaults. | | | | | |
| (| add your own custo | m tracks (configure | tracks and display (clear po | osition | |

Since we're looking for **RS11**, we'll type "RS11" into the "position" window (as shown above) and hit the **Submit** button. After a few seconds the Browser will have found sequences that match the string you've entered, and a new window will appear:



It's usually best to take the appropriate link under the first heading that appears. That may be either the "Known Genes" heading, or "RefSeq Genes," as shown below for the "Clock" gene:

12 11.2 11.2 12

92 Mh



Click on either the "Known" or "RefSeq" link, and a window will open showing the location of the gene you've chosen. Here's the window for RS11:

| position | chr19:54,691,499-54,694,754 | 4 jump clear Si | ze 3,256 bp. configure |
|--------------|--|---|---|
| chr19 | (q13.33) 19p13.3 p13.2 | 19p12 19q12 | q13.2 |
| Base Posit | ion 54692000 5469250 Known Genes (Nov | 0 54693000 54693500 22, 04) Based on SWISS-PROT, | 54694000 54694500 TrEMBL, mRNA, and RefSeq |
| RP: AK130 | \$11 324 | | ************************************ |

Don't be surprised if your window looks <u>a lot</u> more complex than the one shown above. The Browser allows you to display so much information that it can be truly confusing. So, try the following. First, hit the "Hide All" button:

hide all

This will close all of the data display options. And now, open just two of them that you're going to use: The "Known Genes" button and the "Base Position" option:



Hit the "Refresh" button, and your window should now look just like the one above, displaying a manageable amount of information. As you can see, there's a map of our gene (actually called RPS11) showing its exact position (on chromosome 19), and several little blocks showing that it consists of 5 exons and is transcribed from left to right. The window also allows you to move to the left or the right along the chromosome, and to zoom in or out to look at the "neighborhood" around the RPS11 gene.

For detailed information, click on RPS11 and a new window will open with a brief, but informative description of the gene and (in this case) its protein product:

| Human Gene RPS11 Description and Page Index | | | | | | |
|--|------------------|-------------------|------------|---------------|--|--|
| Description: 40S ribosomal protein S11. Alternate Gene Symbols: NM_001015 Representative mRNA: <u>BC007283</u> Protein: <u>P62280</u> (aka RS11_HUMAN) RefSeq Summary: Ribosomes, the organelles that catalyze protein synthesis, consist of a small 40S subunit and a large 60S subunit. Together these subunits are composed of 4 RNA species and approximately 80 structurally distinct proteins. This gene encodes a ribosomal protein that is a component of the 40S subunit. The protein belongs to the S17P family of ribosomal proteins. It is located in the cytoplasm. The gene product of the E. coli ortholog (ribosomal protein S17) is thought to be involved in the recognition of termination codons. This gene is co-transcribed with a small nucleolar RNA gene, which is located in its third intron. As is typical for genes encoding ribosomal proteins, there are multiple processed pseudogenes of this gene dispersed through the genome. | | | | | | |
| Page Index Quick Links U | IniProt Comments | Sequence | Microarray | RNA Structure | | |
| Protein Structure Other Species G | O Annotations | mRNA Descriptions | Pathways | Methods | | |

As you can see, the description on this page gives us the basic facts about the protein, which turns out to be one of the proteins in the small ribosome subunit. To find even more about RPS11, click on any of the other links on the page.

The "Sequence" links give the DNA sequence of the gene, a comparison of the human version of the gene with its corresponding gene in the mouse ("Comparative"), the mRNA sequence, and the Amino Acid sequence of the protein.

To get detailed information about the gene, click on one of the "Tools and Databases" links:

Quick Links to Tools and Databases

| Genome Browser | Gene Sorter | Proteome Browser | UniProt | Entrez Gene | PubMed |
|----------------|--------------|------------------|---------|-----------------|------------|
| OMIM | GeneLynx | GeneCards | CGAP | Stanford SOURCE | ExonPrimer |
| Ensembl | Jackson Labs | H-INV | | | |

The "OMIM" (On-Line Mendelian Inheritance in Man) or the Stanford "SOURCE" links are especially useful, and will link you to a wealth of information about the gene. Here's what the OMIM window for our RS11 search:



Either of these will tell you the scientific background of the gene, list any genetic disorders associated with the gene, and refer you directly to scientific publications on the gene or its protein product.

The "Structure" link will even show you the three-dimensional structure of the protein product (if known):



Happy Gene Hunting!