Citation

How to cite GREAT

- If you use GREAT in the context of human or mouse, please cite:

- If you use GREAT in the context of zebrafish, please cite:

Why Cite GREAT

Over the long run, the best way to assure the continued availability and development of GREAT is for us to be able to show its relevance and contribution to biomedical research. We thank you for your support!

Publications & Reprints

The Nature Biotechnology paper is currently the inaugural GREAT publication. Reprints are available at http://bejerano.stanford.edu/publications.html#ref27.

Some of the papers that cite GREAT

See Google Scholar for a more up to date list.

We really appreciate people’s supportive e-mails and re-cap a handful of them here to encourage folks to take the plunge.

Date: April 21, 2011
Hi Gill,

Just wanted to let you know that we tried GREAT on a new kind of genomic data set after your talk. It worked amazingly well and made sense with respect to known biology. Keep up the good work!

Subject: your tool is cool

Date: October 19, 2010
Dear GREAT developers

I find this tool very useful in analyzing my ChIP-seq peaks, especially after being quite frustrated using other tools to get insight into the functions/pathways associated with my TF. ...

Subject: Re: great

Date: May 5, 2010
Hey guys, I had a chance to try out GREAT for a bit with our ... chip-seq data. It looks like an amazing tool and it’s way ahead of anything else out there for chipseq-related functional analysis. My favorite part is being able to click through each annotation and seeing the actual genes that were assigned.. very fancy and cool. That and the integration with the mouse phenotype ontologies--a big thing that other tools don’t have. Anyway we’ll definitely be using it and I’ll be sure to spread the word!
Hi ... 
This is not a ... ad. GREAT is awesome, we have been playing around with it quite a bit.

... 
you certainly will be referenced in a paper that we want to submit by the end of this month. And I suspect many more after that.