



Phenosaurus

Getting started guide

Fishtail plot of single screen

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FISHTAIL PLOT

GENELOT

LIST ALL GENES

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Select Screen

Select action on click:

Textsize
(px):

P-value cutoff (can also be written as 1E-xx):

o Label all significant hits:

List all significant genes in table:

IRF1

[Link to Genecards](#)

11px

5E-2

Submit

IRF1 using CST Antibody #8478 (induced with 10 ng/ml IFN gamma 24hr)

Select the screen
you want to plot

Here you can
define the
behavior of what
happens when
you click on a
gene in the graph

Optionally you can manually change the p-value cutoff to change which genes are colored and/or labeled

Label all genes
that meet the p-
value threshold

Hover over a gene
to see its p-value

P-Value: 7.755e-27
Gene: IRF1

Insertions [10log]

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Select Screen

IRF1

Select action on click:

Link to Genecards

Textsize (px):

11px

P-value cutoff (can also be written as 1E-xx):

5E-2

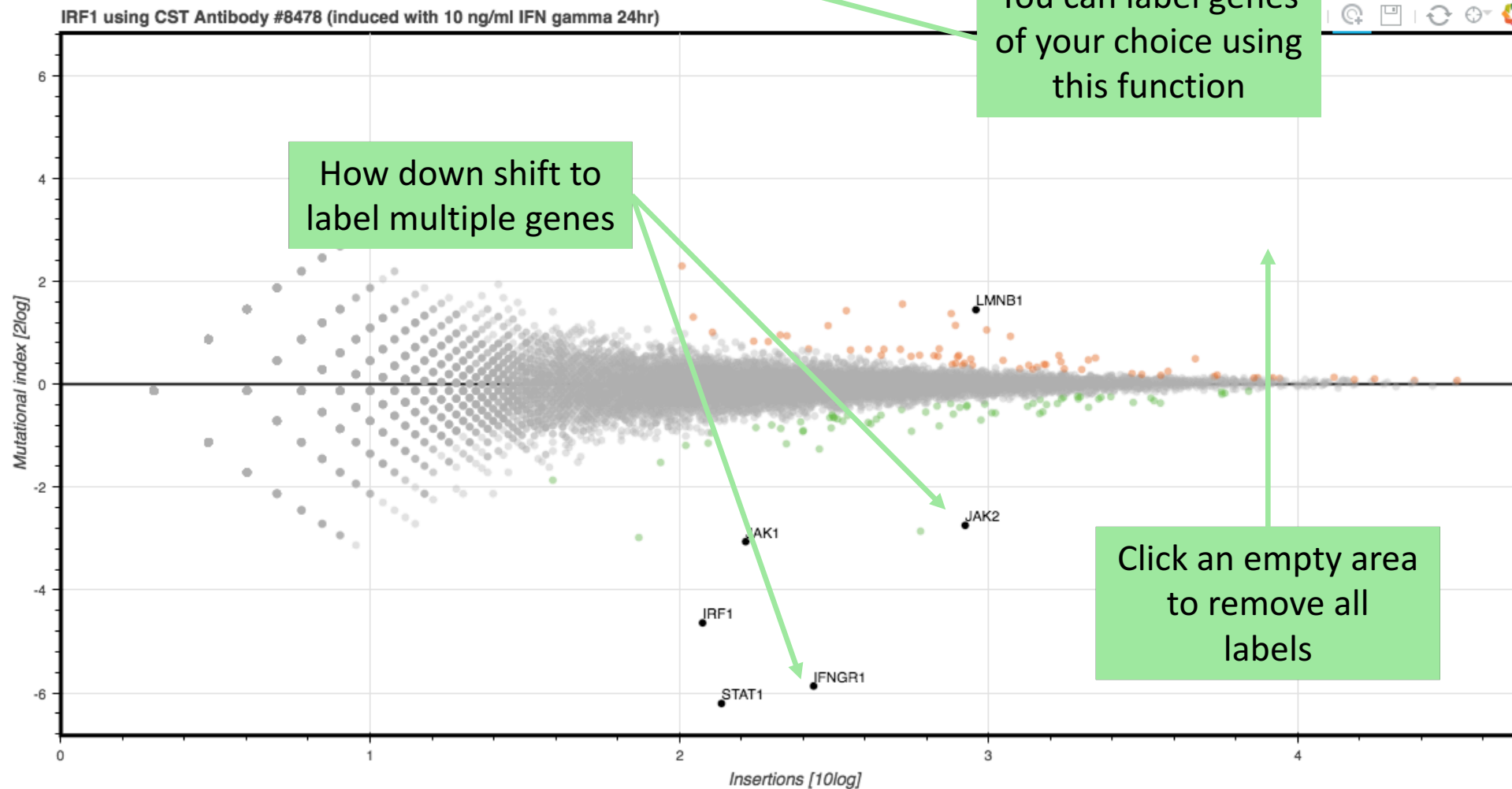
Label all significant hits:

☐

List all significant genes in table:

☐

Submit



Gene Finder

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Select screen(s)

glycosylated LAMP1
GNB1
GNB1 in KCTD5 knockout
Histone H2A(K119) crotonylation
Histone H3(K27) trimethylation
IRF1
non-phospho-beta-Catenin
phospho-AKT
phospho-AKT in KCTD5 knockout
phospho-ERK
phospho-p38
spliced XBP1

Enter genename(s), space separated:

KRAS

P-value cutoff (can also be written as 1E-xx):

0,05

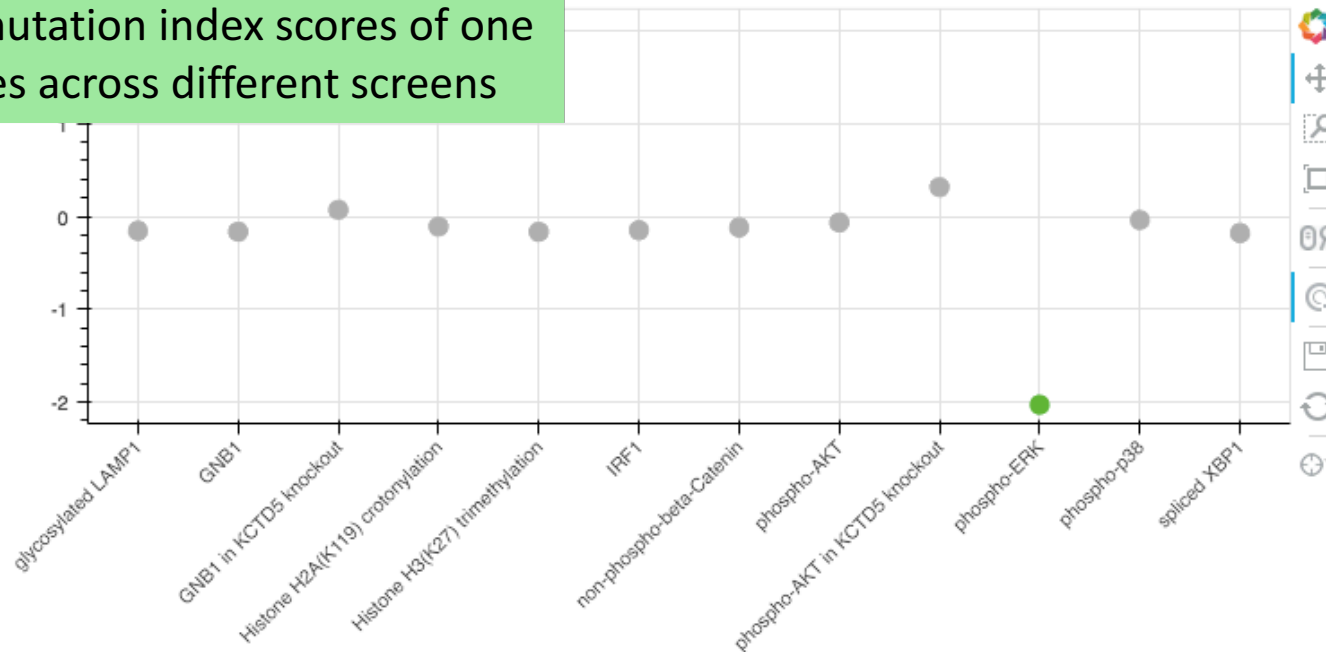
Plot width:

small: 800px

Enter one ore more gene names (space separate
and CAPITALIZED)

Select one or more screens,
hold down SHIFT to select multiple
or use ⌘-A / CTRL-A to select all

The 'Geneplot' functions allows you to
compare the mutation index scores of one
or more genes across different screens



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Genes

name	chromosome	orientation
A1BG	19	-
A1CF	10	-
A2M	12	-
A2ML1	12	+
A3GALT2	1	-
A4GALT	22	-
A4GNT	3	-
AAAS	12	-
AACS	1	+
AADAC	1	+
AADACL2	5	+
AADACL3	1	+
AADACL4	1	+
AADAT	4	-
AAED1	9	-
AAGAB	15	-
AAK1	2	-
AAMDC	11	+
AAMP	2	-
AANAT	17	+
AAR2	20	+
AARD	8	+
AARS	16	-
AARS2	6	-
AARSD1	17	-
AASDH	4	-
AASDHPPT	11	+
AASS	7	-
AATF	17	+
AATK	17	-

Here you can find all genes in the database
and identify their name

On the bottom right of the graph is a bar that lets you zoom, scale, reset the zoom-level and save the graph



Drag a certain area of the graph to the center with your arrow

Draw a box around a region of interest and zoom into it

Resize the whole window of the graph

The button lets you zoom in using the scroll function of your mouse. If you put your arrow OUTSIDE the graph next to an axis it allows you to zoom in only over that axis

Save the graph to a PNG file. SVG files are unfortunately not yet supported.

Reset the layout of the graph to its default view.

Questions?

- We are more than curious about your experience with the tool, so please feel free to contact us with any questions or suggestions you might have.
- Found a bug? Please let us know!
- Contact: Elmer Stickel e.stickel@nki.nl