

Firehose Reports | At-a-Glance

→ Reports are compatible with Firefox 4+, Chrome 12+, Safari 5+, Opera 11+ and Internet Explorer 9+.

Navigation: UP, DOWN, LEFT, RIGHT arrows for navigating between reports or the overview page.

Expand/Collapse: EXPAND ALL, COLLAPSE ALL buttons to manage report sections.

Printability: PRINT REPORT button to load a printable version.

Feedback: REPORT A PROBLEM button to contact the report maintainer.

Responsive Design: SET AUTO WIDTH button to fit the report to the browser window.

Significance: Red markers and red boxes highlight statistically significant results.

Interactivity: Clickable figures and table headers (e.g., GET FULL TABLE) for further exploration.

Supplementary Information: Clickable underlined results in tables to view associated supplementary data.

Download Results: Download Results section for downloading analysis files from the TCGA Data Coordination Center.

Table 1: Amplifications Table

Cytoband	Q value	Residual Q value	Wide Peak Boundaries	# Genes in Wide Peak
7p11.2	0		chr7:34954372-54968011	0 [EGFR]
12q14.1	5.19228	6.2028-113	chr12:56411663-5644647	5
4q12	6.7649e-85	6.7649e-85	chr4:34727006-34861623	1
12q15	1.5248e-57	1.7421e-57	chr12:200564385-202813140	2
12q16	3.8162e-70	4.0322e-71	chr12:67427108-67531544	2
3p06.33	4.5642e-09	4.5642e-09	chr3:18284087-183044402	2
7q31.2	9.0818e-09	1.7003e-08	chr7:16161324-16167511	1
12p13.33	2.4873e-08	2.4873e-08	chr12:3839333-430326	3
10q14	2.0116e-07	4.0273e-07	chr10:4105213-242804011	5
7p12.1	1.2098e-06	3.7782e-06	chr7:91966270-9268284	5
11p15.21	1.7964e-05	1.7964e-05	chr11:1732335-1429524	1
2p24.3	4.5845e-05	4.5845e-05	chr2:1933336-16904271	2
19p13	0.03487	0.03487	chr19:10856148-109682638	3
19q12	0.059145	0.059145	chr19:34867390-35007574	2

Table 2: Deletions Table - 62 significant deletions found.

Table S1: Genes in Wide Peak

Genes	# Genes in Wide Peak
CDK4	1
CYP27B1	1
TSPAN31	1
MARCH9	1
AGAP2	1