

RNAseq ID: ARHGAP11B =
Survival: OS
Auto select best cutoff: checked
Follow up threshold: all
Censore at threshold: checked
Compute median over entire database: false
Cutoff value used in analysis: 2.86
Expression range of the probe: -1 - 5
Probe set option: user selected probe set
Invert HR values below 1: not checked

Restrictions

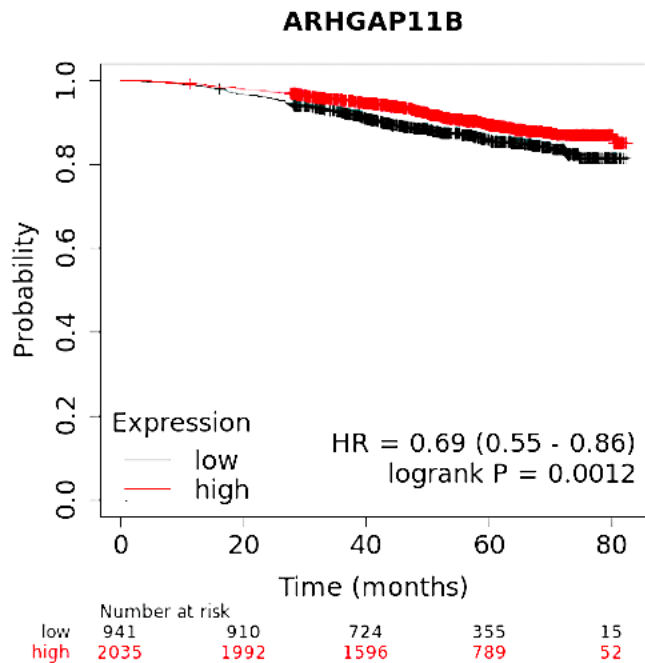
Lymph node status: all
 ER status: all
 PGR status: all
 HER2 status: all
 KI67 status: all
 Nottingham histologic grade: all
 PAM50 subtype: all

Cohorts

Endocrine treated: all
 Chemo treated: all

Results

P value: 0.0012
FDR: over 50%



RNAseq ID: ARHGAP11B
Survival: OS
Auto select best cutoff: checked
Follow up threshold: all
Censore at threshold: checked
Compute median over entire database: false
Cutoff value used in analysis: 3.17
Expression range of the probe: 2 - 5
Probe set option: user selected probe set
Invert HR values below 1: not checked

Restrictions

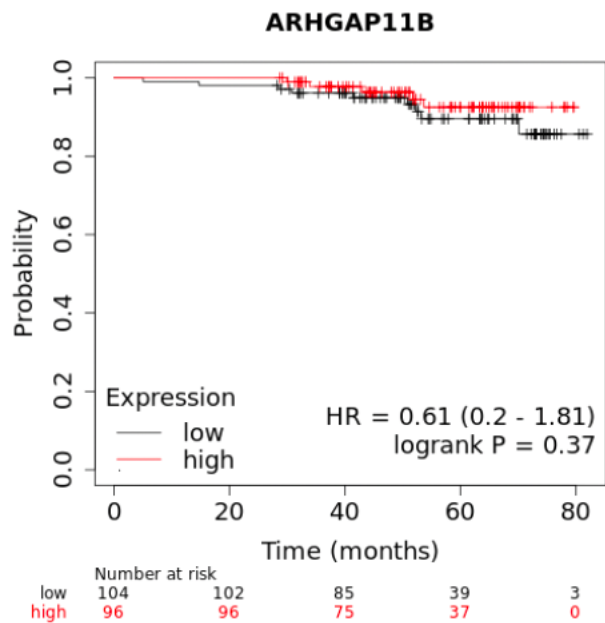
Lymph node status: all
 ER status: all
 PGR status: all
 HER2 status: all
 KI67 status: all
 Nottingham histologic grade: all
 PAM50 subtype: Normal

Cohorts

Endocrine treated: all
 Chemo treated: all

Results

P value: 0.3658
FDR: 100%



RNAseq ID: ARHGAP11B
Survival: OS
Auto select best cutoff: checked
Follow up threshold: all
Censore at threshold: checked
Compute median over entire database: false
Cutoff value used in analysis: 3.03
Expression range of the probe: 1 - 4
Probe set option: user selected probe set
Invert HR values below 1: not checked

Restrictions

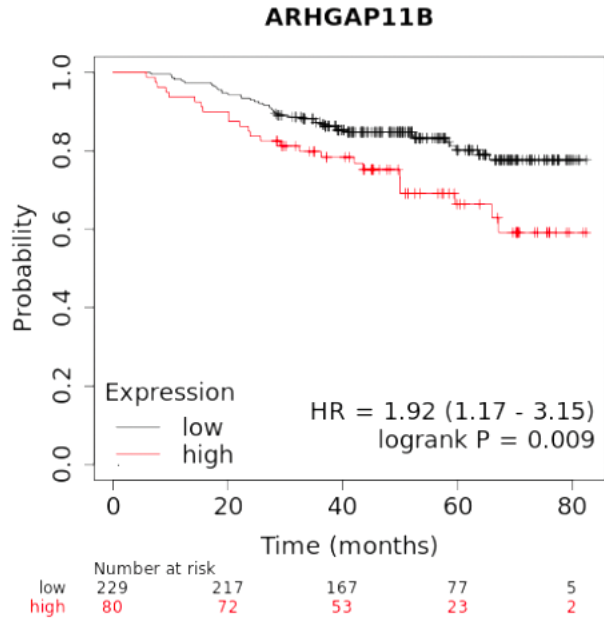
Lymph node status: all
 ER status: all
 PGR status: all
 HER2 status: all
 KI67 status: all
 Nottingham histologic grade: all
 PAM50 subtype: Basal

Cohorts

Endocrine treated: all
 Chemo treated: all

Results

P value: 0.009
FDR: over 50%



RNAseq ID: ARHGAP11B
Survival: OS
Auto select best cutoff: checked
Follow up threshold: all
Censore at threshold: checked
Compute median over entire database: false
Cutoff value used in analysis: 3.24
Expression range of the probe: 1 - 5
Probe set option: user selected probe set
Invert HR values below 1: not checked

Restrictions

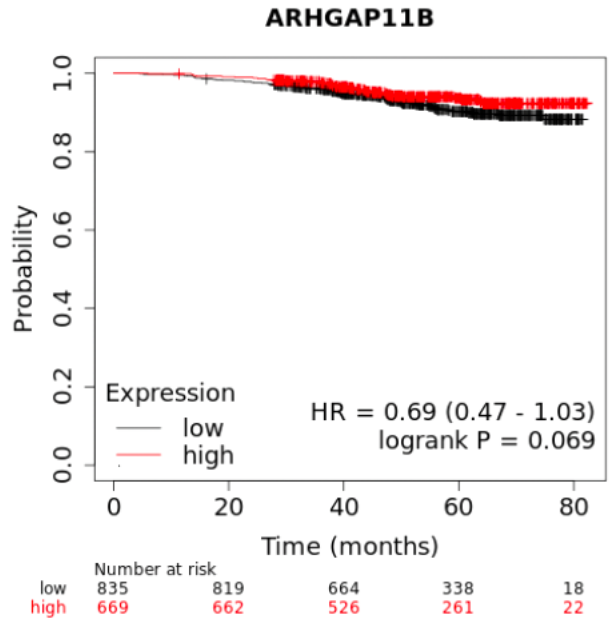
Lymph node status: all
 ER status: all
 PGR status: all
 HER2 status: all
 KI67 status: all
 Nottingham histologic grade: all
 PAM50 subtype: LuminalA

Cohorts

Endocrine treated: all
 Chemo treated: all

Results

P value: 0.0686
FDR: 100%



RNAseq ID: ARHGAP11B
Survival: OS
Auto select best cutoff: checked
Follow up threshold: all
Censore at threshold: checked
Compute median over entire database: false
Cutoff value used in analysis: 2.77
Expression range of the probe: -1 - 5
Probe set option: user selected probe set
Invert HR values below 1: not checked

Restrictions

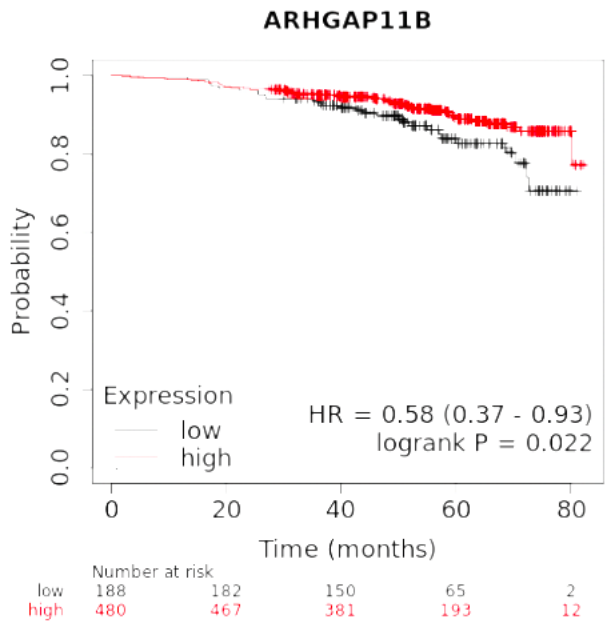
Lymph node status: all
 ER status: all
 PGR status: all
 HER2 status: all
 KI67 status: all
 Nottingham histologic grade: all
 PAM50 subtype: LuminalB

Cohorts

Endocrine treated: all
 Chemo treated: all

Results

P value: 0.0215
FDR: over 50%



RNAseq ID: ARHGAP11B
Survival: OS
Auto select best cutoff: checked
Follow up threshold: all
Censore at threshold: checked
Compute median over entire database: false
Cutoff value used in analysis: 3.21
Expression range of the probe: 2 - 5
Probe set option: user selected probe set
Invert HR values below 1: not checked

Restrictions

Lymph node status: all
 ER status: all
 PGR status: all
 HER2 status: all
 KI67 status: all
 Nottingham histologic grade: all
 PAM50 subtype: Her2

Cohorts

Endocrine treated: all
 Chemo treated: all

Results

P value: 0.0354
FDR: over 50%

