

https://hgserver1.amc.nl/cgi-bin/r2/main.cgi

< > ↻ 🏠 ↶ ☆ 📄 🔒 https://hgserver1.amc.nl/cgi-bin/r2/main.cgi 🔍

**Go to: [Main](#)** **R2: Genomics Analysis and Visualization Platform**

673699 (623521 unique) samples available

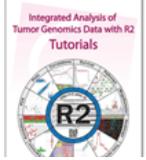
**1** Choose single or multiple dataset analysis  
Single Dataset ▾ ⓘ

**2** Select a dataset for analysis  
Tum or Neuroblastom a public  
Versteeg - 88 - MAS5.0 - u133p2 ⓘ [Change Dataset](#)

**3** Select type of analysis  
View a Gene ▾ ⓘ

**4** Proceed  
[Next](#) [Reset](#)

- Main
- Time series
- Kaplan-Meier**
- Sample maps
- Small Tools
- DataGrabber
- Genome Browser
- ChIP data
- TAR literature
- Change Data Scope ▶
- [Login/Register](#)
- Help ▶
- Contact / About R2



Go to: [Main](#)

R2: Kaplan-Meier

- Main
- Time series
- Kaplan-Meier
- Sample maps
- Small Tools
- DataGrabber
- Genome Browser
- ChIP data
- TAR literature
- Change Data Scope ▶
- Login/Register
- Help ▶
- Contact / About R2

**Kaplan-Meier analysis using a data set**

Current dataset:  
Tumor Neuroblastoma public - Versteeg - 88 - MAS5.0 - u133p2 ⓘ

Separate by:  ▼

Type here or use the dropdown menu to select a dataset

Select	Species	Data type	Category	Tissue/Tumor	Platform	Normalizat...	N	Author	Accessi...	Release d...	R2 date	Access
	Sele	Select Filte	Selec	neurob		Select Filt						Select
Select	hs	miRNA data	Tumor	Neuroblastoma	mirbase22	tmm	97	Bell		2000-01-01	2019-12-15	public
Select	hs	Expression da...	Tumor	Neuroblastoma	ag44kcwolf	custom	649	Kocak	gse45547	2013-03-28	2013-05-08	public
Select	hs	Expression da...	Tumor	Neuroblastoma	u95a	custom	101	Maris	GSE3960	2006-01-04	2000-01-01	public
Select	hs	Expression da...	Tumor	Neuroblastoma	amexp255	custom	251	Oberthuer	E-TABM-38	2000-01-01	2010-01-29	public
Select	hs	Expression da...	Tumor	Neuroblastoma	ag44kcwolf	custom	498	SEQC	gse49710	2014-10-21	2014-11-03	public
Select	hs	Expression da...	Tumor	Neuroblastoma	seqcnb1	RPM	498	SEQC	gse62564	2014-10-22	2014-11-03	public
Select	hs	Expression da...	Tumor	Neuroblastoma (Stage IV)	ag44kcwolf	custom	27	Capasso	gse79910	2016-04-05	2017-05-10	public
Select	hs	Expression da...	Tumor	Neuroblastoma ALT	gencode19	tpm	144	Westermann		2020-10-28	2020-10-30	public
Select	hs	aCGH data	Tumor	Neuroblastoma HR	gent005	custom	556	de Preter		2000-01-01	2020-02-11	public
Select	hs	Expression da...	Tumor	Neuroblastoma Primary	huex10t	rma_sketch ...	283	NRC	gse85047	2018-07-02	2018-09-05	public

Clear filters

Type here or use the dropdown menu to select a dataset

Sel...Y	Spec...Y	Data type Y	Categ...Y	Tissue/Tumor Y	Platform Y	Normalizat...Y	N Y	Author Y	Accessi...Y	Release d...Y	R2 date Y	Access Y
	Sele ▾	Select Filte ▾	Selec ▾	neurobl		Select Filt ▾						Select ▾
Select	hs	miRNA data	Tumor	Neuroblastoma	mirbase22	tmm	97	Bell		2000-01-01	2019-12-15	public
Select	hs	Expression da...	Tumor	Neuroblastoma	ag44kewolf	custom	649	Kocak	gse45547	2013-03-28	2013-05-08	public
Select	hs	Expression da...	Tumor	Neuroblastoma	u95a	custom	101	Maris	GSE3960	2006-01-04	2000-01-01	public
Select	hs	Expression da...	Tumor	Neuroblastoma	amexp255	custom	251	Oberthuer	E-TABM-38	2000-01-01	2010-01-29	public
Select	hs	Expression da...	Tumor	Neuroblastoma	ag44kewolf	custom	498	SEQC	gse49710	2014-10-21	2014-11-03	public
Select	hs	Expression da...	Tumor	Neuroblastoma	seqcnb1	RPM	498	SEQC	gse62564	2014-10-22	2014-11-03	public
Select	hs	Expression da...	Tumor	Neuroblastoma (Stage IV)	ag44kewolf	custom	27	Capasso	gse79910	2016-04-05	2017-05-10	public
Select	hs	Expression da...	Tumor	Neuroblastoma ALT	gencode19	tpm	144	Westermann		2020-10-28	2020-10-30	public
Select	hs	aCGH data	Tumor	Neuroblastoma HR	gent005	custom	556	de Preter		2000-01-01	2020-02-11	public
Select	hs	Expression da...	Tumor	Neuroblastoma Primary	huex10t	rma_sketch ...	283	NRC	gse85047	2018-07-02	2018-09-05	public

Clear filters

## R2: Kaplan-Meier

**Kaplan-Meier analysis using a data set**

Current dataset:  
Tumor Neuroblastoma - Kocak - 649 - custom - ag44kcwlf 

Separate by:

- a categorical track ▼
- a categorical track
- a numerical track
- a single gene**
- multiple genes

Kaplan-Meier  on data

### R2: Kaplan-Meier Scanner

**Kaplan Meier Scanner:**

Gene / Reporter:

Cut-off mode:

**Sample Filter**

Subset track:

Selected sample subset: None

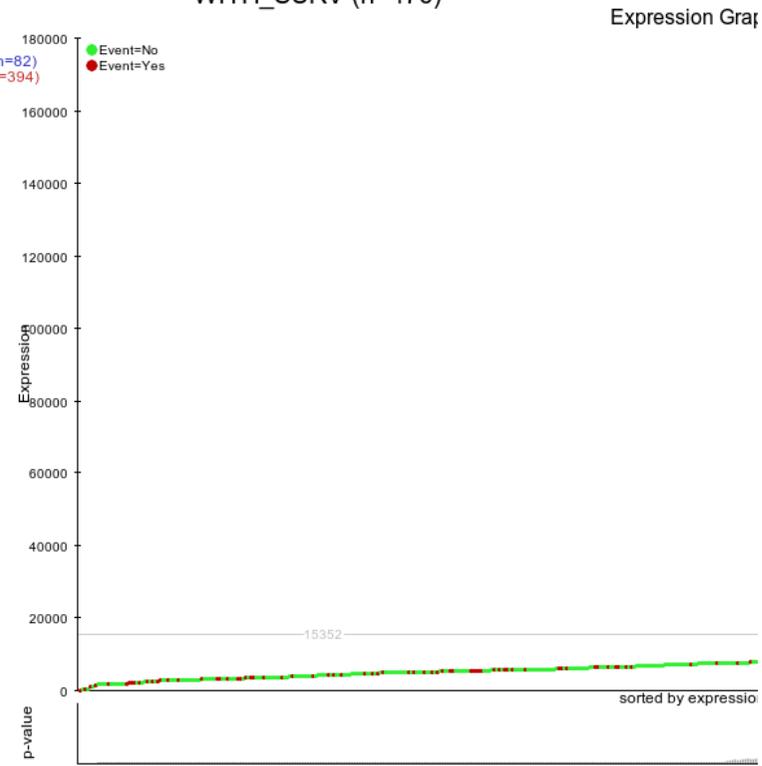
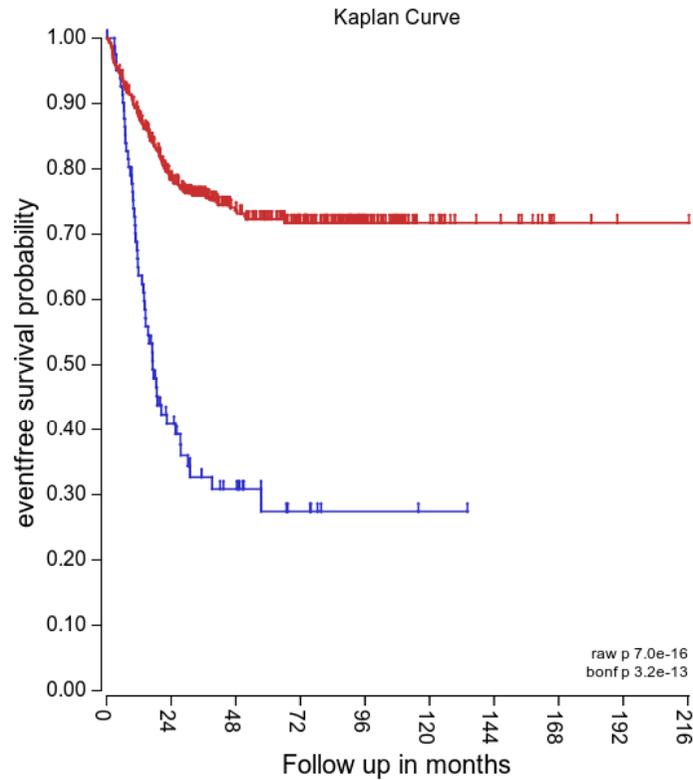
**Other settings**

Type of Survival:

Minimal group size:

先输入，之后在下拉菜单中选择

**Tumor Neuroblastoma**  
Kocak - 649 - custom - ag44kcwof  
MYCN (UKv4\_A\_24\_P94402)  
Expression cutoff: 15352.3 (min.grp=8)  
WITH\_SURV (n=476)



chi=65.13 df=1 p=7.0e-16

**high** is worse

ST:

[View MYCN in 2geneview](#)

更改图片样式

**Adjustable settings:**

Gene / Reporter:	MYCN	UKv4_A_24_P94402	advanced search
cutoff_modus:	scan		
Cutoff:	15352.3	7.0e-16	5.28e-13

Subset track: ⚙️ ❌ i

Number of samples in s  
n=476

Selected sample subset: **WITH\_SURV**  
no subset selected

---

**Other settings**

Only draw up to: 120 months

Minimal groupsize: 8

LineWidth: 3

Expression Graph: no

Color for Group low: 3232C8

Color for Group high: C83232

fontsize\_ruler: 30

fontsize\_y: 30

fontsize\_t1: 30

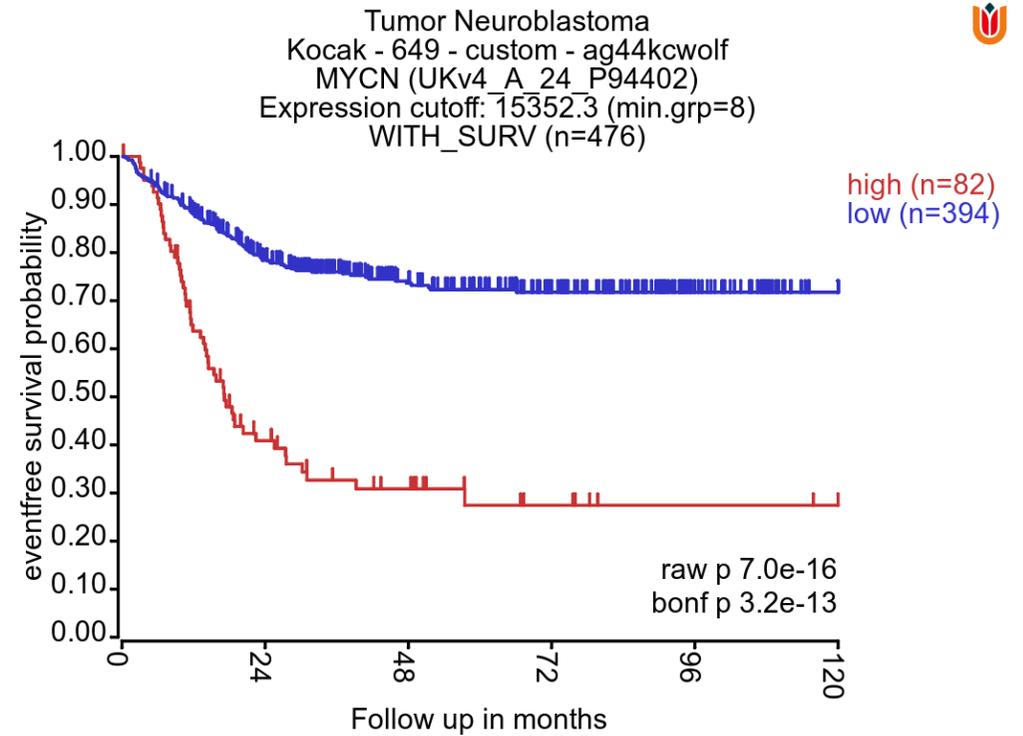
fontsize\_tsub: 30

fontsize\_inline: 30

axis\_width: 3

Vector (SVG) output: false i

Redraw Graph   Reset



View [MYCN](#) in 2geneview

**Adjustable settings:**

Gene / Reporter: MYCN UKv4\_A\_24\_P94402 advanced search

cutoff\_modus: scan

Cutoff: 394 - : raw p: 7.0e-16 (bonf: 3.2e-13)

**Sample Filter**

Subset track: mycn (2 cat + nd)

Selected sample subset: mycn (2 cat + nd)

Only draw up to:

Minimal groupsize: 8

LineWidth: 3

Expression Graph: no

Color for Group low: 3232C8

Color for Group high: C83232

fontsize\_ruler: 30

fontsize\_y: 30

fontsize\_t1: 30

也可以选择 组内绘图。

chi=65.13 df=1 p=7.0e-16  
**high** is worse  
ST:  
View [MYCN](#) in 2geneview

**Adjustable settings:**

Gene / Reporter: MYCN UKv4\_A\_24\_P94402 advanced search

cutoff\_modus: scan

Cutoff: 394 - : raw p: 7.0e-16 (bonf: 3.2e-13)

**Sample Filter**

Subset track: mycn (2 cat + nd)

Number of samples in n=476

Selected sample subset: WITH\_SURV no subset selected

Other set

Only draw up to: 120 months

Minimal groupsize: 8

LineWidth: 3

Expression Graph: no

Color for Group low: 3232C8

Color for Group high: C83232

fontsize\_ruler: 30

fontsize\_y: 30

fontsize\_t1: 30

fontsize\_tsub: 30

fontsize\_inline: 30

axis\_width: 3

mycn - WITH\_SURV (n=476)

amp (66)

na (5)

non\_amp (405)

check-all

OK Cancel