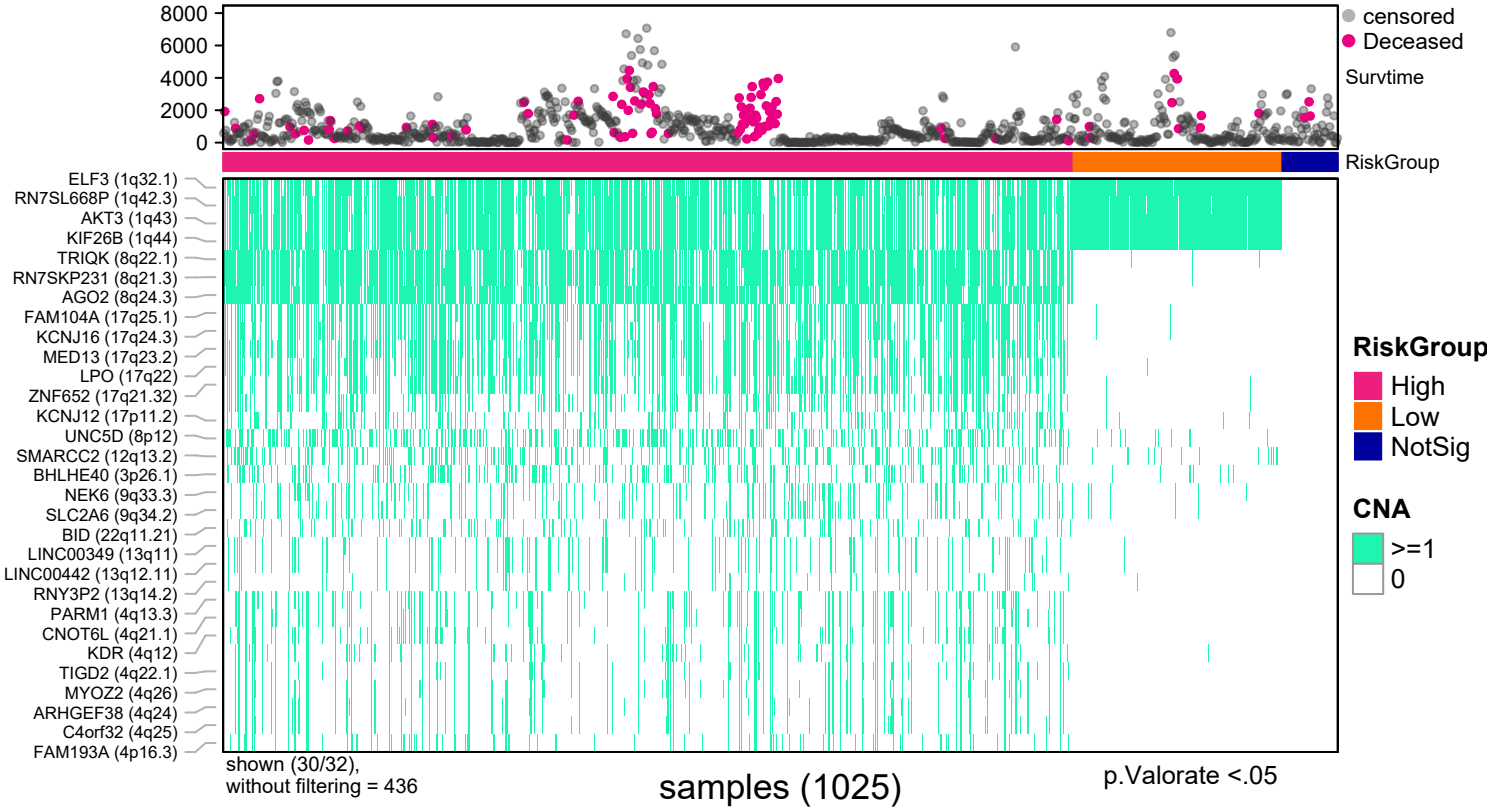
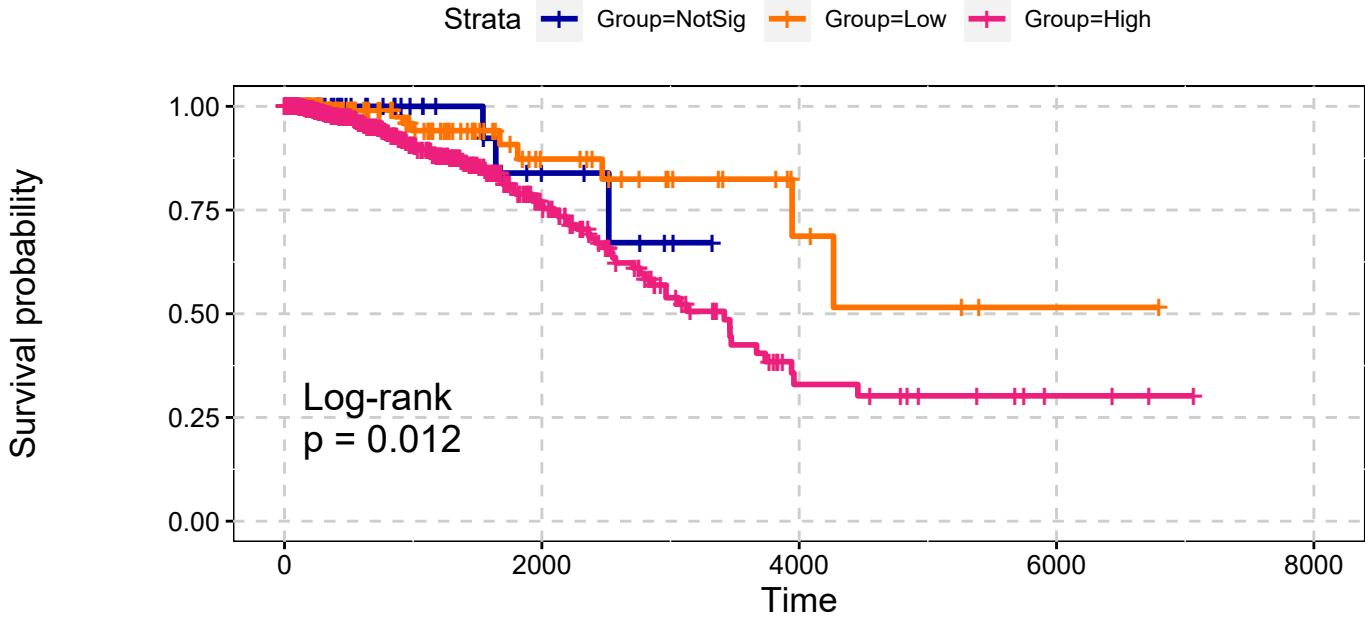


BRCA
All Amplifications
Single Data Signature



BRCA
All Amplifications
Single Data Signature

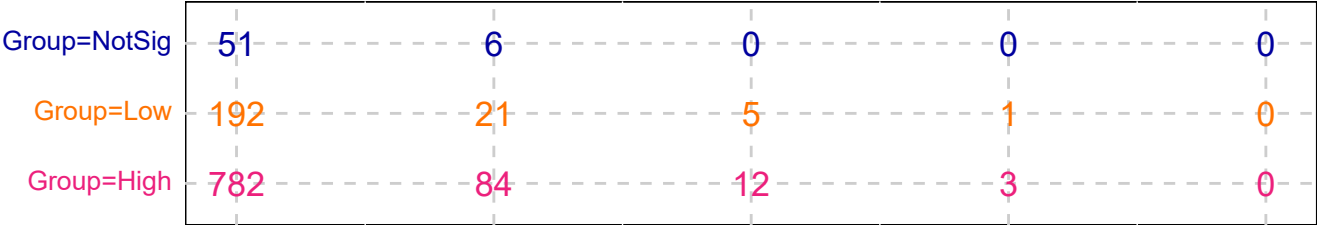


p.Valorate <.05

explanatory	beta	HR	L95	U95	p
Low	-0.32	0.72	0.19	2.68	0.63
High	0.63	1.87	0.59	5.92	0.29

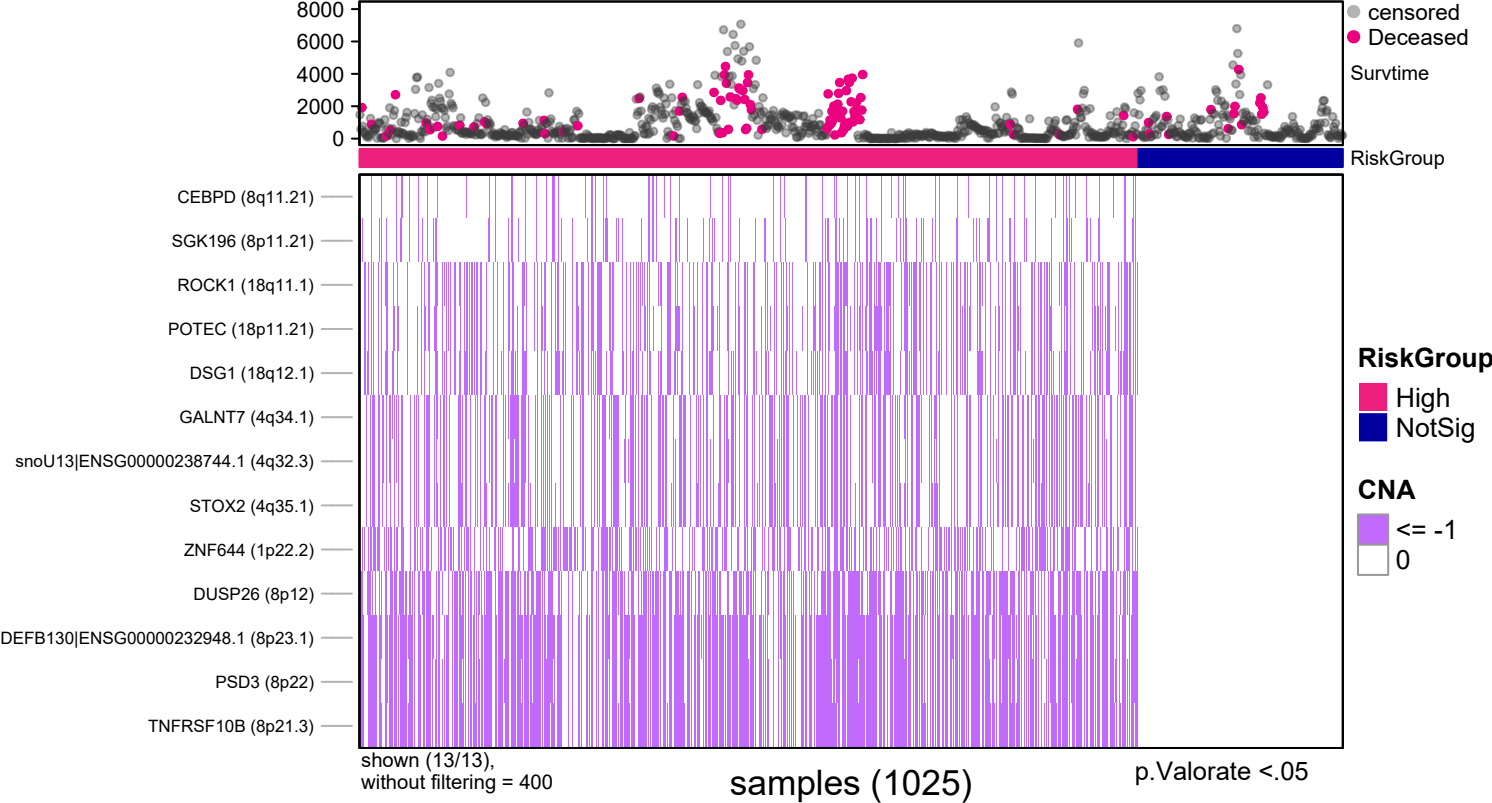
n= 1025, number of events =101
Score(logrank) test = 0.012

Number at risk

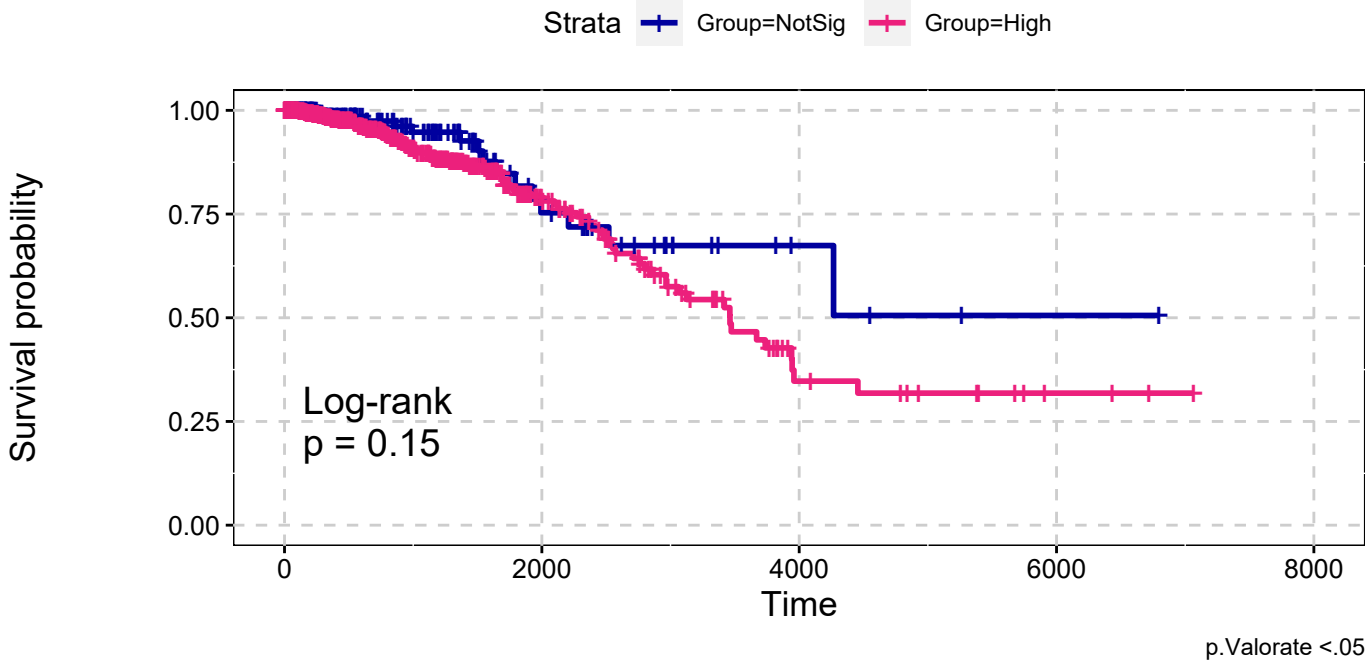


p.Valorate <.05

BRCA
All Deletions
Single Data Signature



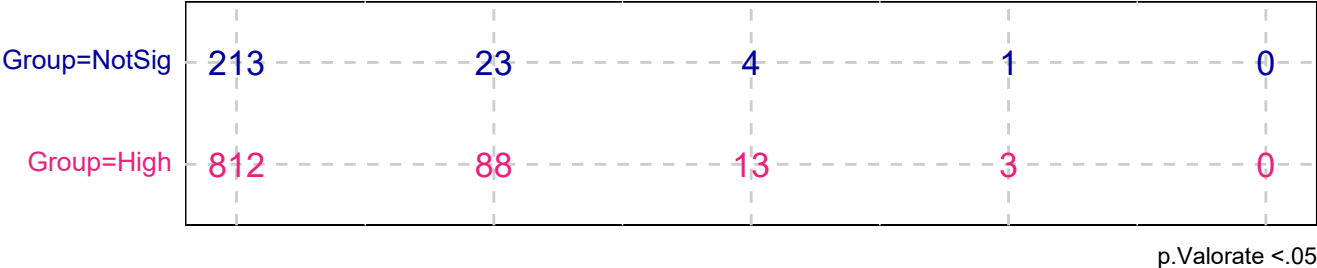
BRCA
All Deletions
Single Data Signature



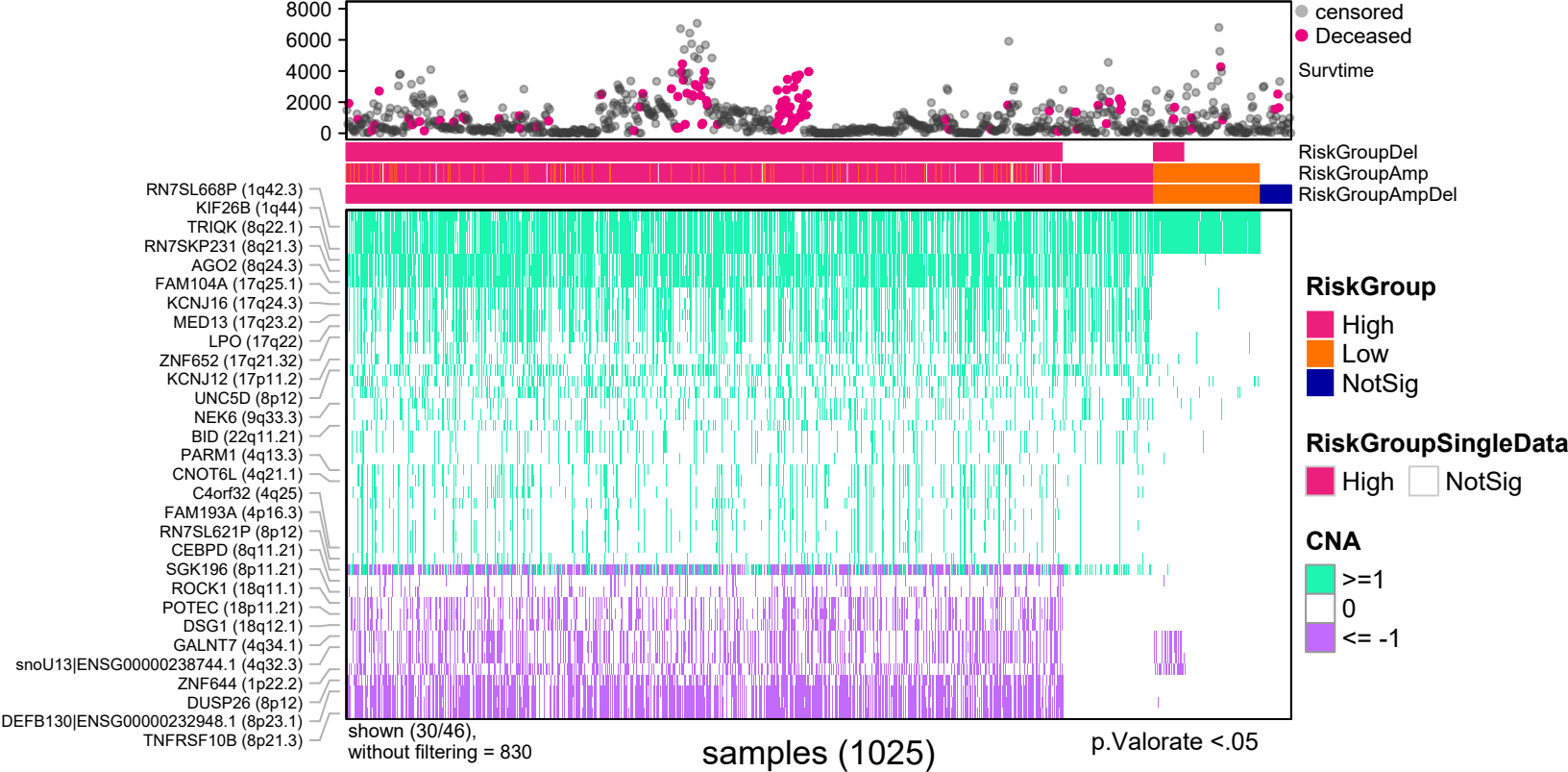
explanatory	beta	HR	L95	U95	p
High	0.40	1.49	0.86	2.58	0.15

n= 1025, number of events =101
Score(logrank) test = 0.152

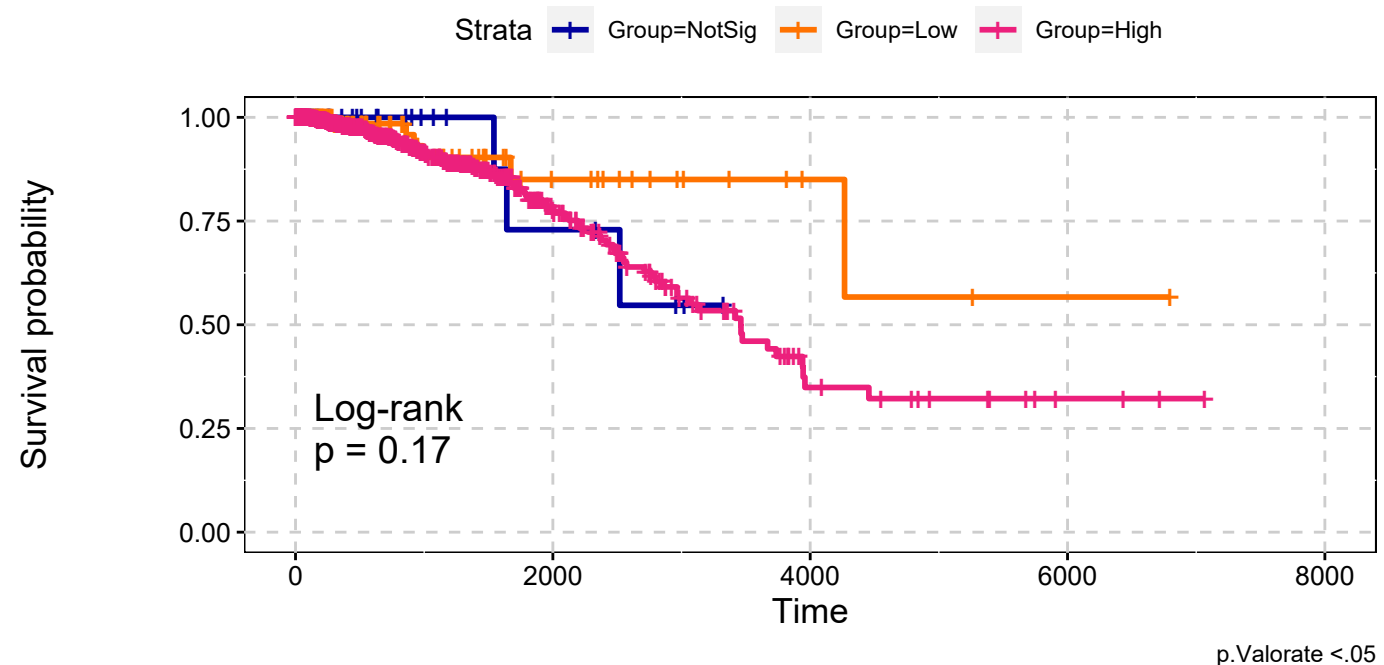
Number at risk



BRCA
All Amplifications & All Deletions
Max Sum Significance Signatures

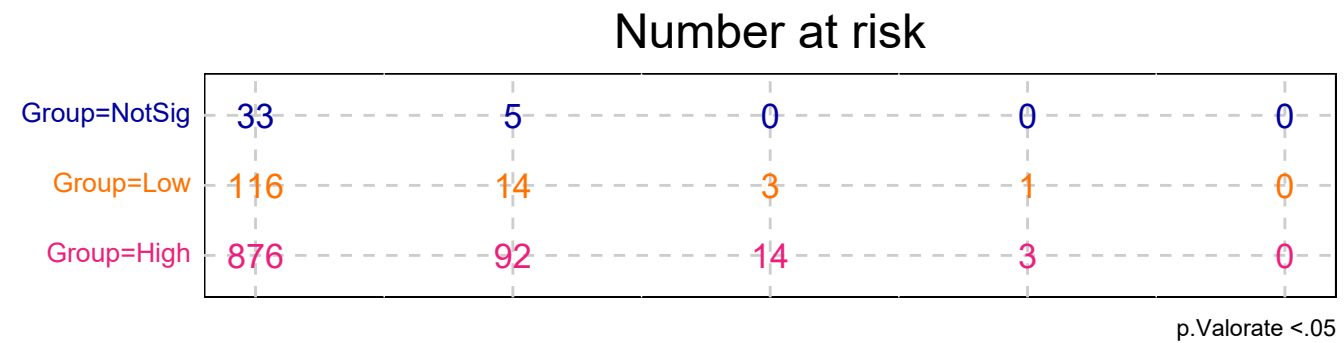


BRCA
All Amplifications & All Deletions
Max Sum Significance Signatures

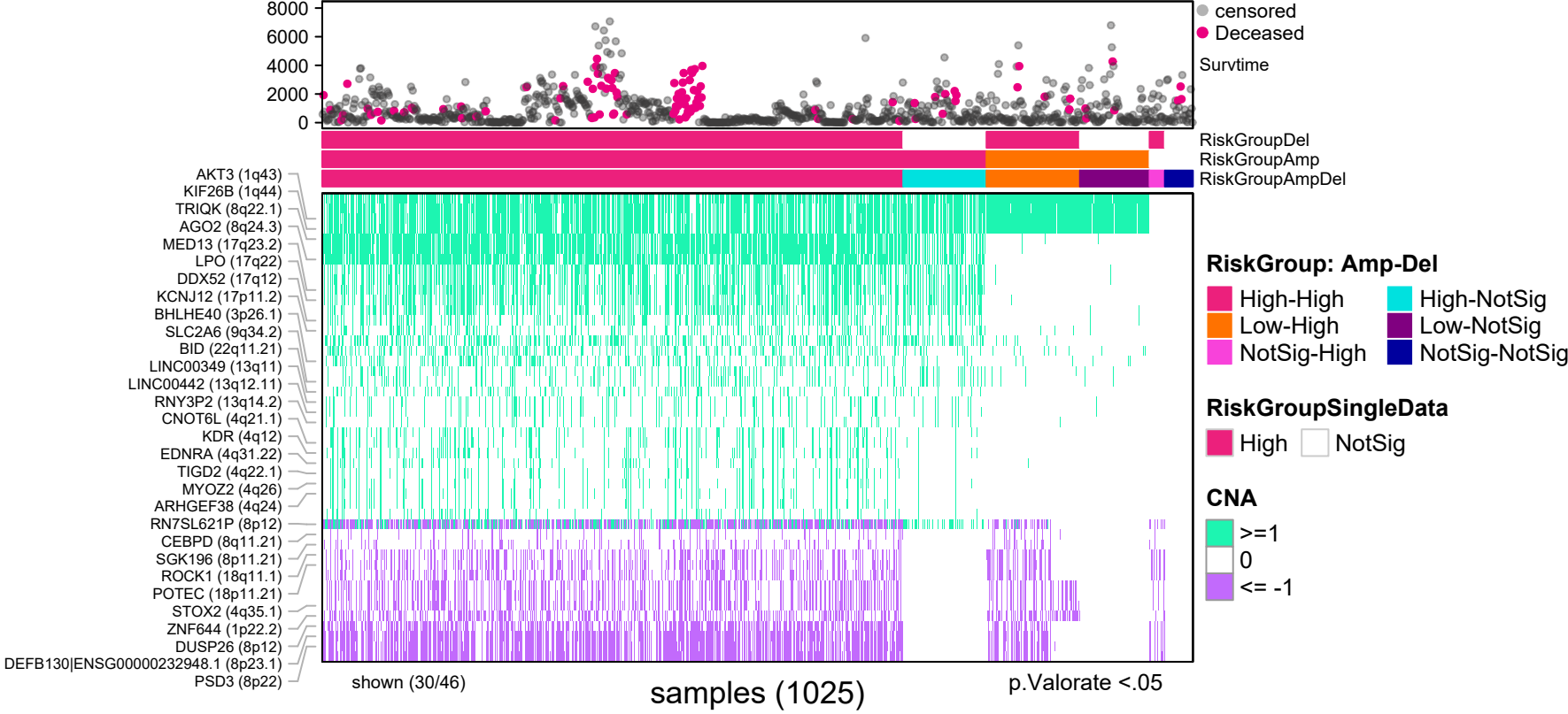


explanatory	beta	HR	L95	U95	p
Low	-0.60	0.55	0.14	2.20	0.40
High	0.16	1.18	0.37	3.73	0.78

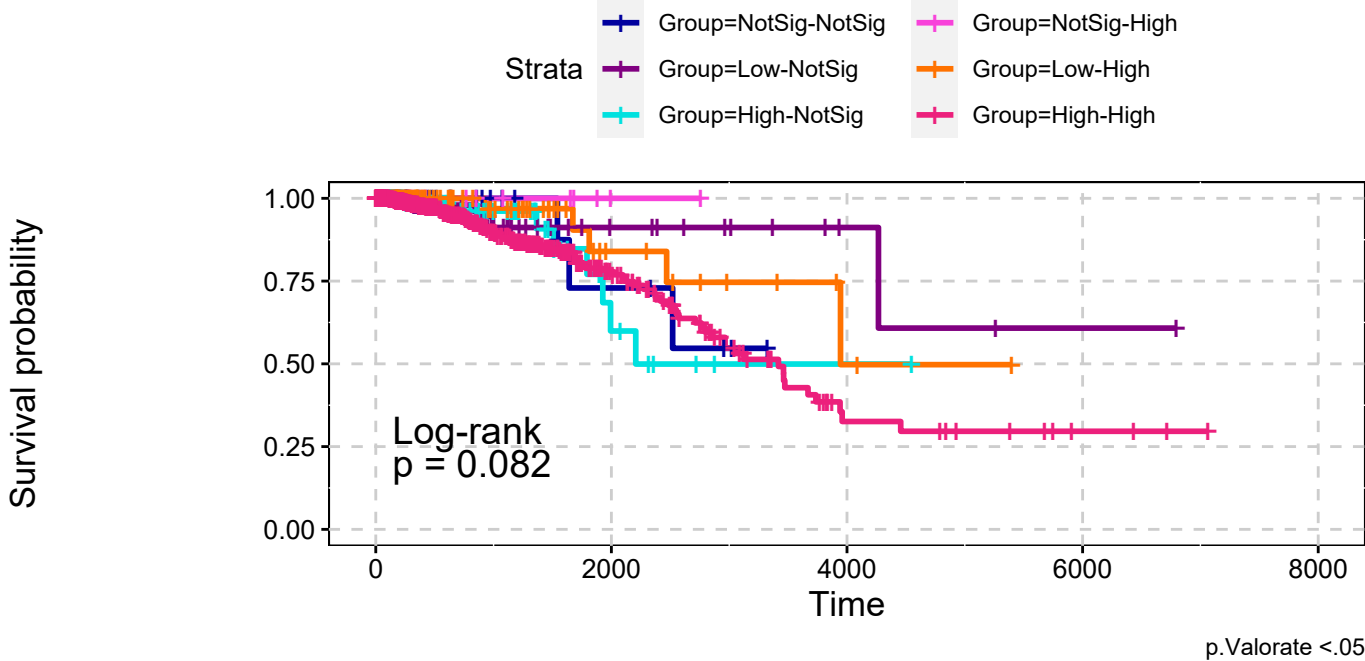
n= 1025, number of events =101
Score(logrank) test = 0.173



BRCA
All Amplifications & All Deletions
combining signatures



BRCA
All Amplifications & All Deletions
combining signatures



explanatory	beta	HR	L95	U95	p
Low-NotSig	-0.83	0.43	0.10	1.95	0.28
High-NotSig	0.16	1.18	0.31	4.45	0.81
NotSig-High	-15.95	0.00	0.00	Inf	0.99
Low-High	-0.59	0.56	0.13	2.34	0.42
High-High	0.25	1.29	0.41	4.09	0.67

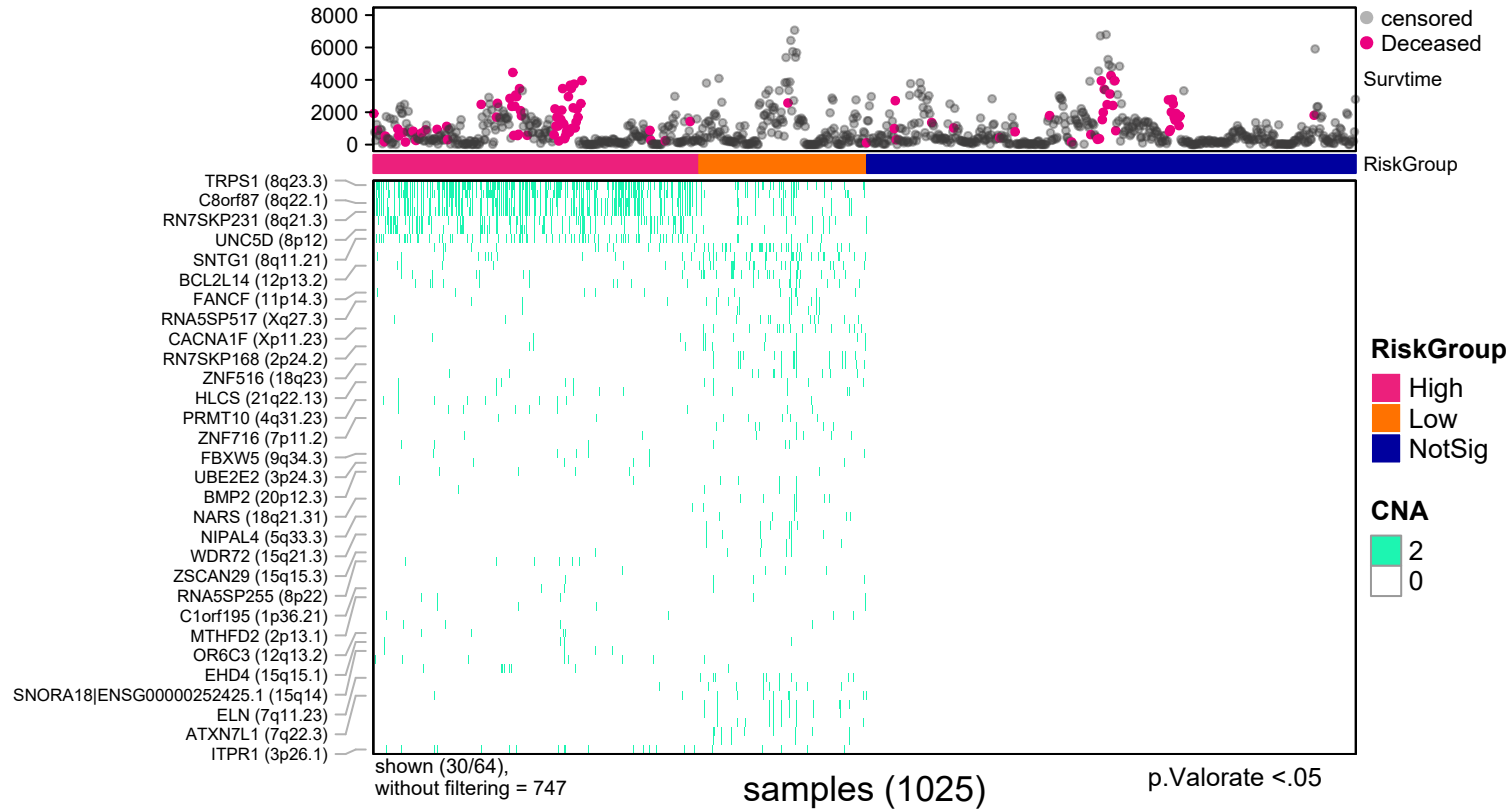
n= 1025, number of events =101
Score(logrank) test = 0.082

Number at risk

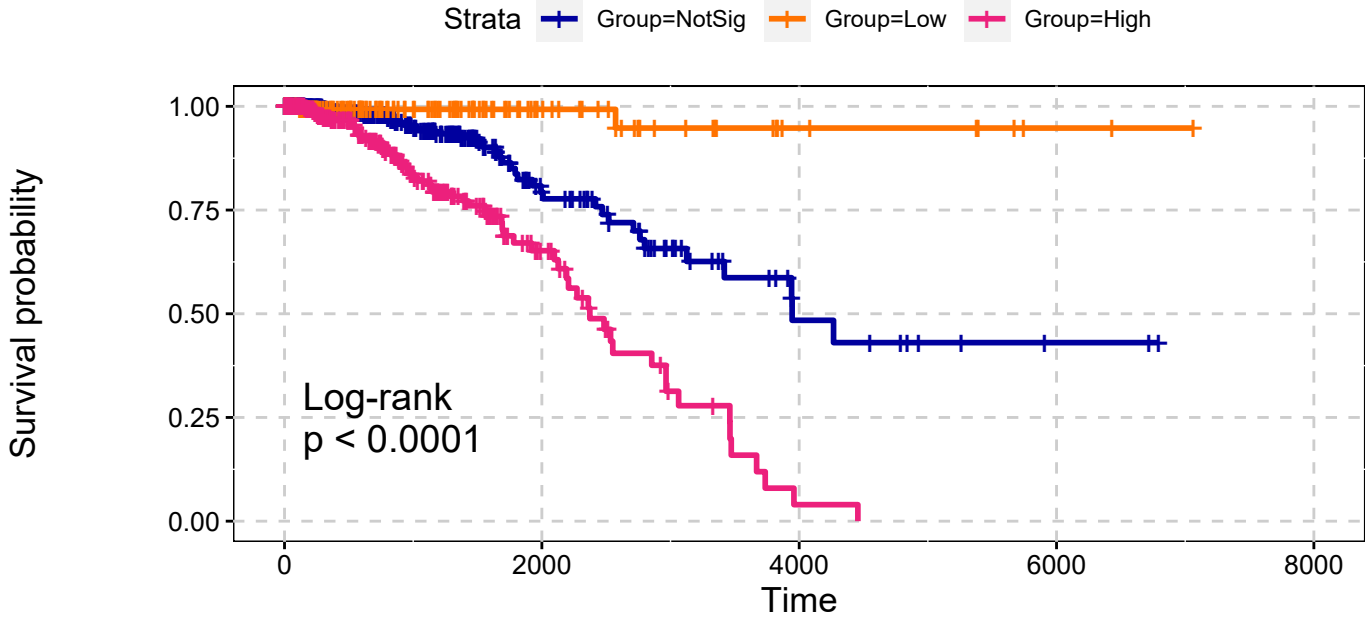
Group=NotSig-NotSig	33	5	0	0	0
Group=Low-NotSig	82	11	3	1	0
Group=High-NotSig	98	7	1	0	0
Group=NotSig-High	18	1	0	0	0
Group=Low-High	110	10	2	0	0
Group=High-High	684	77	11	3	0

RiskGroup: Amp-Del, p.Valorate <.05

BRCA
Deep Amplifications
Single Data Signature



BRCA
Deep Amplifications
Single Data Signature

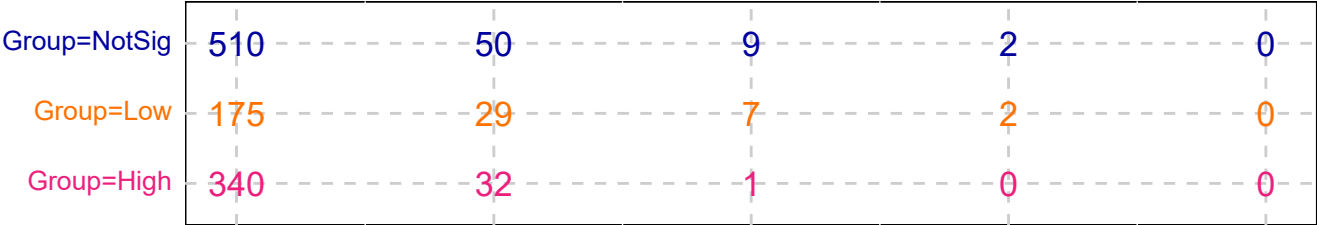


p.Valorate <.05

explanatory	beta	HR	L95	U95	p
Low	-2.25	0.11	0.03	0.44	0.00
High	1.12	3.08	2.03	4.67	0.00

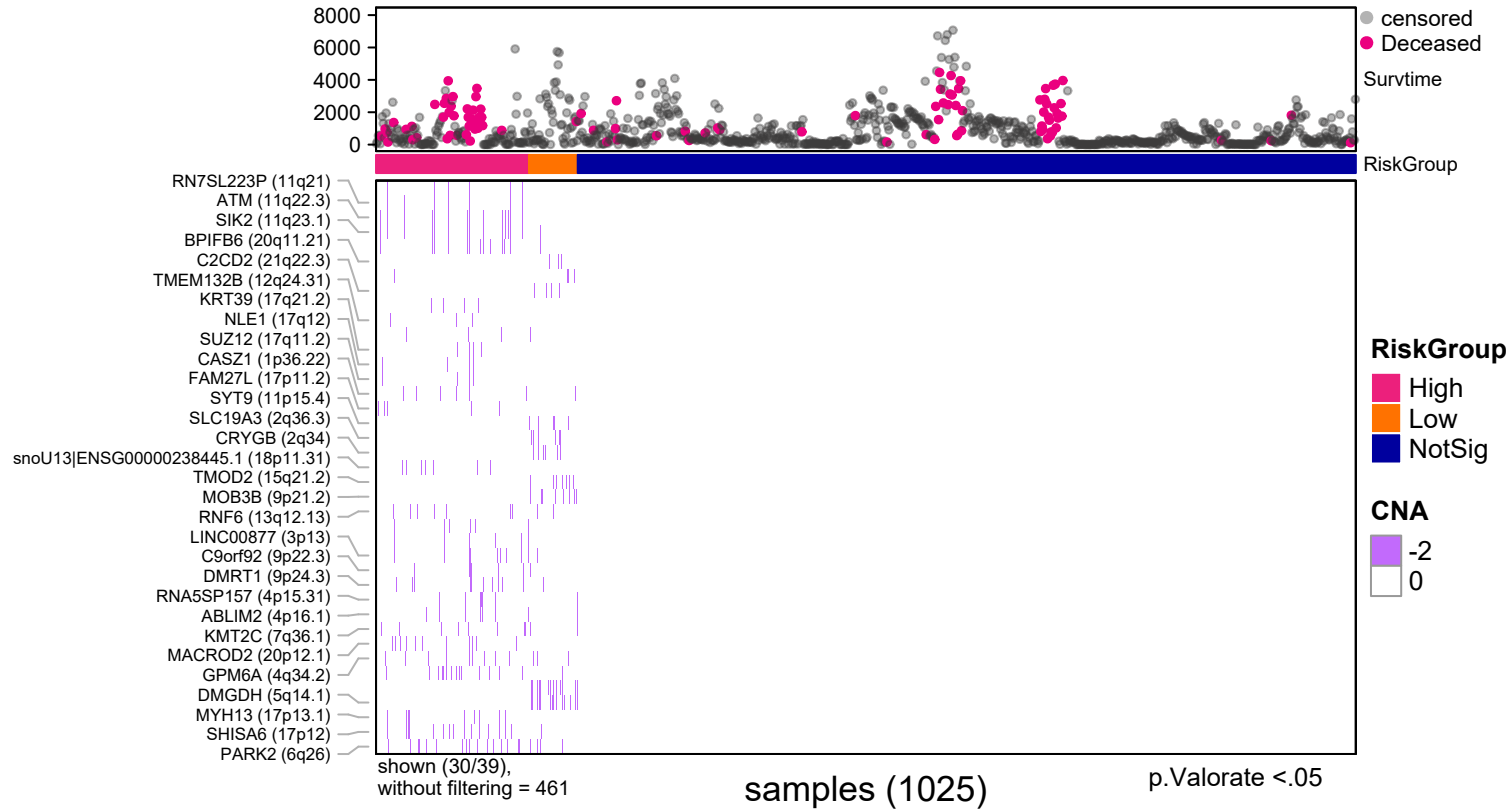
n= 1025, number of events =101
Score(logrank) test = p <.0001

Number at risk

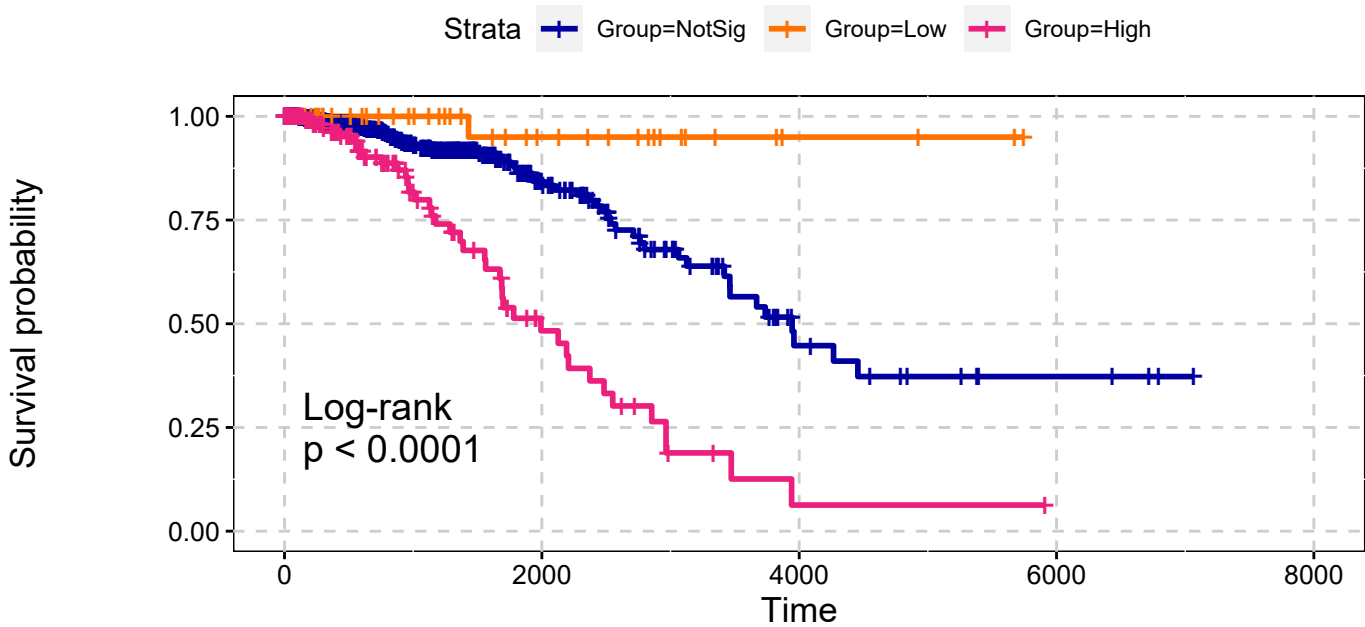


p.Valorate <.05

BRCA
Deep Deletions
Single Data Signature



BRCA
Deep Deletions
Single Data Signature

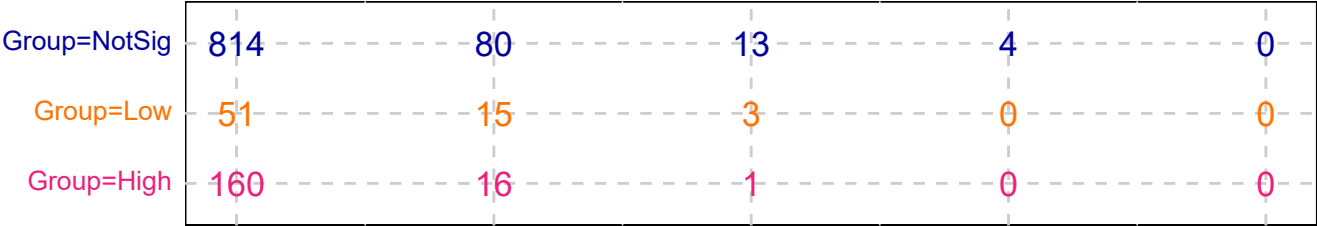


explanatory	beta	HR	L95	U95	p
Low	-2.19	0.11	0.02	0.81	0.03
High	1.27	3.56	2.38	5.32	0.00

n= 1025, number of events =101
Score(logrank) test = p <.0001

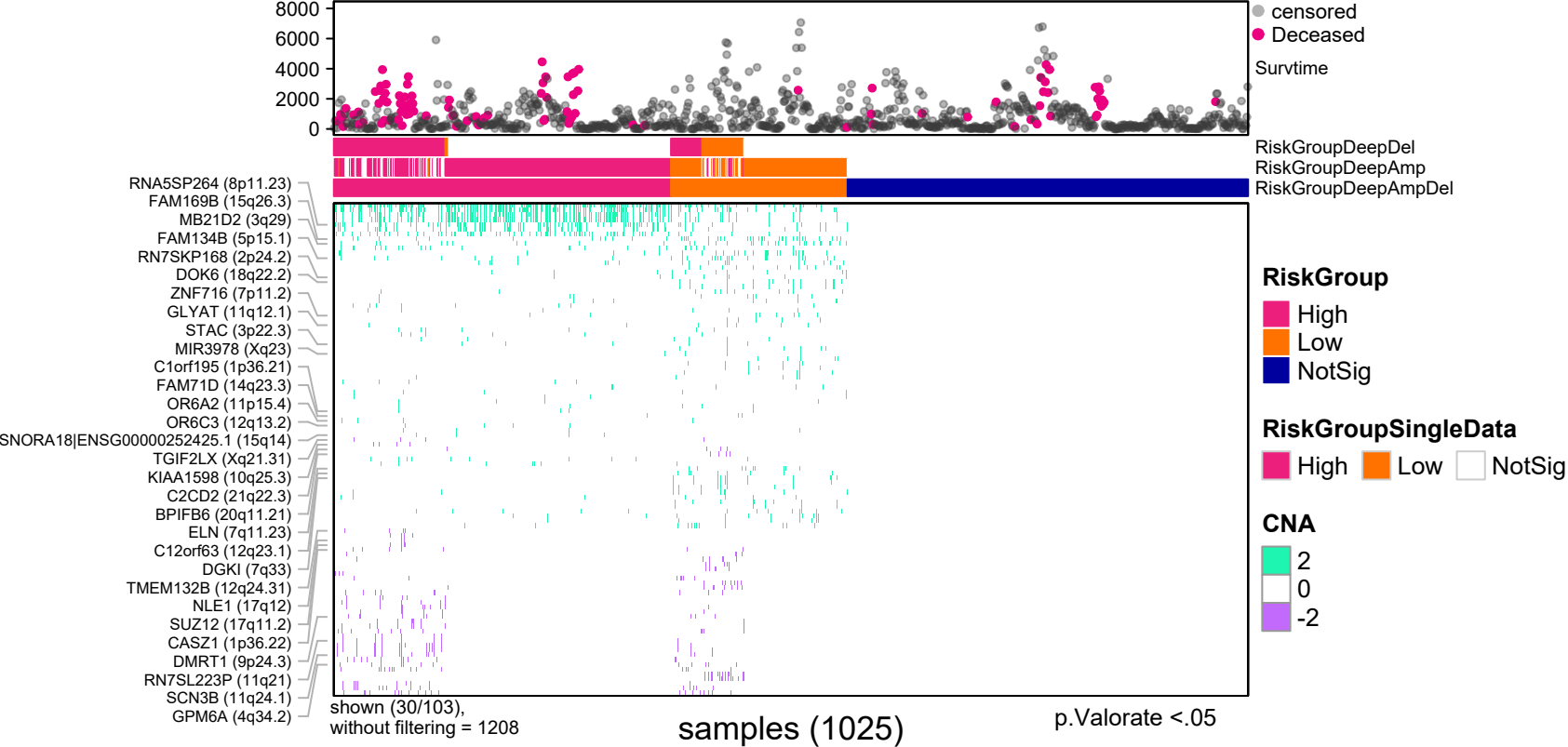
p.Valorate <.05

Number at risk



p.Valorate <.05

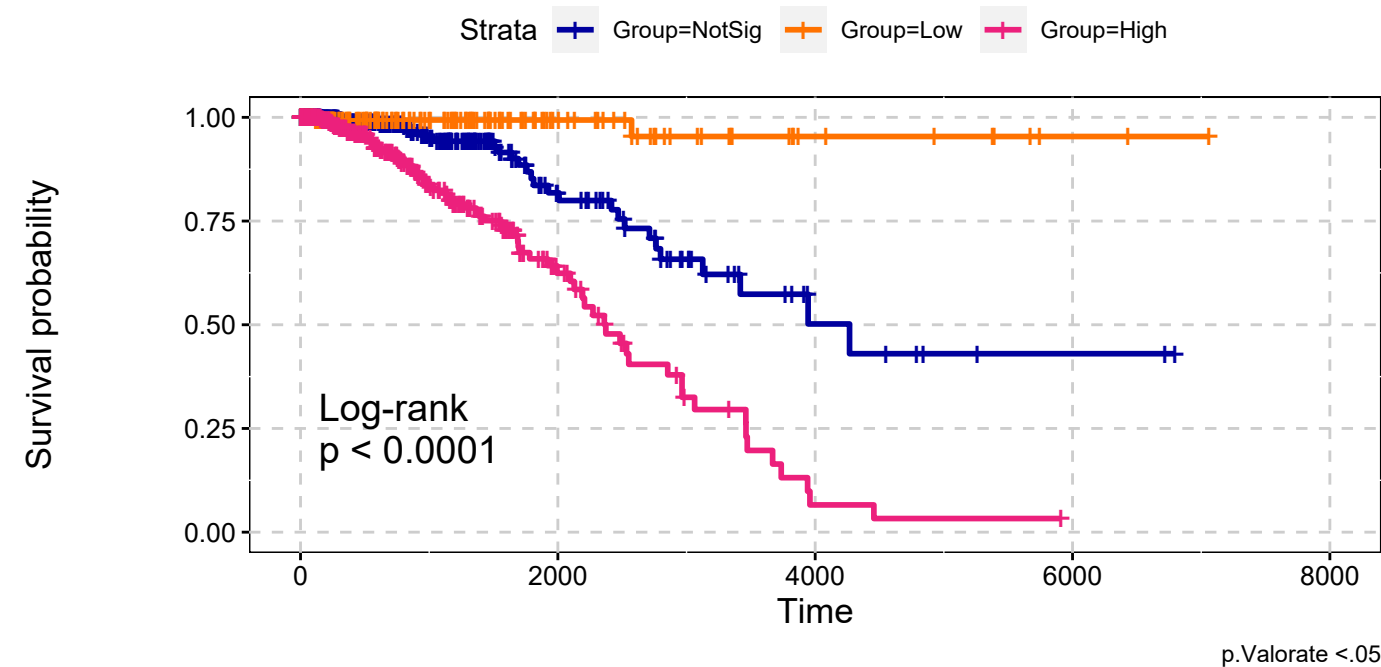
BRCA
Deep Amplifications & Deep Deletions
Max Sum Significance Signatures



BRCA

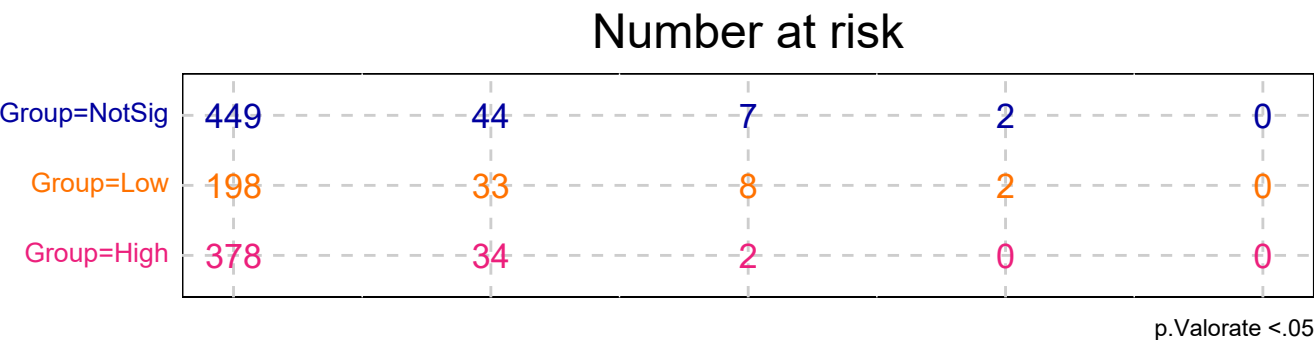
Deep Amplifications & Deep Deletions

Max Sum Significance Signatures

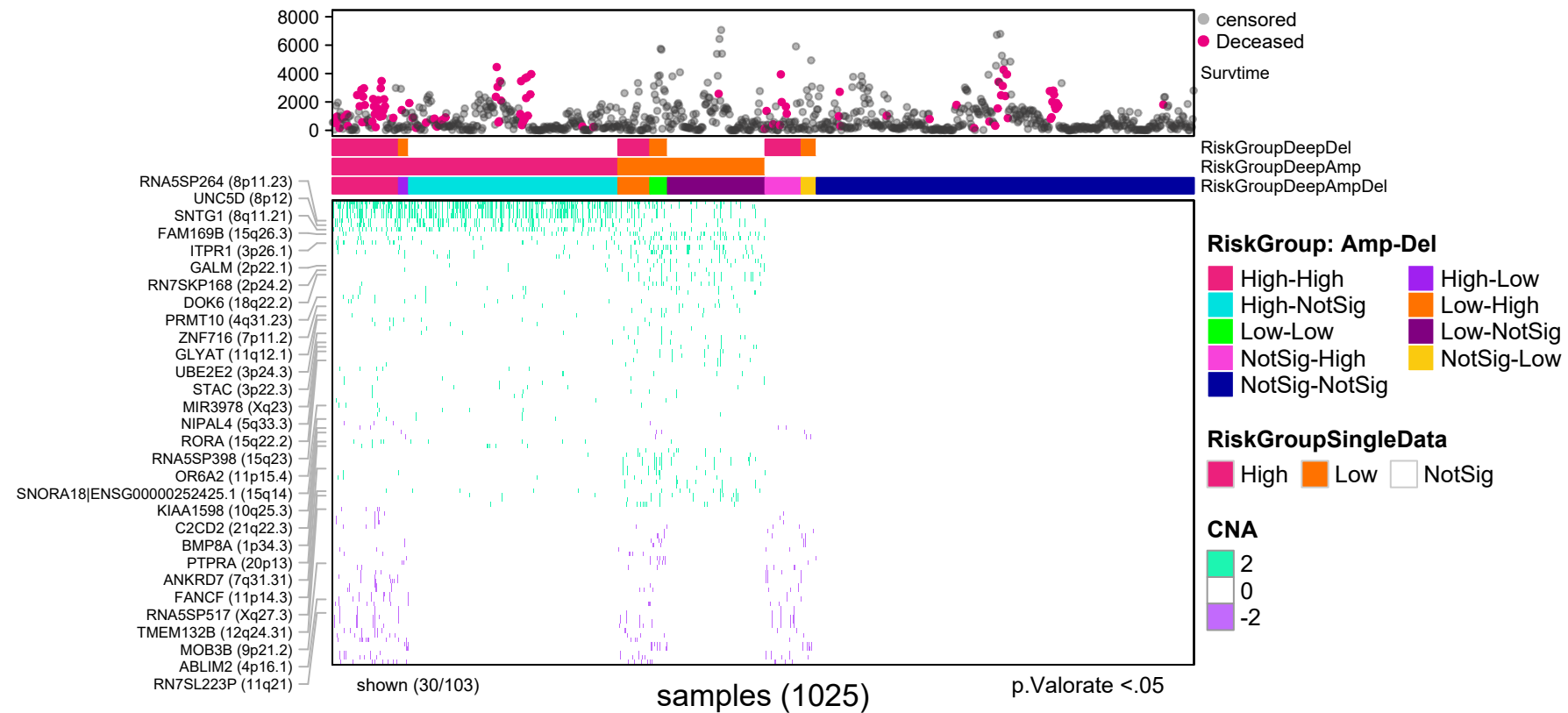


explanatory	beta	HR	L95	U95	p
Low	-2.31	0.10	0.02	0.42	0.00
High	1.17	3.21	2.08	4.97	0.00

n= 1025, number of events =101
Score(logrank) test = p <.0001

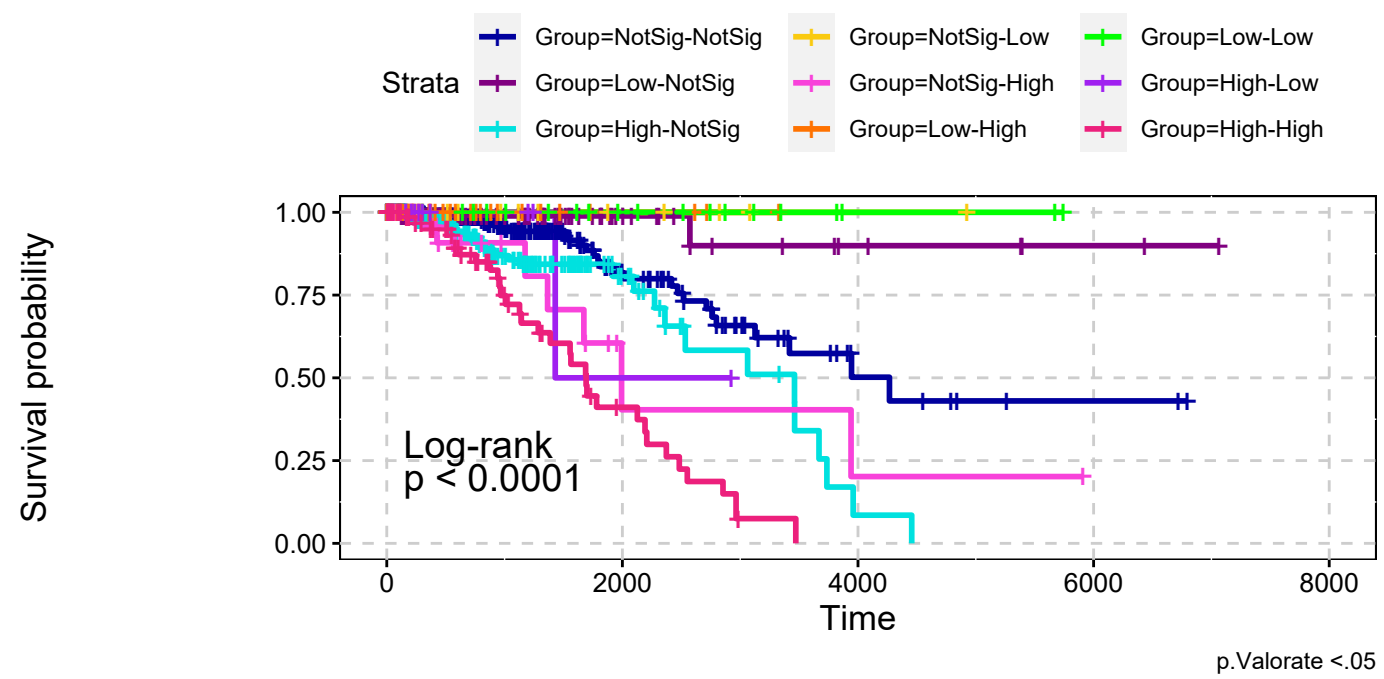


BRCA
Deep Amplifications & Deep Deletions
combining signatures



BRCA

Deep Amplifications & Deep Deletions combining signatures



explanatory	beta	HR	L95	U95	p
Low-NotSig	-1.70	0.18	0.04	0.77	0.02
High-NotSig	0.80	2.22	1.32	3.73	0.00
NotSig-Low	-17.68	0.00	0.00	Inf	1.00
NotSig-High	0.96	2.62	1.14	6.01	0.02
Low-High	-17.85	0.00	0.00	Inf	1.00
Low-Low	-17.69	0.00	0.00	Inf	1.00
High-Low	0.54	1.72	0.23	12.70	0.59
High-High	1.81	6.08	3.65	10.13	0.00

n= 1025, number of events =101
Score(logrank) test = p <.0001

