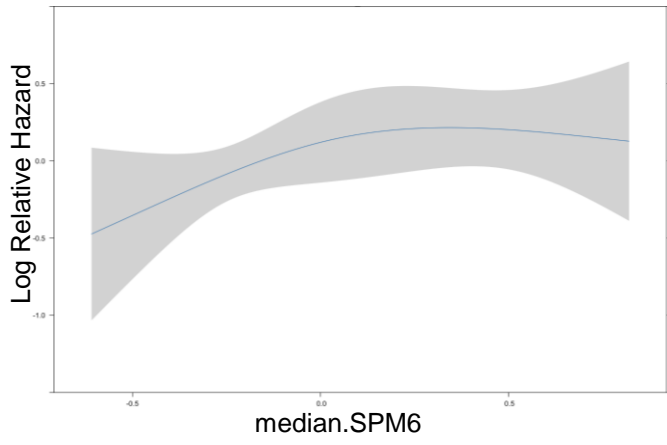
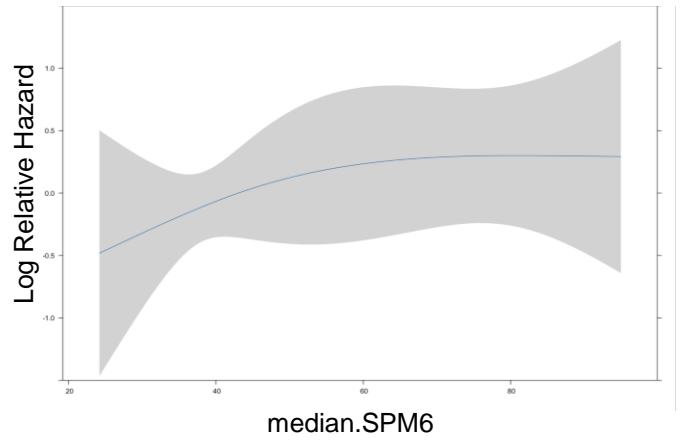


A

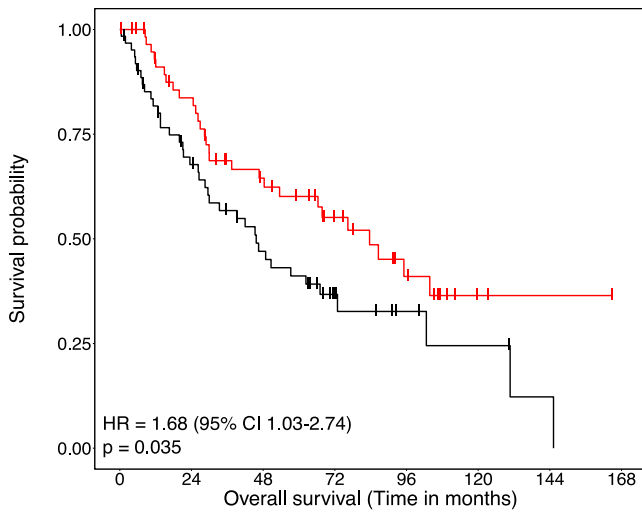
Proteomics cohort

**B**

TCGA cohort

**C**

Proteomics cohort

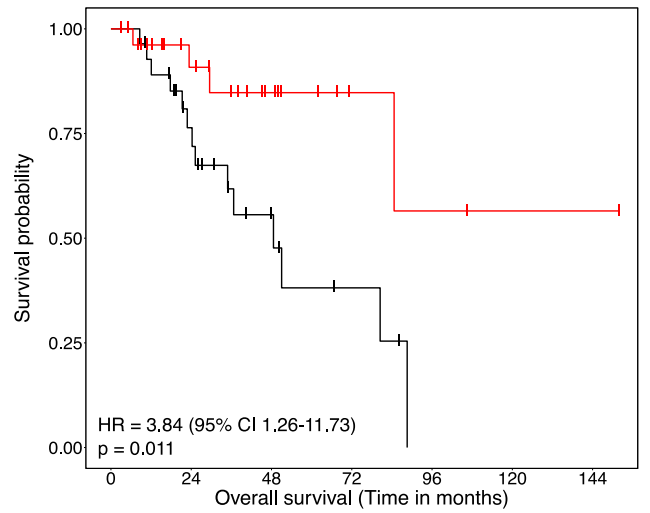


Number at risk

SPM6 high	62	38	24	12	5	3	1	0
SPM6 low	61	45	30	19	10	2	1	0

D

TCGA cohort



Number at risk

SPM6 high	28	17	7	3	0	0	0
SPM6 low	28	17	9	3	2	1	1