

**Supplementary table 2.** Comparison of different prognostic parameters with survival in 30 uveal melanoma patients. An Univariate Cox regression analysis was performed here, with the unknown deaths not censored, but attributed to death to metastasis.

<i>Parameters</i>		<i>Univariate analysis- unknown cases attributed to death due to metastasis</i>			
		95% confidence interval			
<b>Standard clinical/histological parameters</b>	<b><i>n</i></b>	<b>HR</b>	<b>Lower</b>	<b>Upper</b>	<b>p-value</b>
Gender					
Male	17	-	-	-	-
Female	13	1.2	0.5	3.3	0.65
Ciliary Body involvement					
No	17	-	-	-	-
Yes	13	3.8	1.4	10.4	0.008
Stage group					
Stage I & IIA	9	-	-	-	-
Stage IIB	10	2.4	0.6	10.2	0.24
Stage IIIA	9	8.1	1.9	33.7	0.004
Stage IIIB	2	2.5	0.3	25.1	0.43
Age at enucleation (years; low to high; continuous)	30	1.0	1.0	1.1	0.14
Largest basal diameter (mm; low to high; continuous)	30	1.3	1.1	1.6	0.008
<b>Chromosomal aberrations</b>					
FISH on isolated nuclei, cut-off at 5%					
Disomy chromosome 3	11	-	-	-	-
Monosomy chromosome 3	19	15.1	2.0	115.1	0.009
FISH on isolated nuclei, cut-off at 30%					
Disomy chromosome 3	15	-	-	-	-
Monosomy chromosome 3	15	6.4	2.0	20.5	0.002
SNP on tumour DNA					
Disomy chromosome 3	14	-	-	-	-
Monosomy chromosome 3	16	8.3	2.4	29.5	0.001
Disomy chromosome 8q	11	-	-	-	-
Gain chromosome 8q	19	4.8	1.4	17.1	0.015
Disomy chrom 3 + disomy chrom 8q	10	-	-	-	-
Monosomy chrom 3 + gain chrom 8q	15	16.7	2.2	129.6	0.007
<b>Gene expression</b>					
15-gene expression assay class					
Class 1	14	-	-	-	-
Class 2	14	6.7	1.9	23.9	0.003
<i>BAP1</i> gene expression (dichotomized at median)					
High	14	-	-	-	-
Low	14	8.0	2.2	28.8	0.001
<i>BAP1</i> gene expression (high to low; continuous)	28	4.4	1.8	10.8	0.001
<b>Immunohistochemistry</b>					
Positive <i>BAP1</i> immunostaining	14	-	-	-	-
Negative <i>BAP1</i> immunostaining	14	5.0	1.6	15.9	0.007

*HR = hazard ratio.*