- II -			0.00/ 01	_ 1
Predictor	Number affected	Hazard ratio	95% CI	<i>P</i> -value
Presence of metastatic disease	168	4.05	3.05, 5.37	<0.001***
Pathway mutation in				
P1★ (KRAS, NRAS)	53	1.61	1.14, 2.29	0.007**
P21★ (TP53, MTOR, AKT1)	74	1.38	0.99, 1.92	0.058
P110★ (ARID1A, PBRM1, ARID2	2) 136	0.79	0.60, 1.04	0.088
P108★ (ATM, BRCA2, BRCA1	.) 39	0.67	0.42, 1.08	0.10

<sup>&</sup>lt;sup>1</sup> \*p<0.05; \*\*p<0.01; \*\*\*p<0.001

N = 374 patients (266 events). 15 pathways included in the model with p > 0.1 not shown.