Suppl. Fig. S12

A

Endpoint	Subgroup		HR	CI	p
DSS	Age_group:Older Adults		1.34	(0.88–2.03)	0.167
	Tumor_Size:Intermediate (10–		1.62	(1.09–2.39)	0.016
	Tumor_Size:High (> 20 cm)		4.80	(2.31–9.95)	<0.001
	Mitotic_Rate:Intermediate (10		1.98	(1.04–3.79)	0.038
	Mitotic_Rate:High (> 20)		2.37	(1.26–4.46)	0.007
	Necrosis:Present		2.25	(1.18–4.32)	0.014
	Genomic.risk.group:Intermedi		1.40	(0.85–2.31)	0.192
	Genomic.risk.group:High		2.22	(1.35–3.64)	0.002

Univariate Cox regression analysis

B

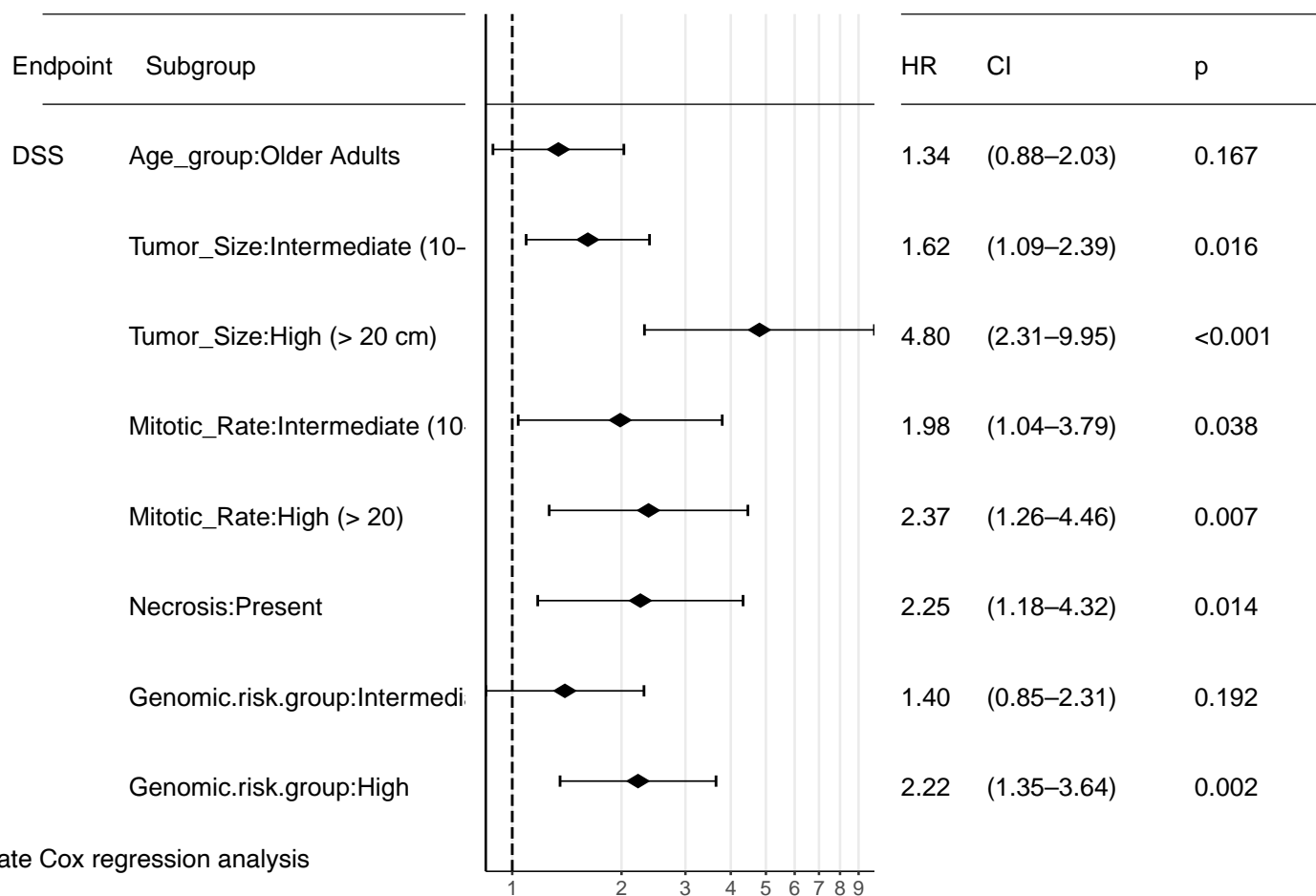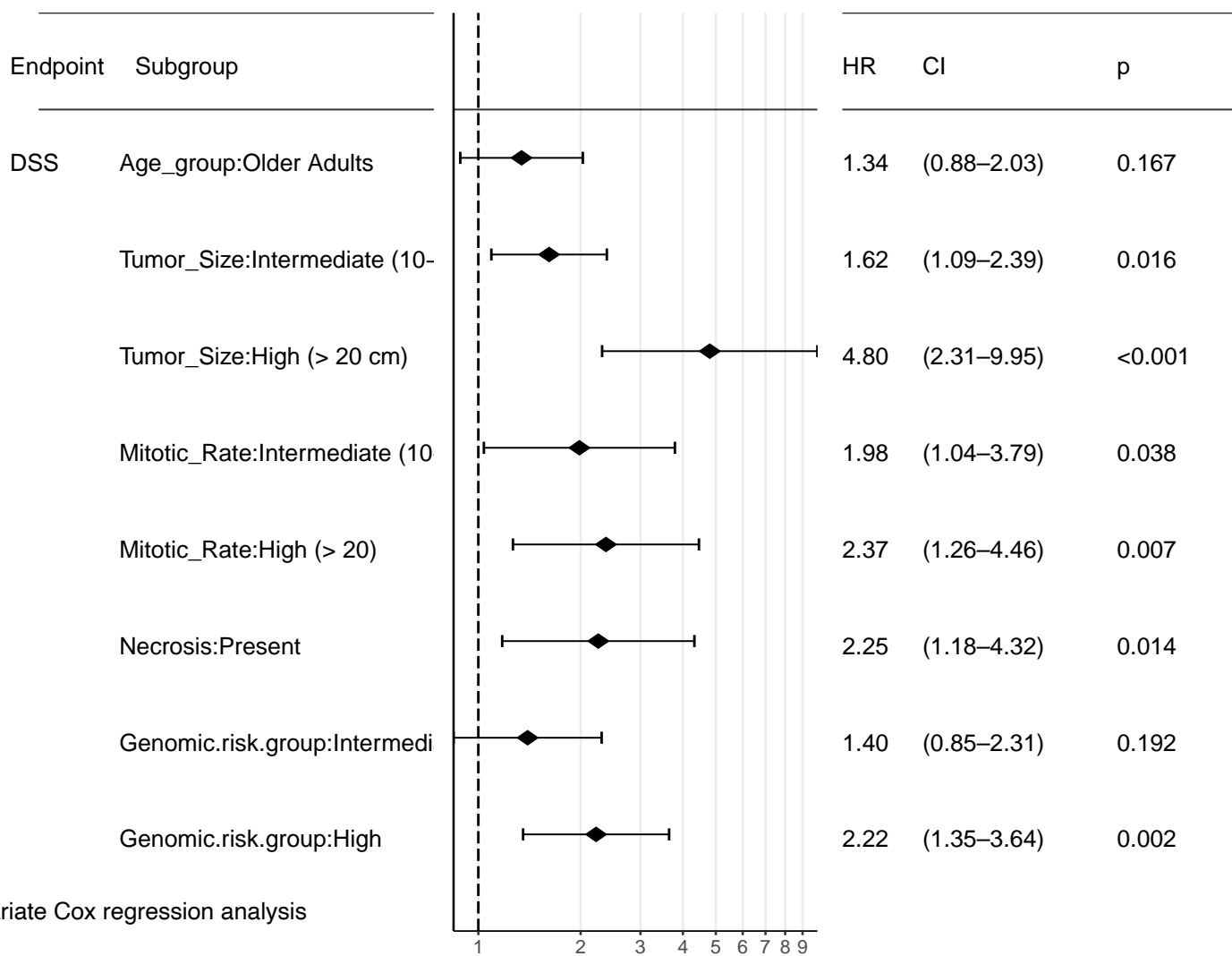| Endpoint	Subgroup		HR	CI	p
DSS	Age_group:Older Adults		1.34	(0.88–2.03)	0.167
	Tumor_Size:Intermediate (10–		1.62	(1.09–2.39)	0.016
	Tumor_Size:High (> 20 cm)		4.80	(2.31–9.95)	<0.001
	Mitotic_Rate:Intermediate (10		1.98	(1.04–3.79)	0.038
	Mitotic_Rate:High (> 20)		2.37	(1.26–4.46)	0.007
	Necrosis:Present		2.25	(1.18–4.32)	0.014
	Genomic.risk.group:Intermedi		1.40	(0.85–2.31)	0.192
	Genomic.risk.group:High		2.22	(1.35–3.64)	0.002

Multivariate Cox regression analysis

Supplementary Figure S12. A. Forest plot depicting univariate Cox regression analysis for clinicopathologic and genomic risk variables for DSS in ULMS patients. B. Forest plot depicting multivariate Cox regression analysis comparing tumor size, mitotic rate and necrosis with high genomic risk group for DSS in ULMS patients.