OS: N = 49 with 43 events

	HR (95% CI)	P–value
HGG, H3 wildtype	1.00 (NA–NA)	
HGG, H3 wildtype, TP53	3.32 (0.9–12.0)	P = 0.07
DMG, H3 K28	6.53 (1.7–25.5)	P < 0.01
DMG, H3 K28, TP53	9.22 (2.6–32.1)	P < 0.01
DHG, H3 G35	1.34 (0.3–7.0)	P = 0.73
DHG, H3 G35, TP53	1.75 (0.4–8.5)	P = 0.49
HGG, IDH, TP53	7.95 (0.6–107.4)	P = 0.12
HGG, PXA	8.96 (1.3–63.3)	P = 0.03
Cluster 2	1.00 (NA–NA)	
Cluster 4	3.23 (0.6–18.7)	P = 0.19
Cluster 6	1.95 (0.7–5.3)	P = 0.19
Cluster 8	0.84 (0.1–4.9)	P = 0.85
spliceosome gsva score	0.18 (0.1–0.6)	P < 0.01
age at diagnosis days	1.00 (1.0–1.0)	P = 0.31

Hazard Ratio ± 95% CI