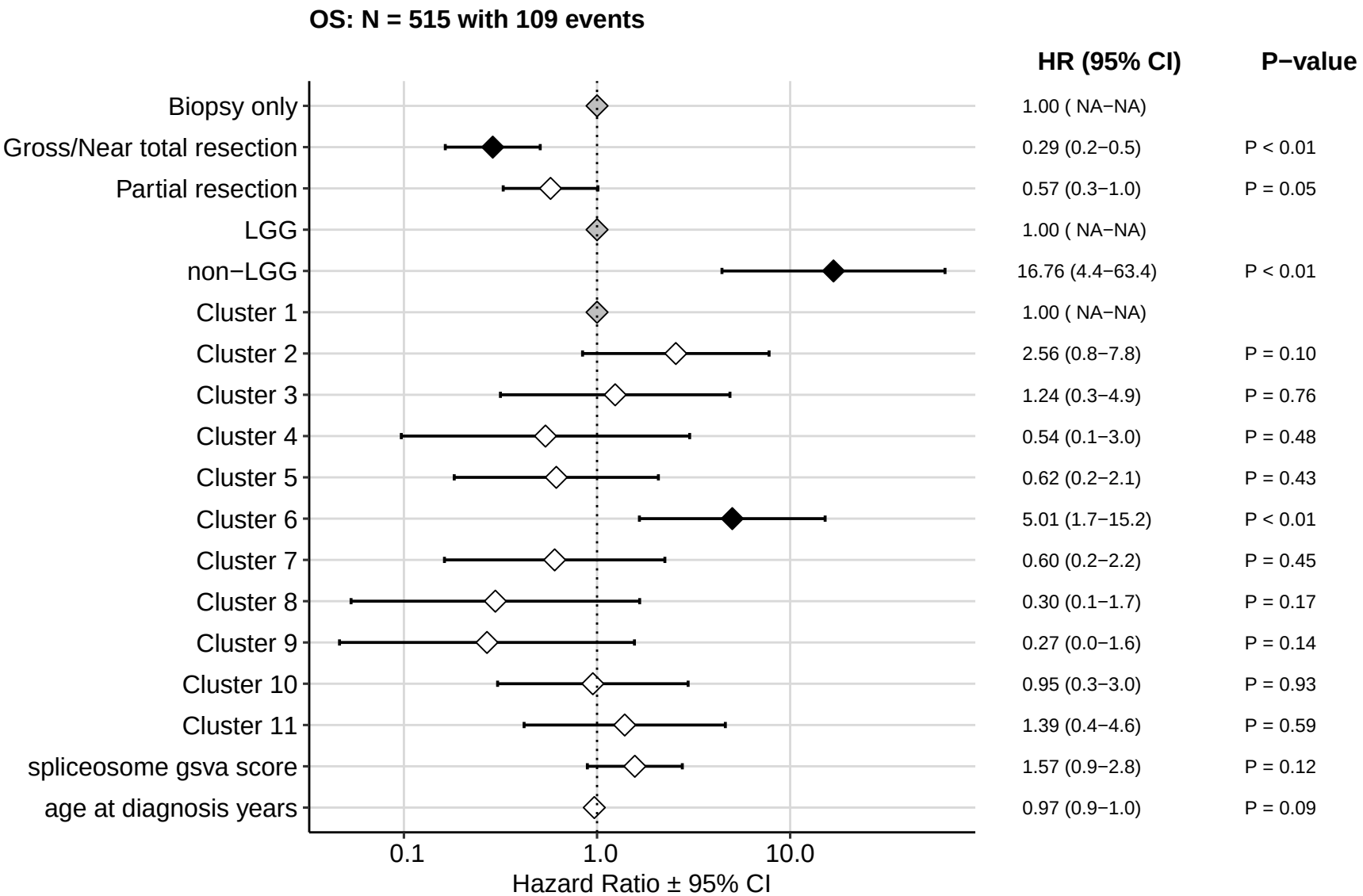
OS: N = 515 with 109 events
HR (95% CI)
P−value
Biopsy only
1.00 (NA−NA)
Gross/Near total resection
0.29 (0.2−0.5)
P < 0.01
Partial resection
0.57 (0.3−1.0)
P = 0.05
LGG
1.00 (NA−NA)
non−LGG
16.76 (4.4−63.4)
P < 0.01
Cluster 1
1.00 (NA−NA)
Cluster 2
2.56 (0.8−7.8)
P = 0.10
Cluster 3
1.24 (0.3−4.9)
P = 0.76
Cluster 4
0.54 (0.1−3.0)
P = 0.48
Cluster 5
0.62 (0.2−2.1)
P = 0.43
Cluster 6
5.01 (1.7−15.2)
P < 0.01
Cluster 7
0.60 (0.2−2.2)
P = 0.45
Cluster 8
0.30 (0.1−1.7)
P = 0.17
Cluster 9
0.27 (0.0−1.6)
P = 0.14
Cluster 10
0.95 (0.3−3.0)
P = 0.93
Cluster 11
1.39 (0.4−4.6)
P = 0.59
spliceosome gsva score
1.57 (0.9−2.8)
P = 0.12
age at diagnosis years
0.97 (0.9−1.0)
P = 0.09
0.1
1.0
10.0
Hazard Ratio ± 95% CI