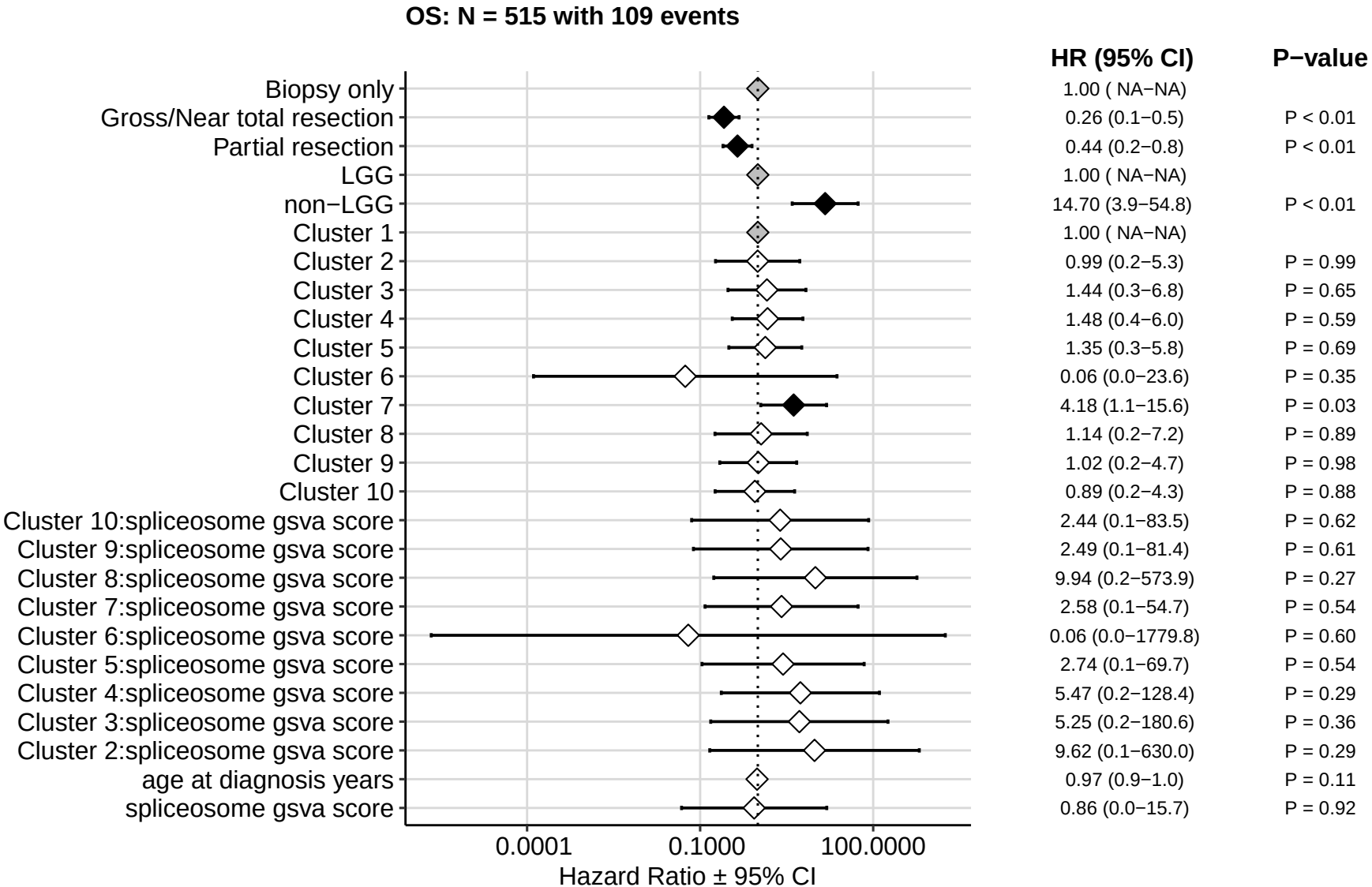
OS: N = 515 with 109 events
HR (95% CI)
P−value
Biopsy only
1.00 (NA−NA)
Gross/Near total resection
0.26 (0.1−0.5)
P < 0.01
Partial resection
0.44 (0.2−0.8)
P < 0.01
LGG
1.00 (NA−NA)
non−LGG
14.70 (3.9−54.8)
P < 0.01
Cluster 1
1.00 (NA−NA)
Cluster 2
0.99 (0.2−5.3)
P = 0.99
Cluster 3
1.44 (0.3−6.8)
P = 0.65
Cluster 4
1.48 (0.4−6.0)
P = 0.59
Cluster 5
1.35 (0.3−5.8)
P = 0.69
Cluster 6
0.06 (0.0−23.6)
P = 0.35
Cluster 7
4.18 (1.1−15.6)
P = 0.03
Cluster 8
1.14 (0.2−7.2)
P = 0.89
Cluster 9
1.02 (0.2−4.7)
P = 0.98
Cluster 10
0.89 (0.2−4.3)
P = 0.88
Cluster 10:spliceosome gsva score
2.44 (0.1−83.5)
P = 0.62
Cluster 9:spliceosome gsva score
2.49 (0.1−81.4)
P = 0.61
Cluster 8:spliceosome gsva score
9.94 (0.2−573.9)
P = 0.27
Cluster 7:spliceosome gsva score
2.58 (0.1−54.7)
P = 0.54
Cluster 6:spliceosome gsva score
0.06 (0.0−1779.8)
P = 0.60
Cluster 5:spliceosome gsva score
2.74 (0.1−69.7)
P = 0.54
Cluster 4:spliceosome gsva score
5.47 (0.2−128.4)
P = 0.29
Cluster 3:spliceosome gsva score
5.25 (0.2−180.6)
P = 0.36
Cluster 2:spliceosome gsva score
9.62 (0.1−630.0)
P = 0.29
age at diagnosis years
0.97 (0.9−1.0)
P = 0.11
spliceosome gsva score
0.86 (0.0−15.7)
P = 0.92
0.0001
0.1000
100.0000
Hazard Ratio ± 95% CI