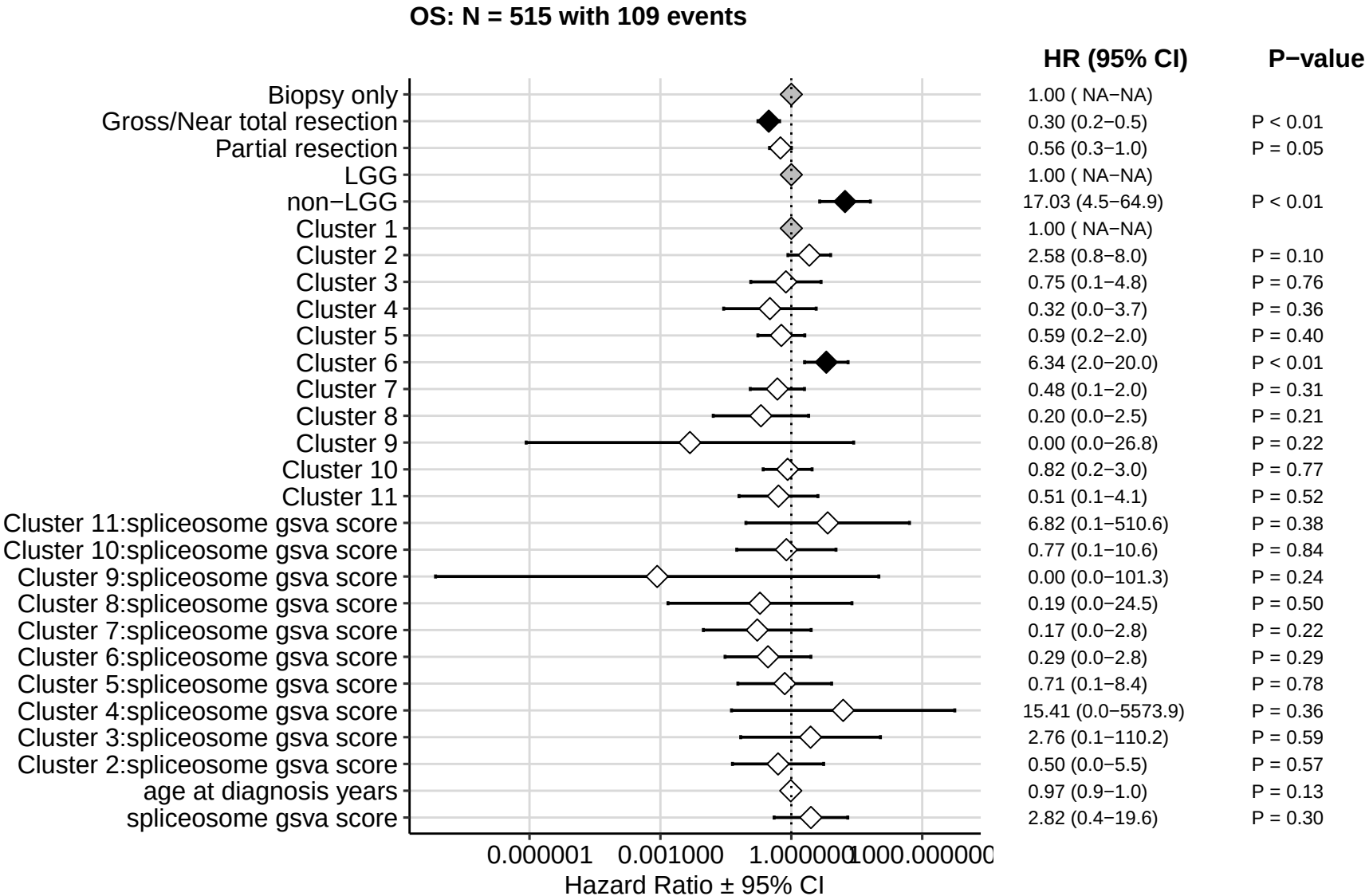
OS: N = 515 with 109 events
HR (95% CI)
P−value
Biopsy only
Gross/Near total resection
Partial resection
LGG
non−LGG
Cluster 1
Cluster 2
Cluster 3
Cluster 4
Cluster 5
Cluster 6
Cluster 7
Cluster 8
Cluster 9
Cluster 10
Cluster 11
Cluster 11:spliceosome gsva score
Cluster 10:spliceosome gsva score
Cluster 9:spliceosome gsva score
Cluster 8:spliceosome gsva score
Cluster 7:spliceosome gsva score
Cluster 6:spliceosome gsva score
Cluster 5:spliceosome gsva score
Cluster 4:spliceosome gsva score
Cluster 3:spliceosome gsva score
Cluster 2:spliceosome gsva score
age at diagnosis years
spliceosome gsva score
1.00 (NA−NA)
0.30 (0.2−0.5) P < 0.01
0.56 (0.3−1.0) P = 0.05
1.00 (NA−NA)
17.03 (4.5−64.9) P < 0.01
1.00 (NA−NA)
2.58 (0.8−8.0) P = 0.10
0.75 (0.1−4.8) P = 0.76
0.32 (0.0−3.7) P = 0.36
0.59 (0.2−2.0) P = 0.40
6.34 (2.0−20.0) P < 0.01
0.48 (0.1−2.0) P = 0.31
0.20 (0.0−2.5) P = 0.21
0.00 (0.0−26.8) P = 0.22
0.82 (0.2−3.0) P = 0.77
0.51 (0.1−4.1) P = 0.52
6.82 (0.1−510.6) P = 0.38
0.77 (0.1−10.6) P = 0.84
0.00 (0.0−101.3) P = 0.24
0.19 (0.0−24.5) P = 0.50
0.17 (0.0−2.8) P = 0.22
0.29 (0.0−2.8) P = 0.29
0.71 (0.1−8.4) P = 0.78
15.41 (0.0−5573.9) P = 0.36
2.76 (0.1−110.2) P = 0.59
0.50 (0.0−5.5) P = 0.57
0.97 (0.9−1.0) P = 0.13
2.82 (0.4−19.6) P = 0.30
0.000001
0.001000
1.000000
1000.000000
Hazard Ratio ± 95% CI