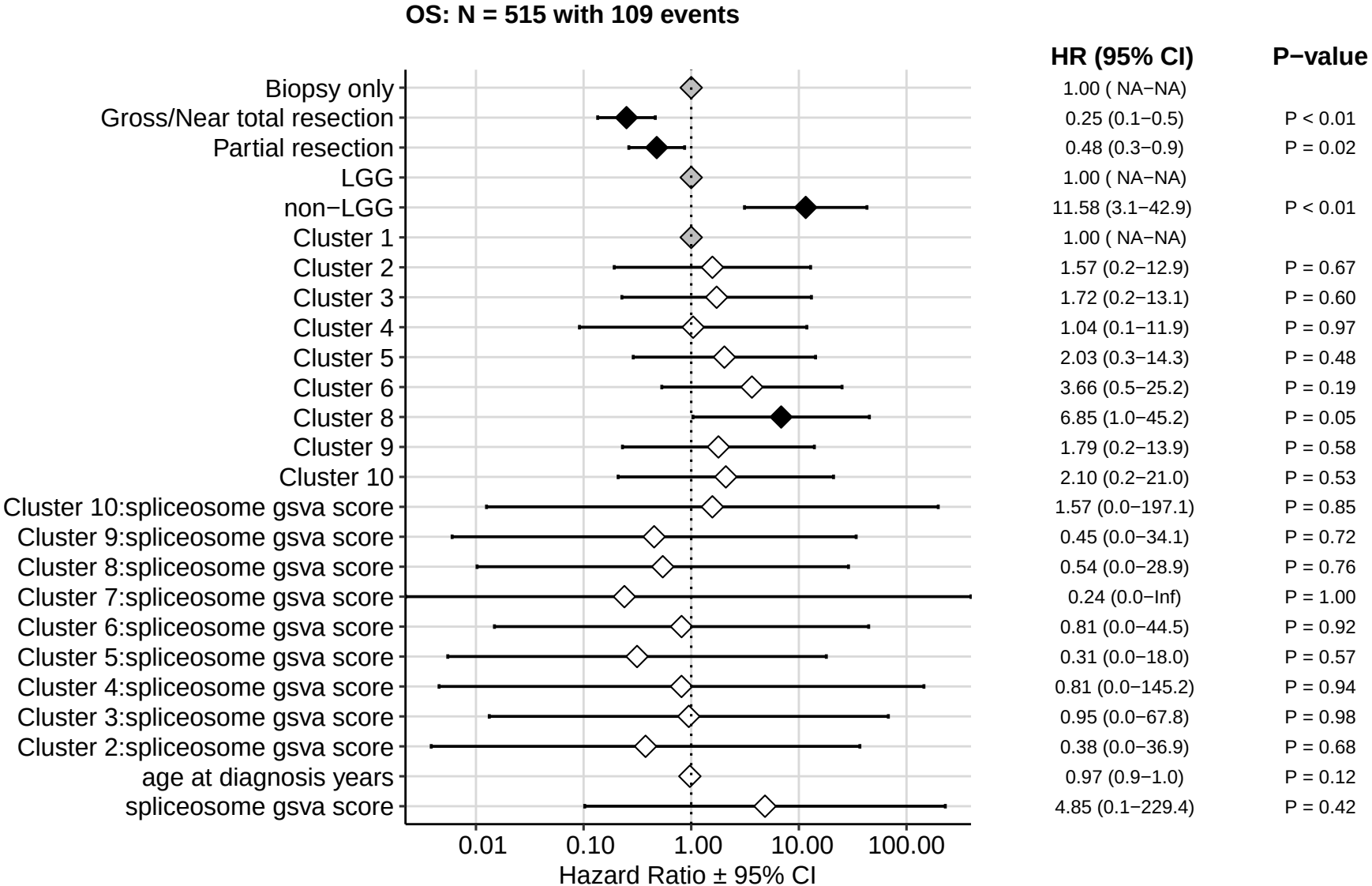
OS: N = 515 with 109 events
HR (95% CI)
P−value
Biopsy only
1.00 (NA−NA)
Gross/Near total resection
0.25 (0.1−0.5)
P < 0.01
Partial resection
0.48 (0.3−0.9)
P = 0.02
LGG
1.00 (NA−NA)
non−LGG
11.58 (3.1−42.9)
P < 0.01
Cluster 1
1.00 (NA−NA)
Cluster 2
1.57 (0.2−12.9)
P = 0.67
Cluster 3
1.72 (0.2−13.1)
P = 0.60
Cluster 4
1.04 (0.1−11.9)
P = 0.97
Cluster 5
2.03 (0.3−14.3)
P = 0.48
Cluster 6
3.66 (0.5−25.2)
P = 0.19
Cluster 8
6.85 (1.0−45.2)
P = 0.05
Cluster 9
1.79 (0.2−13.9)
P = 0.58
Cluster 10
2.10 (0.2−21.0)
P = 0.53
Cluster 10:spliceosome gsva score
1.57 (0.0−197.1)
P = 0.85
Cluster 9:spliceosome gsva score
0.45 (0.0−34.1)
P = 0.72
Cluster 8:spliceosome gsva score
0.54 (0.0−28.9)
P = 0.76
Cluster 7:spliceosome gsva score
0.24 (0.0−Inf)
P = 1.00
Cluster 6:spliceosome gsva score
0.81 (0.0−44.5)
P = 0.92
Cluster 5:spliceosome gsva score
0.31 (0.0−18.0)
P = 0.57
Cluster 4:spliceosome gsva score
0.81 (0.0−145.2)
P = 0.94
Cluster 3:spliceosome gsva score
0.95 (0.0−67.8)
P = 0.98
Cluster 2:spliceosome gsva score
0.38 (0.0−36.9)
P = 0.68
age at diagnosis years
0.97 (0.9−1.0)
P = 0.12
spliceosome gsva score
4.85 (0.1−229.4)
P = 0.42
0.01
0.10
1.00
10.00
100.00
Hazard Ratio ± 95% CI