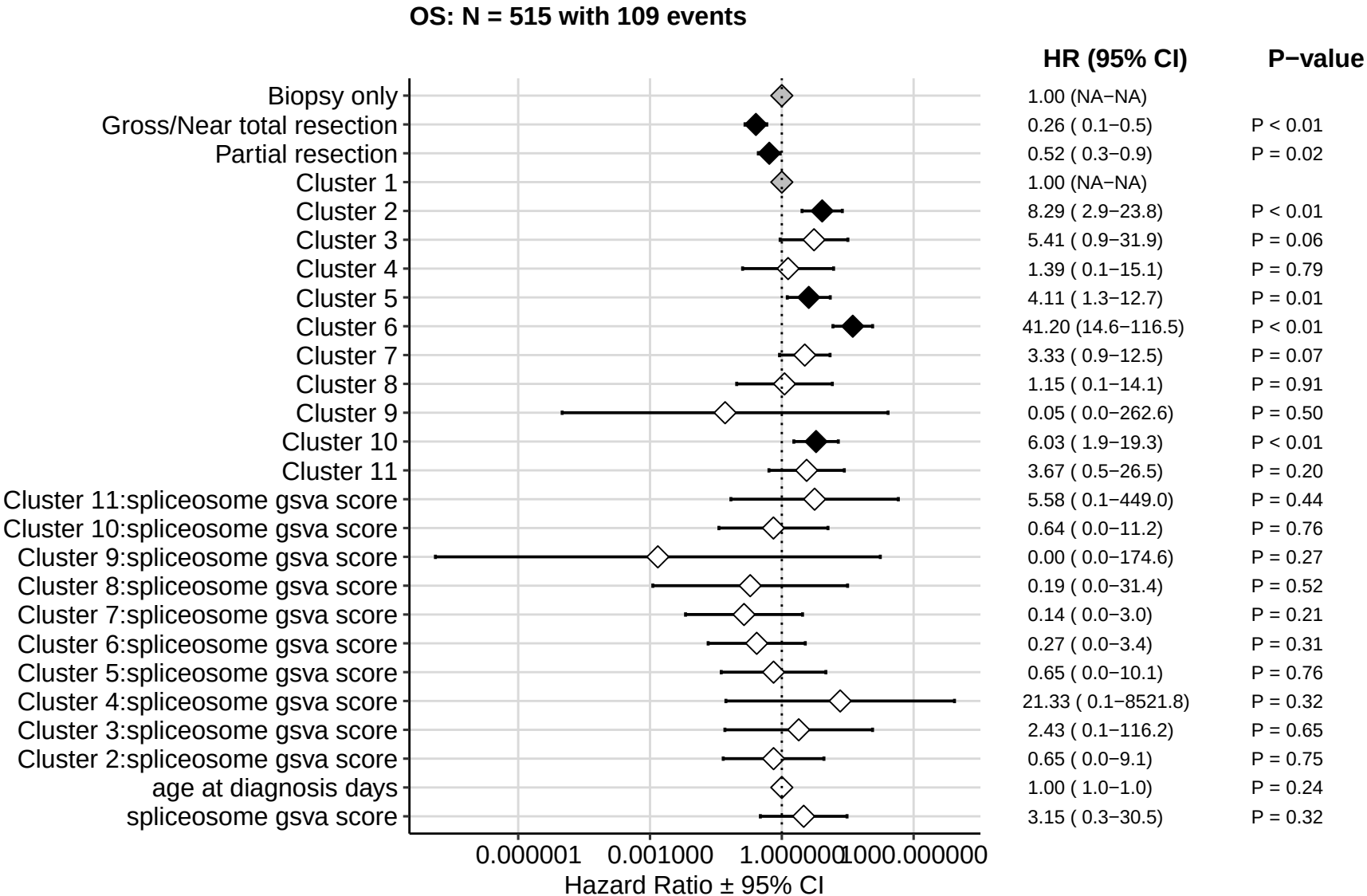
OS: N = 515 with 109 events
HR (95% CI)
P−value
Biopsy only
Gross/Near total resection
Partial resection
Cluster 1
Cluster 2
Cluster 3
Cluster 4
Cluster 5
Cluster 6
Cluster 7
Cluster 8
Cluster 9
Cluster 10
Cluster 11
Cluster 11:spliceosome gsva score
Cluster 10:spliceosome gsva score
Cluster 9:spliceosome 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 days
spliceosome gsva score
1.00 (NA−NA)
0.26 (0.1−0.5) P < 0.01
0.52 (0.3−0.9) P = 0.02
1.00 (NA−NA)
8.29 (2.9−23.8) P < 0.01
5.41 (0.9−31.9) P = 0.06
1.39 (0.1−15.1) P = 0.79
4.11 (1.3−12.7) P = 0.01
41.20 (14.6−116.5) P < 0.01
3.33 (0.9−12.5) P = 0.07
1.15 (0.1−14.1) P = 0.91
0.05 (0.0−262.6) P = 0.50
6.03 (1.9−19.3) P < 0.01
3.67 (0.5−26.5) P = 0.20
5.58 (0.1−449.0) P = 0.44
0.64 (0.0−11.2) P = 0.76
0.00 (0.0−174.6) P = 0.27
0.19 (0.0−31.4) P = 0.52
0.14 (0.0−3.0) P = 0.21
0.27 (0.0−3.4) P = 0.31
0.65 (0.0−10.1) P = 0.76
21.33 (0.1−8521.8) P = 0.32
2.43 (0.1−116.2) P = 0.65
0.65 (0.0−9.1) P = 0.75
1.00 (1.0−1.0) P = 0.24
3.15 (0.3−30.5) P = 0.32
0.000001
0.001000
1.000000
1000.000000
Hazard Ratio ± 95% CI