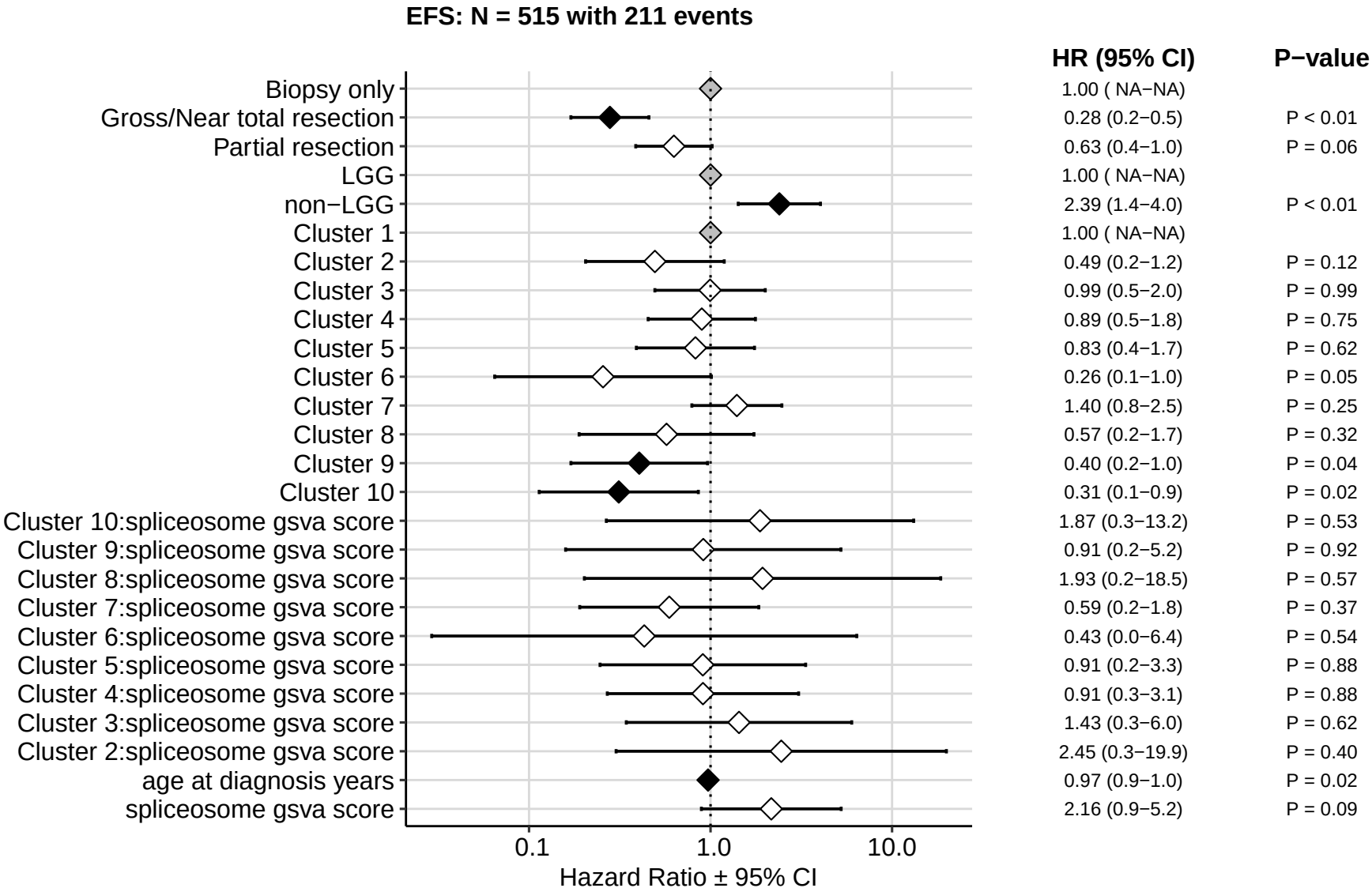
EFS: N = 515 with 211 events
HR (95% CI)
P−value
Biopsy only
1.00 (NA−NA)
Gross/Near total resection
0.28 (0.2−0.5)
P < 0.01
Partial resection
0.63 (0.4−1.0)
P = 0.06
LGG
1.00 (NA−NA)
non−LGG
2.39 (1.4−4.0)
P < 0.01
Cluster 1
1.00 (NA−NA)
Cluster 2
0.49 (0.2−1.2)
P = 0.12
Cluster 3
0.99 (0.5−2.0)
P = 0.99
Cluster 4
0.89 (0.5−1.8)
P = 0.75
Cluster 5
0.83 (0.4−1.7)
P = 0.62
Cluster 6
0.26 (0.1−1.0)
P = 0.05
Cluster 7
1.40 (0.8−2.5)
P = 0.25
Cluster 8
0.57 (0.2−1.7)
P = 0.32
Cluster 9
0.40 (0.2−1.0)
P = 0.04
Cluster 10
0.31 (0.1−0.9)
P = 0.02
Cluster 10:spliceosome gsva score
1.87 (0.3−13.2)
P = 0.53
Cluster 9:spliceosome gsva score
0.91 (0.2−5.2)
P = 0.92
Cluster 8:spliceosome gsva score
1.93 (0.2−18.5)
P = 0.57
Cluster 7:spliceosome gsva score
0.59 (0.2−1.8)
P = 0.37
Cluster 6:spliceosome gsva score
0.43 (0.0−6.4)
P = 0.54
Cluster 5:spliceosome gsva score
0.91 (0.2−3.3)
P = 0.88
Cluster 4:spliceosome gsva score
0.91 (0.3−3.1)
P = 0.88
Cluster 3:spliceosome gsva score
1.43 (0.3−6.0)
P = 0.62
Cluster 2:spliceosome gsva score
2.45 (0.3−19.9)
P = 0.40
age at diagnosis years
0.97 (0.9−1.0)
P = 0.02
spliceosome gsva score
2.16 (0.9−5.2)
P = 0.09
0.1
1.0
10.0
Hazard Ratio ± 95% CI