| **Characteristic** | **N** N = 17*1* | **Y** N = 18*1* | **p-value***2* |
| --- | --- | --- | --- |
| sex |  |  | 0.2 |
| F | 10 (59%) | 7 (39%) |  |
| M | 7 (41%) | 11 (61%) |  |
| age\_at\_surgery | 42 (32, 53) | 59 (43, 67) | 0.062 |
| race |  |  | >0.9 |
| Asian | 0 (0%) | 1 (5.6%) |  |
| Hispanic | 1 (5.9%) | 1 (5.6%) |  |
| Other | 1 (5.9%) | 2 (11%) |  |
| White | 15 (88%) | 14 (78%) |  |
| steroids\_at\_surgery |  |  | <0.001 |
| N | 10 (59%) | 2 (11%) |  |
| Unknown | 2 (12%) | 0 (0%) |  |
| Y | 5 (29%) | 16 (89%) |  |
| grade\_category |  |  | <0.001 |
| High | 7 (41%) | 17 (94%) |  |
| Low | 10 (59%) | 1 (5.6%) |  |
| shotgun\_seq\_batch |  |  | 0.4 |
| Batch 1 | 4 (24%) | 2 (11%) |  |
| Batch 2 | 6 (35%) | 10 (56%) |  |
| Batch 6 | 7 (41%) | 6 (33%) |  |
| glioma\_idh\_mutation |  |  | 0.001 |
| Mutant | 11 (65%) | 2 (11%) |  |
| Wild | 6 (35%) | 16 (89%) |  |
| glioma\_fu\_time | 618 (318, 901) | 238 (155, 419) | 0.019 |
| glioma\_recurrent\_status\_at\_surg |  |  | 0.088 |
| N | 16 (94%) | 12 (67%) |  |
| Y | 1 (5.9%) | 6 (33%) |  |
| Tumor size | 4.70 (3.70, 5.40) | 4.90 (3.00, 6.70) | 0.6 |
| Unknown | 2 | 0 |  |
| *1*n (%); Median (Q1, Q3) | | | |
| *2*Pearson's Chi-squared test; Wilcoxon rank sum exact test; Fisher's exact test; Wilcoxon rank sum test | | | |