|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | **Immune gene cluster** | | | | |
| **Variable** | **T1** | **T2** |  | **NT1** | **NT2** |
| Patient # | 14 | 16 |  | 13 | 20 |
| Age (mean±SD) | 66.79±10.45 | 66.13±10.40 |  | 66.92±8.95 | 68.2±10.38 |
| Gender (M/F) | 9/5 | 12/4 |  | 9/4 | 11/9 |
| HCC nodules (median, range) | 1 (1-2) | 1 (1-5) |  | 1 (1-2) | 1 (1-5) |
| Nodules size (cm, mean±SD) | 46.15±18.61 | 58.25±26.63 |  | 42.5±16.5 | 53.8±25.1 |
| HBsAg (# pos) | 1 | 1 |  | 0 | 0 |
| Anti-HCV (# pos) | 9 | 15 |  | 9 | 19 |
| Alcohol use (# pos) | 4 | 2 |  | 4 | 2 |
| Microvascular invasion (# pos) | 6 | 11 |  | 4 | 11 |
| Lymphomononuclear infiltrate (# pos) | 5 | 3 |  | 3 | 4 |
| Ishak fibrosis stage (median, range) | 6 (3-6) | 6 (2-6) |  | 6 (2-6) | 6 (3-6) |

**Supplementary Table 2**. Clinico-pathological features of patients included in different immune gene expression clusters.

Comparison between groups (T1 vs T2, NT1 vs NT2) did not show statistical differences (Mann-Whitney test, Fisher's exact test).