Supplementary Table 2. Sample information of RNA-based TCR repertoire sequencing

| Sample names                        | cell events | RNA extraction (ng/ml) |
|-------------------------------------|-------------|------------------------|
| 7D after injury CD44<sup>high</sup> Tregs (BH) | 352,000     | 14.1                   |
| 7D after injury CD44<sup>low</sup> Tregs (BL) | 485,000     | 10.7                   |
| Uninjured CD44<sup>high</sup> Tregs (SH) | 73,000      | 5.6                    |
| Uninjured CD44<sup>low</sup> Tregs (SL) | 288,000     | 12.2                   |