

Establishing Molecular Subgroups of CD8+ T Cell-Associated Genes in the Ovarian Cancer Tumour Microenvironment and Predicting the Immunotherapy Response

Supplementary Table S2. Primer sequences for qRT-PCR analysis.

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|----------|--------------------------|
| TXK-F | GATGGGCAGTTGTTTACCAGCC |
| TXK-R | ATGGACCACTCCAAACTGACCG |
| STX7-F | CGAGTAAGAGCCAGTTCCAGAG |
| STX7-R | TGAATAAGACGGAGGTCATCCTC |
| SEMA4F-F | GCAGGATGTTGCTGTGCTTCGA |
| SEMA4F-R | CTGTCCGAATGTCTTGTTGCTCG |
| HEMGN-F | GAACCTGGAGTTTGAGAAACAGAG |
| HEMGN-R | CCTGTTCTCTGCTGCTTGCGTT |

Supplementary Table S3. Clinical information on 10 patients with OC used for immunohistochemistry

| Clinical characteristics | N (%) |
|----------------------------------|--------------|
| Age | |
| <60 | 7 (70%) |
| >60 | 3 (30%) |
| Pathological type | |
| high grade serous ovarian cancer | 10 (100%) |
| Clinical Stages | |
| III | 7 (70%) |
| IV | 3 (30%) |