## Supplementary Table 1

| Table S1 Clinicopathological | characteristics of the | patients in this study |
|------------------------------|------------------------|------------------------|
|------------------------------|------------------------|------------------------|

| Characteristic  | Chemotherapy resistant<br>group <sup>†</sup> | Chemotherapy sensitive<br>group <sup>†</sup> |
|---|--|--|
| Sample counts   | 12   | 27   |
| Age(median(range))                                    | 52 (40-76)                                   | 58 (34-72)                                   |
| Tumor size>5cm (yes/no)                               | 10/2   | 19/8   |
| Massive Ascites >500ml (yes/no)                       | 6/6  | 7/20   |
| Liver metastasis (yes/no)                             | 3/9  | 1/27   |
| Preoperative CA-125 level<br>(mIU/ml, median (range)) | 1016 (158-5856)                              | 539 (94-6455)                                |
| FIGO stage (IIIC/IV)                                  | 5/7  | 24/3   |
| Residual disease (R0/R1/R2) <sup>‡</sup>              | 1/5/6  | 6/15/6                                       |

Notes: Chemotherapy resistance was defined as recurrence within 6 months of achieving complete recovery after the initial treatment or disease progression during chemotherapy. Chemotherapy sensitive patients were defined as having recurrence more than 6 months after achieving complete recovery after the initial treatment.

 $\ddagger$ R0 refers to no residual disease; R1 refers to residual disease with tumors  $\leq$  1cm; R1 refers to residual disease with tumors  $\geq$  1cm

## Supplementary Table 2

Table S2 Primers for reverse transcription-quantitative polymerase chain reaction.

|           | Primer sequence (5'-3')   |                           |  |
|-----------|---------------------------|---------------------------|--|
| Gene Name | Forward                   | reverse                   |  |
| actin     | GAAATCGTGCGTGACATTA       | AGGCAGCTCGTAGCTCTT        |  |
| DUSP6     | GAGTCTGACCTTGACCGAGACCCC  | TTCCTCCAACACGTCCAAGTTGGT  |  |
| cyclinD1  | GTCCTACTTCAAATGTGTGCAGAAG | GGGGATGGTCTCCTTCATCTTAGAG |  |
| cyclinD3  | GACCATCGAAAAACTGTGCATCTAC | CATGGCAAAGGTATAATCTGTAGCA |  |
| cyclinE2  | AACTACCCAGGATGTCAAAAAAAGA | ATCAGGCAAAGGTGAAGGATTAATA |  |