

## Supplementary data

### Could redox biomarkers be useful in patients with high-grade ovarian cancer?

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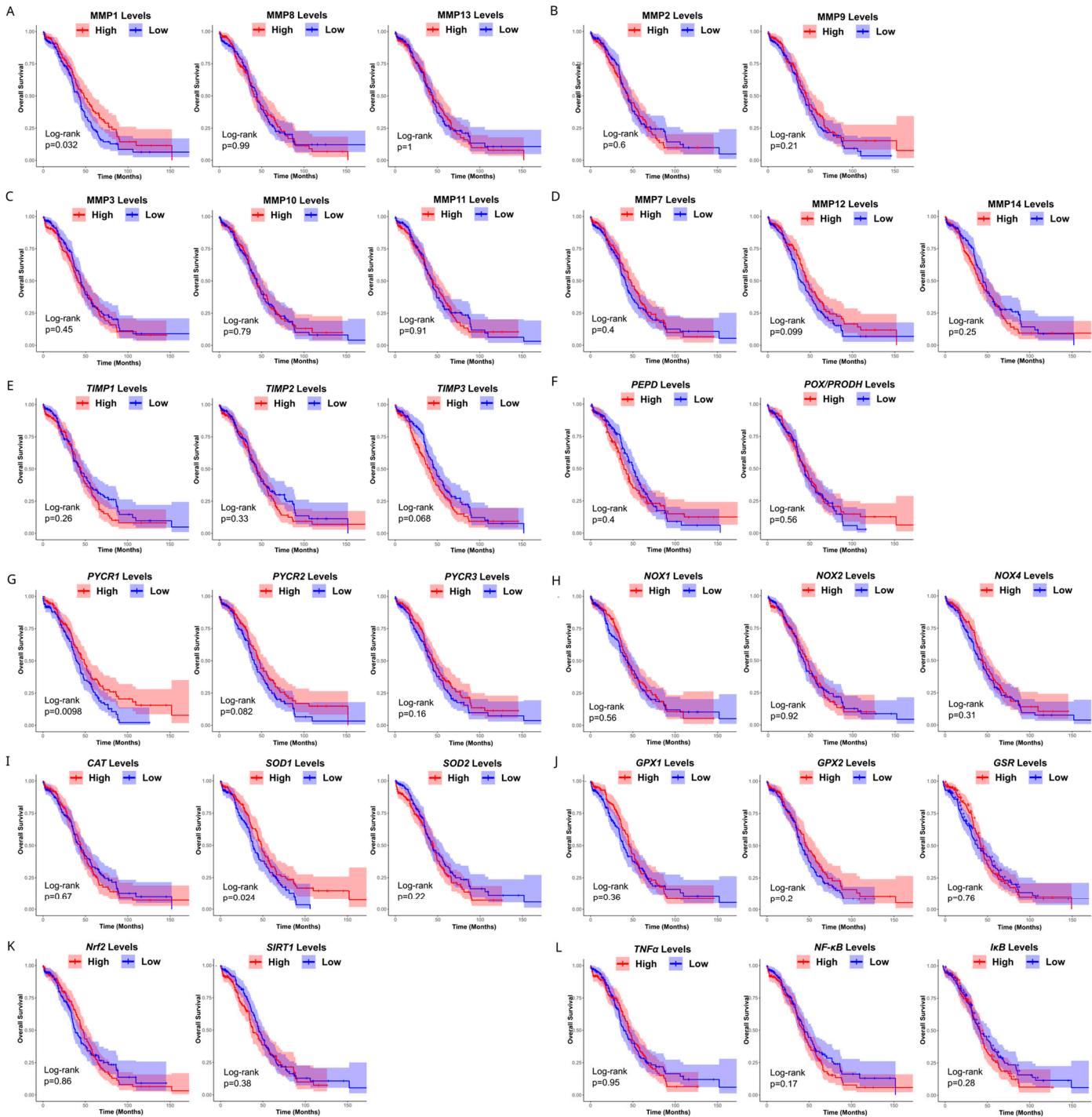
**Table S1.** List of primers used for real-time PCR in the study.

| Target Gene | Accession number | Forward Primer (5'-3')        | Reverse Primer (5'-3')    | Amplicon Length [bp] |
|-------------|------------------|-------------------------------|---------------------------|----------------------|
| MMP1        | NM_00242<br>1    | ACAGCCCAGTACTTATTCCCTTG       | GGGCTTGAAGCTGCTTACGA      | 74                   |
| MMP2        | NM_00453<br>0    | AGCGAGTGGATGCCGCCTTAA         | CATTCCAGGCATCTGCGATGAG    | 138                  |
| MMP3        | NM_00242<br>2    | AAATCCTGATCTTAAAGACAGGCA<br>C | GAGATCAAATGCAATTCAAGGTTCA | 76                   |
| MMP7        | NM_00242<br>3    | TCGGAGGAGATGCTCACTTCGA        | GGATCAGAGGAATGTCCCATAACC  | 127                  |
| MMP8        | NM_00242<br>4    | CCAAGTGGGAACGCACTAACATTGA     | TGGAGAATTGTCACCGTGATCTCTT | 200                  |
| MMP9        | NM_00499<br>4    | GCACGACGTCTTCCAGTACC          | CAGGATGTCATAGGTACGTAGC    | 124                  |
| MMP10       | NM_00242<br>5    | CAAAAGAGGAGGACTCCAACA         | TTCACATCCTTTCGAGGTTG      | 76                   |
| MMP11       | NM_00594<br>0    | AAGAGGTTCGTGCTTCTGG           | CCATGGGAACCGAAGGAT        | 72                   |
| MMP12       | NM_00242<br>6    | TGCTGATGACATACGTGGCA          | AGGATTGGCAAGCGTTGG        | 70                   |
| MMP13       | NM_00242<br>7    | CCTTGATGCCATTACCAGTCTCC       | AAACAGCTCCGCATCAACCTGC    | 97                   |
| MMP14       | NM_00499<br>5    | GGATGGACACAGAGAACTTCGTG       | CGAGAGGTAGTTCTGGGTTGAG    | 118                  |
| TIMP1       | NM_00325<br>4    | TGTTGTTGCTGTGGCTGATAGC        | TCTGGTGTCCCCACGAACCTT     | 118                  |
| TIMP2       | NM_00325<br>5    | ACCCTCTGTGACTTCATCGTGC        | GGAGATGTAGCACGGATCATG     | 129                  |
| TIMP3       | NM_00036<br>2    | TACCGAGGCTTCACCAAGATGC        | CATCTGCCATCATAGACGCGAC    | 132                  |
| PEPD        | NM_00028<br>5    | GAGAACTCAGCCGTGCTACACT        | AAGGAGCAGGTGATGTCGGAAG    | 131                  |
| POX/PRODH   | NM_01633<br>5    | AGAGTCAGCGATGACGGCTTCA        | CTTGCTCCACAGCCATTGGTG     | 127                  |
| PYCR1       | NM_00690<br>7    | AAGATGGCAGGCTTGTGGAGCA        | CAGAGCATCCAGGGCTGTGAAA    | 134                  |
| PYCR2       | NM_01332<br>8    | TGCAAGCCAGACACATCGTGGT        | GTGTTGGTCATGCAGCGAACATC   | 115                  |
| PYCR3       | NM_02307<br>8    | GTGGAAGCTCAGCACATACTGG        | CTTGGTGGCAAAGATGACGAGC    | 135                  |
| Nrf2        | NM_00616<br>4    | CCAAAAGGAGCAAGAGAAAGCC        | GGCAACCTGGGAGTAGTTGG      | 133                  |

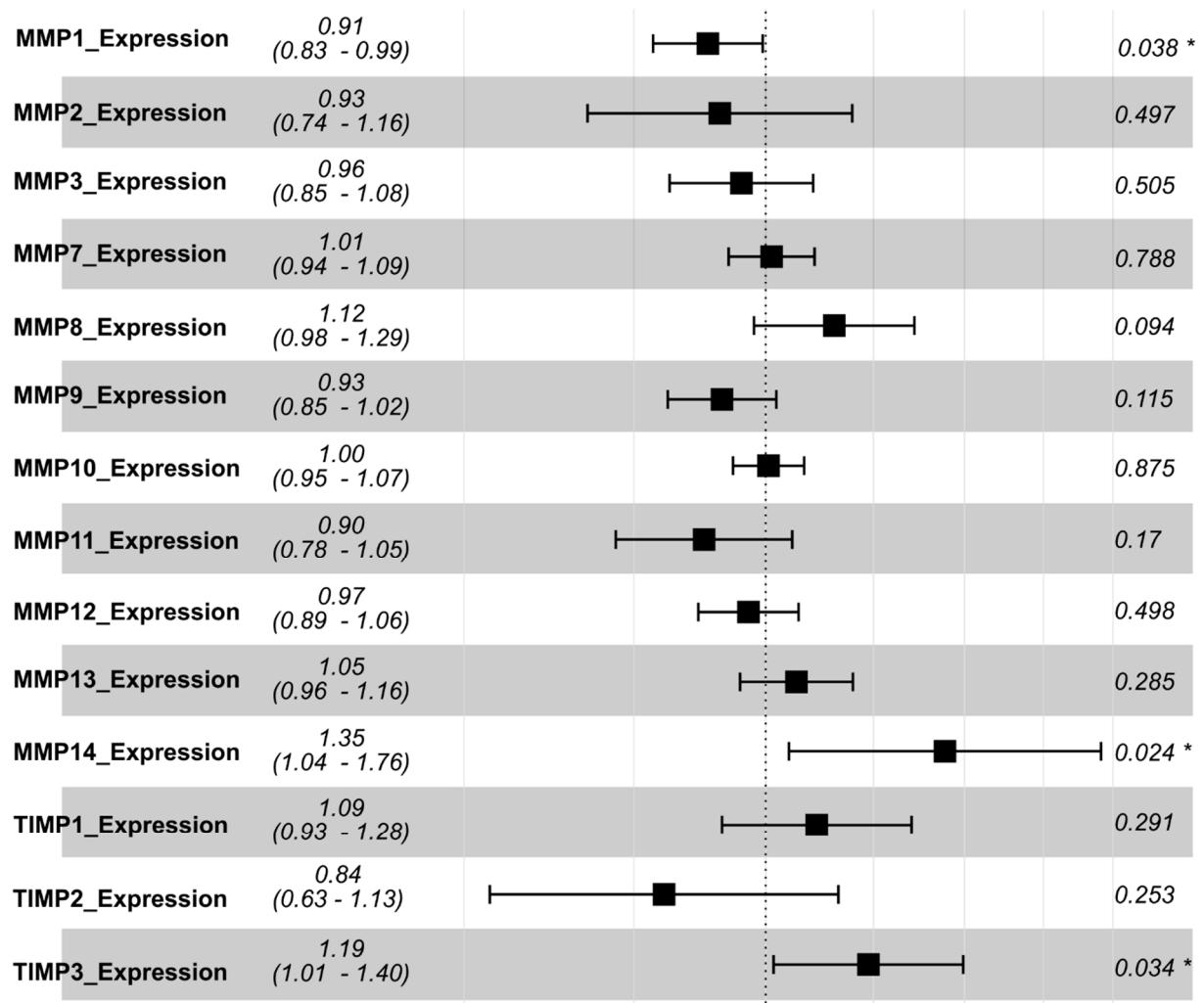
|                                |               |                        |                          |     |
|--------------------------------|---------------|------------------------|--------------------------|-----|
| <i>Sirt1</i>                   | NM_01223<br>8 | TAGACACGCTGGAACAGGTTGC | CTCCTCGTACAGCTTCACAGTC   | 117 |
| <i>GPx</i>                     | NM_00208<br>5 | ACAAGAACGGCTGCGTGGTGAA | GCCACACACTTGTGGAGCTAGA   | 100 |
| <i>GSR</i>                     | NM_00063<br>7 | TGGCACTTGCCTGAATGTTG   | CACATAGGCATCCGCTTTTC     | 157 |
| <i>TNF<math>\alpha</math></i>  | NM_00059<br>4 | CTGGGGCCTACAGCTTGAT    | GGCTCCGTCTCAAGGAAG       | 170 |
| <i>NF-<math>\kappa</math>B</i> | NM_00399<br>8 | GCAGCACTACTTCTTGACCACC | TCTGCTCCTGAGCATTGACGTC   | 130 |
| <i>I<math>\kappa</math>B</i>   | NM_02052<br>9 | CAGCAGACTCCACTCCACTT   | GAGAGGGTATTCCTCGAA       | 114 |
| $\beta$ -actin                 | NM_00110<br>1 | AGTCGGTTGGAGCGAGCATC   | GGACTTCCTGTAACAACGCATCTC | 115 |

**Table S2.** The receiver operating characteristic (ROC) analysis in HGSOC patients for lymph node metastasis.

| Parameter                    | AUC    | p value | 95%<br>confidence<br>interval | Cut-off  | Sensitivity [%] | Specificity [%] |
|------------------------------|--------|---------|-------------------------------|----------|-----------------|-----------------|
| <i>MMP11</i>                 | 0.7143 | 0.0495  | 0.5231 - 0.9055               | > 1.488  | 73.33           | 71.43           |
| <i>I<math>\kappa</math>B</i> | 0.7188 | 0.0417  | 0.5249 - 0.9126               | < 0.6655 | 68.75           | 71.43           |

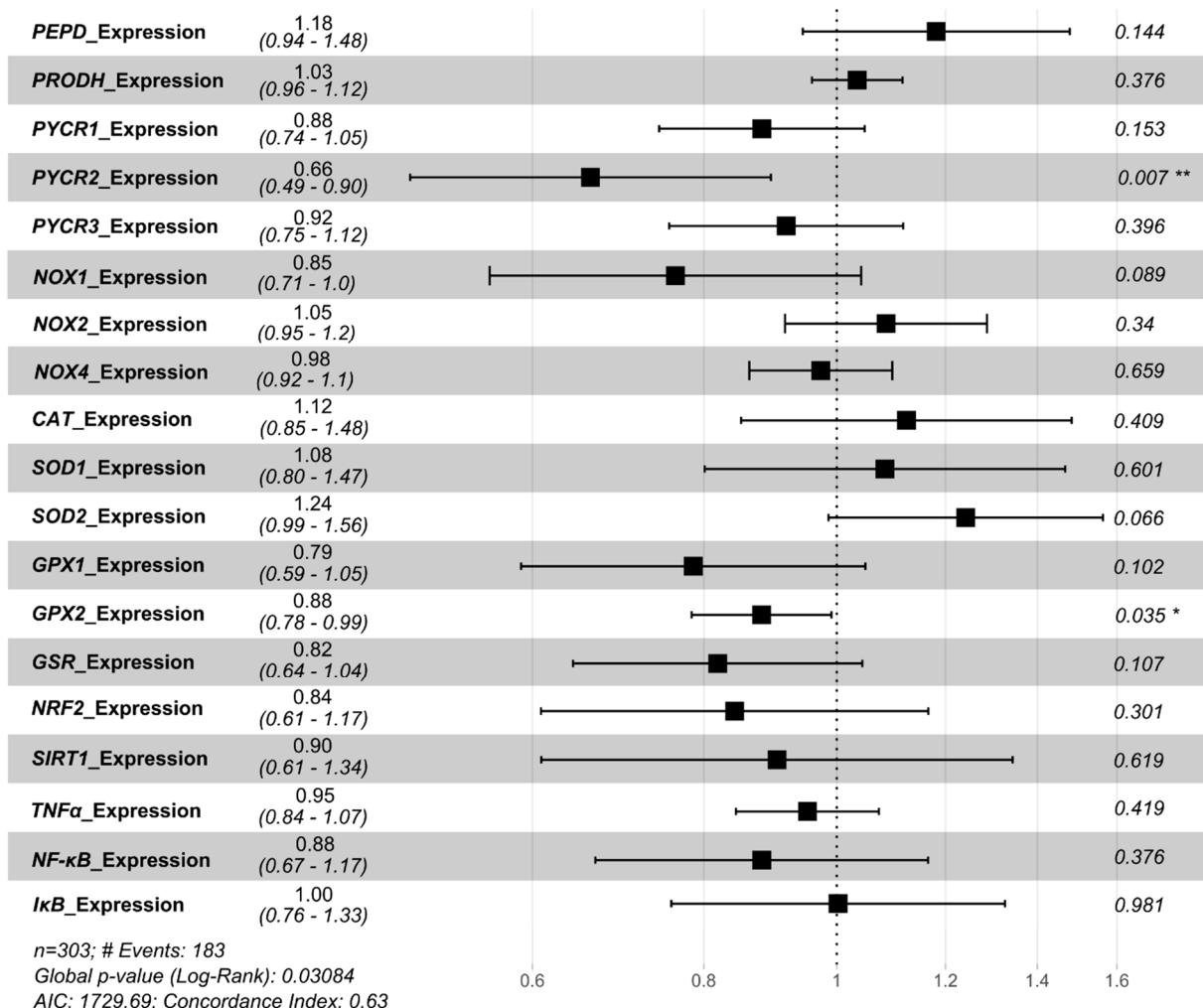


**Figure S1.** Kaplan-Meier overall survival curves of the matrix- (A-G) and oxidative stress-associated (H-L) genes based on TCGA dataset. OC samples were assigned into two separate groups depending on whether target expression of each sample is higher (high expression) or lower (low expression) than the median.

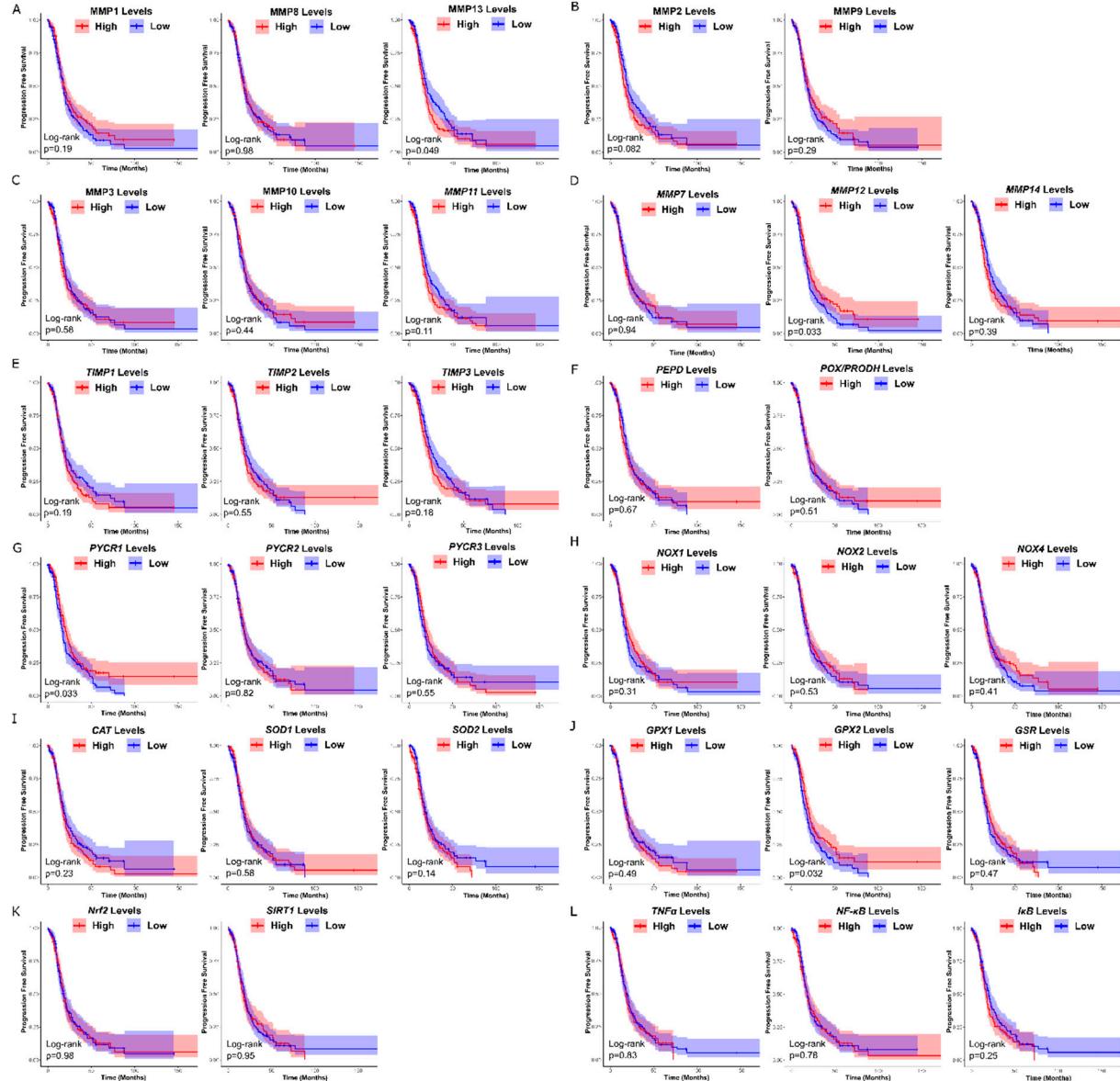


$n=3$ ; # Events: 183  
 Global p-value (Log-Rank): 0.0082427  
 AIC: 1724.03; Concordance Index: 0.62

**Figure S2.** MMPs and TIMPs hazard ratio for overall survival in TCGA ovarian cancer patients. \*  $p < 0.05$



**Figure S3.** Oxidative stress-associated hazard ratio for overall survival in TCGA ovarian cancer patients. \* p<0.05, \*\* p<0.01



**Figure S4.** Kaplan-Meier progression free survival curves of the matrix- (A-G) and oxidative stress-associated (H-L) genes based on TCGA dataset. OC samples were assigned into two separate groups depending on whether target expression of each sample is higher (high expression) or lower (low expression) than the median.