

SUPPLEMENTARY TABLES

Supplementary Table 3. A list of significantly prognostic genes identified by univariate Cox regression analysis.

| Gene | HR | CI5 | CI95 | P-value |
|----------|------|------|------|-------------|
| CXCL14 | 0.92 | 0.86 | 0.98 | 0.008001475 |
| WLS | 0.88 | 0.79 | 0.99 | 0.028963359 |
| SDC1 | 1.21 | 1.05 | 1.39 | 0.009853082 |
| TNN | 0.89 | 0.82 | 0.96 | 0.001691972 |
| PDLIM4 | 0.87 | 0.77 | 0.99 | 0.030797728 |
| ITPRIPL2 | 1.15 | 1 | 1.32 | 0.047845533 |
| EMP1 | 1.23 | 1.05 | 1.43 | 0.009450169 |
| CERCAM | 1.24 | 1.03 | 1.49 | 0.020515589 |
| MARVELD1 | 1.24 | 1.04 | 1.48 | 0.017340631 |
| CYTH3 | 1.23 | 1.02 | 1.49 | 0.027050121 |
| PRKG1 | 1.16 | 1.01 | 1.33 | 0.030374849 |
| XG | 1.15 | 1.02 | 1.3 | 0.026768892 |