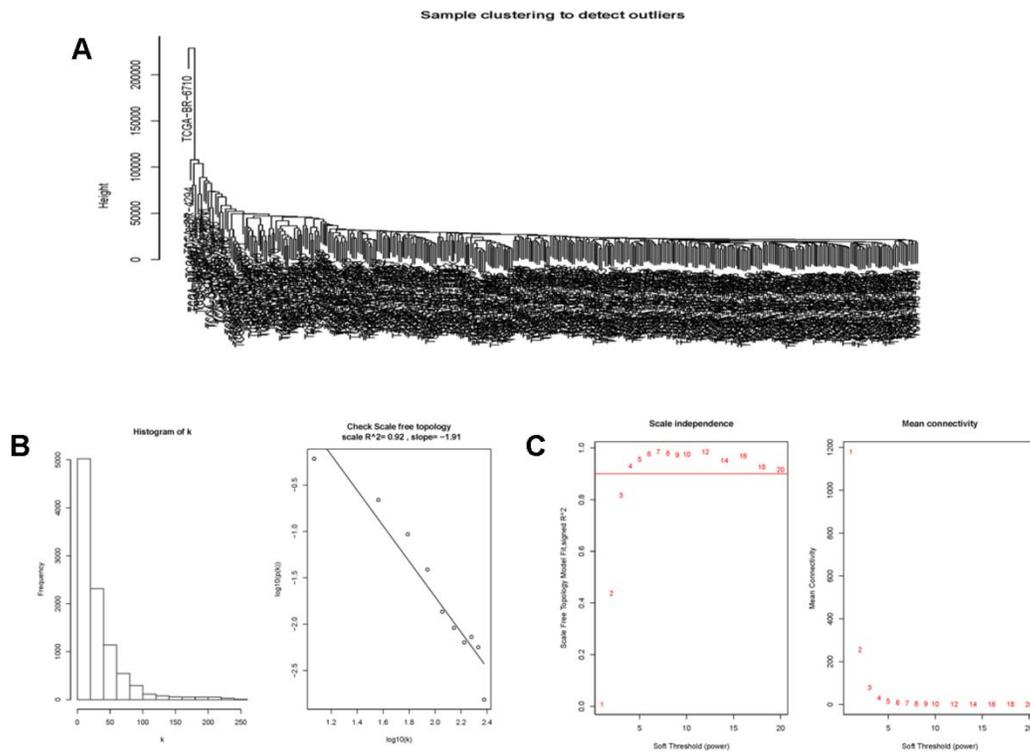
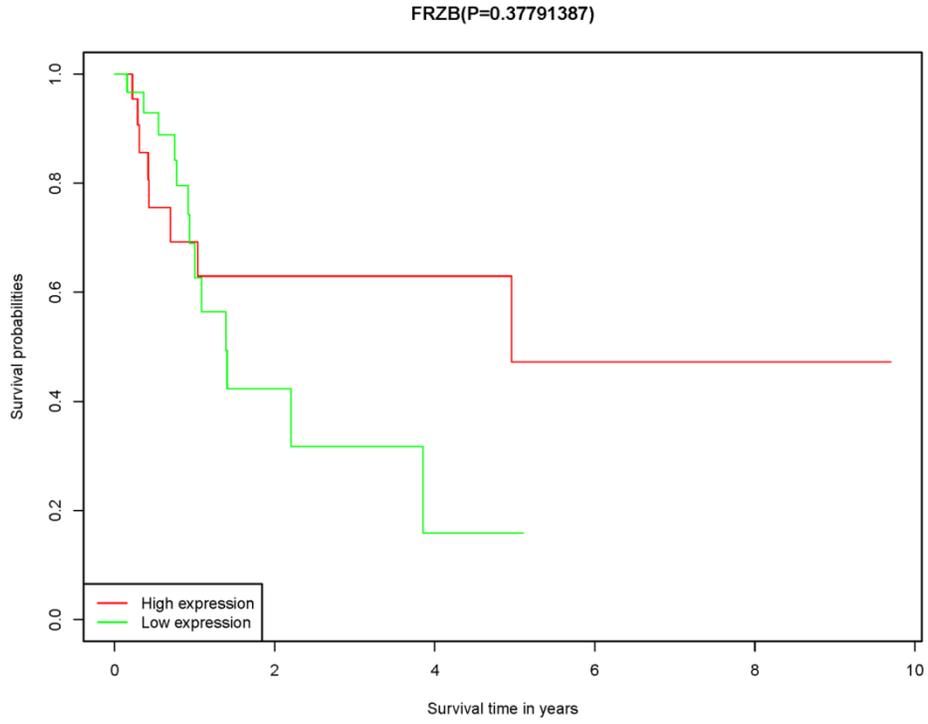


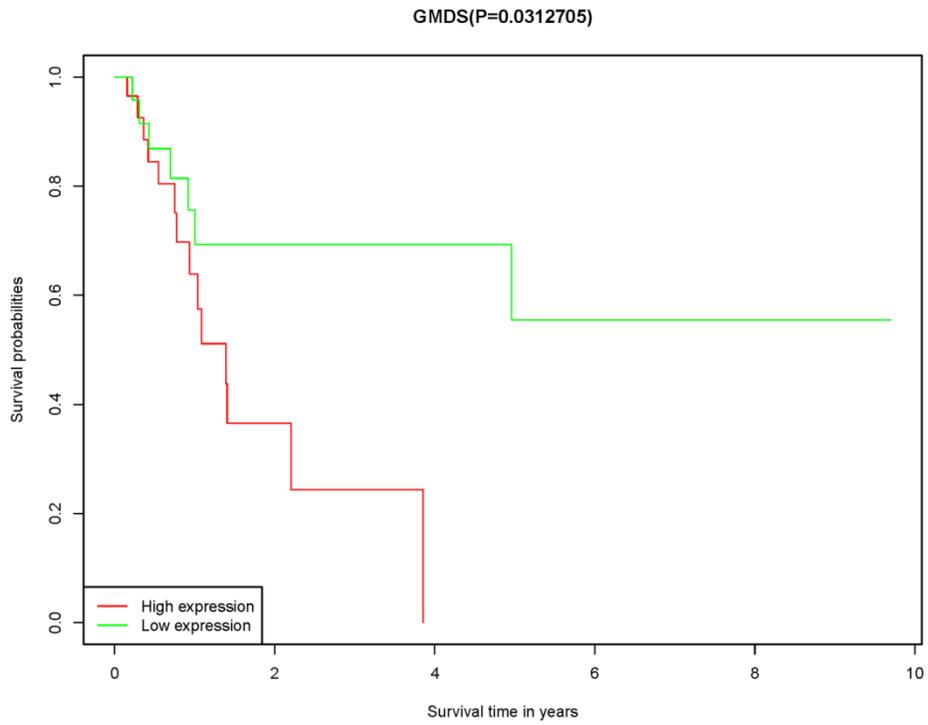
SUPPLEMENTARY FIGURES



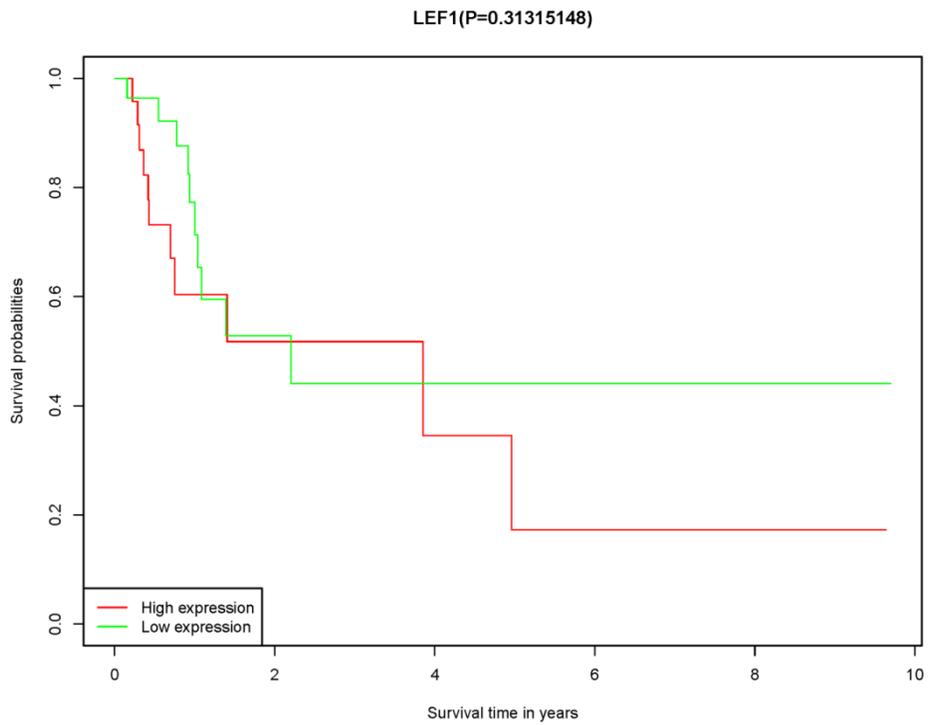
Supplementary Figure 1. (A) Sample clustering to detect outliers. (B, C) Determination of soft-thresholding power in the co-expression network analysis.



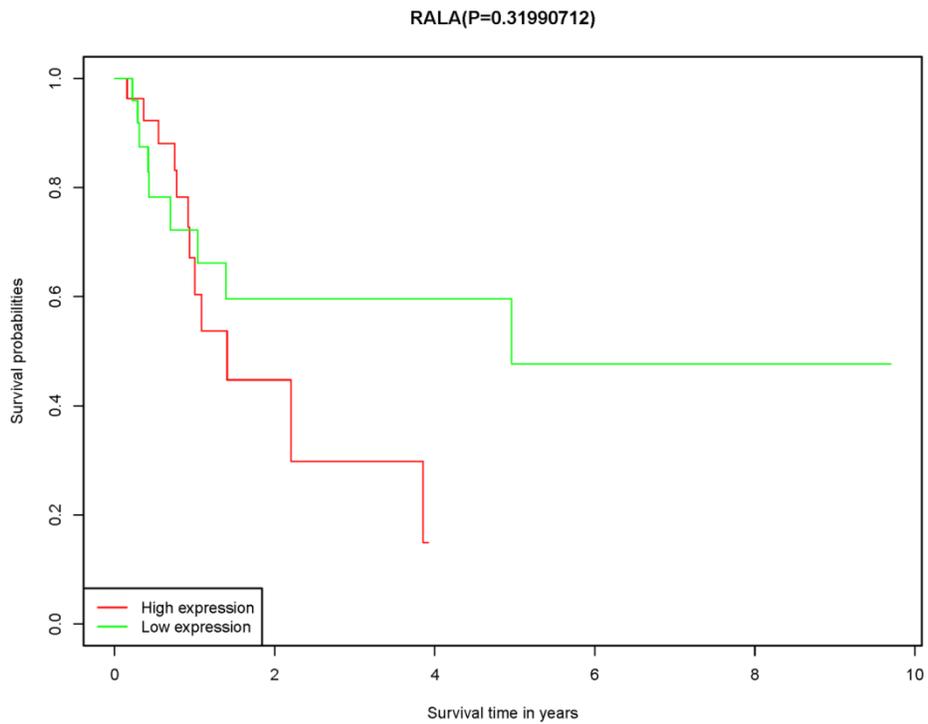
Supplementary Figure 2. The relation of high- and low expression of 10 hub genes for OS.



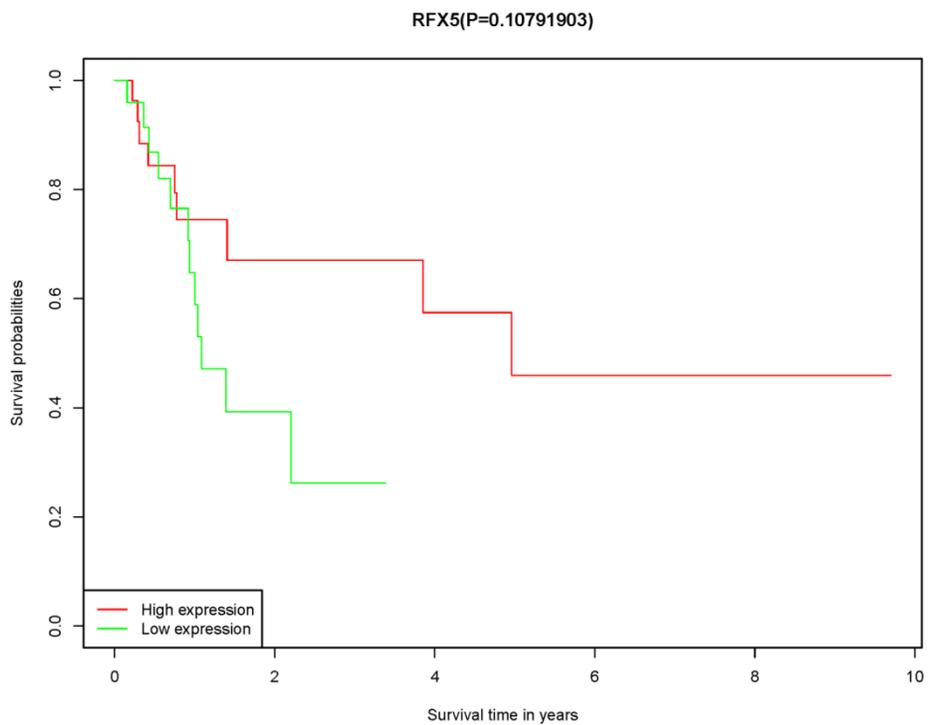
Supplementary Figure 3. The relation of high- and low expression of 10 hub genes for OS.



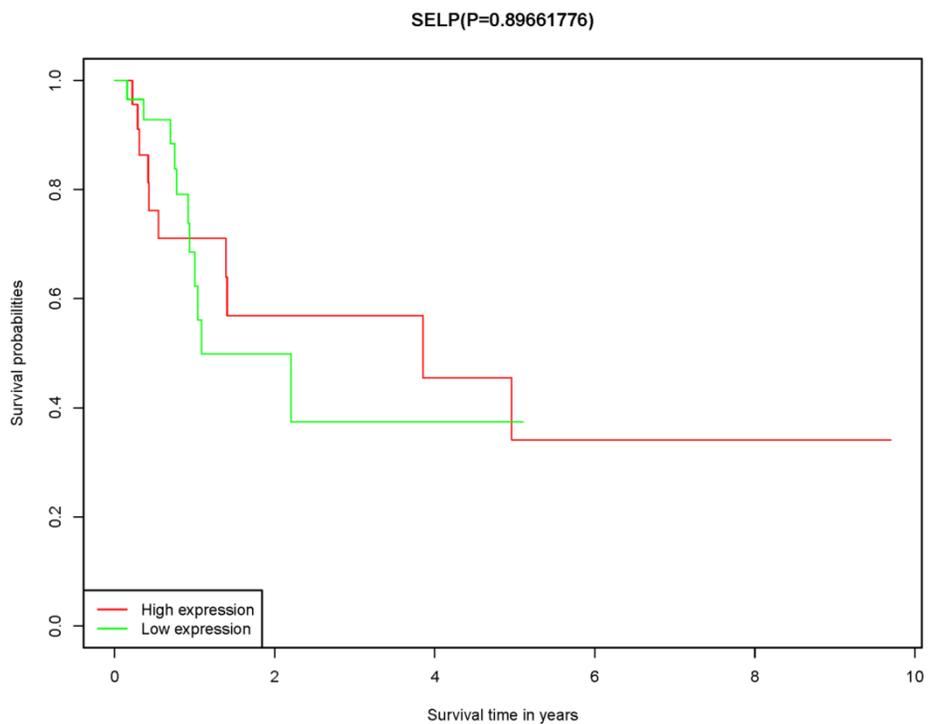
Supplementary Figure 4. The relation of high- and low expression of 10 hub genes for OS.



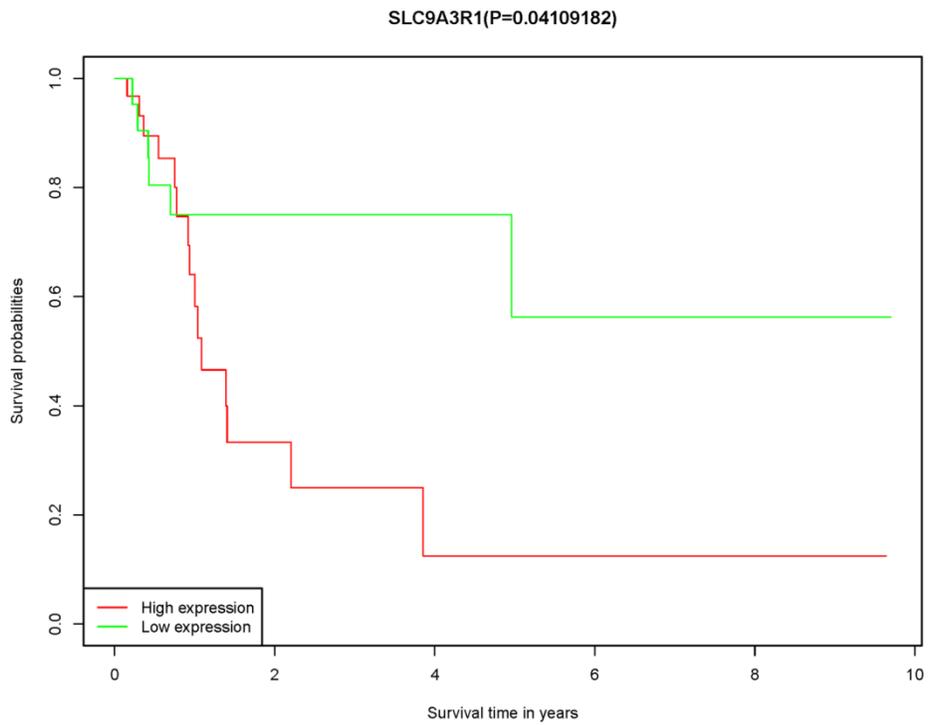
Supplementary Figure 5. The relation of high- and low expression of 10 hub genes for OS.



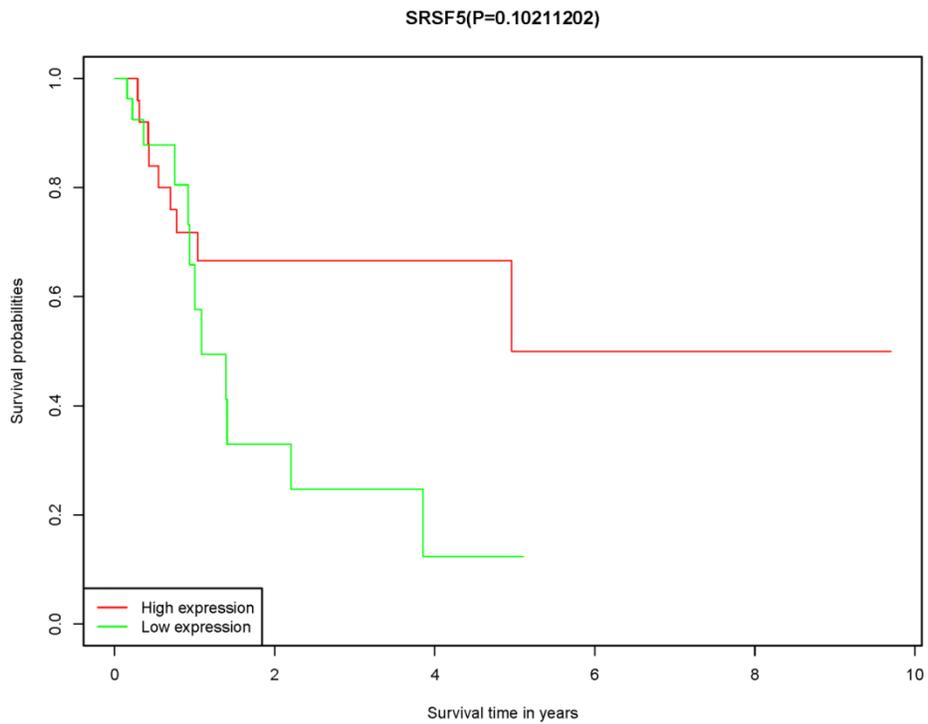
Supplementary Figure 6. The relation of high- and low expression of 10 hub genes for OS.



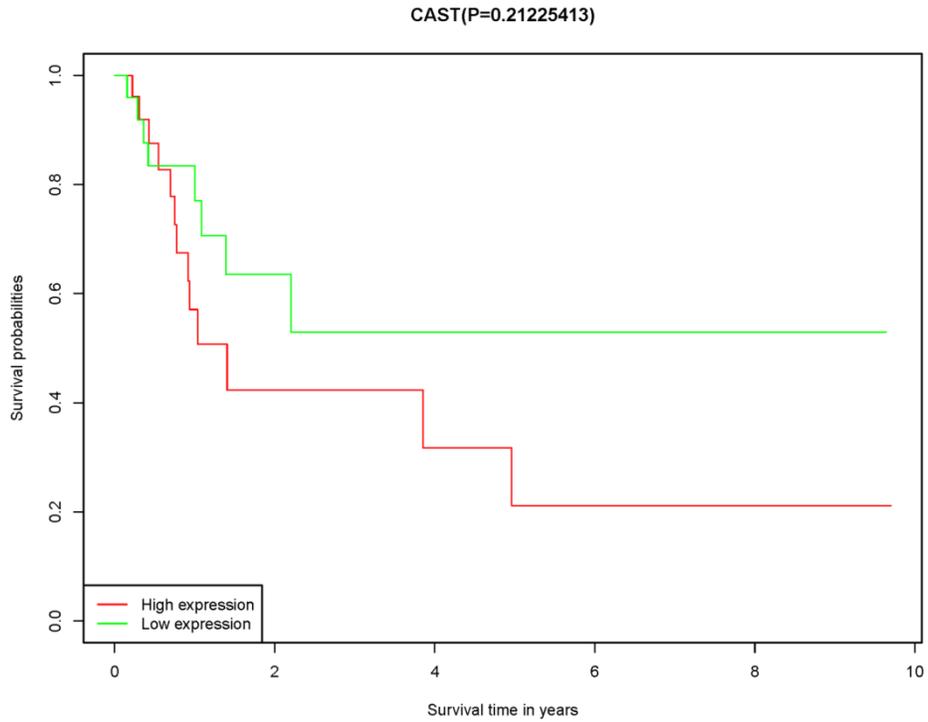
Supplementary Figure 7. The relation of high- and low expression of 10 hub genes for OS.



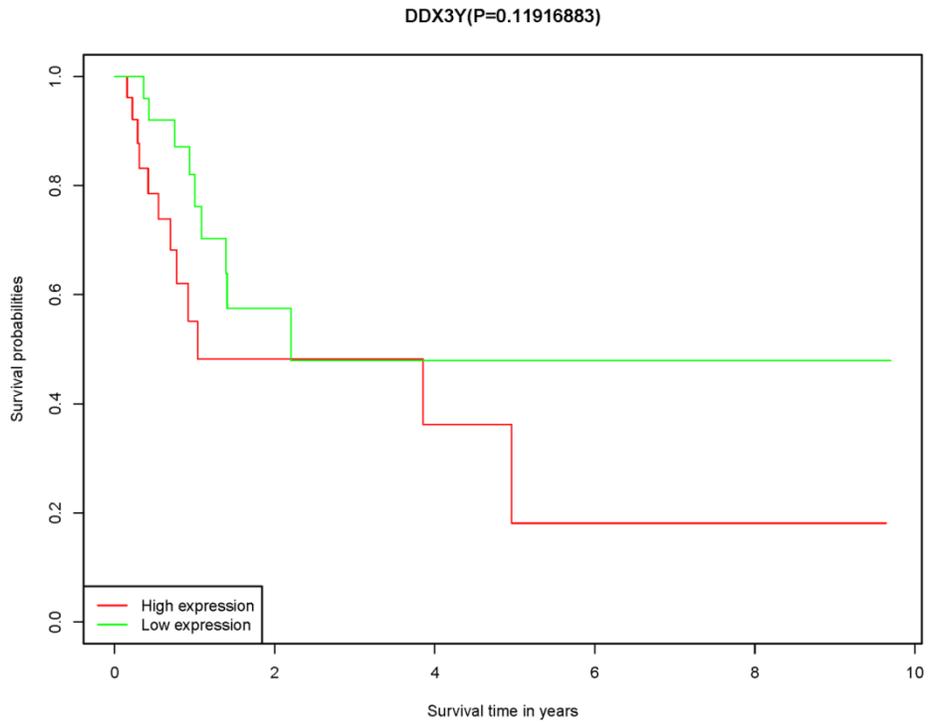
Supplementary Figure 8. The relation of high- and low expression of 10 hub genes for OS.



Supplementary Figure 9. The relation of high- and low expression of 10 hub genes for OS.



Supplementary Figure 10. The relation of high- and low expression of 10 hub genes for OS.



Supplementary Figure 11. The relation of high- and low expression of 10 hub genes for OS.