



**Supplementary Figure S1.** Bioinformatic analysis indicated that KIF14 is enhanced in BLCA patients and correlated with the prognosis. **A** – KIF14 expression level in BLCA tissues and normal tissues was analyzed from bioinformatic analysis (for tumor tissues and normal tissues,  $n = 404$  and  $28$  respectively, from TCGA dataset). **B, C** – KIF14 expression was correlated with the DFS of BLCA patients in different numbers of cases

\* $p < 0.05$ .