

**Figure 1** Bioinformatics analyses of the expression of NTNG gene in ovarian cancer. Volcano plots of gene expression profile in GSE45553 and GSE73935 datasets (A, B). The mRNA level of NTNG1 was higher in cisplatin-sensitive than that in cisplatin-resistant cells (C, D). Progression-free survival (PFS) in overall cases (E), and in the subgroup receiving cisplatin treatment (F): patients with a high NTNG1 level in cancer tissues had shorter PFS compared with those with a low level. \*: p < 0.05.

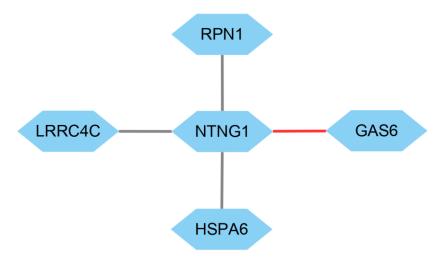
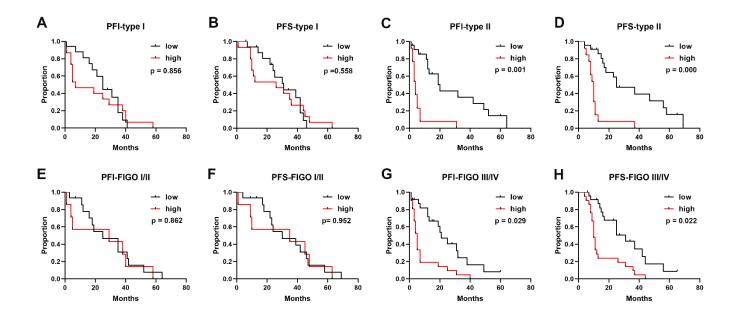


Figure 2 Proteins interacted with NTNG1 in the BioGRID database.



**Figure 3** NTNG1 level correlated with prognosis in EOC patients. Platinum-free interval (PFI) and Progression-free survival (PFS) in type I (n = 32; low-grade serous: 24, clear cell: 7, endometrioid: 1) (A, B) and type II cancers (n = 35; high-grade serous) (C, D). PFI and PFS in FIGO I/II (n = 22) (E, F) and FIGO III/IV cancers (n = 45) (G, H): type II or FIGO III/IV cancer patients with a high NTNG1 level in cancer tissues had shorter PFI and PFS compared with those with a low level.