

Figure S1. OS data in all CTSL abnormally expressed malignancies. OS, overall survival; CTSL, cathepsin L.

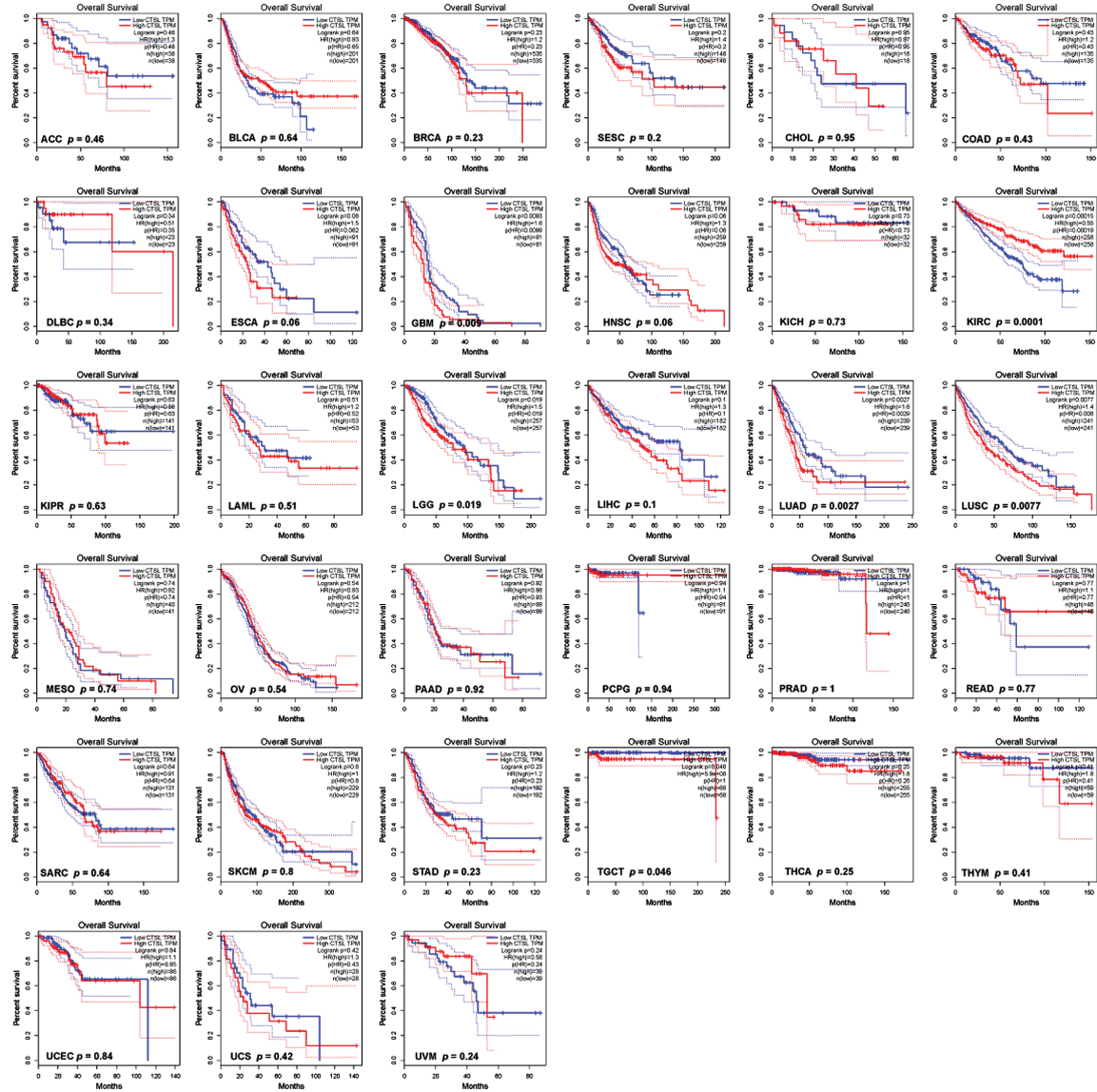


Figure S2. DNA methylation aberration of CTSL in all tumors. CTSL, cathepsin L.

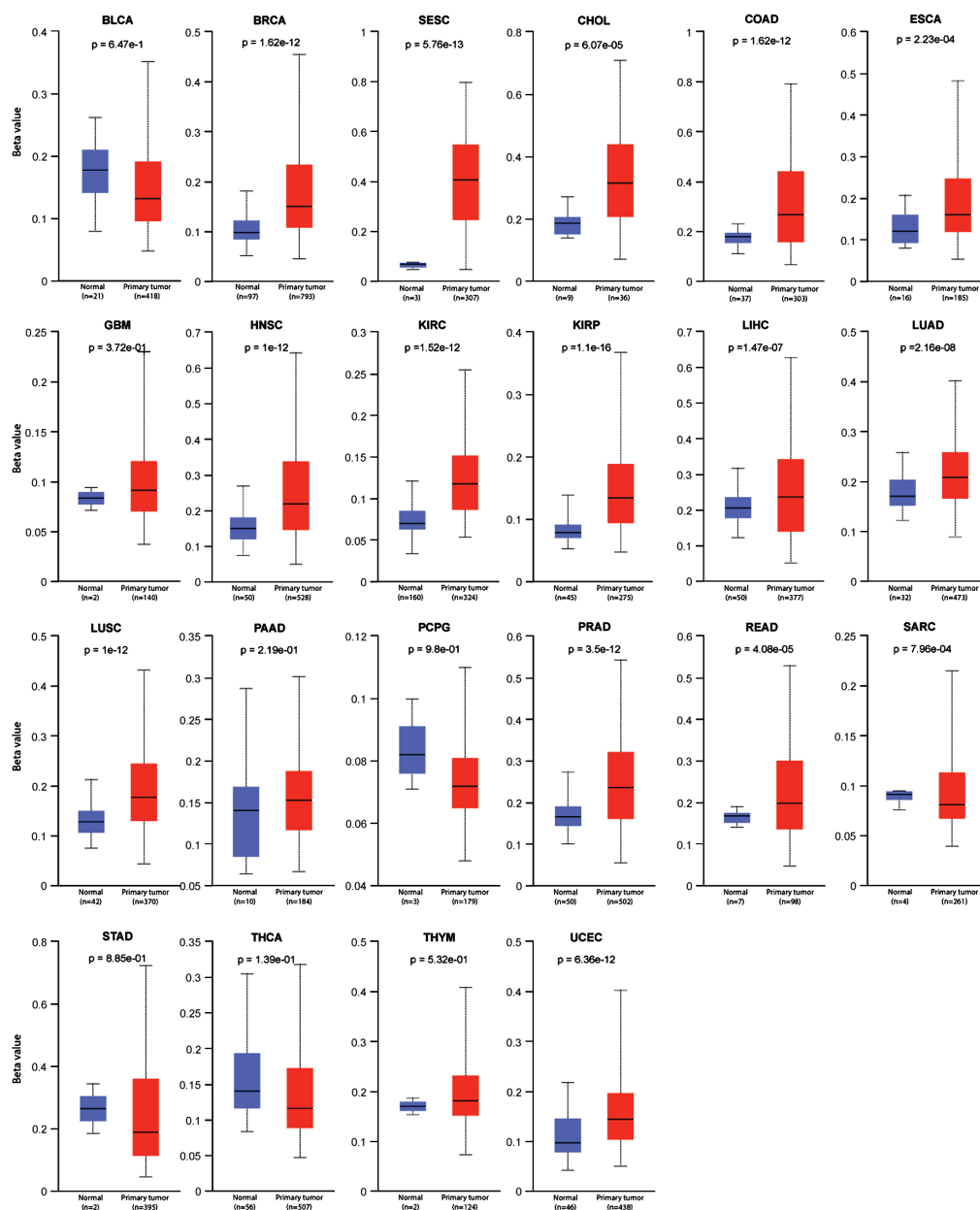


Figure S3. OS data in all TMRSS2 abnormally expressed malignancies. OS, overall survival; TMRSS2, transmembrane protease serine 2.

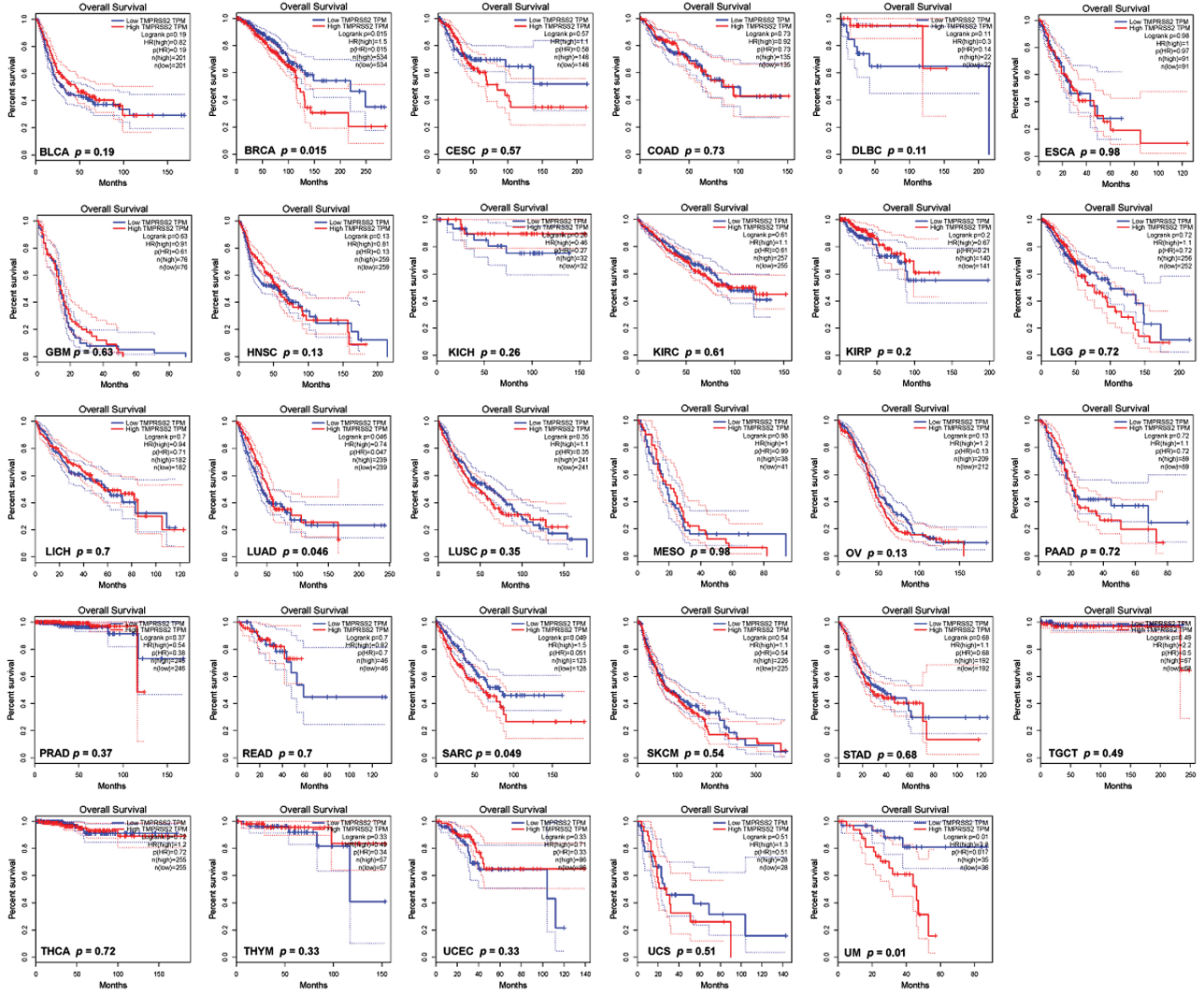


Figure S4. DNA methylation aberration of TMPRSS2 in all tumours. TMPRSS2, transmembrane protease serine 2.

