**Supplementary Figure S1.**
**A,** Das et al. cohort showing *TTP* mRNA levels at prostatectomy in patient-matched normal adjacent tissues versus primary prostate tumor tissues (***p < .001, paired t-test). **B,** Kaplan-Meier curve showing the time to BCR in low-grade primary prostate cancer patients with PSA ≤10 ng/ml from the Das et al. cohort separated into *TTP*-High and *TTP*-Low subtypes based on the median *TTP* RNA level (p-value, Mantel-Cox log-rank test). **C** and **D,** Kaplan-Meier curves showing the time to BCR in primary prostate cancer patients separated into *TTP*-High and *TTP*-Low subtypes based on median *TTP* RNA levels for pathologically high-grade patients from the Taylor et al. cohort (**C**) and the Das et al. cohort (**D**) (p-values, Mantel-Cox log-rank test).