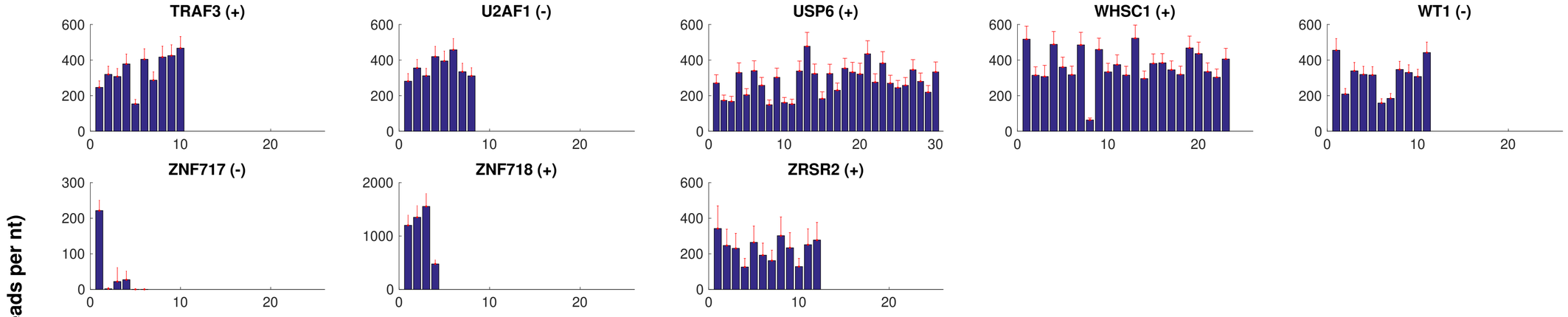
TRAF3 (+)
U2AF1 (-)
USP6 (+)
WHSC1 (+)
WT1 (-)
ZNF717 (-)
ZNF718 (+)
ZRSR2 (+)
Standardised coverage (reads per nt)
Captured coding exons are ranked as per target bed file with orientation reversed for genes on the (-) genomic strand