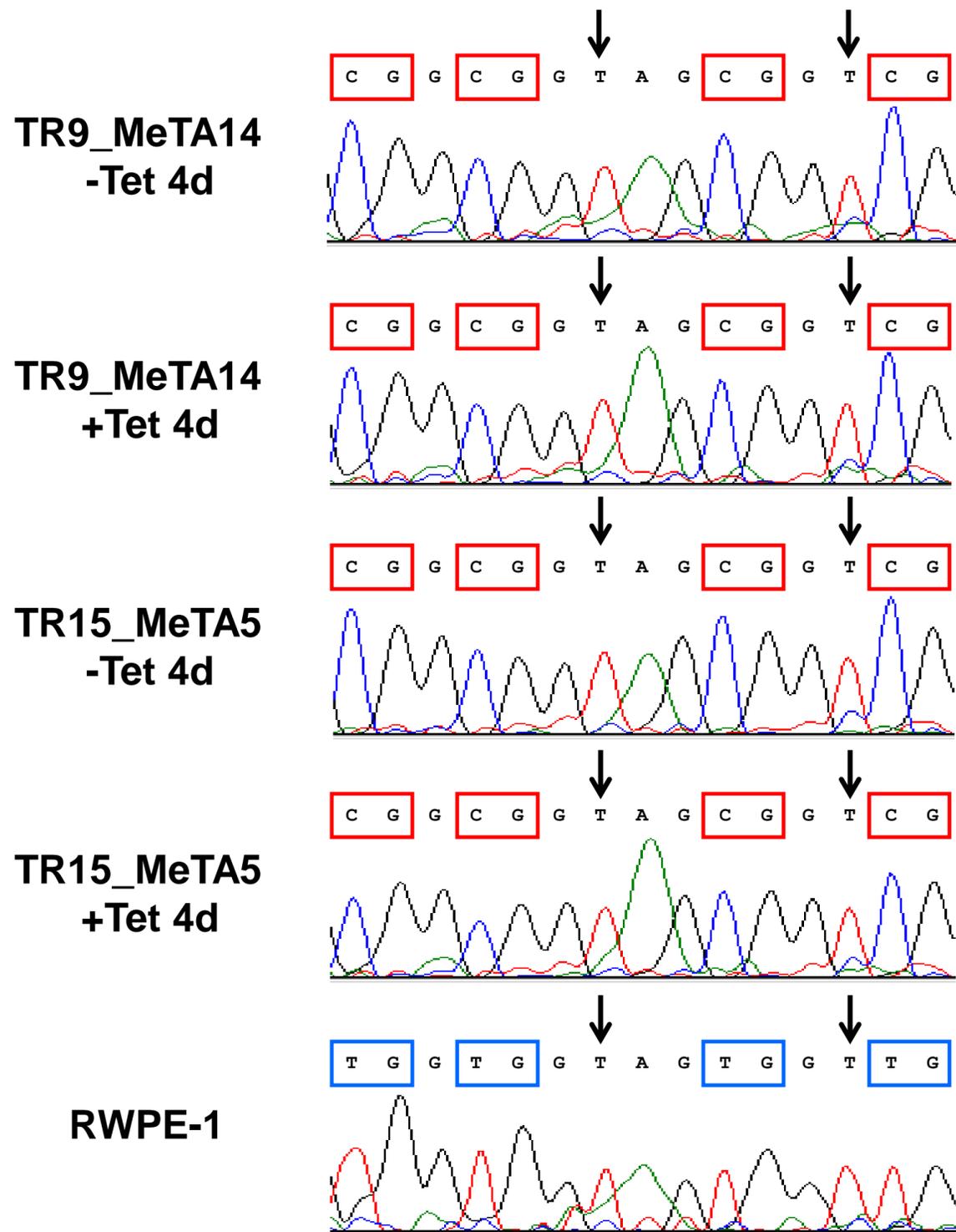


PYCARD



TNFRSF25

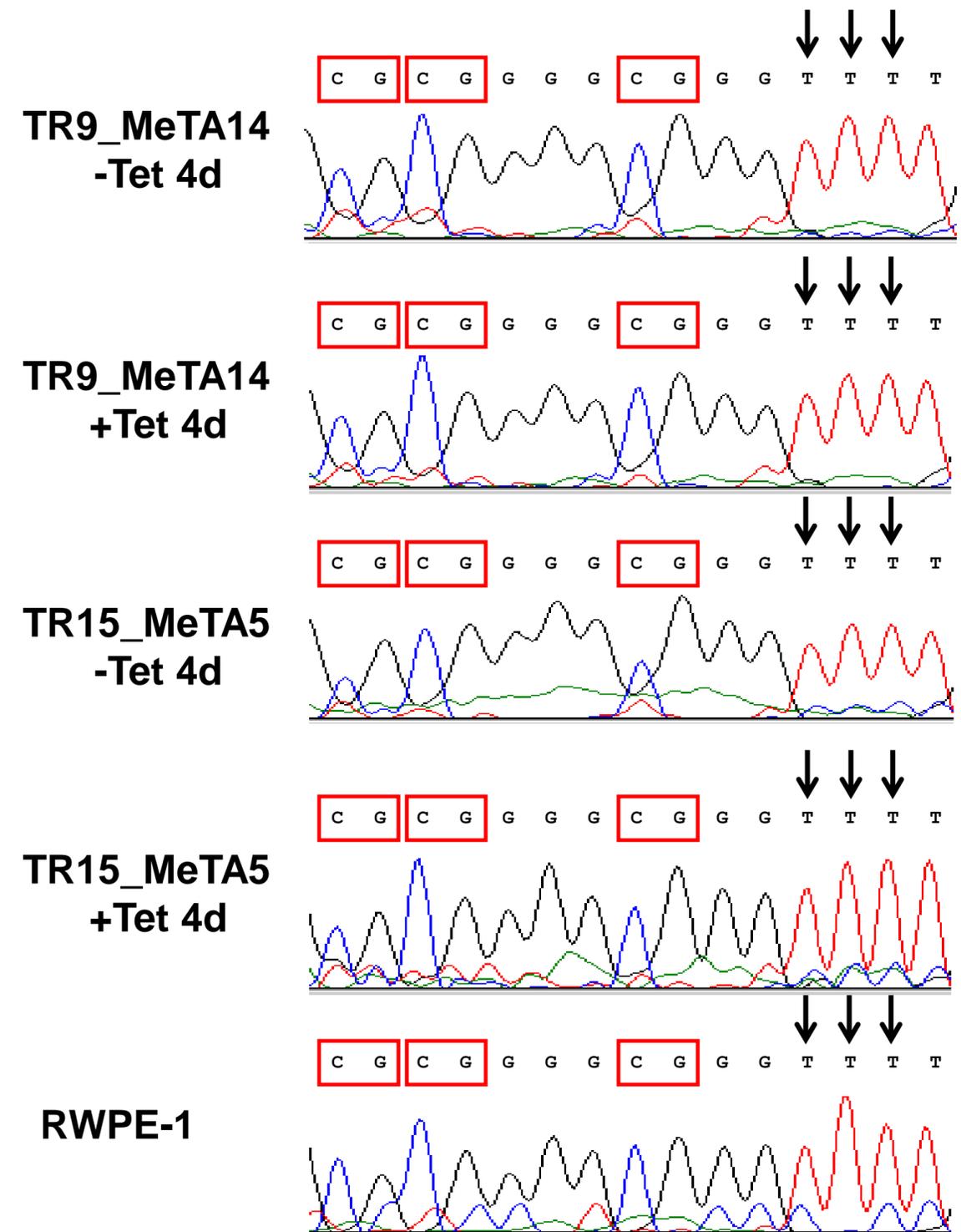


Figure S2. Representative bisulfite sequencing results of *PYCARD* (left) and *TNFRSF25* (right) promoter regions. *PYCARD* promoter region was highly methylated in LNCaP-derived prostate cancer cells but not in RWPE-1 normal prostate epithelial cells. Genomic DNAs were treated with sodium bisulfite and the promoter CGIs of *PYCARD* were sequenced. *TNFRSF25* promoter region was highly methylated both in LNCaP-derived prostate cancer cells and in RWPE-1 normal prostate epithelial cells. Arrows indicate the positions of cytosine residue converted by sodium bisulfite. Methylated CpG sites are boxed with red, and unmethylated CpG sites are boxed with blue.