Supplemental Fig. S2. Repair of 7meG lesions shows no obvious strand bias among yeast genes in WT cells. Fraction of 7meG lesions remaining following 2hr repair in WT cells relative to a matched 0hr mag1Δ control. Data is shown for DNA regions surrounding the transcription start site (TSS) of 5762 yeast genes. The transcribed strand (TS) is depicted in blue and the non-transcribed strand (NTS) is depicted in black.