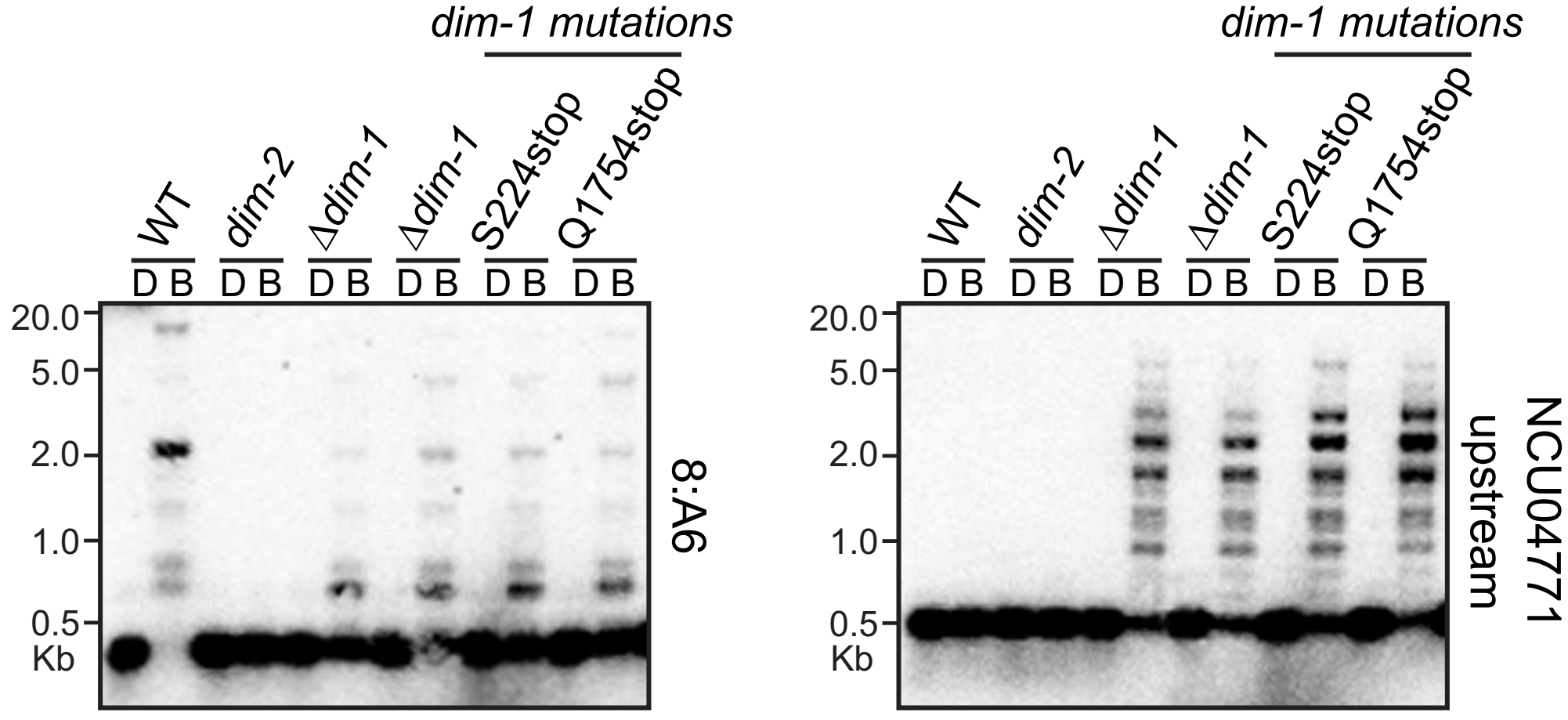

Figure S1: Aberrant cytosine methylation in* dim-1 *strains. Southern hybridizations of genomic DNA from the indicated strains digested with either DpnII (D) or its 5^{m}C-sensitive isoschizomer BfuCI (B) and probed for a constitutive heterochromatic region (8:A6, top)(Selker et al. 2003) or an intergenic region (NCU04771 upstream, bottom), as in Figures 1A, 2B.