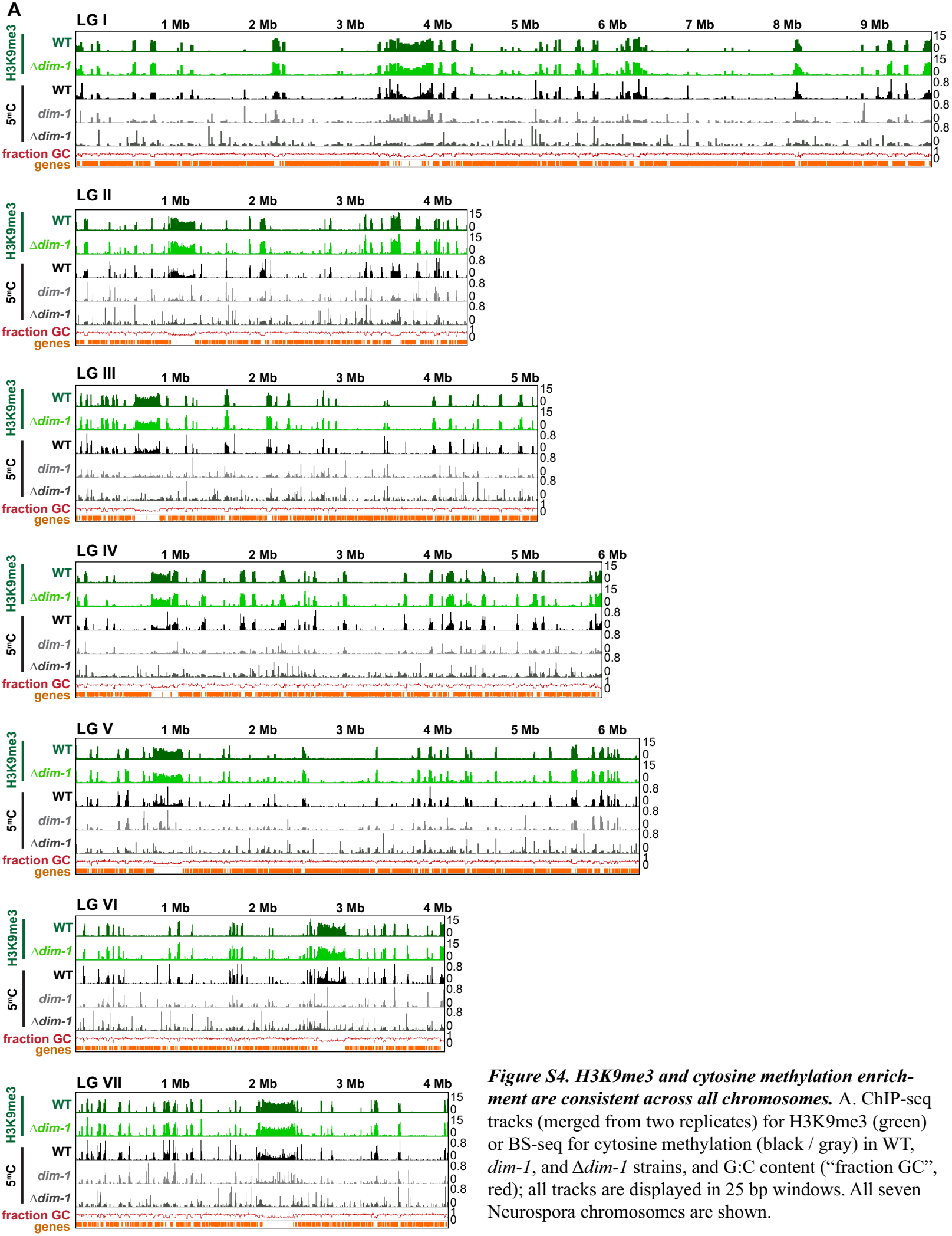

Figure S4. H3K9me3 and cytosine methylation enrichment are consistent across all chromosomes. A. ChIP-seq tracks (merged from two replicates) for H3K9me3 (green) or BS-seq for cytosine methylation (black / gray) in WT, *dim-1*, and Δ*dim-1* strains, and G:C content ("fraction GC", red); all tracks are displayed in 25 bp windows. All seven Neurospora chromosomes are shown.