

Genome aberrations revealed by coverage depth changes

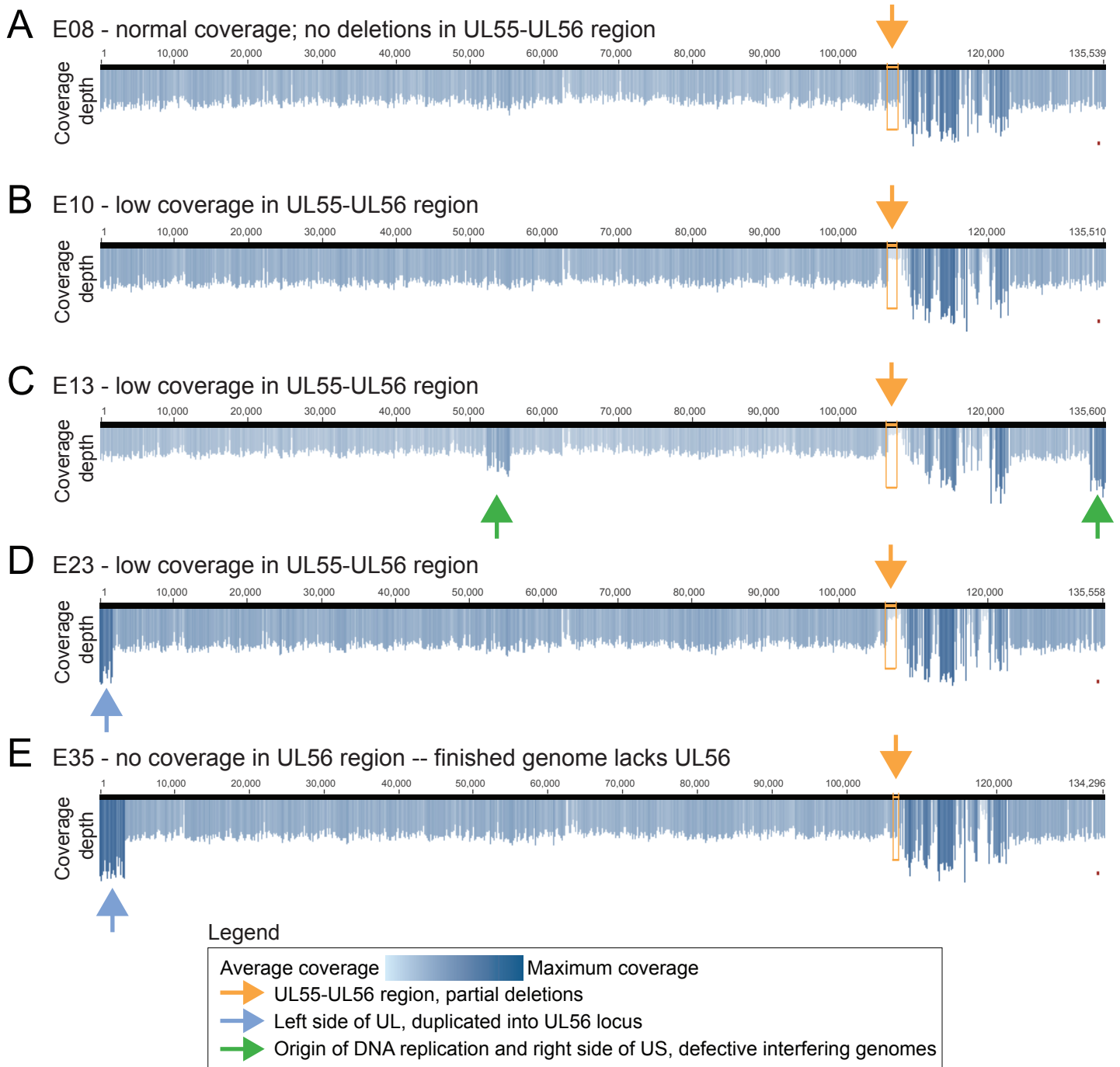


Figure S2. Read coverage scan from four HSV-1 strains depicting likely deletions in the UL55-UL56 region.