Duplication percentage (R1 trimmed) H3K27ac H3K27me3 60 -40 histone\_mark H3K27ac H3K27me3 H3K36me3 20 **-**H3K9me3 H3K36me3 H3K9me3 ind 1 2 3 60 **-**40 -20 interaction(other\_time, trt) 144r.VEH <sup>-</sup> 24t.DOX -144r.VEH <sup>-</sup> 24t.5FU <sup>-</sup> 144r.5FU <sup>-</sup> 24r.DOX -24r.5FU 24t.VEH 144r.5FU 24t.DOX 24r.VEH 24t.5FU 24r.5FU

percent duplication