Duplication percentage (R1 no trim) VEH VEH VEH VEH 1 2 3 4 VEH 5FU 5FU 5FU 5 3 2 histone\_mark H3K27ac H3K27me3 H3K36me3 H3K9me3 5FU 5FU DOX DOX ind 4 5 2 1 2 3 5 H3K27me3 H3K36me3 H3K9me3 DOX DOX DOX H3K27ac 3 5

