



Figure S14. Nucleosome shifting and deacetylation may explain NDR disappearance (biological replicate). **(A)** Delta metagene plots showing the average read depth of the indicated modifications after subtracting the interphase depths from the mitotic depths. **(B)** A Scatter plot showing the normalized read counts at H3K9ac peaks on cells treated with TSA in mitosis versus interphase. **(C)** Metagene plot for H3K9ac data after TSA treatment. **(D)** Delta metagene plot for H3K9ac data before (dotted line) and after (solid line) TSA treatment.