



Figure S3. MNase analysis of chromatin from cells treated or not with 5FU. **(A)** Nucleosome profile after digestion of chromatin with 0, 15, 60 and 240 U/ml of MNase. Samples were separated by gel electrophoresis and quantified by densitometry. **(B)** DNA protected by nucleosomes obtained from the control or 5FU-treated cells. Values represent mean \pm standard deviation of five independent replicates. No significant differences were found between controls and 5FU-treated cells.