Supplementary Figures and Legends



Fig. S1. Confirming the COMMD4 and H2B binding pose. (a). An ELISA showing the binding of H2BWTpep and recombinant COMMD4. An EC50 of 70 nM for COMMD4 is shown. **(b)**. The binding of overexpressed FLAG-H2BWT and FLAG-H2BMUT1 with GST-COMMD4 in control and COMMD4 siRNA depleted cells. COMMD4 levels are shown, and IgG heavy chain shows the loading.



Fig. S2. Viability of NSCLC cells. (a-d). Luminescent cell viability assay of H460, H1299 and CRL5889 NSCLC cells and HBEC3-KT cells, transfected with FLAG-H2BWT, FLAG-H2BMUT1 or mock-transfected and treated with varying doses of IR is shown. Asterix (*) denotes P < 0.05, n.s; not significant. Error bars represent mean \pm SD from three independent experiments.



Fig. S3. Induction of γ H2AX foci in H460 cells. (a). Immunofluorescence staining of H460 NSCLC cells transfected with control or COMMD4 siRNA #2 and FLAG-H2BWT or FLAG-H2BMUT1, treated with 6 Gy IR followed by recovery at time points shown. (b). Plot of γ H2AX foci numbers for (a). **; *P* < 0.005. Error bars represent mean ± SD from three independent experiments where 50 cells were quantified per condition. γ H2AX is shown in green and DAPI shows the nucleus. Scale bar denotes 5 µm.



Fig. S4. Induction of γ H2AX foci in CRL5889 cells. (a). Immunofluorescence staining of CRL5889 NSCLC cells transfected with control or COMMD4 siRNA #2 and FLAG-H2BWT or FLAG-H2BMUT1, treated with 6 Gy IR followed by recovery at time points shown. (b). Plot of γ H2AX foci numbers for (a). n.s; not significant and *; *P* < 0.05. Error bars represent mean ± SD from three independent experiments where 50 cells were quantified per condition. γ H2AX is shown in green and DAPI shows the nucleus. Scale bar denotes 5 µm.