Figure S5. Binding of hnRNP C to transcripts of HR genes. A. Genome browser view of *PALB2* and *BARD1* genes displaying RNA-Seq data (overlapping reads per nucleotide; blue) from control and hnRNP C knockdown HeLa cells, that were independently transfected with two different siRNAs (KD1 and KD2), as well as hnRNP C iCLIP data (crosslink events per nucleotide; purple). RefSeq transcript annotations (blue) and *Alu* elements in antisense orientation to the shown strand (orange) are depicted below. No *Alu* exonization events were found in these two genes. B. “Weblogo” showing the base composition at the hnRNP C crosslink sites (position 0) within *BRCA1, BRCA2, PALB2, RAD51, BARD1* and *BRIP1* gene transcripts as well as the surrounding sequence. The y-axis indicates the informational content for each position in bits. The graph shows the aggregate of all the crosslink sites in the 6 genes.