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**Figure S4 - Read alignments of multiples of mRNA-seq samples visualised in IGB.** Alignments from different RNA-seq experiments were obtained from untreated, control and SNW1 siRNA treated HeLa and mSnw1-LAP cells, and visualised at the sororin locus. I and II indicate two biological replicates in the same sequencing run, 50 and 100 bp indicate 50 bp single-end and 100 bp paired-end sequencing, respectively.