Figure S7: An example of FOXA1 deletion detected by RNA-seq but missed by WES due to low coverage in DNA sequencing. A 95bp deletion is detected in exon 2 of FOXA1 in SU2C subject 1115161 at chr14:38,061,136-38,061,230 (hg19 coordinate) by RNA-seq with 33% variant allele fraction. This deleted region is covered by only 9 reads in whole exome DNA-seq data.