

Figure S2. The dynamically generated bubble chart from Seten WI, showing the comparison of significantly enriched MalaCards Disease Ontology terms for DDX6 in K562 and HepG2 cell lines. Only gene sets which had more than 5% of the total genes and exhibited a minimal p-value of 0.05 in one of the cell lines are included in this comparison. The radius of bubbles is computed as negative Log10(corresponding p-value).

