Figure 7-figure supplement 1

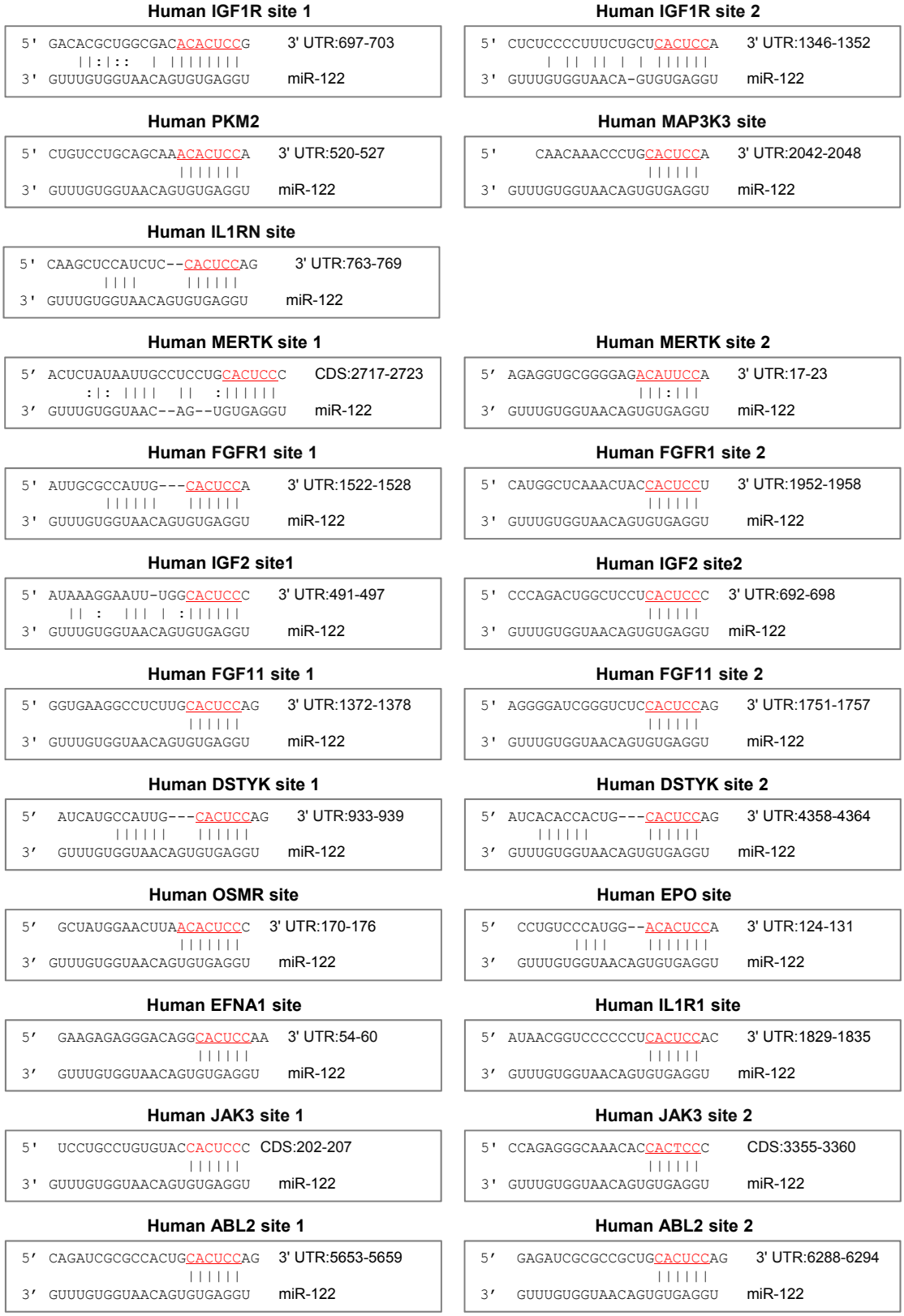

Figure 7-figure supplement 1. miR-122 binding sites in the 3′UTR or the CDS of candidate genes, predicted by TargetScan or RegRNA. The binding sites of IGF1R (S1 and S2), PKM2 and MAP3K3 have been reported to be functional.