

Target gene		Free energy	Go Function	Go Process
<i>Sec23b</i>	target 5' A UUU G G 3' GUG GAAU AUACCCU CGC CUUA UAUGGGGA tRF 3' U G 5'	-25,3	binding: protein	developmental process: neurocranium morphogenesis; cartilage development
<i>Ptpa</i>	target 5' A A CUC U 3' GC GG AAUCAUACCCU CG UC UUAGUAUGGGGA tRF 3' C 5'	-24,6	catalytic activity	cellular process: metabolic
<i>Myst3</i>	target 5' A U A A 3' GC GAGAG CGUACCC CG CUCUU GUAUGGGG tRF 3' A A 5'	-26,7	binding: DNA	developmental process: embryonic patterning; cartilage and skeletal development
<i>Parva</i>	target 5' U U UU A 3' GC G AAUUAUACCCU CG C UUAGUAUGGGGA tRF 3' UC 5'	-23,8	unknown	unknown
<i>Exoc6</i>	target 5' A CA G 3' GUGAGG AUACCUU CGCUCU UAUGGGG tRF 3' UAG A 5'	-19,2	unknown	cellular process: exocytosis
<i>Pgm1</i>	target 5' C UU A 3' G UCAUACCC UC AGUAUGGGG tRF 3' CGC UU A 5'	-23,1	catalytic activity	Cellular process: metabolic