

**Table 1 – miR-181 targets identification in RGC *Xenopus laevis* isolated axons**

Table 1: Predicted miR-181-5p targets, detectable within the axonal transcriptome, ranked by TotalScore. The final candidate selection is based on the 20% best ranked candidates which were conserved as targets in human and mice, filtered by the Reactome axon guidance pathway (R-HSA-422475) and the Reactome integrin cell surface interactions pathway (R-HSA-216083).

Column header	Comment
TCS	Total Context Score from miR-181a-5p targets predicted using TargetScan 6
8mer-1a	Type of miR-181-5p MRE (miRNA responsive element)
7mer-m8	
7mer-1a	
TS.a-5p	Total sites of miR-181a-5p MREs in <i>Xenopus</i>
TS.a-1-3p	Total sites of miR-181a-1-3p MREs in <i>Xenopus</i>
TS.a-2-3p	Total sites of miR-181a-2-3p MREs in <i>Xenopus</i>
TS.miR	Total sites of miR-181a MREs in <i>Xenopus</i>
TS.3p	Total sites of miR-181a-3p MREs in <i>Xenopus</i>
h-TS-miR	Human total number of miR-181 MREs
h-TS-5p	Human total number of miR-181-5p MREs
h-TS-3p	Human total number of miR-181-3p MREs
tc-h-ID	miR-181 TargetScantargeting conserved in human based on entrezID > yes or no
tc-h-Gn	miR-181 TargetScantargeting conserved in human based on gene name > yes or no
m-TS-miR	Mice total number of miR-181 MREs
m-TS-5p	Mice total number of miR-181-5p MREs
m-TS-3p	Mice total number of miR-181-3p MREs
tc-m-ID	miR-181 TargetScantargeting conserved in mice based on entrezID > yes or no
tc-m-Gn	miR-181 TargetScantargeting conserved in mice based on gene name > yes or no
Integrin_Reactome	Transcripts reported in the Reactome integrin cell surface interactions pathway (R-HSA-216083) >yes (if it is reported in the selected Reactome pathway)
Axon-guidance_Reactome	Transcripts reported in the Reactome axon guidance pathway (R-HSA-422475) >yes (if it is reported in the selected Reactome pathway)













