ESM Table 1.

Predicted miR-153 targets	Targetscan Context+ score	Targetscan aggregate P_{CT}	Brain expression/function [reference]	Pancreas expression/function [reference]
BSN	-0,05	0.85	[1]	[2]
PCLO	-0,10	0,89	[3]	[4]
SNAP25	-0,14	0,78	[5]	[6]
SNCA	-0,18	0,85	[7]	[8]
SYT1	>-0,02	< 0,1	[9]	[10]
SYT4	-0,04	0,73	[11]	[12]
VAMP2	-0,26	0.93	[13]	[14]

References to ESM Table 1.

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ESM Table 1. **Candidate miR-153 target genes that correlate with IA-2 beta function at sites of IA-2beta expression**. The first column shows the names of the predicted miR-153 targets. Targetscan total context+ scores (looking at the context of the sites within the UTR; the lower the score, the better) and aggregate P_{CT} (Probability conserved targeting) preferential conservation of the sites; score between 0 and 1.0; the higher the score, the better) are shown in the second and third column, respectively. References that demonstrated correlation between expression and/or function of IA-2beta and target genes in brain and pancreas are shown in the fourth and fifth column, respectively. The list of relevant references is shown below the table.