

NEUROBIOLOGY. For the article “Identification of genes expressed with temporal-spatial restriction to developing cerebellar neuron precursors by a functional genomic approach,” by Qing Zhao, Alvin Kho, Anna Marie Kenney, Dong-in Yuk, Isaac Kohane,

and David H. Rowitch, which appeared in number 8, April 16, 2002, of *Proc. Natl. Acad. Sci. USA* (99, 5704–5709), column headings in Table 1 did not appear correctly due to a printer’s error. The corrected table appears below.

Table 1. Expression of genes/ESTs *in vivo* in the neonatal mouse cerebellum

Genes	Accession no.	PN 1	PN 7				PN 15			
		EGL	EGLa	EGLb	PK	IGL	EGLa	EGLb	PK	IGL
Training set: EGL expression—positive*										
CycD1	W08016	+++	+++	–	–	–	–	–	–	–
STK1	D21099	+++	++	++	–	–	–	–	–	–
Cdc20	AA00468	+++	+++	+++	–	–	+	+	+	+
CKS2	AA11263	++	++	–	–	–	–	–	–	–
AYK1	U80932	+++	++	–	–	–	+	+	–	–
IPL1	AA050055	+	++	++	–	–	–	–	–	–
SET	W78604	+++	+++	+++	–	–	+	+	+	+
BM28	AF004105	+++	+++	–	–	–	–	–	–	–
RanBP1	X56045	+++	+++	–	–	–	–	–	–	–
<u>DP1</u>	X72310	++	+++	++	–	–	–	–	–	++
Training set: EGL expression—negative*										
Stra13	Y07836	–	–	–	–	–	–	–	–	++
EST69 [†]	C76791	–	–	+/-	++	+	–	–	–	–
Test set: EGL expression—positive [‡]										
E2F1	L21973	+	++	++	–	–	–	–	–	++
RNM2	M14233	++	+++	+++	–	–	–	–	–	–
CycB2	X66032	+++	++	++	–	–	–	–	–	–
N-myc	X03919	+++	+++	+++	–	–	–	–	–	–
CycA2	Z26580	+++	+++	+++	–	–	–	–	–	–
CycB1	AA426917	++	+++	+++	–	–	–	–	–	–
mMIS5	AA689977	+++	+++	+++	–	–	–	–	–	–
<u>EIF4A</u>	AA166088	++	+++	+++	+	+	–	–	–	–
<u>HMG1</u>	Z11997	++	+++	+++	–	–	–	–	–	–
PAL31	W48027	++	+++	+++	+	+	–	–	–	–
Test set: EGL expression—negative [‡]										
AA84	AA688784	–	+	+	–	–	–	–	–	–
AA96	AA267296	–	+	+	–	–	–	–	–	–
FKBP-13 [†]	AA072278	–	+/-	+/-	–	–	–	–	–	–
BC73	AA16900	–	+/-	+/-	+	–	–	–	–	–
RCC	AA051583	–	+	+	–	–	–	–	–	–
AA88	AA275288	–	+	+	–	–	–	–	–	–
QK15A	AA174970	–	–	–	+++	+++	–	–	–	–
FIN-14	U42386	–	–	–	++	++	–	–	–	–

PK, Purkinje cell; IGL, internal granule layer; EGLa, actively proliferating granule cells; EGLb, granule cells that recently have left the cell cycle.

*A training set of 12 genes was tested for EGL-specific expression in PN 1, 7, and 15 mouse cerebella by ISH. Ten of these (*CycD1*, *STK1*, *CDC20*, *CKS2*, *AYK1*, *IPL1*, *SET*, *BM28*, *RanBP1*, and *DP1*) showed expression in the PN 7 EGL in contrast to *Stra13* and *EST69*.

[†]Entries identified by the TRP-EGL that failed to show expression in the EGL *in vivo* (false positives). False negatives, not identified by the TRP-EGL but having EGL-specific expression, are underlined.

[‡]Test data set expression *in vivo*. Of an additional 18 genes tested, 10 were expressed in the PN 7 EGL, and 8 genes were not expressed specifically in the EGL. The relative intensity of expression is shown as strong (+++), moderate (++) or weak (+) or absent (–).