

Table 5. Probes for *in situ* hybridization

Gene name	NCBI accession no.	pBSA #	Insert	Species
<i>CNTNAP2</i>	NM_014141	2	bps 548-4300	Human
<i>CNTNAP2</i>	NM_014141	13	bps 3550-4100	Human
<i>CNTNAP2</i>	NM_014141.4	83	bps 2318-1867	Human
<i>Cntnap2</i>	52138537	98	bps 191-667	Mouse
<i>Cntnap2</i>	52138537	57	bps 3600-4900	Mouse
<i>COL1A1</i>	NM_000088	61	bps 1 - 4748	Human
<i>CD19</i>	BC006338	82	bps 1826-1384	Human
<i>CG033</i>	NP_660347.1	92	bps 594-1049	Human
<i>GLIS3</i>	NM_152629.2	87	bps 2219-1734	Human
<i>HOXB13</i>	AY937237	1	bps 32-824	Human
<i>LDB1</i>	BC028170.1	4	bps 46-623	Human
<i>MAB21L1</i>	BC028170	22	bps 75-664	Human
<i>MGC13024</i>	NM_152288	25	bp 802-1305	Human
<i>NR4A2</i>	BC009288	3	bps 4-2175	Human
<i>NR4A2</i>	NM_173172	52	bps 2700-3300	Human
<i>Nr4a2</i>	NM_013613	55	bps 1950-2450	Mouse
<i>P15-2</i>	NM_018698	29	bps 208-847	Human
<i>PCDH17</i>	NM_014459.2	100	bps 4025-4566	Human
<i>PCDH17</i>	NM_014459.2	86	bps 3374-3906 *	Human
<i>PIASXB</i>	AF077954.1	8	bps 79-599	Human
<i>TTR</i>	BC020791	14	bps 27-616	Human

*An additional 80 bp (below), presumably representing a splice variant, were present between base pairs 3541-3542 of NM_014459.2.

>Additional sequence within cloned cDNA insert.

tgccacagatgaatgccgagtgcttggctcattctgacaggtgctggatgccacagttcctgcagccaatcaggctgaaaatgc