

**Table S3. DNA sequence flanking deletions in RT-PCR products**

No.	5' sequence	del	3' sequence
<b>from Kanadia and Cepko (2010)</b>			
ECO	... ACTGACTGCAC   GTGAGTCGCTCCGTC	[ 639 ]	TGTTTTGTAG   GTTTCATGAGTGGACATC ...
InC	... CCCTCCGGCGGGAGCT   CGCGGCGCGC	[ 71 ]	CGCGGCGCGCAGGCGTCTGGCGGCCAAC ...
E12P20	... TCGCGGCGCGCCCCCGT   GCGCGGGCG	[ 85 ]	GCGCGAGCGGCGCCGCATGCAGGGGCTG ...
32	... TCGCGGCGCGCCCCCG   TCGCGGGCG	[ 134 ]	TCGCGAGGTTGGTGCCGCAGTGGGGCCA ...
9	... GGCCCTCCGGCGGGAGCTCGC   GGC GC	[ 122 ]	GG   GGC TGAACACGGCGTTCGACCGGCTG ...
<b>from E14.5 retina RT-PCR (<math>\leq 2X</math> MasterAmp™, 40 cycles)</b>			
1	... AAGTCGGCCTGCAAACCCAC   GGCC	[ 70 ]	GGCCGGGCGGCTGGAGAGCGCGGCGCG ...
2*	... ATGAAGTCGGCCTGCAA   ACCCCACG	[ 235 ]	CTGCAG   ATGGCGCTCAGCTACATCATCG ...
<b>from RT-PCR of IVT RNA (<math>\leq 2X</math> MasterAmp™)</b>			
3	... CGTGGATGAAGTCGGCCTG   CAAACC	[ 746 ]	TAAATCC TAGCGTCATTCAGGAGGTGGA ...
4	... GAAAGCTTTCTAT   CCCC GACCC	[ 605 ]	CCCC TACCTCCCTTTCCGGGTGCTAG ...
5	... CTCCGGCGGGAGCTC   GCGGCGCGC	[ 106 ]	GCGGCGCGCATGCAGGGGCTGAACAC ...
6	... CTCCGGCGGGAGCTC   GCGGCGCGC	[ 106 ]	GCGGCGCGCATGCAGGGGCTGAACAC ...
7	... CTCCGGCGGGAGCTC   GCGGCGCGC	[ 70 ]	GCGGCGCGCAGGCGTCTGGCGGCCAA ...
8	... CGCGCCCCCGT GCGCGG   GCGCAGCC	[ 85 ]	GCGCGA   GCGGC GCCGCATGCAGGGGCT ...
9	... CTCGCGGCGCGCCCCCG   TCGCGGGC	[ 103 ]	TCGAGGGCTGAACACGGCGTTCGACC ...
10	... CCCACGGCCCTCCG GCGG   GAGCTCG	[ 110 ]	GCGC   GAGCGGCGCCGCATGCAGGGGCT ...
11	... CCCTCCGGCGGGAGCT   CGCGGCGCGC	[ 71 ]	CGCGGCGCGCAGGCGTCTGGCGGCCAA ...
12	... TGCAAACCCACGGCCCT   CCGGC GG	[ 155 ]	CCGGC TCGCAGGGTGGTGCCGCAGTG ...

Spurious RT-PCR products are numbered to match Figure 5. The highlighted text shows areas of micro-homology, which can promote RT template switching. The size of the deletion [bp] and breakpoints (|) are indicated.

\*Product is likely due to mis-priming of LP6.