Table S2. PCR conditions in this study

	<u>Primers</u>					
Experiment	FOR	REV	# cyclest	Product	Size	Figure
RT-PCR (gPCR)	LP8	LP4	35 [§]	Math5 full length ECO	1087 448	3a, 3b, 4b
RT-PCR (gPCR)	LP6	LP7	35 §		486	3c, 3d, 4c
RT-PCR	LP13	LP4	35	unspliced Cb splice	567 368	Suppl 3d
RT-PCR	LP15	LP4*	33	Cb splice	228	Suppl 3e
3'RACE pA1	LP5 LP10	UAP UAP	15 20	initial RACE nested RACE	>591‡ >236‡	2đ
3'RACE pA6	LP11 LP12	UAP UAP	15 20	initial RACE nested RACE	>379‡ >342‡	2đ
Triplex RT-PCR [¶]	LP8 LP13	LP4*	33	unspliced ECO	567 448	6b, 6c
Triplex RT-PCR [¶]	LP8 LP14	LP4*	33 33	unspliced ECO	301 448	6b, 6c
Triplex RT-PCR [¶]	LP14 LP15	LP4*	33	unspliced Cb splice	301 228	Suppl 3e

Notes:

^{*} End-labeled with 6-FAM (carboxyfluorescein)

tAll PCRs had an initial denaturation step (94°C x 3 min); followed by # cycles of 94°C x 30 sec denaturation, 57°C x 45 sec annealing, and 72°C x 60-70 sec extension; plus a final extension step (72°C x 7 min)

 $[\]S$ To generate deletion products, 40 cycles were used in these reactions, with basic Taq polymerase (Invitrogen cat. 10342)

[‡]Products include a variable polyA tract

 $[\]P$ Triplex PCRs utilized 3 primers, with 2 FOR primers at 0.1 μM each and 1 REV primer at 0.2 μM .