

Transcript id	Type	Length (nt)	Other transcript ID	Distance (bp)
STRG.1411.1	Antisense	2948	SPAC4H3.02c	358
STRG.2665.1	Intergenic	639	SPBC2F12.05c	123
STRG.2999.1	Antisense	998	SPBC6B1.07	198
STRG.4201.1	Antisense	1675	SPCC584.13	177

Table S4. Novel transcripts predicted to have originated from bidirectional promoters. The transcripts are located in antisense orientation to other transcripts and the distance between the two transcription start sites is less than 400 bp.