

| 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.