**Table S1**

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| --- | --- | --- | --- |
| ***Gene*** | **Primer labeling in** **Fig 2F** & **Fig S3C** | **Sequence (5’ to 3’)** | **Putative Su(H) binding sites covered** |
| *E(spl)m8*-*HLH* | *E(spl)m8* | F: GGAGCGGAGCGAAGCATGTGR: GCGTGGGAACCGAGCTGAAA | GTGAGAA |
| *bantam*  | *bantam S1* | F: GTTCTTCGCTTCTCTGTGGTCTR: ACCGGTTTCGTAGTCAAATCAT | ATGTGAG |
| *bantam*  | *bantam S2* | F: ATTGGAGAAATACGGGCGCR: TATTTTCACACGCGCCGC  | GTGTGAA |
| *bantam* | *bantam con* | F: CCAATGCAATTGGAGAAATACGGR: ACACAATTAGTGGCGCTGAG | - |
| *RP49 (RpL32)* | *rp49* | F: CCCTGATAAGGATTGATGCTGR: CAGACCATTGCGGTCTTTCT | - |