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PVKW010072700.1
PVKW010000146.1

[illegible]

[illegible]

| | Copy number | Proportion |
|------------------------------------|-------------|------------|
| T ⁺ copies | 22 | 30 % |
| Conventional T ⁺ copies | 6 | 8% |
| T ⁻ copies | 45 | 62% |
| Total | 73 | |

Figure S11. Random Dip_b2 copies from the *A. bullata* genome. Nucleotide sequences are highlighted as follows: PASS, strong pol III terminators, medium terminators, rudimentary terminators, and oligo(A). Conventional T⁺ copies do not contain either a terminator or an oligo(A).