

Appendix Table 3. Recombination estimates

	γ / base pair					
	<i>al</i>	<i>bab2</i>	<i>Gpdh</i>	<i>ple</i>	<i>Sod</i>	<i>up</i>
<i>D. malerkotliana</i>	0.035	0.052	0.107	0.031	0.063	0.020
<i>D. bipectinata</i>	NE	0	0.159	0.022	0.116	NE
<i>D. parabiptinata</i>	NE	NE	0	0	0.080	NE
<i>D. pseudoananassae</i>	NE	NE	0.049	NE	0.028	NE
<i>D. malerkotliana</i> + <i>D. bipectinata</i>	0.006	0.051	0.052	0.025	0.080	0.040
<i>D. bipectinata</i> + <i>D. parabiptinata</i>	0	0.020	0.071	0.010	0.075	NE
<i>D. malerkotliana</i> + <i>D. bipectinata</i> + <i>D. parabiptinata</i>	0.003	0.029	0.065	0.016	0.093	0.056
	Minimum number of recombination events					
	<i>al</i>	<i>bab2</i>	<i>Gpdh</i>	<i>ple</i>	<i>Sod</i>	<i>up</i>
<i>D. malerkotliana</i>	1 530	8 2237	7 1125	3 961	10 1037	1 549
<i>D. bipectinata</i>	0 518	1 2245	8 1116	1 960	12 1001	0 553
<i>D. parabiptinata</i>	0 517	0 2247	0 1123	0 959	3 1020	0 553
<i>D. pseudoananassae</i>	0 526	0 2252	2 1107	0 961	1 667	0 476
<i>D. malerkotliana</i> + <i>D. bipectinata</i>	1 530	9 2275	12 1176	3 968	17 1058	1 583
<i>D. bipectinata</i> + <i>D. parabiptinata</i>	0 530	6 2275	13 1176	2 968	14 1058	0 583
<i>D. malerkotliana</i> + <i>D. bipectinata</i> + <i>D. parabiptinata</i>	1 530	14 2275	16 1176	4 968	20 1058	1 583