

Supplementary Material SM17

A) Duplication breakpoints

| name | chrom | start | end | sequence (AgamP4, from Assogba 2016) | window (Lucas et al. 2019) |
|---------------------------|-------|---------|---------|---------------------------------------|----------------------------|
| upstream_breakpoint_out | 2R | 3436893 | 3436926 | TCTTATCATGTGCTTGGTGACAATTTTGTTC | 3436800 + 300bp |
| upstream_breakpoint_in | 2R | 3436927 | 3436964 | GAATCCACGGAGCGGTGCGATTTAGGGCAGGACCGAT | 3436800 + 300bp |
| downstream_breakpoint_in | 2R | 3639803 | 3639835 | ATATTTTCCCATTCTATTTTATCCGACGAAGC | 3639600 + 300bp |
| downstream_breakpoint_out | 2R | 3639836 | 3639873 | AAACCTGCAAAGTGAACAATAAGCTCTATCTGCAAC | 3639600 + 300bp |



B) Normalised coverage along the *Ace1* duplication

--- Major duplication breakpoints
--- Minor duplication (Guinea)
█ *Ace1* gene body

