Figure S4. Window-based genetic differentiation ($F_{ST}$) in quantiles ($Q$) between a lightly pigmented population (Zambia) and three melanic populations: (A) Uganda, (B) Cameroon and (C) Ethiopia at two pigmentation-associated QTLs. Dotted lines refer to boundaries of pigmentation candidate genes (cf. Table 1) with pigmentation candidate genes delimited with dashed lines: yellow ($y$), black ($b$), and CG15023 ($Pfk$). Coordinates are given in kb with respect to release 5 of the *D. melanogaster* genome.