**S8 Table.** Pairwise genetic distance ($F_{ST}$) and gene flow ($N_m$) between subpopulations of *An. balabacensis* based on the combined sequence. Values above the diagonal are for $N_m$, while values below the diagonal are for $F_{ST}$. Values marked with asterisk indicate the genetic distances between two subpopulations are significant: *p<0.05, **p<0.01, ***p<0.001.

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