

Figure S2. Sequence alignment of the ITS2 of *Anopheles algeriensis*. The five sequences shown in the alignment are representative of the *Anopheles algeriensis* samples examined for each geographic area. The colored nucleotides indicate the position of single nucleotide polymorphisms. Red asterisks positioned in correspondence with the dash (-) highlight the position of insertions or deletions in the alignment. Start end of the presumptive ITS2 region are indicated by red arrows.

