

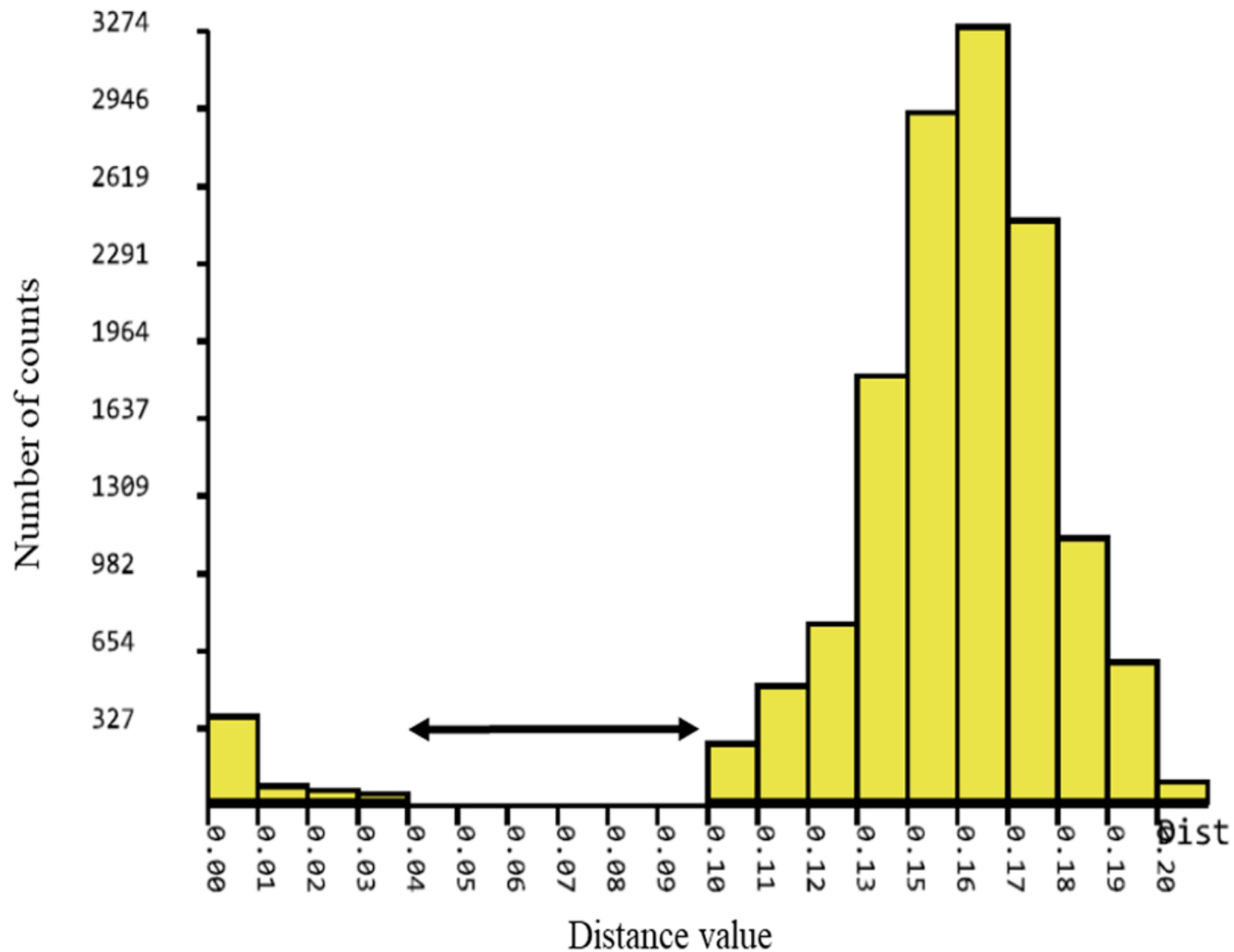


Figure S32. Histogram showing the distribution of inter- (right) and intra-specific K2P genetic distances (left) for the COI region between each pair of 169 sand fly sequences. The arrow indicates the barcode gap.