

Table 4 Maximum composite likelihood estimate of the pattern of nucleotide substitution for *P. xylostella* mitochondrial *cox1* gene

	A	T	C	G
A	-	3.22	1.25	17.45
T	2.49	-	8.72	1.24
C	2.49	22.37	-	1.24
G	35.07	3.22	1.25	-