

Table3 Abbreviation: n: number of samples; Ht: number of haplotypes; S: number of polymorphic sites; Hd: haplotype diversity; Pi: nucleotide diversity; k: average nucleotide difference. * P< 0.05;

	n	Ht	S	Hd	Pi	Tajima's	Fu and Li's D	Fu and Li's F	Fu's Fs
CF	29	27	749	0.99261	0.18248	-0.01536	0.36967	0.28470	-0.661
EEDS	26	20	264	0.96308	0.06074	-1.10881	-2.63032*	-2.51768	-0.978
SZWQ	25	23	742	0.99333	0.09745	-0.63235	-2.38827	-2.14566	1.027
HLBE	29	21	676	0.94089	0.08740	-0.38827	-1.91491	-1.66059	0.438
TOTAL	109	82	990	0.98318	0.11369	-1.67931	-3.12716*	-2.96282*	-0.688*