Table S4: Intra- and inter-specific uncorrected pair-wise nucleotide distances (*p*-dist) between six *Cryptolestes* species based on 13 concatenated mitochondrial protein coding genes (11,024bp). Draft mitogenomes of the six *Cryptolestes* species are available from Tay et al. (2024). Specimen MK614530 is the outgroup Cucujidae beetle. Intra-specific *p*-dist estimates are indicated by orange cells. Green coloured cells showed genetic distances expected at the inter-specific level between closely related species (i.e., *C. pusillus*/*C*. sp. WTT-2013; *C. ferrugineus*/*C. pusilloides*). Values in the top triangle are *p*-dist estimates from the full mtCOI gene (1,524 bp). Values in the bottom triangle are *p*-dist estimates from the concatenated 13 PCGs of 11,024bp.

	Species	1	2	3	4	5	6	7	8	9
1	MK614530	-	18.42	18.55	16.80	16.86	16.60	16.47	16.28	16.73
2	CspWTT2016-Viet	22.86	-	0.13	15.50	15.50	15.76	16.80	16.93	15.89
3	CspWTT2016-Msia	22.90	0.05	-	15.63	15.63	15.89	16.93	17.06	16.02
4	T6-1_Cpld	20.79	17.35	17.39	-	0.26	7.65	15.50	14.92	13.94
5	N4-2_Cpld	20.91	17.39	17.44	0.32	-	7.39	15.37	14.79	13.81
6	Cfer-06	20.24	17.55	17.60	6.73	6.77	-	14.66	14.4	14.07
7	WTT2013-Chn10	21.65	17.70	17.73	16.60	16.60	16.31	-	6.36	15.30
8	Cpls_T6-11	21.63	17.63	17.67	16.29	16.27	15.91	6.28	-	14.27
9	Ctur-v05	21.57	18.60	18.62	16.93	16.95	16.97	16.71	16.61	-