Table S3: Estiates of pairwise nucleotide distances between *Cryptolestes* species as assigned through the phylogenetic analysis. Within species (in traspecific) nucleotide distances (range) are as shown in the diagonal grey cells. Note that only one *C. capensis* sequence was available and therefore no range was able to be calculated. All sequences are 613bp long and sequences with missing data have been removed. Lower triangle: Inter -specific nucleotide differences (top values = range, bottom values = mean ± s.e.) estimated by the uncorrected *p*-dist method. Mean ± s.e. esti mated based on 500 bootstrap replications using MEGA7. Top triangle values are interspecific nucleotide difference based on the Kimura 2-parameter (K2P) method with gamma shape parameter = 0.2377 with four categories (mean evolutionary rates: 0, 0.06, 0.47, 3.47) and included all codon positions. Note that *C. ferrugineus*-A and *C. ferrugineus*-B are two sister clades within the *C. ferrugineus* species clade with high (97%) bootstrap confidence. The intraspecific nucleotide difference ranges fro m 0-0.359 (i.e., 0% - 3.59%; average = 2.8 ± 0.6%). Closely related species (e.g., *C. ferrugineus*/*C. pusilloides*; *C. pusillus*/*C*. sp. 'WTT-2013') exhibited an average of 7.7 ± 0.1% and 8.1 ± 0.11% nucleotide difference (boxed).

Species (no. sequences)	*C. pusillus*	*C* . sp. WTT-2013	*C. turcicus*	*C. capensis*	*C. ferrugineus* -A	*C. ferrugineus* -B	*C. pusilloides*	*C. corticinus* /*C. spartii*	*C. duplicatus*	*C* . sp. WTT-2016
C. pusillus (n=13)	0 - 0.0253 (0.011 ± 0.002)	0.087 ± 0.012	0.165 ± 0.017	0.163 ± 0.017	0.169 ± 0.017	0.173 ± 0.017	0.184 ± 0.018	0.191 ± 0.018	0.183 ± 0.019	0.185 ± 0.018
C . sp. WTT-2013 (n=9)	0.0767 - 0.0914 (0.081 ± 0.011)	0 - 0.0196 (0.009 ± 0.002)	0.185 ± 0.019	0.171 ± 0.018	0.167 ± 0.017	0.159 ± 0.017	0.186 ± 0.019	0.194 ± 0.019	0.185 ± 0.019	0.182± 0.019
C. turcicus (n=6)	0.1419 - 0.1648 (0.147 ± 0.013)	0.1566 - 0.1746 (0.163 ± 0.014)	0 - 0.031 (0.019 ± 0.003)	0.159 ± 0.018	0.159 ± 0.017	0.158 ± 0.017	0.169 ± 0.018	0.174 ± 0.017	0.170 ± 0.017	0.190 ± 0.019
C. capensis (n=1)	0.1468 - 0.1509 (0.146 ± 0.013)	0.1498 - 0.1547 (0.152 ± 0.014)	0.1501 (0.142 ± 0.014)	-	0.173 ± 0.020	0.168 ± 0.019	0.175 ± 0.019	0.173 ± 0.018	0.176 ± 0.019	0.152 ± 0.016
C. ferrugineus -A (n=14)	0.1485 - 0.1648 (0.151 ± 0.013)	0.1501 - 0.1644 (0.150 ± 0.014)	0.1552 (0.143 ± 0.014)	0.1533 - 0.1648 (0.154 ± 0.015)	0 - 0.0277 (0.011 ± 0.002)	0.028 ± 0.006	0.084 ± 0.012	0.184 ± 0.018	0.175 ± 0.019	0.173 ± 0.018
C. ferrugineus -B (n=6)	0.1436 - 0.168 (0.154 ± 0.014)	0.1436 - 0.1533 (0.143 ± 0.014)	0.1475 (0.142 ± 0.014)	0.1517 (0.150 ± 0.015)	0.0147 - 0.0359 (0.028 ± 0.006)	0 - 0.0131 (0.005 ± 0.002)	0.082 ± 0.012	0.175 ± 0.018	0.163 ± 0.018	0.172 ± 0.018
C. pusilloides (n=13)	0.1599 - 0.1778 (0.163 ± 0.014)	0.1615 - 0.179 (0.164 ± 0.015)	0.1605 (0.151 ± 0.014)	0.1452 - 0.155 (0.156 ± 0.015)	0.0767 - 0.0946 (0.079 ± 0.011)	0.0767 - 0.0901 (0.077 ± 0.010)	0 - 0.0147 (0.007 ± 0.002)	0.184 ± 0.018	0.158 ± 0.017	0.167 ± 0.018
C. corticinus /*C. spartii* (n=6)	0.1648 - 0.1794 (0.168 ± 0.014)	0.1664 - 0.1778 (0.170 ± 0.014)	0.1664 (0.155 ± 0.014)	0.155 - 0.1599 (0.154 ± 0.014)	0.155 - 0.1741 (0.162 ± 0.014)	0.1485 - 0.1631 (0.155 ± 0.014)	0.1582 - 0.1713 (0.168 ± 0.014)	0 - 0.0457 (0.035 ± 0.005)	0.162 ± 0.017	0.155 ± 0.016
C. duplicatus (n=4)	0.1599 - 0.1746 (0.162 ± 0.014)	0.1582 - 0.1688 (0.163 ± 0.014)	0.1615 (0.152 ± 0.014)	0.1599 - 0.1615 (0.157 ± 0.015)	0.1477 - 0.1656 (0.155 ± 0.014)	0.1445 - 0.1526 (0.146 ± 0.014)	0.1396 - 0.1494 (0.142 ± 0.013)	0.1436 - 0.155 (0.145 ± 0.013)	0 - 0.0049 (0.003 ± 0.001)	0.181 ± 0.019
C . sp. WTT-2016 (n=4)	0.1648 - 0.1713 (0.163 ± 0.014)	0.1582 - 0.1656 (0.160 ± 0.014)	0.1737 (0.167 ± 0.015)	0.137 - 0.1403 (0.137 ± 0.014)	0.1501 - 0.1627 (0.153 ± 0.014)	0.1533 - 0.1566 (0.152 ± 0.015)	0.1452 - 0.155 (0.149 ± 0.015)	0.1403 - 0.1485 (0.140 ± 0.014)	0.1631 (0.161 ± 0.015)	0 - 0.0082 (0.004 ± 0.002)