

Figure S12. P-distances of synonymous and non-synonymous substitutions among the Noctuids caspase-1 and -2 sequences using the Nei-Gojobori algorithm with pairwise deletion of gaps and missing data.

Nonsynonymous substitutions per nonsynonymous site (d_N)										
	Ha-C1	Hv-C1	Hs-C1	Se-C1	Mb-C1	Ha-C2	Hv-C2	Hs-C2	Se-C2	Mb-C2
Ha-C1		0.030	0.020	0.060	0.040	0.300	0.310	0.310	0.290	0.300
Hv-C1	0.271		0.030	0.070	0.050	0.300	0.310	0.310	0.300	0.300
Hs-C1	0.260	0.140		0.070	0.040	0.300	0.310	0.310	0.300	0.300
Se-C1	0.658	0.650	0.659		0.060	0.290	0.300	0.300	0.290	0.300
Mb-C1	0.532	0.538	0.528	0.564		0.290	0.300	0.300	0.290	0.300
Ha-C2	0.718	0.728	0.723	0.747	0.692		0.080	0.070	0.220	0.220
Hv-C2	0.728	0.749	0.698	0.750	0.710	0.271		0.030	0.230	0.210
Hs-C2	0.701	0.776	0.738	0.788	0.748	0.260	0.175		0.220	0.210
Se-C2	0.815	0.743	0.753	0.800	0.732	0.682	0.740	0.701		0.210
Mb-C2	0.786	0.783	0.804	0.806	0.765	0.628	0.688	0.670	0.680	
Synonymous substitutions per Synonymous site (d_S)										