

Table S2 Accession numbers of amino acid sequences of Beclin-1.

Species	Genbank No.	Identity(%)
<i>Takifugu rubripes</i>	NP_001032963.1	100.00%
<i>Epinephelus lanceolatus</i>	XP_033488150.1	95.30%
<i>Salarias fasciatus</i>	NP_001139639.1	95.75%
<i>Acanthopagrus latus</i>	XP_029953648.1	96.20%
<i>Mus musculus</i>	AAH05770.1	81.47%
<i>Loxodonta africana</i>	XP_023409488.1	81.29%
<i>Oryctolagus cuniculus</i>	XP_002719455.1	81.03%
<i>Xenopus laevis</i>	NP_001085751.1	81.21%
<i>Xenopus tropicalis</i>	NP_001029112.1	81.21%
<i>Crotalus tigris</i>	XP_039183605.1	81.03%
<i>Homo sapiens</i>	KAI4049683.1	81.11%
<i>Sceloporus undulatus</i>	XP_042327832.1	82.14%
<i>Gallus gallus</i>	NP_001006332.1	80.09%
<i>Patagioenas fasciata monilis</i>	OPJ74544.1	80.36%
<i>Mizuhopecten yessoensis</i>	OWF44040.1	64.82%
<i>Octopus sinensis</i>	XP_029649639.1	59.65%
<i>Plakobranthus ocellatus</i>	GFN82960.1	54.17%
<i>Homalodisca vitripennis</i>	KAG8287127.1	64.08%