

1 **Molecular cloning, recombinant protein expression and purification of a novel**
2 **carboxylesterase from *Liposcelis bostrychophila***

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13 **Summary**

14 Booklice are tiny insect pests commonly found in an indoor environment and characterized by
15 extremely high proliferation ability. Booklouse contamination in stored foods causes serious food
16 loss; therefore, they are recognized as so-called stored-food pests. Furthermore, since booklice in
17 house dust have been reported to be a potent allergen, it should be also considered that the
18 accidental ingestion of booklice possibly causes allergic symptom. Therefore, safe extermination of
19 booklice would contribute to the solution of both food-loss and allergic concerns.
20 Organophosphates (OPs) are compounds generally used as harmless insecticides; nonetheless, the
21 cases of OP poisoning are sometimes reported worldwide. Therefore, considering usage for stored
22 food products, OP doses should be low as far as possible. It is known that
23 carboxylesterase-mediated inactivation of OPs decreases the efficacy of OPs, suggesting that an
24 inhibitor of carboxylesterase could be a synergist to reduce the dose of OP. Herein, I report the
25 molecular cloning and characterization of a novel carboxylesterase highly expressed in a
26 representative species of indoor booklice, *Liposcelis bostrychophila*.

27
28 **Introduction**

29 Losses in the food chain from production to harvest, storage, distribution, processing, sales, and
30 consumption have recently been recognized as a problem because they not only cause possible food
31 shortages but also indirectly contribute to global warming. It was reported that global food loss and
32 waste during 2010–2016 were estimated to equal 8–10% of total anthropogenic greenhouse gas

33 emissions and cost about 1 trillion USD per year [1]. Further, it is estimated that more than 40% of
34 global food loss is "post-harvest loss" [1]; thus, countermeasures to address this issue are an urgent
35 task. Biotic post-harvest food losses include the infestation of microorganisms, pests, and rodents.
36 Pests, which include some insects and mites, occasionally infest food products, *e.g.*, grains and
37 cereals, during storage and distribution. Particularly, insect pests, including cockroaches, ants,
38 molds, warehouse beetles, and booklice (psocids), are considered a nuisance and cause huge losses
39 in the grains [2]. The amounts of stored grain lost due to insect infestation are estimated to be 5% to
40 10% in developed countries and as much as 35% in developing countries [3]. Among the insect
41 pests, booklice did not attract much attention for a long period because solid evidence as to
42 quantitative and qualitative food losses caused by them was missing. However, the status of
43 booklice in terms of food pests began to change in the late 1980s as booklouse infestation in stored
44 grains in diverse locations worldwide was reported [4]. Further, Kučerová reported a quantitative
45 data that the average weight of grain samples (broken wheat kernels) infested with booklice
46 decreased by 9.7% after 3 months of infestation [5]. These reports clearly indicate that food loss
47 caused by booklouse infection should be seriously considered.

48 Booklice are tiny insects measuring 1-4 mm in length. The proliferation of booklice is extremely
49 rapid. Therefore, once the outbreak of booklice occurs, it is extremely difficult to get rid of them.
50 Booklice include numerous species, and some of them are often found in food factories and
51 warehouses. Booklice generally prefer to feed on molds; thus, they can infest stored foods not only
52 directly but also indirectly through food-infesting molds. Among the booklouse species, *Liposcelis*
53 (*L.*) *bostrychophila*, *L. decolor*, *L. entomophila* and *L. paeta* are common food-infesting booklice.
54 Particularly, *L. bostrychophila* is a representative species of indoor booklice, which are commonly
55 found in food facilities, as well as ordinary houses; therefore, it has been studied more commonly
56 than the other species [5].

57 In addition, it has recently been reported that *L. bostrychophila* in house dusts are an allergen to
58 cause allergic asthma [6,7], suggesting that accidental ingestion of foods infested with booklice may
59 induce allergic symptoms. Although it is possible to suppress the proliferation of booklice by
60 maintaining proper temperature and humidity, providing an environment suitable for food storage is
61 sometimes difficult for various reasons such as cost issues. Thus, safe extermination of booklice
62 using less harmful insecticides would contribute to the solution of food-loss problems, which could
63 lead to the achievement of SDGs. A lot of effective and less harmful insecticides have been
64 developed so far. However, their effectiveness to booklice is not sufficient to extirpate booklice,

65 which is attributed to their extremely high proliferative capacity and the emergence of
66 insecticide-resistant booklice [8]. Of insecticides, acetylcholinesterase (AChE) inhibitors such as
67 organophosphates (OPs) have been reported to exert a relatively high anti-booklouse effect [9].
68 Therefore, the usage of high-dose OPs may potentially extirpate booklice. However, OPs should be
69 carefully used for food control, as cases of OP poisoning often occur even though OPs are
70 considered to have low toxicity to humans. In addition, as with many other insects, acquisition of
71 resistance to OPs has been reported for booklice [10]. So far, several mechanisms underlying the
72 acquisition of resistance to OPs have been documented [11,12], which includes carboxylesterase
73 (CE)-mediated OP inactivation in the insect body. Notably, Correy *et al.* showed that a specific
74 inhibitor of α -carboxylesterase (α E7), which was designed based on its protein structure,
75 significantly suppressed OP degradation, and increased the efficacy of OPs, indicating that the
76 combination of OPs and CE inhibitors possibly enables to extirpate booklice at lower insecticide
77 concentrations [13].

78 Herein, we report the molecular cloning of a novel CE expressed at a high level in *L.*
79 *bostrychophila*, designated LBCE1, and the generation of recombinant LBCE1 protein with an
80 esterase activity.

81

82 **Methods**

83

84 **Analysis of the RNA sequencing data of *L. bostrychophila***

85 Reads obtained by our RNA sequencing analysis performed previously, which are available from
86 Sequence Read Archives (NCBI) under the identifier DRR086994, were subjected to contig
87 construction as previously described [7]. The predicted open reading frame in these contigs were
88 then analyzed using the BLAST tool to identify possible homologous proteins in the NCBI protein
89 database.

90

91 **PCR-based molecular cloning of *L. bostrychophila* carboxylesterase 1 (LBCE1)**

92 Primers used to amplify a region encompassing the predicted coding sequence of LBCE1 are as
93 follows. First, total RNA extracted from *L. bostrychophila* bodies were subjected to reverse
94 transcription using a PrimeScript RT Master Mix (Perfect Real Time) (Takara-bio, Japan) and the
95 LBCE1-reverse primer. This was followed by PCR using KOD-plus2 DNA polymerase (Takara)
96 and the primers for LBCE1. The PCR products were purified, inserted in pGEM-Teasy (Promega),

97 and subjected to Sanger sequencing to verify their nucleotide sequences. This was then followed by
98 the subcloning of the PCR product into the expression vector pET22b to construct pET22b-LBCE1.

99

100 **Expression and purification of LBCE1 using an *E. coli* expression system**

101 *E. coli* cells of the strain BL21 (DE3) were transformed with pET22b-LBCE1, and the expression
102 of LBCE1 tagged with 6x His at the N-terminus was induced by culturing the cells in an
103 autoinduction medium, OvernightExpress TB medium (Sigma), for 8 h. Since the expressed protein
104 was shown to be secreted or leaked into the culture medium at a detectable level, the His-tagged
105 LBCE1 was purified from the culture medium as follows. First, 100 ml of the culture medium was
106 concentrated and buffer-changed to the equilibration buffer (20 mM phosphate, 500 mM NaCl,
107 pH7.4) using an Amicon Ultra-15 filtration unit (10-kDa cutoff) (Merck-Millipore). The concentrate
108 was then applied to an affinity column filled with TALEN Metal Affinity Resin (Takara-bio),
109 washed with the equilibration buffer, and resin-bound proteins were eluted with the same buffer
110 containing 10 mM imidazole.

111

112 **LBCE1 activity assay**

113 The purified His-tagged LBCE1 was assayed for esterase activity using *p*-nitrophenol acetic acid as
114 a substrate as described earlier [14]. The hydrolysis of the substrate was monitored by measuring
115 OD₄₅₀.

116

117 **Western blotting**

118 The purified His-tagged LBCE1 was subjected to SDS-PAGE, transferred to a nylon membrane,
119 and probed with the anti-His tag monoclonal antibody (Clone 6C4) (MBL, Tokyo, Japan).

120

121 **Results and Discussion**

122

123 **Identification of a novel carboxylesterase homologue in *L. bostrychophila***

124 First, we searched for contigs that encompass full ORFs coding for CE homologues in *L.*
125 *bostrychophilla*. Several contigs were identified to have ORFs encoding proteins that exhibit
126 significant homology with CEs registered in the NCBI database. Of these contigs, one contig
127 designated Comp59220 encompasses an ORF encoding a CE-homologous protein consisting of 555
128 amino acids (Figure 1). The Comp59220-encoded protein exhibits significant identities with CEs

129 from various insects, with the highest (47.2%) identity with EF4 esterase from *Pediculus humanus*
 130 (human louse) (Table 1); therefore, it was designated *L. bostrychophila* carboxylesterase 1 (LBCE1).
 131 Of proteins registered in the Protein Data Bank (PDB), a database for the three-dimensional

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atg cag ttc ggc tcc gac ctt cga acg agt tac aat cgc gaa aaa aaa aaa atg act gaa
M Q F G S D L R T S Y N R E K K K M T E
agc caa ccg att atc cgc atc gcg gat ggt tcg atc cga ggg gaa aaa ttg gat tca att
S Q P I I R I A D G S I R G E K L D S I
cgc gga ggt tct tat tac agc ttt aag ggg atc cct tat gcc aaa cct cct gtt ggg gat
R G G S Y Y S F K G I P Y A K P P V G D
ttg agg ttt aag gcc ccg gta ccg gtg gaa cct tgg aca ggt gta aga gat gcc tta aaa
L R F K A P V P V E P W T G V R D A L K
cat gga agc gaa gcc ccg gca aag gac atg ttg aaa cat gaa tat atg gaa aat acg agt
H G S E A P A K D M L K H E Y M E N T S
gag gat tgc ctg ttc atc aac gtc tac acg cca gaa ctt ccg aaa agc aaa aat gat cca aaa
E D C L F I N V Y T P E L P K S K N D K
ttg aaa tca gtc ctc gtc tgg gtg cac gga gga gga ttc tcc atg gga tct gga aac tct
L K S V L V W V H G G G F S M G S G N S
gaa atc tac ggc ccc gac tac acg gaa gac gtc gtc ctg gtc act ttc aac tat
E I Y G P D Y L I T E D V V L V T F N Y
cgg ttg gga gtt ttg gga ttc ctc agt ctc gga aca gtc gaa tgc ccc ggg aac ttc ggt
R L G V L G F L S L G T V E C P G N F G
ttg aag gat atg gtc ctt gcc tta aaa tgg gtt caa aag aac att gcc gct ttc ggc gga
L K D M V L A L K W V Q K N I A A F G G
gat ccg aac aac gtc acg att ttc ggt gaa agc gcg gga gga gcc gcc gtt cag tac ctt
D P N N V T I F G E S A G G A A V Q Y L
ttg att tcg aaa gcg acc aga gga ttg ttc cat aag gcc att tcc caa tcg gga acc act
L I S K A T R G L F H K A I S Q S G T T
ttg gac ccg tgg gcg cat aga ctg aat ccc aga gat ttc gcg ttt gct ttg ggg gaa gag
L D P W A H R L N P R D F A F A L G E G
ttg gga tgc aaa aca acc gac gac aaa gtg ctt ctc gac ttc ttg aaa aaa gca tcg cca
L G C K T T D D K V L L D F L K K A S P
aaa gat ttc gta gaa aaa gaa ggg gag ttg ccg aag aaa ctg tac ccc gac agg att ttt
K D F V E K E G E L P K K L Y P D R I F
ctc cag tta tcg ttt gtt ccc gta gtc gaa ccc gaa cac gaa ggg gcc ttt tta acc aaa
L Q L S F V P V V E P E H E G A F L T K
agc cca agg gaa att att caa agc ggg gat ttc aat gat gtc ccg tat att atc gga gga
S P R E I I Q S G D F N D V P Y I I G G
ggt agc ttg gaa ggc ctt att att atc tac aga aat ttc gaa tat aaa gaa tcg acg gcg
V S L E G L I I Y R N F E Y K E S T A
gat gag gat ttg gaa caa gtc ctc cct ctg gga aca tta aac att caa aag gga tcg gga
D E D L E Q V L P L G T L N I Q K G S K
gaa tcc aag gaa ata acg aag aaa att cgg gac ttt tac ttc ccc aac gga tat gag aag
E S K E I T K K I R D F Y F P N G Y E K
gag aaa cta gta gct gtt ctc tcc gcc att tat ttt ctg aac gga atc ggc aaa acc tgc
E K L V A V L S A I Y F L N G I G K T C
gat tgg atc ggc aga tta aag aac aga aat tct ccc act tat ctg tac cat ttc ctg ttc
D W I G R L K N R N S P T Y L Y H F L F
gac gga acc aag gcc ttc ctt aag cat ctt ata ggc tac ggg gat tgg aaa gga act tgc
D G T K A F L K H L I G Y G D W K G T C
cat gct gac gag ctc ggc tat ctc ttc cac atg ccc atg ctc caa gct aaa ctc gag ccg
H A D E L G Y L F H M P M L Q A K L E P
aac acc cct gaa tat acg aca gtt caa cgc atg acc aaa tta tgg acc gat ttt gcg aaa
N T P E Y T T V Q R M T K L W T D F A K
acc gga aac ccg acg ccg aag gat aac tcc tgg aaa ccg ata tct gag aat gac aac acg
T G N P T P K D N S W K P I S E N D N T
tat ctg gaa atc gaa aaa gaa tta act ctc aag aag aat ttc aac gag aaa gag gcg aaa
Y L E I E K E L T L K K N F N E K E A K
ttg tgg aat gaa att tac aaa tcc gtt tgc aca aga cac aag taa
L W N E I Y K S V C T R H K -

```

132 **Figure 1** Nucleotide sequences of the longest ORF of Comp59220 and deduced amino acid
 133 sequence
 134

135 structural data of large biological molecules such as proteins, LCαE7, a CE from *Lucilia cuprina*,
 136 exhibits the highest (30%) identity with LBCE1. LCαE7 was previously reported to be a CE that
 137 potentially inactivates OPs, which leads to the increased OP resistance of *Lucilia cuprina*.
 138 Consistently, it was shown that the use of *in silico*-designed inhibitors of LCαE7 as synergists
 139 significantly increased the efficacy of OPs. Therefore, in this study, we focused on LBCE1 as a
 140 potential target of OP synergists.

141

Table 1 BLAST search hits identified as LBCE1-homologous proteins in the NCBI protein database

Description	Query Cover	E value	Per. Ident	Accession
Esterase FE4 precursor, putative [Pediculus humanus corporis]	96%	5e-171	47.19%	XP_002424560.1
esterase E4 isoform X2 [Cryptotermes secundus]	94%	2e-153	45.49%	XP_023716388.1
esterase E4 isoform X1 [Cryptotermes secundus]	94%	4e-153	45.67%	XP_023716379.1
esterase E4-like isoform X2 [Zootermopsis nevadensis]	95%	5e-152	45.32%	XP_021912901.1
esterase E4-like isoform X1 [Zootermopsis nevadensis]	95%	2e-150	44.49%	XP_021912894.1
esterase FE4 [Cryptotermes secundus]	94%	8e-148	44.07%	XP_023716405.1
uncharacterized protein LOC111869379 [Cryptotermes secundus]	94%	9e-147	45.80%	XP_023716645.1
uncharacterized protein LOC111869395 [Cryptotermes secundus]	96%	2e-142	45.12%	XP_023716667.1
PREDICTED: venom carboxylesterase-6 [Amyeloidis transitella]	94%	7e-142	44.48%	XP_013183727.1
venom carboxylesterase-6 [Manduca sexta]	95%	5e-141	43.68%	XP_030024687.1
venom carboxylesterase-6-like [Aphantopus hyperantus]	95%	2e-139	42.91%	XP_034834181.1
venom carboxylesterase-6-like [Hyposmocoma kahamanoa]	95%	3e-138	43.48%	XP_026315216.1
venom carboxylesterase-6 [Bicyclus anynana]	95%	6e-138	42.75%	XP_023935921.1

151

152 **Enzyme assay of the recombinant LBCE1**

153 The *E. coli* expression system yielded 30 μg of the recombinant His-tagged LBCE1 after
 154 purification (Figure 2A and 1B). The esterase activity of this enzyme was examined using a
 155 *p*-nitrophenol acetic acid as a substrate. As shown in Figure 2C, the substrate was shown to be
 156 hydrolyzed in a time-dependent manner, indicating that this protein is has an esterase activity.

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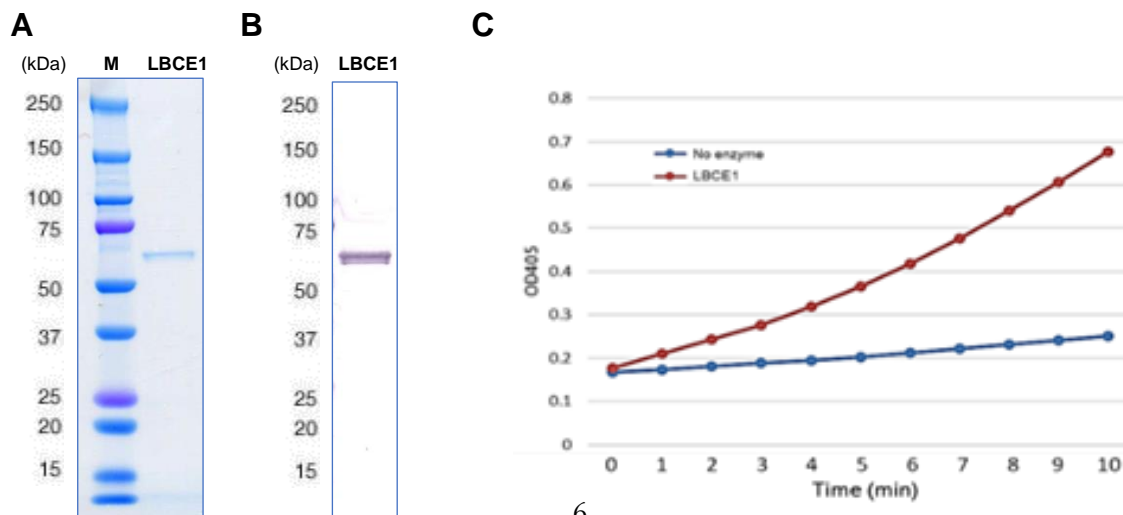
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Figure 2 (A) SDS-PAGE of the purified His-tagged LBCE1. (B) Western blotting of the purified His-tagged LBCE1 using an anti-His tag antibody. (C) Time-dependent hydrolysis of *p*-nitrophenol acetic acid by the recombinant LBCE1. As a control, the substrate was also incubated without LBCE1. M: Size marker.

166

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170

171 **Conflict of interest**

172

173 The authors declare no conflict of interest.

174

175 **References**

176

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