DATE: Day 20 Month April Year 2018

## SUMMARY of

## 2017 RESEARCH RESULTS REPORT

## For International Collaborative Research with IPR, Osaka University

Research Title		Structures of LC3B in complex with LIR motifs of RavZ
Applicant	Name	Hyun Kyu Song
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	Present Title	Crystal structures of LC3B in complex with LIR motifs of RavZ
Research Collaborator (Host PI)		Atsushi Nakagawa

## **Summary**

LC3-family member proteins play a critical role in autophagy, a cellular process responsible for the degradation of massive cellular components including intracellular pathogens. A variety of molecules involved in the autophagic pathway engage in specific interactions with a unique sequence motif referred to as the LIR (LC3-interacting region) motif. Although identification of conserved structural features of LIR motifs in complex with LC3-family members has established a canonical LIR motif, atypical conformations of LIR motifs have recently been revealed. We determined the high resolution crystal structures of LC3B in complex with three different LIR motifs of RavZ from *Legionella pneumophila*, an intracellular pathogen that can manipulate the host autophagy system. The tandem LIR motifs located in the N-terminal region of RavZ adopt a novel β-sheet conformation and thus provide specific ionic interactions with LC3B in addition to canonical hydrophobic plugged-in interactions. Consequently, these motifs possess higher binding affinity to LC3-family members than canonical LIR motifs, although the tandem repeats can only bind to one LC3 molecule. These findings broaden our understanding of the functional repertoire of LIR motifs in autophagy.

<sup>\*</sup>Deadline: May 18, 2018

<sup>\*</sup>Please submit it to E-mail: tanpakuken-kyoten@office.osaka-u.ac.jp.

<sup>\*</sup>We accept only PDF file. Please file it after converting WORD to PDF.

<sup>\*</sup>Please describe this summary within 1 sheet. Please DON'T add some sheets.

<sup>\*</sup>This summary will be published on the web.