

Autophagy and Bacterial Infection: a Competition for Survival

PhD student: Ying, YANG

Supervisor: Professor Margaret IP

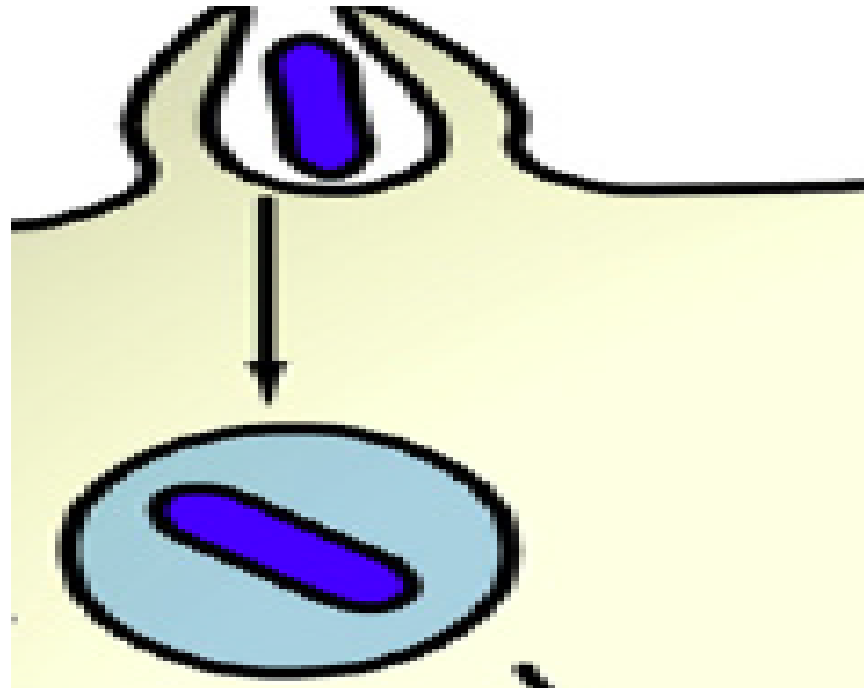
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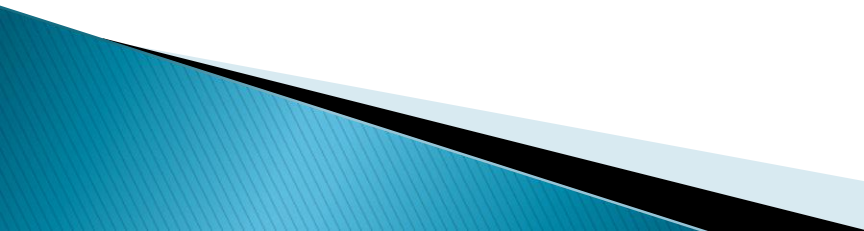


Autophagy: an essential conserved lysosomal degradation pathway

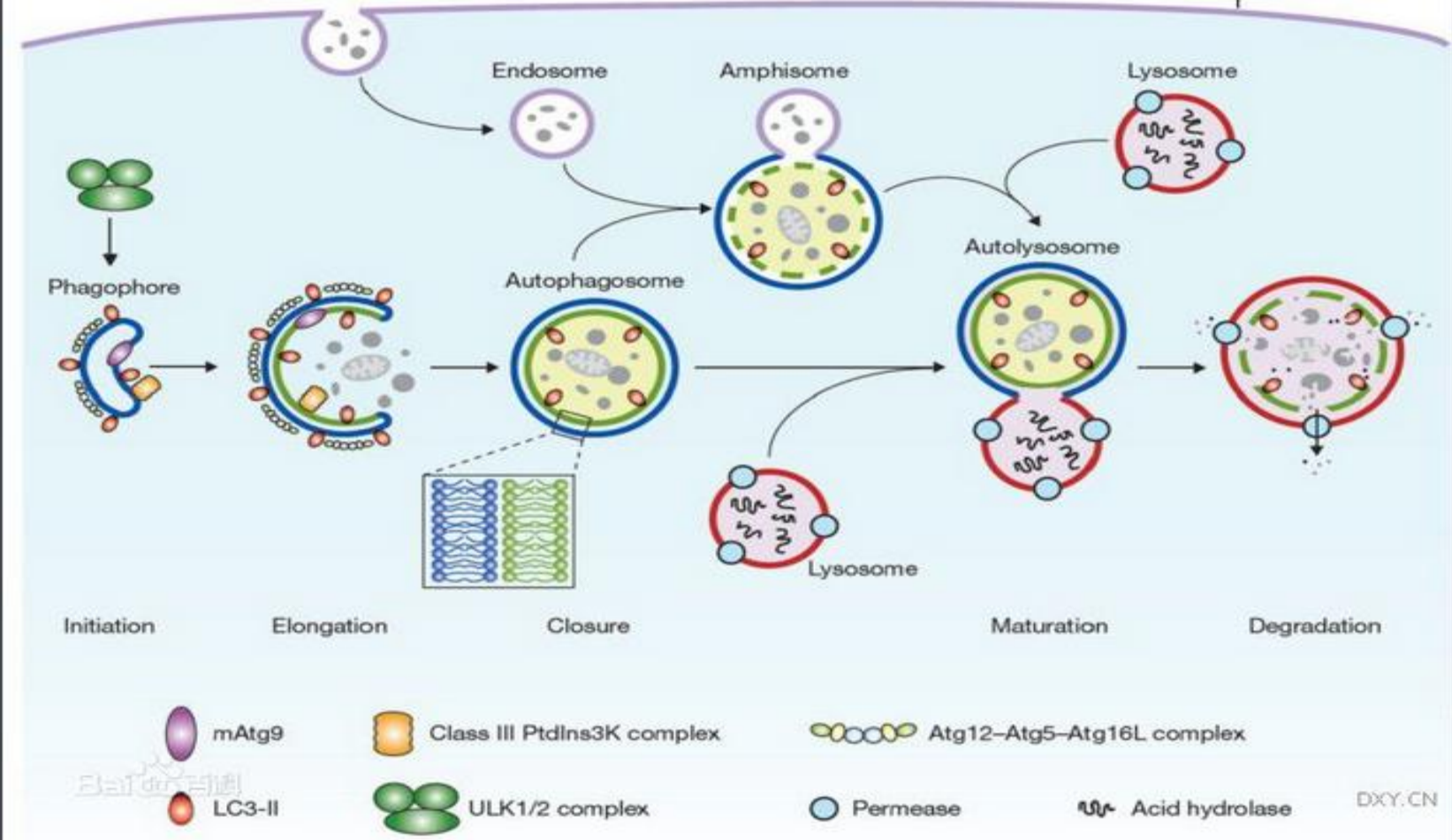
Numerous pathogens have developed the capacity to invade host cells to be protected from components of the systemic immune system.



Autophagy: an essential conserved lysosomal degradation pathway

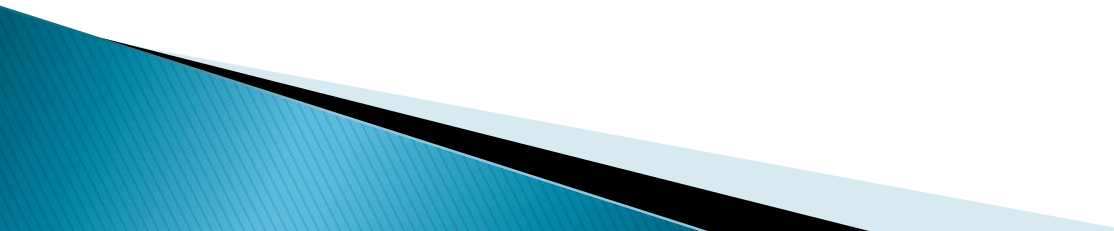
- ▶ Autophagy is an intracellular process delivering cytoplasmic material to the lysosome for degradation.
 - ▶ The cellular events of this ancient and highly conserved process: phagophore- autophagosome- autolysosome and degrade the enclosed material.
 - ▶ Autophagy acts as a cytoplasmic quality control mechanism, eliminating protein aggregates, damaged organelles and intracellular microbes to maintain cellular homeostasis.
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Autophagy: an essential conserved lysosomal degradation pathway

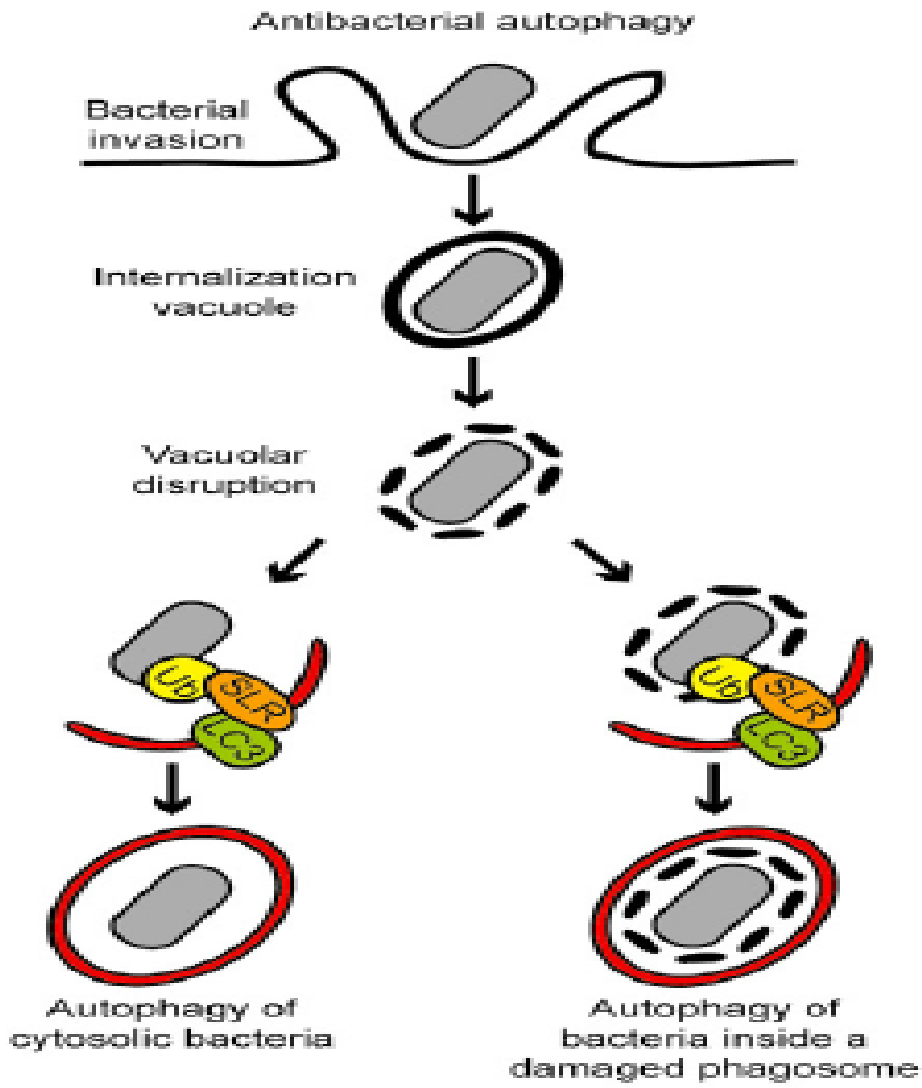


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Autophagy: different roles during different bacterial infections

- ▶ Antibacterial autophagy: restrict bacterial replication.
 - ▶ Non-bacterial autophagy: cross-talk with innate immune signaling pathways.
 - ▶ Pro-bacterial autophagy: support bacterial replication.
- 

Antibacterial autophagy: restrict bacterial replication



Mostowy S. Autophagy and bacterial clearance: a not so clear picture[J]. Cellular microbiology, 2013, 15(3): 395-402.

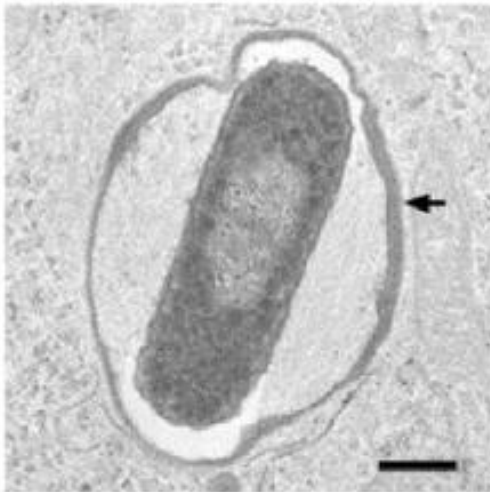
Antibacterial autophagy: Shigella clearance

- ▶ Autophagy of Shigella is triggered by ATG5 recognition of IcsA and is mediated by TECPR1, a Tectonin domain-containing protein, which binds to ATG5 and promotes autophagosome–lysosome fusion.
- ▶ To restrict bacterial motility and autophagy escape, septins are guanosine triphosphate (GTP) binding proteins recruited to sites of IcsA-induced actin polymerization, and form cage-like structures.

Antibacterial autophagy: *Shigella* clearance

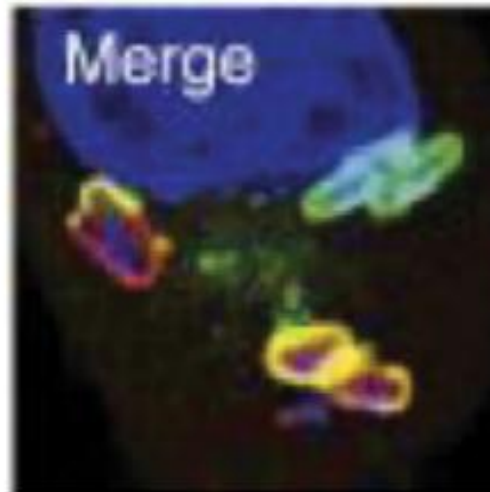
A

Shigella
autophagosome



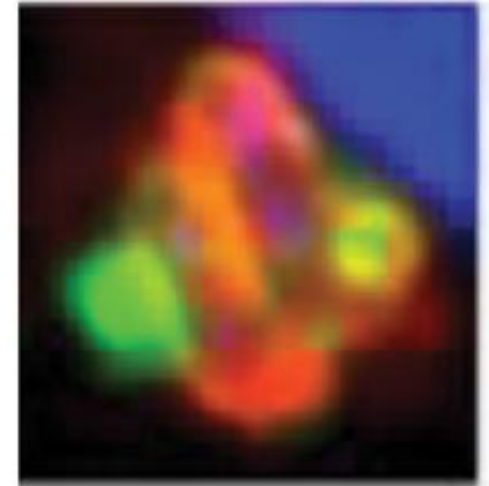
B

ATG5 binding
(TECPR1 and LC3)



C

The septin cage
(SEPT2 and LC3)

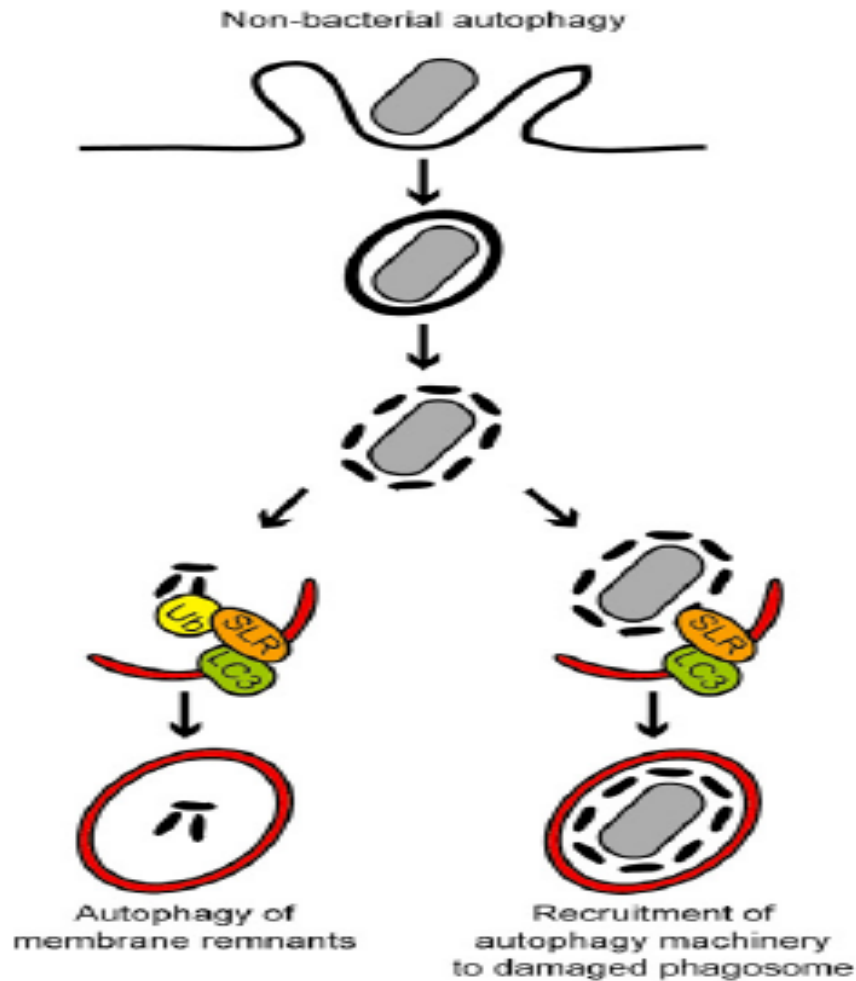


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Antibacterial autophagy: *Mycobacterium tuberculosis* clearance

- ▶ *Mycobacterium tuberculosis* is a vacuolar pathogen that survives within macrophages by arresting phagosomal maturation.
- ▶ Autophagy induction counteracts the maturation block imposed by mycobacterium.
- ▶ A variety of studies have shown that the induction of autophagy by starvation, inhibition of mTOR (a suppressor of autophagy), vitamin D and interferon-gamma (IFN γ) may help restrict mycobacterial replication. p62 appears to be crucial for this process, and provides mycobacterial autophagolysosomes with enhanced antimicrobial capacities relative to conventional phagolysosomes.

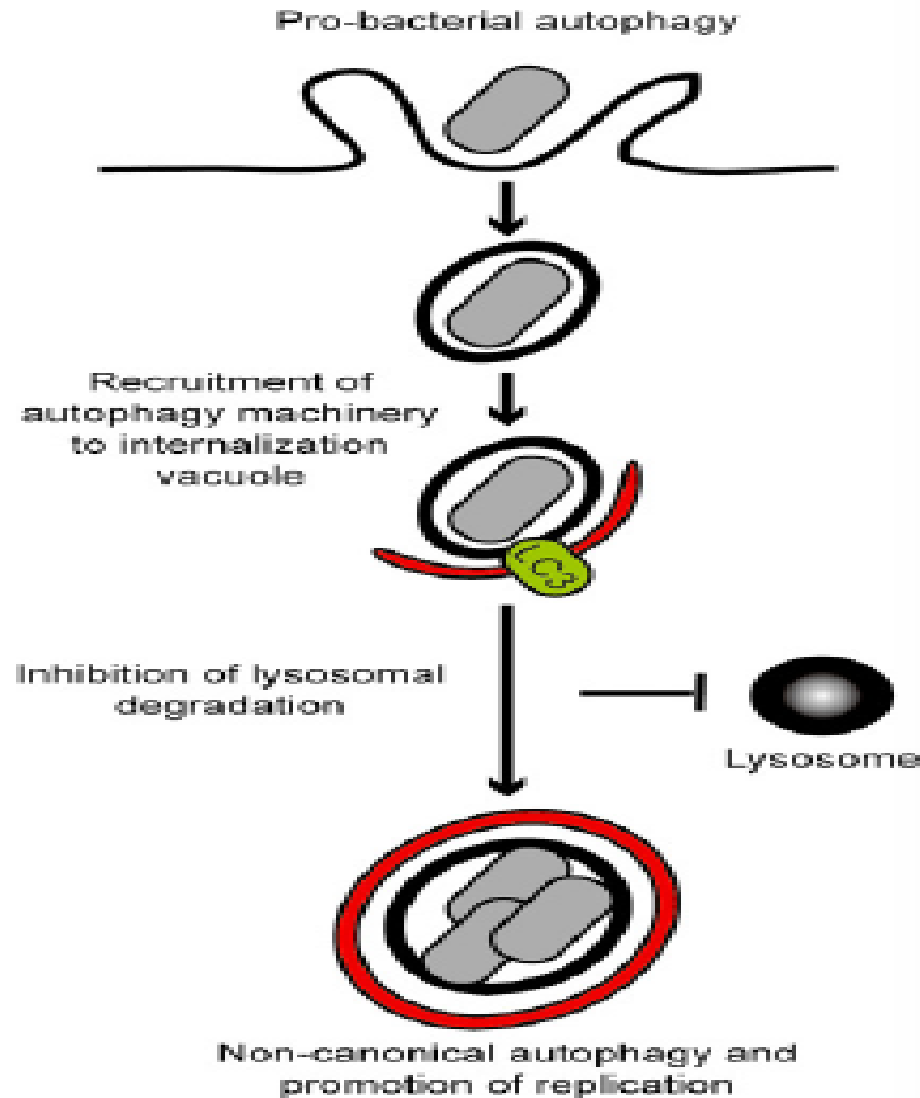
Non-bacterial autophagy : cross-talk with innate immune signaling pathways



Non-bacterial autophagy : cross-talk with innate immune signaling pathways

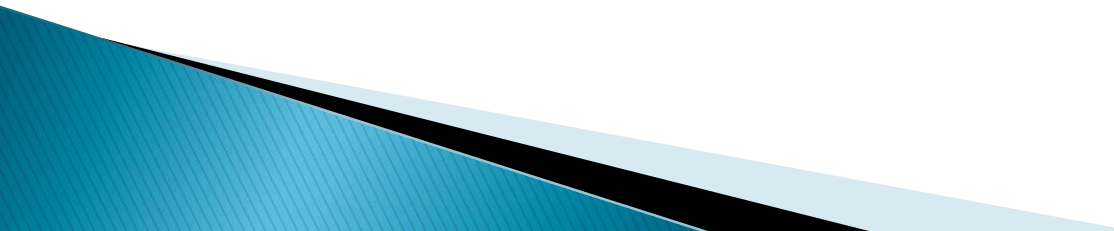
- ▶ Studies using *Shigella* have shown that in the cytosol of infected cells, membrane remnants induced by invading *Shigella* are ubiquitinated and recognized by p62, NBR1 and NDP52 for delivery to autophagosomes.
- ▶ Inflammasome components, localized to damaged membranes, are also ubiquitinated and recognized by p62 for autophagy.
- ▶ The studies with *V.cholerae* cytolysin (VCC) the pore-forming toxin indicate that the toxin triggers an autophagic response in the target cell, and demonstrate that autophagy acts as a cellular defense pathway against a secreted bacterial toxin.

Pro-bacterial autophagy: support bacterial replication



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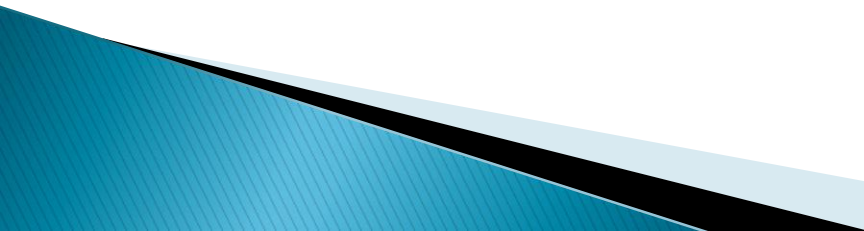
Pro-bacterial autophagy: support the replication of *Brucella*

- ▶ **Brucella** may co-opt autophagosome initiation factors ATG1 (ULK1), Beclin1 and ATG14 to convert *Brucella*-containing vacuoles (BCVs) into autophagosome-like compartments called autophagic BCVs (aBCVs).
 - ▶ A non-canonical autophagy pathway here promotes bacterial replication and survival since autophagosome elongation factors ATG4B, ATG5, ATG7, LC3B and ATG16L1 are not required for biogenesis of aBCVs.
- 

Pro-bacterial autophagy: support the replication of *S. aureus*

- ▶ ***S. aureus*** is a Gram-positive bacterium that can invade cells and replicate in autophagosome-like vacuoles. Hla (α -haemolysin), a pore-forming toxin secreted by *S. aureus*, is required for the recruitment of LC3, suggesting the recruitment of autophagy components by Hla.
- ▶ Hla-induced autophagy requires ATG5, but does not require Beclin1 (ATG6) nor PI3 kinase activity, highlighting the benefit of a non-canonical autophagy pathway for *S. aureus* replication.

Perspectives

- ▶ Autophagy acts as an innate cytoplasmic surveillance mechanism: enables the host cell to remove pathogens, and to subsequently deliver them to the degradative lysosomal compartment.
 - ▶ Pathogens have developed different strategies to avoid this pathway.
 - ▶ Many key points remain to be solved, the search for virulence factors and the elucidation of how they control the autophagy machinery will be the basis of novel therapeutic intervention against intracellular pathogens.
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Perspectives



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Rapamycin, Autophagy, and Alzheimer's Disease

Zhiyou Cai^{a,*} and Liang-Jun Yan^b

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Abstract

Alzheimer's disease (AD) is a neurodegenerative disorder characterized by multiple pathological lesions. At the molecular level, AD is characterized by amyloid-beta (Aβ) production and tau hyper-phosphorylation. Hence, phagosome accumulation and tau hyper-phosphorylation have potentiated the inhibition of mammalian target of rapamycin (mTOR), its downstream effectors. It is neuroprotective in neurodegenerative diseases. Enhancement of autophagy, a biological process that not only facilitates the clearance of mutant proteins but also significantly reduces the build-up of toxic protein aggregates such as Aβ. Since rapamycin enhancement of autophagy has been associated with abrogation of AD pathological processes such as clearance of Aβ and neurofibrillary tangles (NTFs) as well as reduction of tau hyper-phosphorylation and

| Compounds | Company | Structure | Mechanism of action |
|---------------------------|---------------|-----------|--|
| <i>Autophagy inducers</i> | | | |
| Rapamycin | Sigma-Aldrich | | Induces autophagy by inhibiting mTORC1 |

Perspectives

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Rilmenidine attenuates toxicity of polyglutamine expansions in a mouse model of Huntington's disease

Claudia Rose¹, Fiona M. Menzies¹, Maurizio Renna¹, Abraham Acevedo-Arozena², Silvia Corrochano², Oana Sadiq¹, Steve D. Brown² and David G. Klapper^{1*}

¹Department of Medical Genetics, University of Cambridge, Cambridge Hospital, Hills Road, Cambridge CB2 0XY, UK and ²Medical Research Council, Oxfordshire, UK

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Huntington's disease (HD) is an autosomal dominant neurodegenerative disorder caused by a CAG repeat expansion in huntingtin. There are no treatments that are known to modify the progression of this mutation. Mutant huntingtin causes disease via a toxic gain of function that leads to aggregation and formation of intraneuronal inclusions. One therapeutic approach for HD is to enhance the degradation of the mutant protein. We have shown that this can be achieved by upregulating autophagy, using the drug rapamycin. In order to find safer ways of inducing autophagy for clinical purposes, we previously screened United States Food and Drug Administration-approved drugs for their autophagy-stimulating potential. This screen suggested that rilmenidine, a well tolerated, safe, centrally acting anti-hypertensive drug, could induce autophagy in cell culture via a pathway that was independent of the mammalian target of rapamycin. Here we have shown that rilmenidine induces autophagy in mice and in primary neuronal culture. Rilmenidine administration attenuated the signs of disease in a HD mouse model and reduced levels of the mutant huntingtin fragment. As rilmenidine has a long safety record and is designed for chronic use, our data suggests that it should be considered for the treatment of HD and related conditions.

Rilmenidine*

Tocris Bioscience



Lowers cAMP levels

Perspectives

NATURE | ARTICLE



日本語要約

Identification of a candidate therapeutic autophagy-inducing peptide **Tat beclin1**

Sanae Shoji-Kawata, Rhea Sumpter, Matthew Leveno, Grant R. Campbell, Zhongju Zou, Lisa Kinch, Angela D. Wilkins, Qihua Sun, Kathrin Pallauf, Donna MacDuff, Carlos Huerta, Herbert W. Virgin, J. Bernd Helms, Ruud Eerland, Sharon A. Tooze, Ramnik Xavier, Deborah J. Lenschow, Ai Yamamoto, David King, Olivier Lichtarge, Nick V. Grishin, Stephen A. Spector, Dora V. Kaloyanova & Beth Levine

[Affiliations](#) | [Contributions](#) | [Corresponding author](#)

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Thank you!



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