Figure EV1. Evaluation of cell death after P.

A–D A549 cells were either left uninfected (NI) or infected for 4 h with WT and different

P. aeruginosa mutants. Cells were then stained with Sytox-green (A, C) or propidium iodide

(PI) (B, D) and the percentage of dead cells was

analyzed by flow cytometry. Data are mean +/- SD from three independent experiments;

one-way ANOVA analysis followed Dunn's multiple-comparison posttest t, \* $P \le 0.05$ .

aeruginosa infection.



## **Expanded View Figures**



### Figure EV2. Induction of autophagy was attenuated by P. aeruginosa T3SS ExoS.

- A A549 cells were treated for 16 h with DMSO or rapamycin. Cells were then left uninfected (NI) or infected for 4 h with WT or *ApscD P. aeruginosa* and cell lysates were evaluated by immunoblotting.
- B A549 cells were transfected, for 24 h, with a plasmid containing an empty vector (EV), GFP-ExoS or an inactive mutant (GFP-ExoS<sup>G-A-</sup>). Cells were then treated for 16 h with DMSO or rapamycin and cell lysates were evaluated by immunoblotting.

Source data are available online for this figure.



#### Figure EV3. ExoS ADP-ribosyltransferase has the same effect on autophagy in primary epithelial cells; Single cytotoxin ExoT or ExoY does not affect mTOR or autophagy.

A-C A549 cells (A) and primary NHBE cells (B, C) were infected for 4 h, with WT or P. aeruginosa mutants ( $\Delta$ STY,  $\Delta$ T,  $\Delta$ SY,  $\Delta$ Y,  $\Delta$ ST S<sup>WT</sup>, S<sup>G-A-</sup>,  $S^{G-A^{+}}\!\!,$  and  $S^{G+A^{-}}\!\!)\!,$  and cell lysate were evaluated by immunoblotting using indicated antibodies. P. aeruginosa mutant strains legend: S<sup>WT</sup> (P. aeruginosa contains only active ExoS), S<sup>G-A+</sup> (P. aeruginosa contains only ExoS with loss-of-function mutations in the GTPaseactivating domain), S<sup>G+A-</sup> (P. aeruginosa contains ExoS with loss-of-function in the ADP ribosylation domain), S<sup>G-A-</sup> (*P. aeruginosa* contains a nonfunctional ExoS); $\Delta$ STY (mutant deficient for the three cytotoxins),  $\Delta T$  (deficient for ExoT but still express ExoS and ExoY cytotoxins),  $\Delta$ SY (deficient for ExoS and ExoY only express ExoT),  $\Delta Y$  (deficient for ExoY but still express ExoS and ExoT cytotoxins),  $\Delta$ ST (deficient for ExoS and ExoT only express ExoY).

Source data are available online for this figure.





# Figure EV4. ExoS reduced autophagosome and autolysosome formation.

- A A549 cells, transfected with RFP-GFP-LC3 vectors for 24 h, were infected with different *P. aeruginosa* mutant strains. Representative images form fluorescence microscopy assays were shown. Scale bars: 10 μm.
- B Quantitative analysis of yellow puncta generated from overlapping GFP and RFP puncta (represent autophagosome) and RFP-LC3 puncta (represent autolysosome). The puncta from more than 100 cells were counted and the ratios of these puncta per cell are shown. Data are mean +/– SD from three independent experiments. The significance of differences between treatments was determined using two-tailed Student's *t*-test with Welch's correction, \* $P \leq 0.05$ .

Source data are available online for this figure.



#### Figure EV5. Evaluation of cellular activity of Ras mutants and Working model of ExoS facilitates intracellular survival of *P. aeruginosa* by inhibiting autophagy and mTOR.

- A Schematic of the generation of Ras-G12V&R41K from Ras-G12V.
- B A549 cells were transfected, for 24 h, with plasmids containing empty vector (EV), Ras-G12V or Ras-G12V&R41K and cell lysates were evaluated by immunoblotting.
- C Working model of *P. aeruginosa* ExoS role in inhibiting host cells autophagy and mTOR to facilitate intracellular bacterial survival. *P. aeruginosa* injects toxins including ExoS, ExoY, and ExoT through the T3SS. ExoS-mediated ADP ribosylation of RAS and leads to inhibition of mTOR. In addition, ExoS's ADP ribosylation activity suppresses autophagy initiation complex Atg14L-Vps34 kinase activity preventing autophagosome formation. The inhibition of autophagy by ExoS ADP ribosylation protects bacteria from elimination by autolysosome digestion.

Source data are available online for this figure.