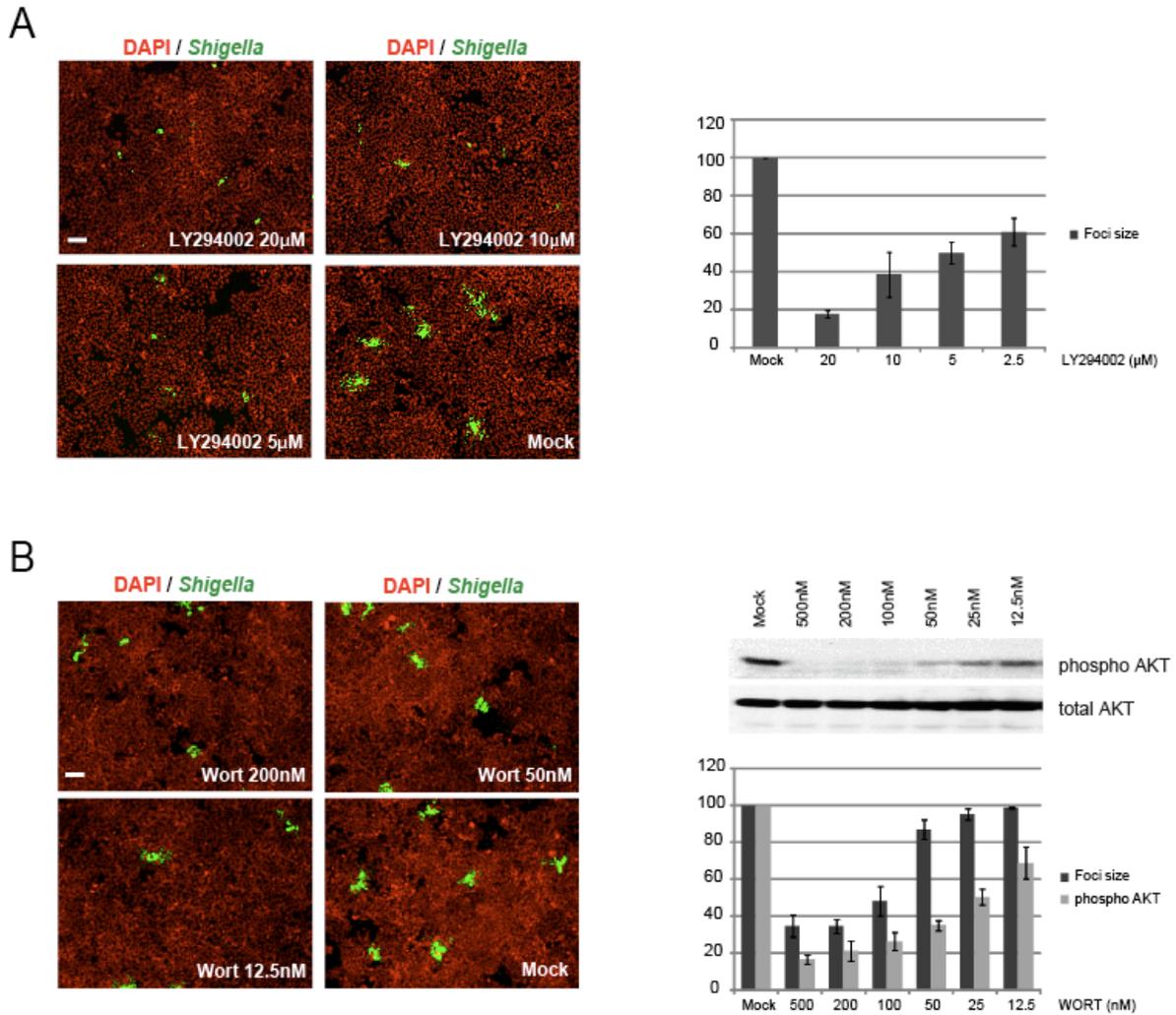


1 **Supplementary materials**

2

3 **Supplementary figures**

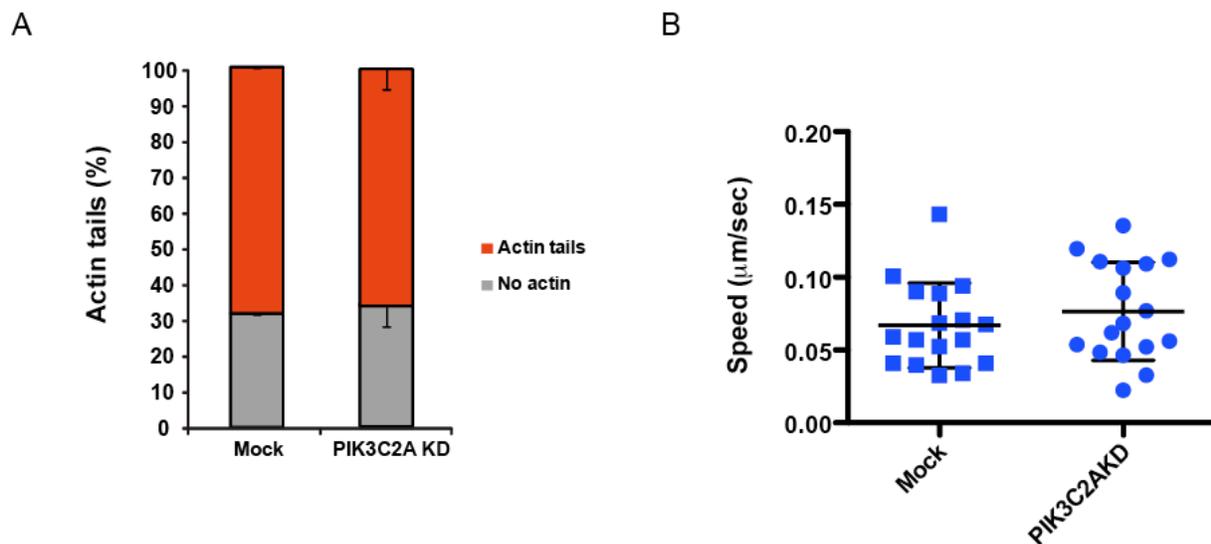


4

5 **Figure S1. Effect of PIK3 inhibitors on *S. flexneri* dissemination in HT-29 cells**

6 (A) Representative images of HT-29 cells pretreated for 1h with various concentrations of
7 Wortmannin and infected with GFP-expressing *S. flexneri* for 8h. (red, nuclei; green, *Shigella*).
8 Graph showing the analysis of focus size in Wortmannin-treated HT-29 cells relative to the size
9 recorded in mock-treated cells set at 100%. Values represent the mean +/- SD of three

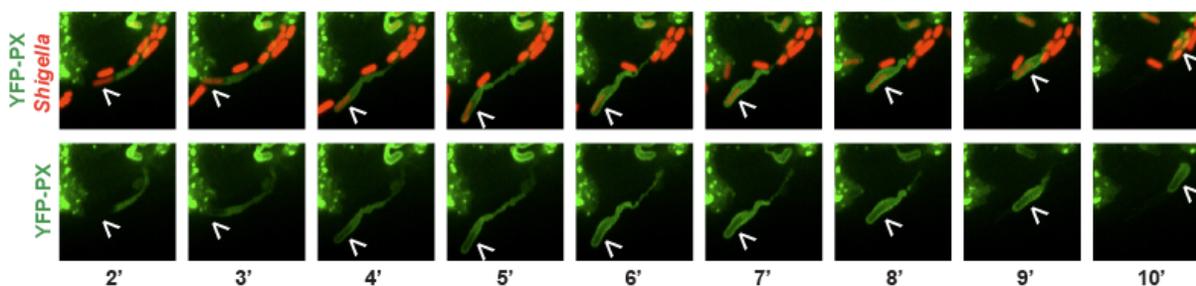
10 independent experiments. (B) Upper panel: western blot analysis of Akt phosphorylation in
 11 mock-treated and Wortmannin-treated cells. Lower panel: Akt phosphorylation levels were
 12 quantified in mock-treated and Wortmannin-treated cells using ImageJ software analysis and
 13 normalized to the total Akt levels. Graph showing Akt phosphorylation level in Wortmannin-
 14 treated cells calculated as percentile level of mock-treated cells (phospho Akt). Statistical
 15 analysis: Mock vs. Wort500 $p < 0.0001$, Mock vs. Wort200 $p < 0.0001$, Mock vs. Wort100
 16 $p < 0.0005$, Mock vs. Wort50 $p < 0.05$, Mock vs. Wort25 NS, Mock vs. Wort12.5 NS. The
 17 experiment shows the inhibitory effect of Wortmannin on Akt phosphorylation in the 50-12.5
 18 nM concentration range, with little to no effect on *Shigella* dissemination.
 19



20
 21 **Figure S2. PIK3C2A is not required for cytosolic motility**
 22 (A) Graph showing counts of actin tails in mock-treated and PIK3C2A-depleted cells. Values
 23 represent the mean +/- SD of three independent experiments. (B) Graph showing measurements
 24 of *S. flexneri* velocity in the cytosol of mock-treated and PIK3C2A-depleted HT-29 cells. The

25 tracking module of the Volocity software was used to score motile bacteria and determine their
26 velocity ($\mu\text{m/s}$).

27

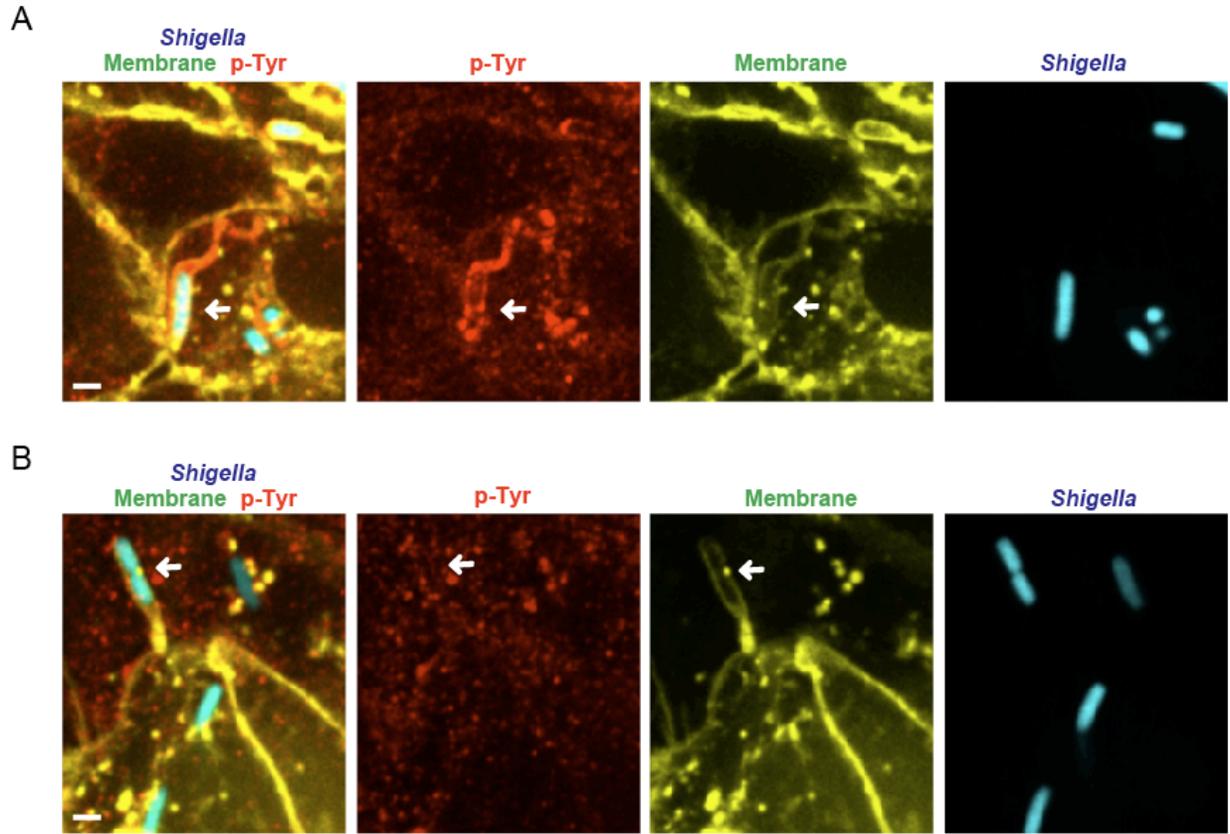


28

29 **Figure S3. Localization of YFP-PX in *S. flexneri* protrusions**

30 Representative time-lapse images of YFP-PX-expressing HT-29 cells (green) infected with RFP-
31 PIK3C2A-expressing *S. flexneri* (red). Note the progressive YFP-PX enrichment in the plasma
32 membrane surrounding protrusions overtime (arrowhead).

33



34

35 **Figure S4. Phospho-tyrosine residues in *S. flexneri* protrusions**

36 Representative images of HT-29 cells infected with CFP-expressing *S. flexneri* (blue) displaying
 37 protrusions (arrows, membrane-YFP, yellow) that scored (A) phospho-tyrosine positive (red) or
 38 (B) phospho-tyrosine negative (red), as counted in Figure 7C.

39

40

41 **Supplementary movies**

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43 **Movie S1. Time lapse imaging of protrusion resolution into vacuole through VLP**
44 **formation**

45 HT-29 cell expressing membrane GFP (yellow) were infected with RFP-expressing *Shigella*
46 *flexneri* and imaged every five minutes. Scale bar: 5 μ m. The movie shows the formation of a
47 protrusion that successfully transitions into a Vacuole-Like-Protrusion (VLP) and resolve into a
48 vacuole.

49

50 **Movie S2. Time lapse imaging of unsuccessful protrusions resolution into vacuole**

51 HT-29 cell expressing membrane GFP (yellow) were infected with RFP-expressing *Shigella*
52 *flexneri* and imaged every five minutes. Scale bar: 5 μ m. The movie shows the formation of a
53 protrusion that fails to transitions into a Vacuole-Like-Protrusion (VLP) and retracts bringing the
54 pathogen back to the primary infected cell.

55

56