



HAL
open science

Diverted recycling- Shigella subversion of Rabs

Noelia López-Montero, Jost Enninga

► **To cite this version:**

Noelia López-Montero, Jost Enninga. Diverted recycling- Shigella subversion of Rabs. Small GTPases, 2018, 9 (5), pp.365-374. 10.1080/21541248.2016.1240494 . pasteur-01388795

HAL Id: pasteur-01388795

<https://pasteur.hal.science/pasteur-01388795>

Submitted on 27 Oct 2016

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution 4.0 International License

1 **Title: Diverted recycling- *Shigella* subversion of Rabs**

2

3 Noelia Lopez-Montero and Jost Enninga

4

5 Institut Pasteur, Research unit “Dynamics of host-pathogen interactions”, 25 rue du Dr.

6 Roux, 75724 Paris, France

7

8 Correspondence: Jost Enninga (jost.eninga@pasteur.fr)

9

Noelia Lopez-Montero (noelia.lopez-montero@pasteur.fr)

10

11

12

13

14

15

16

17

18

19

20

21

22

23

24

25

26

27

28

29 **Keywords**

30 Vesicular traffic, Rab subversion, Rab11, intracellular bacteria, bacterial containing
31 vacuole, vacuolar rupture, *Shigella flexneri*, macropinocytosis.

32

33 **Abbreviations**

34 Bacterial containing vacuoles (BCVs).

35 Early endosome antigen 1 (EEA1).

36 Endoplasmic reticulum (ER).

37 Focused ion beam scanning electron microscopy (FIB/SEM).

38 GDP/GTP exchange factor (GEF).

39 Glyceraldehyde-3-phosphate dehydrogenase (GAPDH).

40 GTP-hydrolysis activating proteins (GAPs).

41 Kinesin superfamily proteins, kinesins (KIFs).

42 Lipoarabinomannan (LAM)

43 Lysosome-related organelles (LROs).

44 Mannose phosphate receptors (MPRs).

45 Multi-vesicular bodies (MVBs).

46 Myosin V (MyoV).

47 Phosphatidylinositol 3-kinase catalytic subunit type 3 (PI3KC3).

48 Phosphatidylinositol 3-phosphate (PI3P).

49 Phosphatidylinositol 4,5-bisphosphate (PI(4,5)P2).

50 Phosphatidylinositol 5-phosphate (PI5P).

51 Pleckstrin homology domain-containing family M member 2 (SKIP).

52 Rab GDP dissociation inhibitors (GDIs).

53 Rab-interacting lysosomal protein (RILP).

54 Rab11 family-interacting protein (FIP).

55 *Salmonella* induced filaments (SIFs).

56 Trans-Golgi network (TGN).

57 Type-3 secretion system (T3SS).

58

59

60 **Abstract**

61 Small GTPases of the Rab protein family control intracellular vesicular trafficking to allow
62 their communication and maintenance. It is a common strategy for intracellular bacteria to
63 exploit these pathways to shape their respective niches for survival. The subversion of
64 Rabs for the generation of an intracellular environment favoring the pathogen has been
65 described almost exclusively for intracellular bacteria that reside within bacterial containing
66 vacuoles (BCVs). However, less is known about Rab subversion for bacteria that rupture
67 the BCV to reach the host cytoplasm. Here, we provide recent examples of Rab targeting
68 by both groups of intracellular bacteria with a special focus on *Shigella*, the causative
69 agent of bacillary dysentery. *Shigella* recruits Rab11, the hallmark of the perinuclear
70 recycling compartment to *in situ* formed macropinosomes at the entry foci via the bacterial
71 effector IpgD. This leads to efficient BCV rupture and cytosolic escape. We discuss the
72 concept of diverted recycling through host Rab GTPases that emerges as a novel
73 pathogen strategy.

74

75

76 **Main text**

77 **Introduction**

78 Intracellular bacteria have evolved molecular weapons to subvert host cells for their
79 own benefit. They manipulate eukaryotic pathways for the promotion of efficient cellular
80 invasion within a bacterial containing vacuole (BCV), intracellular survival, evasion of
81 immune responses, and for their own propagation.¹ The bacterial weaponry that achieves
82 such complex control is often constituted of an “effector cocktail”, secreted bacterial
83 proteins into the host cytosol through specialized injection devices, like the type-3
84 secretion system (T3SS). These effectors interplay with host factors involved in the
85 pathogenicity pathways. It has been recognized that the subversion of the vesicular
86 trafficking machinery represents a key strategy by the bacteria in order to physically shape
87 their local intracellular environment.²⁻⁴

88 **Rab GTPases in vesicular trafficking**

89 Cell membrane trafficking is regulated by the Rab family of proteins, which is part of the
90 Ras superfamily of small GTPases. In humans, almost 70 different Rab proteins have
91 been identified. They coordinate sequential trafficking steps by switching between their
92 active GTP-bound form and the inactive GDP-bound form, participating at different levels
93 for example during vesicle formation, motility, tethering or docking and fusion.⁵

94 Vesicles carrying internalized cargo from different entry pathways, such as clathrin
95 and caveolae-mediated endocytosis, macropinocytosis and phagocytosis traffic to
96 lysosomes for degradation. The pair Rab5-Rab7 is considered as the master regulator of
97 endosomal trafficking and maturation. Rab5 determines the identity of early endosomal
98 compartments.⁶ One of its effectors, the phosphatidylinositol 3-kinase catalytic subunit
99 type 3 (PI3KC3), produces phosphatidylinositol 3-phosphate (PI3P) on endosome

100 membranes, which in turn exerts a positive feedback loop resulting in increased Rab5
101 recruitment.⁷ Rabex5, a GDP/GTP exchange factor (GEF) for Rab5, contributes to the
102 positive loop by stabilizing Rab5 on the membranes.⁸ More than 60 downstream effectors
103 of Rab5 have been identified,⁹ but their functions remain most often poorly understood.
104 The progressive replacement of Rab5 by Rab7 is essential for proper maturation of early
105 endosomes into lysosomes.¹⁰⁻¹² Upon Rab7 recruitment and activation, Rab5 is released
106 allowing the maturation of early endosomes into late endosomes.¹⁰ Retrograde transport of
107 late endosomes along microtubules is necessary for efficient fusion with lysosomes. It is
108 mediated by the binding of active Rab7 to Rab-interacting lysosomal protein (RILP), which
109 induces the subsequent interaction with dynein-dynactin motor complexes.¹³ In addition,
110 the protein Hook has been shown to be involved in endosome maturation; its N-terminus
111 interacts with microtubules and its C-terminus with membranes.¹⁴

112 Alternatively to catabolic degradation, and as a requirement for cell surface
113 homeostasis, a part of the endosomal membrane and cargo are recycled back to the
114 plasma membrane.¹⁵ The membrane recycling system is highly dynamic; multiple
115 pathways for entry and exit from the endocytic recycling compartment have been
116 described with Rab11 present in most of them. Rab11 attracts effectors responsible for the
117 trafficking of recycling endosomes.¹⁶ It mediates vesicular transport along microfilaments
118 by binding the globular tail domain of myosin V (MyoV).¹⁷ Rab11 also interacts with the C-
119 terminal of MyoV forming a ternary complex through Rab11 family-interacting protein 2
120 (FIP2).¹⁸ In addition, Rab11 employs kinesins (KIFs) for anterograde transport along
121 microtubules. FIP5 drives the interaction of Rab11 with KIF3.¹⁹ Rab11 can bind directly
122 KIF13 promoting the formation of tubules.²⁰ The effector protrudin directs the interactions
123 of Rab11 with KIF5. Intriguingly, protrudin interacts selectively with GDP-Rab11²¹. This
124 changes the concept of Rab11 being switched off in its GDP form, and could explain the

125 large promiscuity of Rab11 functions. FIP3 regulates the retrograde transport of Rab11
126 endosomes along microtubules mediated by dynein. This has been proposed for the
127 sorting of peripheral endosomes to the perinuclear recycling compartment.²² In addition,
128 Rab11 is involved in the tethering and fusion of recycling endosomes with the plasma
129 membrane^{23,24}.

130 Although Rab11 is the hallmark of the slow recycling pathway, its highly versatile
131 nature and its complex localization profile in different cell types have led to numerous
132 hypotheses about its roles. Importantly, only one Rab11 GEF, Crag, has been recently
133 identified in *Drosophila*.²⁵ In mitosis, Rab11 has a role in organizing the mitotic spindle and
134 spindle poles, and centriole distribution.²⁶ It is involved in ciliogenesis, cytokinesis,
135 neuritogenesis, and oogenesis.¹⁶ It controls the maintenance of microvilli and
136 apical/basolateral specialization in epithelial cells,²⁷ and it is a regulator of autophagosome
137 maturation.²⁸

138 Besides the three Rabs, Rab5, Rab7 and Rab11, described in some detail above,
139 other Rabs have been analyzed in the context of several local trafficking pathways.
140 Exemplarily, Rab1 and Rab2 mediate endoplasmic reticulum (ER)-Golgi trafficking, Rab3
141 controls exocytic events, Rab32 is involved in the biogenesis of lysosome-related
142 organelles (LROs) and mitochondrial fission, Rab6 regulates intra-Golgi trafficking, Rab9
143 and Rab7L1 mediate retrograde trafficking from late endosomes and lysosomes to the
144 trans-Golgi network (TGN), Rab4 is involved in the fast recycling pathway, as has been
145 reviewed in detail by Mizuno-Yamasaki and cols.²⁹ In addition, many of the Rabs have
146 been described to be frequently involved in more than one single pathway.

147 **Manipulation of Rabs by intracellular bacteria**

148 Intracellular bacterial pathogens use different mechanisms to hijack Rab proteins
149 (see **table 1** for an overview), especially those that reside within BCVs. They can mimic

150 Rab GEFs and GTP-hydrolysis activating proteins (GAPs) through their effector proteins,
151 either favoring recruitment to or displacement of certain Rabs from their vacuoles. For
152 instance, the phosphatase activity of the *Salmonella enterica* effector SopB is required for
153 Rab5 recruitment to the BCV. SopB produces PI3P at the BCV membrane, which prolongs
154 the association of Rab5 allowing BCV maturation. In addition, active Rab5 associates with
155 the PI3KC3, which is responsible for augmented PI3P formation on the BCV membrane.³⁰
156 SopE, another *Salmonella* effector, has been found to act as a GEF for Rab5, recruiting
157 non-prenylated Rab5 on the BCV.³¹ *Mycobacterium tuberculosis* also interferes with Rab5.
158 The glycolipid lipoarabinomannan (LAM), released from the bacterial membrane into the
159 phagosomal membrane, inactivates the PI3KC3 impairing Rab5 recruitment.³²
160 Concomitantly, the *Mycobacterium* phosphatase SapM depletes PI3P contributing to the
161 arrest of BCV maturation.³³

162 Alternatively, some bacterial effectors bind Rabs and thus prevent subsequent
163 interactions with their cognate host effectors. For instance, the *Legionella* effector VipD
164 forms a complex with Rab5 and Rab22 preventing the interaction with their effectors, such
165 as Rabaptin-5 and early endosome antigen 1 (EEA1).³⁴ This leads to a generalized
166 impairment in endosomal maturation and, consequently, in pathogen degradation.³⁵
167 Another example is the complex formed by the *Salmonella* effector SifA and host protein
168 pleckstrin homology domain-containing family M member 2 (SKIP). The SifA-SIKP
169 complex induces the formation of *Salmonella* induced filaments (SIFs).³⁶ It also binds and
170 recruits Rab9 to the BCV and SIFs. Sequestration of Rab9 by SifA-SKIP avoids the
171 recycling of mannose phosphate receptors (MPRs), which prevents the delivery of
172 lysosomal hydrolytic enzymes to the BCV and SIFs.³⁷ In addition, the C-terminus of SifA,
173 which shows similar structure of other bacterial GEFs, binds RhoA although it does not
174 change its GTPase status.³⁸ SIF formation is indirectly supported through another

175 *Salmonella* effector, SopD2, which binds directly to Rab7. This effector inhibits the
176 nucleotide exchange of this Rab. It impairs the recruitment of the Rab effectors to the BCV,
177 and subsequently avoids degradation within lysosomes, thus allowing BCV maturation.³⁹

178 Other bacterial effectors are able to covalently modify Rab proteins and
179 therefore completely change their properties. It has been described that the *Legionella*
180 effector DrrA, which is a GEF for Rab1,⁴⁰ covalently modifies Rab1 by AMPylation of its
181 Switch II region. Then, the AMPylated Rab1 restricts the access of GAPs, becoming
182 constitutively active at the BCV.⁴¹ With opposed function, SidD reverses Rab1
183 AMPylation.⁴² The effector LidA has also an auxiliary role on Rab1 recruitment through the
184 action of DrrA,⁴³ and the effector LepB functions as a Rab1 GAP.⁴⁴ Another *Legionella*
185 effector named AnkX also covalently modifies Rab1. It transfers a phosphocoline group
186 from CDP-choline to a serine also in the Switch II region, leading to a strong inhibition of its
187 interaction with GEFs and Rab GDP dissociation inhibitors (GDIs).⁴⁵ Therefore, as
188 phosphocolinated Rab1 cannot be solubilized by the GDI, it remains membrane bound
189 even in the GDP form. The *Listeria monocytogenes* enzyme glyceraldehyde-3-phosphate
190 dehydrogenase (GAPDH) ADP-ribosylates Rab5. This covalent modification renders Rab5
191 unresponsive for activation by GEFs, and in turn it blocks further maturation into Rab7.⁴⁶

192 Some effectors can even degrade Rabs by proteolytic cleavage. This is the case of
193 the *Salmonella* effector GtgE, which is expressed in the broad-host bacterium *Salmonella*
194 *enterica* serovar Typhimurium (*S. Typhimurium*) and helps to overcome the host restriction
195 barrier. This bacterial effector is specific for Rab32 and Rab7L1. In mouse macrophages,
196 GtgE avoids Rab32 recruitment to the BCV preventing delivery of antimicrobial molecules
197 to the BCV, and allowing bacterial survival and replication. Conversely, the human specific
198 serovars *S. Typhi* and *S. Paratyphi* do not express GtgE. Rab32 is therefore recruited to
199 the BCVs, bringing antimicrobial capacity to the BCVs and killing intravacuolar bacteria.⁴⁷

200 The *Salmonella* GAP for Rab32, SopD2, contributes to Rab32 removal from the BCV in *S.*
201 *Typhimurium*.⁴⁸ However, in *S. Typhi* SopD2 is a pseudogene.⁴⁹ In addition, the absence
202 of GtgE in the human-adapted *S. Typhi* and *S. Paratyphi* allows Rab7L1 recruitment to the
203 BCV, which is required for the export of typhoid toxin, a unique virulence factor for human-
204 adapted serovars.⁵⁰

205 The different examples of manipulation of Rab GTPases by pathogens described
206 above have been characterized in some detail at the molecular level. However, this has
207 not yet been achieved for many other cases. For example, the *Brucella abortus* effector
208 RicA tethers Rab2 to its BCV by an unknown mechanism. *In vitro*, RicA binds preferentially
209 GDP-Rab2 but does not possess GEF activity.⁵¹ *Chlamydia trachomatis* recruits Rab6,
210 Rab11 and Rab14 to the BCV in order to scavenge sphingomyelin from the Golgi.^{52,53} It
211 also recruits Rab39, which is involved in multi-vesicular bodies (MVBs) trafficking, for
212 inclusion growth and bacterial development.⁵⁴ The specific bacterial effectors responsible
213 of such subversions are still unknown. Finally, *Yersinia pestis* promotes the recruitment of
214 Rab1 to the BCVs.⁵⁵ It is thought that this may be a mechanism to control vacuolar pH but,
215 again, neither bacterial effectors responsible nor precise molecular events have been
216 identified so far.

217 **Subversion of Rabs by *Shigella*: Role in vacuolar rupture**

218 For the causative agent of bacillary dysentery in humans, *Shigella flexneri*, Rab
219 subversion has only been studied very recently. It has been shown that the *Shigella*
220 effector VirA functions as a GAP for Rab1, mediating the disruption of ER-Golgi trafficking
221 and suppressing host autophagy, which contributes to intracellular bacterial persistence.⁵⁶
222 Even though not directly acting on Rabs, IpaJ cleaves the Arf1 and Arf2 N-myristoylated
223 host GTPases, which regulate Golgi trafficking, inhibiting thus the host secretory
224 pathway.⁵⁷

225 The scarce information about the usage of host Rabs by *Shigella* may be due to the
226 intricate lifestyle of this pathogen. *Shigella* is a cytosolic bacterium that ruptures its BCV
227 very rapidly after invasion,⁵⁸ replicates in the cytosol, and propagates into the neighboring
228 cells.⁵⁹ The events occurring at the initial steps of infection are highly transient, making
229 them difficult to be studied in detail.⁶⁰ In fact, in both reports mentioned above, the effects
230 of the *Shigella* effectors were studied at late infection times (around 3 hours post-
231 infection). At that stage bacteria are cytosolic and spread inter-cellularly. Therefore, the
232 possible implication of VirA and IpaJ on Rab subversion at the early stages, such as
233 bacterial entry and vacuolar escape, remains unclear.

234 As mentioned before, the step of vacuolar escape is a key event for *Shigella*
235 virulence, but the implication of vesicular trafficking and Rab subversion in the process
236 was never evaluated. Membrane damage, and thus vacuolar rupture, has been thought to
237 occur mainly through the insertion of the T3SS injectisome on the BCV membrane.^{61,62}
238 Contrary to the established ideas, our group has demonstrated that BCV rupture requires
239 host factors involved in vesicular trafficking, including Rab GTPases, to take place in an
240 efficient manner.^{63,64} In particular, *Shigella* hijacks Rab11 to newly formed
241 macropinosomes leading to efficient BCV rupture. This relation was initially discovered
242 characterizing data from a high-content siRNA library screen of host membrane trafficking
243 factors involved in *Shigella* BCV rupture. BCV escape was monitored by automated
244 microscopy using the CCF4 FRET probe as a reporter of vacuolar rupture.⁶⁵ This screen
245 provided further Rab proteins to be potentially involved in vacuolar escape, such as Rab1,
246 Rab3, Rab4, Rab5, Rab7L1 along with Rab11.⁶³

247 Our live imaging revealed that Rab5, Rab7 and Rab11 were recruited to the
248 *Shigella* entry sites upon bacterial invasion. While Rab5 was transiently recruited, Rab11
249 was massively and permanently accumulated.⁶³ Then, in order to evaluate the specific role

250 of Rab11 in vacuolar rupture, we used fluorescently tagged galectin-3 as live cell marker
251 for loss of endomembrane integrity.⁶⁶ In Rab11 knockdown cells, *Shigella* vacuolar escape
252 is dramatically delayed, but it does not affect bacterial entry. Importantly, the GTPase
253 activity of Rab11 is necessary for efficient BCV rupture, since expression of GDP-locked⁶⁴
254 and GTP-locked Rab11 (non published results) led to a strong delay in vacuolar rupture.
255 We found that Rab11 recruitment to the entry foci was entirely controlled by the bacterial
256 effector IpgD,⁶³ an inositol phosphate phosphatase that converts phosphatidylinositol 4,5-
257 bisphosphate (PI(4,5)P2) to phosphatidylinositol 5-phosphate (PI5P).⁶⁷ In agreement with
258 its implication in Rab11 recruitment, the *Shigella* effector IpgD exhibited a strong delay in
259 vacuolar rupture. In addition, we demonstrated that IpgD was also involved in the
260 regulation of macropinosome formation during bacterial invasion. The IpgD mutant, which
261 entered without any delay into epithelial cells, was dramatically impaired for
262 macropinosome formation during the time of invasion.⁶³ Previously, IpgD, through the
263 action of PI5P production was reported to be involved in the shaping of the entry foci.⁶⁸

264 Canonical macropinocytosis has been described as clathrin-independent non-
265 selective endocytosis (see **figure 1** for a comparison of *Shigella* induced
266 macropinosomes). By macropinocytosis, cells are able to internalize considerable volumes
267 of extracellular fluid for the uptake of soluble molecules as well as particles such as
268 viruses, bacteria and apoptotic cell fragments. Antigen presenting cells are capable of
269 constitutive macropinocytosis, while in macrophages, lymphocytes, fibroblast and epithelial
270 cells macropinosomes appear upon applying a variety of external stimuli⁶⁹.
271 Macropinosomes are directed towards the lysosomal pathway through the Rab5/Rab7
272 cascade, Lamp1 acquisition and acidification.⁷⁰ The recycling of them has only been
273 suggested by a few studies, furthermore the molecular mechanism of this and its
274 regulation remains unclear. Intriguingly, in some non-professional phagocytic cells

275 macropinosomes have been reported to recycle to the plasma membrane, with little or no
276 interaction with endosomal vesicles.⁷¹ Our recent study on *Shigella* induced
277 macropinosomes has been the first that proposed a direct implication of Rab11.

278 Classically, it has been considered that *Shigella* takes advantage of ruffling for its
279 own internalization within macropinosome-like vesicles.⁷² In contrast to this paradigm, our
280 results from multidimensional live cell confocal microscopy showed that the recruited
281 Rab5, Rab7 and Rab11 were located exclusively at the membranes of the surrounding
282 macropinosomes (**figure 1**), but never at the forming BCV.⁶⁴ At the ultrastructural level, the
283 different identities of both compartments was confirmed by focused ion beam scanning
284 electron microscopy (FIB/SEM), an emerging technique for tomography of large volumes,⁷³
285 using a correlative approach. This showed clearly that *Shigella* enters in a tight vacuole
286 distinct from the surrounding macropinosomes. In addition, FIB/SEM revealed the
287 presence of contact sites between the BCV and macropinosomes, with the appearance of
288 smaller intramacropinosomal vesicles right at their interface during vacuolar rupture.⁶⁴
289 Although Rab5 and Rab7 are recruited to *Shigella* induced macropinosomes, they do not
290 recruit Lamp1 nor do they show acidification (non published results). Instead, they are
291 blocked for further maturation and hijack Rab11 for virulence purposes. These
292 observations suggest that BCV rupture may be mediated by direct physical contacts
293 between Rab11-positive macropinosomes and the BCV. Such contacts may be an
294 outcome of “misdirected” trafficking towards the BCV instead of their recycling to the
295 plasma membrane, which could be considered as an “internal recycling” pathway.

296 In the scenario of “internal recycling” the exocyst (a protein complex made of eight
297 distinct proteins) may play a role. Rab11-positive recycling endosomes are delivered to the
298 apical cell surface along radial actin filaments by MyoV.¹⁷ Then, they are tethered to the
299 plasma membrane by binding of MyoV and Rab11 to the exocyst component Sec15 to

300 form a tripartite docking complex. Afterwards, Sec15 interacts with SNARE complexes to
301 mediate vesicle fusion with the plasma membrane.^{23,24} Knocking down constituents of the
302 exocyst inhibited the process of *Shigella* internalization (our own unpublished
303 experiments), but the precise step of inhibition needs to be elucidated.

304 Since the GDP/GTP status of Rab11 is important for BCV destabilization, another
305 possible role for Rab11 in BCV rupture could be as a shuttle, providing the nascent
306 macropinosomes with downstream molecules involved in contact formation. It has been
307 described in *Drosophila* that under fed conditions the protein Hook has a negative
308 regulatory role in autophagosome maturation. Hook anchors late endosomes to
309 microtubules and thus impairs their fusion with amphisomes, delaying their maturation. In
310 this context Rab11 is prominently located at recycling endosomes. In starving conditions,
311 Rab11 regulates the relocalization of Hook protein in a GTP-dependent manner binding
312 and recruiting Hook to autophagic structures. Late endosomes are then not anchored
313 anymore to microtubules, and their fusion with amphisomes is favored, allowing a faster
314 maturation and catabolic degradation.²⁸

315 Interestingly, Rab 11 has been also reported to be involved in expulsion of pore
316 forming toxins from the surface of challenged host cells.⁷⁴ A similar function has been
317 described for the ESCRT machinery, which is involved in local membrane deformation and
318 scission, and in repairing plasma membrane wounds.⁷⁵ It could be considered that Rab11
319 is recruited together with the ESCRT complex to the BCV via macropinosomes to repair
320 the damage caused by the insertion of the T3SS translocon complex, pinching off small
321 intraluminal vesicles at the contact interphase, and inducing in turn vacuolar rupture.

322 Finally, the notion that endosomal compartments function as multifunctional
323 platforms on which unique sets of molecular machines are assembled have emerged
324 recently.⁷⁶ Therefore, *Shigella* induced macropinosomes may represent a platform for

325 localized production of specific signalling lipids and docking of certain protein complexes
326 promoting vacuolar scape.

327 **Conclusion and perspective**

328 In summary, as a result of Rab subversion, bacterial effectors induce or avoid
329 multiple interactions between the BCV and other host cell compartments modifying the
330 BCV or its membrane composition. Therefore, bacteria establish an intracellular replicative
331 niche through the generation of a sort of hybrid organelle.^{3,4}

332 Most of the currently described Rab modifications by pathogens lead to an
333 impairment of BCV degradation for bacteria survival. This is achieved either by blocking
334 phagosome maturation, by avoiding fusion with lysosomes or by redirecting BCV to
335 alternative trafficking pathways such as the ER. In addition, intracellular bacteria also
336 subvert vesicular trafficking in order to gain access to nutrients or scavenge building
337 blocks for intracellular replication.

338 From our findings about the cytosolic bacterium *Shigella* we propose that the
339 concept of Rab subversion is not an exclusive strategy for BCV-contained bacteria, but
340 also for cytosolic ones. We have discovered a new role for recycling endosomes, via
341 Rab11, during early stages of *Shigella* infection. This provides a different concept about
342 bacterial vacuolar rupture. Intriguingly, none of the different Rabs recruited to the entry foci
343 (Rab5, Rab7 and Rab11) were located on BCV membranes, but only on macropinosomes.
344 Further research on Rab manipulation by cytosolic bacteria will help to fully understand the
345 process of vacuolar rupture, a key event for their virulence.

346

347

348

349

350 **Acknowledgement**

351 This work was supported by the LabEx consortium IBEID, and by the Institut Pasteur
352 CARNOT-MIE programme. J.E also acknowledges support of an ERC starting grant
353 (Rupteffects, Nr. 261166).

354

355 **Disclosure of Potential Conflicts of Interest**

356 No potential conflicts of interest were disclosed.

357

358 **References**

- 359 1. Bhavsar AP, Guttman JA, Finlay BB. Manipulation of host-cell pathways by bacterial
360 pathogens. *Nature* 2007; 449:827-834.
- 361 2. Cossart P, Craig RR. Manipulation of host membrane machinery by bacterial
362 pathogens. *Curr Opin Cell Biol* 2010; 22:547-554.
- 363 3. Sherwood RK, Roy CR. A Rab-centric perspective of bacterial pathogen-occupied
364 vacuoles. *Cell Host Microbe* 2013 Sep; 14:256-268.
- 365 4. Stein MP, Müller MP, Wandinger-Ness A. Bacterial pathogens commandeer Rab
366 GTPases to establish intracellular niches. *Traffic* 2012; 13:1565-1588.
- 367 5. Zhen Y, Stenmark H. Cellular functions of Rab GTPases at a glance. *J Cell Sci* 2015;
368 128:3171-3176.
- 369 6. Zeigerer A, Gilleron J, Bogorad RL, Marsico G, Nonaka H, Seifert S, Epstein-Barash H,
370 Kuchimanchi S, Peng CG, Ruda VM, Del Conte-Zerial P, Hengstler JG, Kalaidzidis Y,
371 Koteliansky V, Zerial M. Rab5 is necessary for the biogenesis of the endolysosomal
372 system in vivo. *Nature* 2012; 485:465-470.
- 373 7. Shin HW, Hayashi M, Christoforidis S, Lacas-Gervais S, Hoepfner S, Wenk MR,
374 Modregger J, Uttenweiler-Joseph S, Wilm M, Nystuen A, Frankel WN, Solimena M, De

375 Camilli P, Zerial M. An enzymatic cascade of Rab5 effectors regulates phosphoinositide
376 turnover in the endocytic pathway. *J Cell Biol* 2005; 170:607-618.

377 8. Lippe R., Miaczynska M., Rybin V., Runge A., Zerial M. Functional synergy between
378 Rab5 effector Rabaptin-5 and exchange factor Rabex-5 when physically associated in a
379 complex. *Mol Biol Cell* 2001; 12:2219–2228.

380 9. Mottola G. The complexity of Rab5 to Rab7 transition guarantees specificity of pathogen
381 subversion mechanisms. *Front Cell Infect Microbiol* 2014; 4:180.

382 10. Rink J, Ghigo E, Kalaidzidis Y, Zerial M. Rab conversion as a mechanism of
383 progression from early to late endosomes. *Cell* 2005; 122:735-749.

384 11. Poteryaev D, Fares H, Bowerman B, Spang A. *Caenorhabditis elegans* SAND-1 is
385 essential for RAB-7 function in endosomal traffic. *EMBO J* 2007; 26:301-312.

386 12. Huotari J, Helenius A. Endosome maturation. *EMBO J* 2011; 30:3481-3500.

387 13. Harrison RE, Bucci C, Vieira OV, Schroer TA, Grinstein S. Phagosomes fuse with late
388 endosomes and/or lysosomes by extension of membrane protrusions along microtubules:
389 role of Rab7 and RILP. *Mol Cell Biol* 2003; 23:6494-6506.

390 14. Lairo K, Yliannala K, Ahtiainen L, Maunu H, Järvelä I, Kyttälä A, Jalanko A.
391 Interconnections of CLN3, Hook1 and Rab proteins link Batten disease to defects in the
392 endocytic pathway. *Hum Mol Genet* 2004; 13:3017-27.

393 15. Grant BD, Donaldson JG. Pathways and mechanisms of endocytic recycling. *Nat Rev*
394 *Mol Cell Biol* 2009; 10:597-608.

395 16. Welz T, Wellbourne-Wood J, Kerkhoff E. Orchestration of cell surface proteins by
396 Rab11. *Trends Cell Biol* 2014; 24:407-415.

397 17. Pylypenko O, Attanda W, Gauquelin C, Lahmani M, Coulibaly D, Baron B, Hoos S,
398 Titus MA, England P, Houdusse AM. Structural basis of myosin V Rab GTPase-dependent
399 cargo recognition. *Proc Natl Acad Sci U S A* 2013 Dec; 110:20443-20448.

400 18. Hales CM, Vaerman JP, Goldenring JR. Rab11 family interacting protein 2 associates
401 with Myosin Vb and regulates plasma membrane recycling. *J Biol Chem*; 277:50415-
402 50421.

403 19. Schonteich E, Wilson GM, Burden J, Hopkins CR, Anderson K, Goldenring JR,
404 Prekeris R. The Rip11/Rab11-FIP5 and kinesin II complex regulates endocytic protein
405 recycling. *J Cell Sci* 2008; 121:3824-3833.

406 20. Delevoye C, Miserey-Lenkei S, Montagnac G, Gilles-Marsens F, Paul-Gilloteaux P,
407 Giordano F, Waharte F, Marks MS, Goud B, Raposo G. Recycling endosome tubule
408 morphogenesis from sorting endosomes requires the kinesin motor KIF13A. *Cell Rep*
409 2014; 6:445-454.

410 21. Shirane M, Nakayama KI. Protrudin induces neurite formation by directional membrane
411 trafficking. *Science* 2006; 314:818-821.

412 22. Horgan CP, Hanscom SR, Jolly RS, Futter CE, McCaffrey MW. Rab11-FIP3 links the
413 Rab11 GTPase and cytoplasmic dynein to mediate transport to the endosomal-recycling
414 compartment. *J Cell Sci* 2010; 123:181-191

415 23. Takahashi S, Kubo K, Waguri S, Yabashi A, Shin HW, Katoh Y, Nakayama K. Rab11
416 regulates exocytosis of recycling vesicles at the plasma membrane. *J Cell Sci* 2012;
417 125:4049-4057.

418 24. Guichard A, Nizet V, Bier E. Rab11-mediated trafficking in host-pathogen interactions.
419 *Nat Rev Microbiol* 2014; 12:624-634.

420 25. Xiong B, Bayat V, Jaiswal M, Zhang K, Sandoval H, Charng WL, Li T, David G,
421 Duraine L, Lin YQ, Neely GG, Yamamoto S, Bellen HJ. Crag is a GEF for Rab11 required
422 for rhodopsin trafficking and maintenance of adult photoreceptor cells. *PLoS Biol* 2012;
423 10:e1001438.

424 26. Das S, Hehnly H, Doxsey S. A new role for Rab GTPases during early mitotic stages.
425 Small GTPases 2014; 5:e29565.

426 27. Knowles BC, Weis VG, Yu S, Roland JT, Williams JA, Alvarado GS, Lapierre LA, Shub
427 MD, Gao N, Goldenring JR. Rab11a regulates syntaxin 3 localization and microvillus
428 assembly in enterocytes. J Cell Sci 2015; 128:1617-1626.

429 28. Szatmári Z, Kis V, Lippai M, Hegedus K, Faragó T, Lorincz P, Tanaka T, Juhász G,
430 Sass M. Rab11 facilitates cross-talk between autophagy and endosomal pathway through
431 regulation of Hook localization. Mol Biol Cell 2014; 25:522-531.

432 29. Mizuno-Yamasaki E, Rivera-Molina F, Novick P. GTPase networks in membrane
433 traffic. Annu Rev Biochem 2012; 81:637-659.

434 30. Mallo GV, Espina M, Smith AC, Terebiznik MR, Alemán A, Finlay BB, Rameh LE,
435 Grinstein S, Brumell JH. SopB promotes phosphatidylinositol 3-phosphate formation on
436 Salmonella vacuoles by recruiting Rab5 and Vps34. J Cell Biol 2008; 182:741-752.

437 31. Mukherjee K, Parashuraman S, Raje M, Mukhopadhyay A. SopE acts as an Rab5-
438 specific nucleotide exchange factor and recruits non-prenylated Rab5 on Salmonella-
439 containing phagosomes to promote fusion with early endosomes. J Biol Chem 2001;
440 276:23607-23615.

441 32. Fratti RA, Backer JM, Gruenberg J, Corvera S, Deretic V. Role of phosphatidylinositol
442 3-kinase and Rab5 effectors in phagosomal biogenesis and mycobacterial phagosome
443 maturation arrest. J Cell Biol 2001; 154:631-644.

444 33. Vergne I, Chua J, Lee HH, Lucas M, Belisle J, Deretic V. Mechanism of
445 phagolysosome biogenesis block by viable Mycobacterium tuberculosis. Proc Natl Acad
446 Sci U S A 2005; 102:4033-4038.

447 34. Ku B, Lee KH, Park WS, Yang CS, Ge J, Lee SG, Cha SS, Shao F, Heo WD, Jung JU,
448 Oh BH. VipD of *Legionella pneumophila* targets activated Rab5 and Rab22 to interfere
449 with endosomal trafficking in macrophages. *PLoS Pathog* 2012; 8:e1003082.

450 35. Gaspar AH, Machner MP. VipD is a Rab5-activated phospholipase A1 that protects
451 *Legionella pneumophila* from endosomal fusion. *Proc Natl Acad Sci U S A* 2014;
452 111:4560-4565.

453 36. Brumell JH, Goosney DL, Finlay BB. SifA, a type III secreted effector of *Salmonella*
454 *typhimurium*, directs *Salmonella*-induced filament (Sif) formation along microtubules.
455 *Traffic* 2002; 3:407-415.

456 37. McGourty K, Thurston TL, Matthews SA, Pinaud L, Mota LJ, Holden DW. *Salmonella*
457 inhibits retrograde trafficking of mannose-6-phosphate receptors and lysosome function.
458 *Science* 2012; 338:963-967.

459 38. Zhao W1, Moest T1, Zhao Y1, Guilhon AA1, Buffat C2, Gorvel JP1, Méresse S. The
460 *Salmonella* effector protein SifA plays a dual role in virulence. *Sci Rep* 2015; 5:12979.

461 39. D'Costa VM, Braun V, Landekic M, Shi R, Proteau A, McDonald L, Cygler M, Grinstein
462 S7, Brumell JH. *Salmonella* Disrupts Host Endocytic Trafficking by SopD2-Mediated
463 Inhibition of Rab7. *Cell Rep* 2015; 12:1508-1518.

464 40. Murata T, Delprato A, Ingmundson A, Toomre DK, Lambright DG, Roy CR. The
465 *Legionella pneumophila* effector protein DrrA is a Rab1 guanine nucleotide-exchange
466 factor. *Nat Cell Biol* 2006; 8:971-977.

467 41. Müller MP, Peters H, Blümer J, Blankenfeldt W, Goody RS, Itzen A. The *Legionella*
468 effector protein DrrA AMPylates the membrane traffic regulator Rab1b. *Science* 2010;
469 329:946-949.

470 42. Tan Y, Luo ZQ. *Legionella pneumophila* SidD is a deAMPylase that modifies Rab1.
471 *Nature* 2011; 475:506-509.

472 43. Machner MP, Isberg RR. A bifunctional bacterial protein links GDI displacement to
473 Rab1 activation. *Science* 2007; 318:974–977.

474 44. Gazdag EM, Streller A, Haneburger I, Hilbi H, Vetter IR, Goody RS, Itzen A.
475 Mechanism of Rab1b deactivation by the *Legionella pneumophila* GAP LepB. *EMBO Rep*
476 2013; 14:199-205.

477 45. Tan Y, Arnold RJ, Luo ZQ. *Legionella pneumophila* regulates the small GTPase Rab1
478 activity by reversible phosphorylcholation. *Proc Natl Acad Sci U S A* 2011; 108:21212-
479 21217.

480 46. Alvarez-Dominguez C, Madrazo-Toca F, Fernandez-Prieto L, Vandekerckhove J,
481 Pareja E, Tobes R, Gomez-Lopez MT, Del Cerro-Vadillo E, Fresno M, Leyva-Cobián F,
482 Carrasco-Marín E. Characterization of a *Listeria monocytogenes* protein interfering with
483 Rab5a. *Traffic* 2008; 9:325-337.

484 47. Spanò S, Galán JE. A Rab32-dependent pathway contributes to *Salmonella typhi* host
485 restriction. *Science* 2012; 338:960-963.

486 48. Spanò S, Gao X, Hannemann S, Lara-Tejero M, Galán JE. A Bacterial Pathogen
487 Targets a Host Rab-Family GTPase Defense Pathway with a GAP. *Cell Host Microbe*
488 2016; 19:216-226.

489 49. Parkhill J, Dougan G, James KD, Thomson NR, Pickard D, Wain J, Churcher C,
490 Mungall KL, Bentley SD, Holden MT, Sebahia M, Baker S, Basham D, Brooks K,
491 Chillingworth T, Connerton P, Cronin A, Davis P, Davies RM, Dowd L, White N, Farrar J,
492 Feltwell T, Hamlin N, Haque A, Hien TT, Holroyd S, Jagels K, Krogh A, Larsen TS, Leather
493 S, Moule S, O'Gaora P, Parry C, Quail M, Rutherford K, Simmonds M, Skelton J, Stevens
494 K, Whitehead S, Barrell BG. Complete genome sequence of a multiple drug resistant
495 *Salmonella enterica* serovar Typhi CT18. *Nature* 2001; 413:848-852.

496 50. Spanò S, Liu X, Galán JE. Proteolytic targeting of Rab29 by an effector protein
497 distinguishes the intracellular compartments of human-adapted and broad-host
498 Salmonella. Proc Natl Acad Sci U S A 2011; 108:18418-18423.

499 51. de Barsy M, Jamet A, Filopon D, Nicolas C, Laloux G, Rual JF, Muller A, Twizere JC,
500 Nkengfac B, Vandenhaute J, Hill DE, Salcedo SP, Gorvel JP, Letesson JJ, De Bolle X.
501 Identification of a Brucella spp. secreted effector specifically interacting with human small
502 GTPase Rab2. Cell Microbiol 2011; 13:1044-1058.

503 52. Rejman Lipinski A, Heymann J, Meissner C, Karlas A, Brinkmann V, Meyer TF, Heuer
504 D. Rab6 and Rab11 regulate Chlamydia trachomatis development and golgin-84-
505 dependent Golgi fragmentation. PLoS Pathog 2009; 5:e1000615.

506 53. Capmany A, Damiani MT. Chlamydia trachomatis intercepts Golgi-derived
507 sphingolipids through a Rab14-mediated transport required for bacterial development and
508 replication. PLoS ONE 2010; 5:e14084.

509 54. Gambarte Tudela J, Capmany A, Romao M, Quintero C, Miserey-Lenkei S, Raposo G,
510 Goud B, Damiani MT. The late endocytic Rab39a GTPase regulates the interaction
511 between multivesicular bodies and chlamydial inclusions. J Cell Sci 2015; 128:3068-3081.

512 55. Connor MG, Pulsifer AR, Price CT, Abu Kwaik Y, Lawrenz MB. Yersinia pestis
513 Requires Host Rab1b for Survival in Macrophages. PLoS Pathog 2015; 11:e1005241.

514 56. Dong N, Zhu Y, Lu Q, Hu L, Zheng Y, Shao F. Structurally distinct bacterial TBC-like
515 GAPs link Arf GTPase to Rab1 inactivation to counteract host defenses. Cell 2012;
516 150:1029-1041.

517 57. Burnaevskiy N, Fox TG, Plymire DA, Ertelt JM, Weigele BA, Selyunin AS, Way SS,
518 Patrie SM, Alto NM. Proteolytic elimination of N-myristoyl modifications by the Shigella
519 virulence factor IpaJ. Nature 2013; 496:106-109.

520 58. Ray K, Bobard A, Danckaert A, Paz-Haftel I, Clair C, Ehsani S, Tang C, Sansonetti P,
521 Tran GV, Enninga J. Tracking the dynamic interplay between bacterial and host factors
522 during pathogen-induced vacuole rupture in real time. *Cell Microbiol* 2010; 12:545-556.

523 59. Ray K, Marteyn B, Sansonetti PJ, Tang CM. Life on the inside: the intracellular lifestyle
524 of cytosolic bacteria. *Nat Rev Microbiol* 2009; 7:333–340.

525 60. Mellouk N, Enninga J. Cytosolic Access of Intracellular Bacterial Pathogens: The
526 *Shigella* Paradigm. *Front Cell Infect Microbiol* 2016; 6:35.

527 61. High N, Mounier J, Prévost MC, Sansonetti PJ. IpaB of *Shigella flexneri* causes entry
528 into epithelial cells and escape from the phagocytic vacuole. *EMBO J* 1992; 11:1991-1999.

529 62. Du J, Reeves AZ, Klein JA, Twedt DJ, Knodler LA, Lesser CF. The type III secretion
530 system apparatus determines the intracellular niche of bacterial pathogens. *Proc Natl*
531 *Acad Sci U S A* 2016; 113:4794-4799.

532 63. Mellouk N, Weiner A, Aulner N, Schmitt C, Elbaum M, Shorte SL, Danckaert A,
533 Enninga J. *Shigella* subverts the host recycling compartment to rupture its vacuole.
534 *Cell Host Microbe* 2014; 16:517-530.

535 64. Weiner A, Mellouk N, Lopez-Montero N, Chang YY, Souque C, Schmitt C, Enninga J.
536 Macropinosomes are key players in early *Shigella* invasion and vacuolar escape in
537 epithelial cells. *PLoS Pathog* 2016; 12:e1005602.

538 65. Nothelfer K, Dias Rodrigues C, Bobard A, Phalipon A, Enninga J. Monitoring *Shigella*
539 *flexneri* vacuolar escape by flow cytometry. *Virulence* 2011; 2:54-57.

540 66. Keller C, Mellouk N, Danckaert A, Simeone R, Brosch R, Enninga J, Bobard A. Single
541 cell measurements of vacuolar rupture caused by intracellular pathogens. *J Vis Exp* 2013;
542 12:e50116.

543 67. Niebuhr K, Giuriato S, Pedron T, Philpott DJ, Gaits F, Sable J, Sheetz MP, Parsot C,
544 Sansonetti PJ, Payraastre B. Conversion of PtdIns(4,5)P(2) into PtdIns(5)P by the *S.flexneri*
545 effector IpgD reorganizes host cell morphology. *EMBO J* 2002; 21:5069-5078.

546 68. Niebuhr K, Jouihri N, Allaoui A, Gounon P, Sansonetti PJ, Parsot C. IpgD, a protein
547 secreted by the type III secretion machinery of *Shigella flexneri*, is chaperoned by IpgE
548 and implicated in entry focus formation. *Mol Microbiol* 2000; 38:8-19.

549 69. Lim JP, Gleeson PA. Macropinocytosis: an endocytic pathway for internalising large
550 gulps. *Immunol Cell Biol* 2011; 89:836-843.

551 70. Racoosin EL, Swanson JA. Macropinosome maturation and fusion with tubular
552 lysosomes in macrophages. *J Cell Biol* 1993; 121:1011-1020.

553 71. Hamasaki M1, Araki N, Hatae T. Association of early endosomal autoantigen 1 with
554 macropinocytosis in EGF-stimulated A431 cells. *Anat Rec A Discov Mol Cell Evol Biol*
555 2004; 277:298-306.

556 72. Cossart P, Sansonetti PJ. Bacterial invasion: the paradigms of enteroinvasive
557 pathogens. *Science* 2004; 304:242–248.

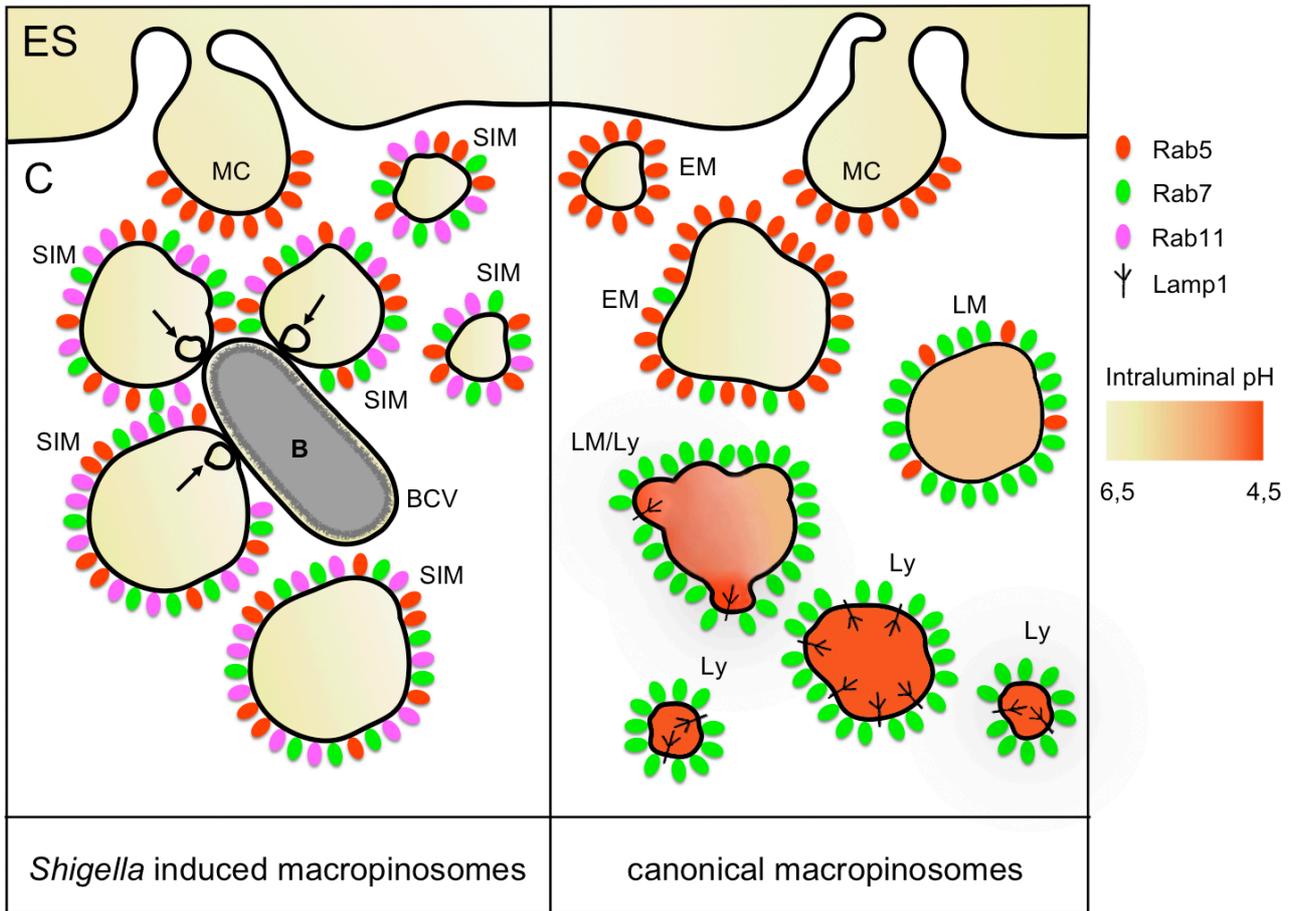
558 73. Murphy GE, Narayan K, Lowekamp BC, Hartnell LM, Heymann JA, Fu J,
559 Subramaniam S. Correlative 3D imaging of whole mammalian cells with light and electron
560 microscopy. *J Struct Biol* 2011; 176:268–78.

561 74. Los FC, Kao CY, Smitham J, McDonald KL, Ha C, Peixoto CA, Aroian RV. RAB-5- and
562 RAB-11-dependent vesicle-trafficking pathways are required for plasma membrane repair
563 after attack by bacterial pore-forming toxin. *Cell Host Microbe* 2011; 9:147-157.

564 75. Jimenez AJ, Maiuri P, Lafaurie-Janvore J, Divoux S, Piel M, Perez F. ESCRT
565 machinery is required for plasma membrane repair. *Science* 2014; 343:1247136.

566 76. Gould GW1, Lippincott-Schwartz J. New roles for endosomes: from vesicular carriers
567 to multi-purpose platforms. *Nat Rev Mol Cell Biol* 2009; 10:287-292.

Pathogen	Effector	Mechanism of subversion	Effect on Rabs	Advantages for infection	References
<i>Salmonella enterica</i>	SopB	Inositol phosphate phosphatase. Converts PI(3) to PI(3)P	Prolongs and increases Rab5 recruitment on the BCV.	Delays interaction of BCVs with late endosomes	Mialo et al. (2008)
	SopE	GEF for Rab5. Acts on non-prenylated Rab5	Prolongs and increases Rab5 recruitment on the BCV.	Delays interaction of BCVs with late endosomes	Mukherjee et al. (2001)
	SfIA	Forms a complex with SKIP, which binds Rab9	Induces recruitment of Rab9 to BCVs and SIFs	Prevents the delivery of hydrolytic enzymes to the BCV	Burwell et al. (2002) McGourry et al. (2012)
<i>Salmonella enterica</i> sv Typhimurium	GIGE	Rab32 protease	Avoids Rab32 recruitment to the BCV	Prevents the delivery of antimicrobial molecules to the BCV	Spano and Galan (2012)
	SopD2	GAP for Rab32	Contributes to Rab32 removal from BCV membranes	Prevents the delivery of antimicrobial molecules to the BCV	Spano et al. (2016)
	SopD2	Binds to Rab7	Stabilizes Rab7 on SCVs and inhibits GTPase cycling.	Contributes to the evasion of lysosomal degradation, promotes the formation of SIFs	D'Costa et al. (2015)
<i>Mycobacterium tuberculosis</i>	LAM	Inactivates sPI3PKC3 inhibiting PIP production	Impairs recruitment of Rab5 to the BCV	Arrests phagosome maturation and degradation	Fraiti et al. (2001)
	SapM	Acid phosphatase. Dephosphorylates PISp	Depletes PISp on BCVs avoiding Rab5 recruitment	Arrests phagosome maturation and degradation	Vergne et al. (2005)
	VipD	Binds to Rab5 and Rab22	Prevents interaction of Rab5 and Rab22 with their effectors	Decreases bacterial degradation by general impairment of endosomal maturation	Ku et al. (2012) Gaepser et al. (2014)
<i>Legionella pneumophila</i>	DrpA	GEF for Rab1. Modifies Rab1 by AMPylation	AMPylated Rab1 is inert to GAPs and becomes constitutively active on BCVs	Establishment of a replicative niche with ER characteristics. Tight modulation of the different effectors is necessary to control BCV trafficking during the different phases of infection	Murata et al. (2006)
	SidD	DeAMPylates Rab1	Opposed function of DrpA		Tan and Luo (2011)
	LdaA	Binds GDI-free Rab1 previously activated by DrpA	Involved in accumulating activated Rab1 on BCVs		Machner and Isberg (2007)
	LeppB	GAP for Rab1	Deactivates Rab1 on the BCV		Gazdag et al. (2013)
	AnkX	Transfers phosphocholine to Rab1	Phosphocholinated Rab1 is inert to GEFs and GDBs on BCV membranes		Tan et al. (2011)
<i>Listeria monocytogenes</i>	GADPH	ADP-ribosylates Rab5	Recruits Rab5 to BCVs and inhibits the activation of GEFs	Blocks maturation of the phagosomes	Alvarez-Dominguez et al. (2008)
<i>Brucella abortus</i>	RicaA	Binds to Rab2	Tethers Rab2 to the BCV	Helps to establish a replicative BCV with ER characteristics	de Batsy et al. (2011)
<i>Chlamydia trachomatis</i>	?	?	Recruitment of Rabs, Rab11, Rab14 and Rab39 to the BCV	Scavenges sphingomyelin for bacterial nutrition from Golgi and MVBs	Felman Ljinski et al. (2009) Castrany and Darian (2010) Gambarte Tudea et al. (2015)
<i>Yersinia pestis</i>	?	?	Recruitment of Rab1 to the BCV	Avoids acidification of BCVs to establish a replicative niche	Connor et al. (2015)
<i>Shigella flexneri</i>	Vf/A	GAP for Rab1	Deactivates Rab1 on Rab1 positive endosomes	Inhibits ER-to-Golgi transport and impairs autophagy	Dong et al. (2012)
	IpgD	Inositol phosphate phosphatase. Converts PI(4,5)P to PI3P	Induces the recruitment of Rab11 to macropinosomes	Enhances vacuolar rupture and bacterial escape into the cytosol	Melouk et al. (2014) Weiner et al. (2016)



569 **Figure 1: Schematic representation of macropinosome-like vesicles induced by**
 570 ***Shigella* infection in comparison to the canonical macropinocytic pathway.** On the
 571 right side, canonical macropinosomes traffic along the endolysosomal pathway, where they are
 572 eventually degraded. Rab5 (red) is first recruited to nascent macropinosomes, similar to its
 573 recruitment to early endosomes. Then, it is replaced by Rab7 (green). Acquisition of Rab7 implies
 574 retrograde transport along microtubules and subsequent fusion with lysosomes. Upon fusion with
 575 lysosomes, macropinosomes acquire lysosomal markers such as Lamp1 and hydrolytic enzymes
 576 leading to their acidification. In contrast, macropinosome-like vesicles induced by *Shigella*
 577 infection block their maturation before their fusion with lysosomes (left side). Rab11 (magenta) is
 578 instead recruited by the bacterial effector IpgD. Bacterial subversion of Rab11, and its recruitment
 579 to *Shigella*-induced macropinosomes, promotes efficient vacuolar escape. ES, extracellular space;
 580 C, cytosol; MC, macropinocytic cup; SIM, *Shigella* induced macropinosome; BCV, bacteria
 581 containing vacuole; B, bacteria; EM, early macropinosome; LM, late macropinosome; LM/Ly, late
 582 macropinosome-lysosome; Ly, lysosome; arrow, intramacropinosomal vesicle.