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Yersinia pseudotuberculosis YopH targets SKAP2-dependent and independent signaling pathways to block neutrophil antimicrobial mechanisms during infection

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Abstract

Yersinia suppress neutrophil responses by using a type 3 secretion system (T3SS) to inject 6-7 Yersinia effector proteins (Yops) effectors into their cytoplasm. YopH is a tyrosine phosphatase that causes dephosphorylation of the adaptor protein SKAP2, among other targets in neutrophils. SKAP2 functions in reactive oxygen species (ROS) production, phagocytosis, and integrin-mediated migration by neutrophils. Here we identify essential neutrophil functions targeted by YopH, and investigate how the interaction between YopH and SKAP2 influence Yersinia pseudotuberculosis (Yptb) survival in tissues. The growth defect of a $\Delta yopH$ mutant was restored in mice defective in the NADPH oxidase complex, demonstrating that YopH is critical for protecting Yptb from ROS during infection. The growth of a $\Delta yopH$ mutant was partially restored in Skap2-deficient (Skap2KO) mice compared to wildtype (WT) mice, while induction of neutropenia further enhanced the growth of the $\Delta yopH$ mutant in both WT and Skap2KO mice. YopH inhibited both ROS production and degranulation triggered via integrin receptor, G-protein coupled receptor (GPCR), and Fcy receptor (FcyR) stimulation. SKAP2 was required for integrin receptor and GPCR-mediated ROS production, but dispensable for degranulation under all conditions tested. YopH blocked SKAP2-independent FcyR-stimulated phosphorylation of the proximal signaling proteins Syk, SLP-76, and PLC_Y2, and the more distal signaling protein ERK1/2, while only ERK1/2 phosphorylation was dependent on SKAP2 following integrin receptor activation. These findings reveal that YopH prevents activation of both SKAP2-dependent and -independent neutrophilic defenses, uncouple integrin- and GPCR-dependent ROS production from FcyR responses based on their SKAP2 dependency, and show that SKAP2 is not required for degranulation.

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Author summary

Pathogenic *Yersinia* species carry a virulence plasmid encoding a type 3 secretion system that translocates 6–7 effector Yops into host cells. We demonstrate that YopH protects *Yersinia pseudotuberculosis* from neutrophil-produced reactive oxygen species (ROS) and degranulation by interfering with signaling pathways downstream of three major receptor classes in neutrophils. We show that a previously identified target of YopH, SKAP2, controls some of the pathways essential for YopH to inactivate during infection. SKAP2 is essential in mediating ROS production downstream of two major receptors; however, it is dispensable for degranulation from the three major receptors tested. Our study illustrates that YopH protects *Y. pseudotuberculosis* by blocking both SKAP2-dependent and independent signaling pathways that regulate several neutrophil functions.

Introduction

The three pathogenic species of Yersinia that are associated with human infections are the causative agent of the pneumonic and bubonic plague, Yersinia pestis, and the gastrointestinal pathogens Y. pseudotuberculosis (Yptb) and Y. enterocolitica [1-3]. These species harbor a 70kb virulence plasmid encoding a Type 3 Secretion System (T3SS) that injects a set of 6–7 effector proteins, called Yops (Yersinia outer proteins), into mammalian cells [1, 4–7]. Yops are essential for virulence and interfere with many antimicrobial responses in immune cells, yet their functions in tissue infections are not completely understood [5, 8-11]. Infection with *Yptb* in tissues is characterized by formation of *Yptb* microcolonies that directly interface with neutrophils that have migrated to sites of infection [12–14]. Consistent with their physical proximity to Yptb microcolonies in tissues, neutrophils are typically the cells that are most frequently injected with Yops [15–17]. Yptb encodes two bacterial adhesins, Invasin and YadA, which bind either directly or indirectly to $\beta 1$ integrins expressed on neutrophils [18, 19]. This binding facilitates Yop-injection into neutrophils during tissue infection, whereas in the absence of Yops, the binding via these adhesins triggers phagocytosis of *Yptb* [15, 20, 21]. In addition to integrin receptor-mediated interactions, Yersinia activates other receptors on neutrophils during infection, including G-protein coupled receptors (GPCRs) [22-25]. Activation of integrin receptors and GPCRs in turn leads to a variety of neutrophil antimicrobial responses including reactive oxygen species (ROS) production and proteolytic enzyme release from granule stores, also known as degranulation.

The effector, YopH, is critical for the colonization and virulence of all three pathogenic *Yersinia* species, is homologous to eukaryotic protein tyrosine phosphatases (PTPase), and is the most active of all known PTPases [5, 26-35]. Growth of a *Yptb ΔyopH* mutant is significantly restored in the absence of neutrophils, indicating that YopH targets antimicrobial functions of neutrophils [36]. In isolated human or murine neutrophils YopH inhibits zymosan-induced ROS production, calcium flux, phagocytosis, and cytokine production [36-39], and together with YopE blocks human neutrophil degranulation [40-42]. While these findings imply potential roles for YopH in modulating neutrophil functions during tissue infection, the specific signal-transduction pathway(s) that are necessary for YopH to inactivate for *Yptb* virulence in animal tissues remains unknown. To begin to address this question, some tyrosine-phosphorylated signal transduction pathway(s) disrupted by YopH during animal infection were previously identified [36]. In neutrophils isolated from spleens of infected mice, the adaptor proteins Src Kinase Associated Phosphoprotein 2 (SKAP2), PML-RARA Regulated Adaptor Molecule 1 (PRAM-1) and SH2 domain containing leukocyte protein of 76kDa (SLP-76), and

the effector proteins, Vav guanine nucleotide exchange factor 1 (Vav1) and Phospholipase C γ 2 (PLC γ 2), are dephosphorylated in the presence of YopH [36]. Previous work in macrophages has shown that SKAP2 and Adhesion and Degranulation Adaptor Protein (ADAP), a homolog of PRAM-1 [43], are co-immunoprecipitated by YopH [44]. These results suggest that SKAP2-controlled pathways may be a critical target of YopH in neutrophils during tissue infection.

SKAP2 is a cytosolic adaptor protein comprised of a N-terminal coiled-coil dimerization domain region, a proximal pleckstrin homology domain (PH), an interdomain containing two tyrosine-based signaling motifs and a C-terminal Src homology 3 (SH3) domain that mediates its binding to PRAM-1 and ADAP [45–48]. SKAP2 is a substrate for Scr kinases, and was recently found to be essential in regulating TNF α –, selectin- and CXCL1-mediated integrin activation [49]. In addition, SKAP2 regulates neutrophil functions including migration to sites of sterile inflammation, neutrophil extracellular trap (NET) formation, and integrin-mediated ROS production [49].

ROS generated by the activated NADPH oxidase complex occurs after stimulation of various surface receptors, including integrin, GPCR, and Fc receptors [49-55] that triggers assembly and activation of the NADPH oxidase complex at the plasma or phagosomal membrane. The gp91^{phox} subunit is required for the assembly of a functional NADPH oxidase complex and subsequent superoxide production [56]. ROS can act directly as an antimicrobial agent by causing oxidative damage to DNA molecules and destroying pathogen membrane structures by lipid peroxidation. It can also activate other neutrophil antimicrobial functions such as phagocytosis [57–59], cytokine release [60–62], NET formation [63–65], and degranulation [57, 66].

Here we investigate whether YopH is required to prevent ROS production during *Yptb* infection of mouse tissues and whether SKAP2 must be inactivated for a $\Delta yopH$ mutant to survive and thus is an important YopH target. We find that while inactivation of SKAP2-regulated pathways is a key function of YopH, YopH has additional functions independent of SKAP2 that enable further inhibition of both neutrophil ROS production and degranulation. Our studies also shed light on SKAP2 function, demonstrating its essential role in neutrophil ROS activation after integrin or GPCR-stimulation, but not in degranulation or migration of neutrophils to infected tissues sites.

Results

Growth of a $\Delta yopH$ mutant is restored in mice lacking a functional NADPH oxidase complex and YopH is sufficient to block ROS production from murine neutrophils

Our previous work [36] demonstrated that several signaling proteins that are important for ROS production downstream of integrin signaling, including SKAP2, PRAM-1, SLP-76, Vav1, and PLC γ 2 [49–54], were hypo-phosphorylated in YopH-injected neutrophils isolated from infected spleens. However, whether YopH plays a role in supporting *Yptb* growth during infection by suppressing ROS production is unknown. To test this, C57BL/6J and *gp91^{phox-/-}* mice were infected intravenously (I.V.) or intratracheally (I.T.) with an equal mixture of IP2666 WT-*Yptb* and a *ΔyopH* mutant. Two days post-infection spleens and lungs were harvested and the competitive index (C.I.) was determined. By contrast with previous work on *ΔyopE* [67], the *ΔyopH* mutant competed significantly better in the spleens and lungs of *gp91^{phox-/-}* mice than C57BL/6J mice (Fig 1A and 1B). The total CFU of *Yptb* recovered from *gp91^{phox-/-}* mice were comparable to numbers recovered from C57BL/6J mice, but the numbers of the recovered *ΔyopH* mutant rose significantly in the *gp91^{phox-/-}* mice (S1A Fig). These results support the idea YopH inactivates one or more pathways leading to the generation of ROS and thereby protects *Yptb* from superoxide-mediated killing during tissue infection.



Fig 1. The growth of a $\Delta yopH$ mutant is restored in absence of ROS *in vivo* and YopH is necessary and sufficient to block extracellular ROS produced by BM neutrophils *in vitro*. (A-B) Female C57BL/6J or C57BL/6J gp91^{phox-/-} mice were inoculated with a 1:1 mixture of IP2666 WT-*Yptb* and $\Delta yopH$ -Kan^R (A) intravenously with 1x10³ CFU or (B) intratracheally with 5x10³ CFU. Spleens and lungs were collected, weighed, homogenized, and plated for CFU two days post- infection. The competitive indices (C.I.) were determined by patching 100 colonies on L-Irgasan and L-Irgasan+Kanamycin plates. Each symbol represents an individual mouse; horizontal bars represent the geometric mean. (C-F) BM neutrophils were infected at a MOI of 20:1 with (C-D) WT-*Yptb*, $\Delta yopH$, or $\Delta 5$ (a mutant lacking 5 effector Yops, H, E, O, M, J), (E-F) WT-*Yptb* +pBAD, $\Delta 5$ +YopH (a strain expressing only YopH at native levels from an arabinose inducible plasmid), or $\Delta 5$ +pBAD, and monitored for extracellular ROS production using isoluminol chemiluminescence assay. (D and F) Total chemiluminescence was determined by computing the area under the curve (AUC) using GraphPad Prism version 7. Mean ±SD of a representative experiment from 2–3 independent experiments done in 2–3 replicates is presented. Statistical significance was calculated using (A) the Mann-Whitney t-test, (B) one-way ANOVA followed by Tukey's Post-test after log₁₀ transformation of the data, or (D, F) one-way ANOVA followed by Tukey's Post-test for area under the curve (AUC).

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To examine the ability of YopH to block ROS production in neutrophils, extracellular ROS production was measured after *Yptb* infection of neutrophils isolated from bone marrow (BM) of BALB/c mice (S1B Fig). BM neutrophils were infected with WT-*Yptb*, $\Delta yopH$, $\Delta 5$ (a mutant lacking 5 effector Yops, H, E, O, M, J) or $\Delta 5$ +pYopH, a strain expressing only YopH at native levels from an arabinose inducible plasmid (S1 Table). There were no differences in amount of YopH secretion between $\Delta 5$ +pYopH and WT-*Yptb* (S2 Fig). YopH was necessary to block ROS production induced by WT-*Yptb* infection (Fig 1C and 1D) because infection with the $\Delta yopH$ strain generated high levels of ROS that were equivalent to infection with the $\Delta 5$ strain (Fig 1D). In addition, YopH was the only Yop needed to block ROS production as much as WT-*Yptb* infection (Fig 1F). Combined, these results indicate that YopH blocks production of extracellular ROS by neutrophils induced by *Yptb* infection.

Growth of a $\Delta yopH$ mutant in the spleen is partially restored in the absence of SKAP2 and is further enhanced in neutropenic mice

SKAP2 is essential for integrin-stimulated ROS production in neutrophils [49], is dephosphorylated in neutrophils injected with YopH during tissue infection [36], and associates with



Fig 2. *Skap2KO* neutrophils cluster to *Yptb* microcolonies and the growth of a *AyopH* mutant is partially restored in competition with WT-*Yptb* in *Skap2KO* mice. (A) WT-BALB/c and *Skap2KO* mice were injected with 1A8 or an isotype control antibody and 16 hrs later I.V. infected with an equal mixture of $1x10^3$ CFU of IP2666 WT-*Yptb* and $\Delta yopH$ -Kan^R. Spleens were collected three days post-infection, weighed, homogenized, and plated for CFU on selective and non-selective agar. The number of bacteria recovered from selective and non-selective plates was used to determine the C.I. Each dot represents a mouse; horizontal bars represent the geometric mean. Significance was calculated using 2-way ANOVA followed by Tukey's Post-test of log₁₀ transformed values, with only significant values shown. The results are a composite of three experiments. (B-I) BALB/c and *Skap2KO* mice were (B-C) uninfected or (D-I) I.V. infected with 10^3 CFU of IP2666 WT-*Yptb*-GFP (green), and sacrificed at (D-E) 24, (F-G) 33, or (H-I) 48 hours post-infection. Frozen sections of spleens were stained using an anti-Ly6G antibody (neutrophils-red) and DAPI (nuclei-blue) and visualized by fluorescence confocal microscopy at 40X magnification. (J-K) Using Volocity software, the size of the (J) bacterial microcolony was determined by generating the summed volume of the individual GFP signal and the (K) number of neutrophils reruited was determined by detecting the signal intensity of the mCherry channel. Each dot represents a microcolony or an area of clustered neutrophils. The horizontal solid bars represent the median neutrophil signal intensity in uninfected controls of BALB/c and *Skap2KO* mice. Scale bars: 50 µm. Statistical significance was determined (J-K) using Mann-Whitney for comparing BALB/c at *Skap2KO* at each time point and Kruskal-Wallis for comparisons between the three time points for each mouse genotype. The results are a composite of 2-3 mice from two independent experiments.

YopH in macrophages [44]. These observations, combined with the observations that *Yptb* binds to neutrophils via integrins [15, 20, 68, 69] and that YopH blocks ROS production (Fig 1), led us to evaluate whether SKAP2 influences *Yptb* survival in the presence and absence of YopH. BALB/c (WT) and BALB/c-*Skap2KO* (*Skap2KO*) mice were co-infected I.V. with an equal mixture of WT-*Yptb* and a Δ *yopH* mutant. Three days post-infection spleens were collected, and the C.I. was determined. The Δ *yopH* mutant colonized *Skap2KO* mice almost 10-fold more efficiently compared to WT mice while overall bacterial burdens were similar (Fig 2A, S3A Fig), suggesting that SKAP2 may function in a YopH sensitive pathway(s) that serves to limit growth of a Δ *yopH* mutant *in vivo*. Although, *Skap2KO* mice have normal levels of circulating neutrophils [70], their extravasation into tissues occurs at significantly lower rates in a sterile model of inflammation [49]. Thus, it was possible that the improved survival of the Δ *yopH* mutant in the *Skap2KO* mice was due to reduced rates of neutrophil recruitment to infected sites. Analysis of the kinetics of neutrophil recruitment in the spleens of *Yptb*-

infected WT and *Skap2KO* mice revealed no differences in the recruitment of neutrophils and bacterial burdens were comparable between the WT and *Skap2KO* mice at the tested time points (Fig 2B–2K, S3B Fig). Collectively, these results support the idea that the restoration of Δ *yopH* mutant growth in *Skap2KO* mice (Fig 2A) is due to a functional defect in *Skap2*-deficient neutrophils that increases the survival of a Δ *yopH* mutant. This suggests that SKAP2 mediates signaling from receptors stimulated by *Yptb* infection and participates in a subset of relevant antimicrobial neutrophil functions normally targeted by YopH.

The $\Delta yopH$ mutant was still defective for survival compared to WT-*Yptb* in *Skap2KO* mice suggesting that *Skap2KO* neutrophils may retain some killing activities against the $\Delta yopH$ mutant. To determine whether YopH targets pathways, which are independent of SKAP2, in neutrophils, additional cohorts of mice from each genotype were subjected to 1A8 antibodymediated depletion of neutrophils. Growth of the $\Delta yopH$ mutant was significantly higher in neutrophil-depleted *Skap2KO* mice compared to *Skap2KO* mice (Fig 2A), indicating that *Skap2KO* neutrophils retain some antimicrobial activities and that YopH targets additional SKAP2-independent pathways in neutrophils. Notably, $\Delta yopH$ growth was comparable in neutrophil-depleted WT and neutrophil-depleted *Skap2KO* mice, highlighting that the lack of SKAP2 activity in other innate immune cells does not enhance survival of the $\Delta yopH$ mutant in the absence of neutrophils. Thus, YopH targets both SKAP2-dependent and SKAP2-independent antimicrobial processes.

YopH blocks SKAP2-dependent and -independent ROS production by neutrophils

Since $\Delta yopH$ growth was restored in $gp91^{phox-/-}$ mice we evaluated the ability of YopH to block ROS production downstream of three major receptor classes (integrin receptors, G- protein coupled receptor (GPCR) and FcyR) known to activate the NADPH oxidase complex, two of which may be stimulated during *Yptb* infection [18, 19, 22–25]. BM WT neutrophils were infected with WT-*Yptb*, $\Delta yopH$, $\Delta 5$, or $\Delta 5+pYopH$ and stimulated with poly-RGD (a polymer of arginine-glutamate-aspartate) to trigger integrin receptors, primed with E. coli lipopolysaccharide (LPS) and then stimulated with formyl-methionyl-leucyl-phenylalanine (fMLP) to trigger GPCR, or stimulated with IgG immune complex (IC) to trigger Fcy receptor (FcyR) (Fig 3A–3C) [49, 55, 71–76]. In all cases, WT-Yptb and $\Delta 5$ +pYopH inhibited ROS production similarly demonstrating that YopH was sufficient for ROS inhibition. In addition, high levels of ROS were detected in BM neutrophils infected with $\Delta 5$ or $\Delta yopH$ (Fig 3A-3C) demonstrating that YopH was also necessary for ROS inhibition when other Yops are present. Unlike stimulation with integrin or Fcy receptors (Fig 3A and 3C), ROS production from $\Delta yopH$ infected from BM neutrophils that were exposed to LPS and fMLP was less than Δ 5-infected cells, indicating that another Yop acts to partially reduce ROS production in the absence of YopH downstream of GPCR stimulation (Fig 3B). Further, ROS production after infection with $\Delta 5$ and exposure to LPS and fMLP was significantly higher than after exposure to LPS and fMLP alone (Fig 3B). This suggests that infection with $\Delta 5$ augments ROS production through activation of other receptors, possibly due to bacteria-mediated crosslinking of and signaling by integrin receptor in the absence of Yops (Fig 1C-1F) [18, 19].

Given our findings that *Skap2KO* mice retain some killing activities against the $\Delta yopH$ mutant (Fig 2A), we next tested the ability of *Skap2KO* neutrophils to produce ROS when stimulated through integrin receptor, GPCR and FcγR (Fig 3D–3F). *Skap2KO* neutrophils failed to generate high levels of ROS after poly-RGD and LPS + fMLP exposure (Fig 3D and 3E, S4A–S4D Fig), but stimulation of FcγR in *Skap2KO* neutrophils resulted in high levels of extracellular ROS, amounting to 80–90% of the ROS produced by WT neutrophils (Fig 3F,





S4E and S4F Fig). To verify that defects in ROS production resulted from a receptor-mediated signaling defect and not an inability of *Skap2KO* neutrophils to activate the NADPH oxidase complex, WT and *Skap2KO* neutrophils were stimulated with phorbol myristate acetate (PMA), which bypasses receptor-mediated activation and directly activates protein kinase C (PKC), to trigger assembly of the NADPH oxidase complex [55, 73, 74]. Indeed, PMA-stimulated WT and *Skap2KO* neutrophils produce equivalent levels of ROS (Fig 3D–3F). Together, these results indicate that a functional NADPH oxidase complex can assemble in *Skap2KO* neutrophils, but that SKAP2 signaling is required for integrin receptor and GPCR-mediated

ROS production in neutrophils. Since these two receptors may be stimulated during infection with *Yptb*, these results suggest that the inability of *Skap2KO* neutrophils to generate ROS contributes to the partial growth restoration of $\Delta yopH$ observed during *in vivo* infections (Fig 2A). While Fc γ R is unlikely to be activated by *Yptb* during infection of a naïve host, this result nonetheless raises the possibility that neutrophils have other receptors that may recognize *Yptb*, signal and function independently of SKAP2, leading to ROS generation in the absence of YopH. This superoxide production could potentially contribute to partial elimination of the $\Delta yopH$ mutant in *Skap2KO* mice.

YopH reduces phosphorylation of signaling proteins upon FcγRstimulation independent of SKAP2

Since YopH is sufficient to block FcyR-mediated ROS production (Fig 3C) but Skap2KO neutrophils produce ROS downstream of FcyR triggering by IC (Fig 3F), there remains an intact SKAP2-independent ROS-producing pathway downstream of FcyR that is targeted by YopH. We therefore evaluated whether YopH reduced tyrosine phosphorylation of signaling molecules important for ROS production after FcyR stimulation, namely Syk (pY352), SLP-76 (pY128), PLCy2 (pY1217) and ERK1/2 (pThr202/pTyr204) [53, 54, 80]. All were robustly phosphorylated in both WT and Skap2KO neutrophils after exposure to IC (Fig 4A-4D) indicating that these signaling proteins are activated upstream and/or independent of SKAP2. However, YopH reduced tyrosine phosphorylation of all these proteins (Fig 4A-4D) showing that YopH can interfere with ROS-generating pathways independent of SKAP2. In comparison, we evaluated whether phosphorylation of Syk, SLP-76, PLCy2, and ERK1/2 required SKAP2 after integrin stimulation, since Skap2KO neutrophils failed to produce ROS downstream of the integrin receptor (Fig 3D). SKAP2 was not required for the phosphorylation of Syk (pY352), SLP-76 (pY128) and PLCγ2 (pY1217) but was required for maximal phosphorylation of ERK1/2 (pThr202/pTyr204) (Fig 5A-5D). WT and Skap2KO BM neutrophils had low levels of SKAP1 indicating that the loss of SKAP2 was not compensated by overexpression of its homologue SKAP1 (S5 Fig). Therefore, SKAP2 acts downstream or independently of Syk, SLP-76 and PLCγ2 phosphorylation after integrin activation, but is important for relaying signals to activate ERK1/2, which is required for NADPH oxidase complex activity [49, 77-80]. Furthermore, YopH can interrupt both SKAP2-dependent and independent signaling pathways.

YopH targets degranulation downstream of integrin receptor, GPCR and FcγR, and degranulation is SKAP2-independent

To investigate the ability of YopH to inhibit additional antimicrobial activities and evaluate whether these activities are controlled by SKAP2, we assessed YopH-dependent inhibition of degranulation in WT and *Skap2KO* neutrophils. In human and murine neutrophils, YopH partially contributes to the inhibition of neutrophil degranulation as mutants infected with either $\Delta yopH$ or $\Delta yopE$ are partially defective in degranulation [40, 41]. The ability of YopH to prevent tertiary degranulation was tested by infecting WT BM neutrophils with WT-*Yptb*, $\Delta yopH$, $\Delta 5$ or $\Delta 5$ +pYopH and stimulating them with poly-RGD, fMLP, or IC (Fig 6A–6C). The amount of matrix metalloproteinase-9 (MMP-9), a component of tertiary granules, released in the cell-free supernatants was quantified by ELISA. Western blot analysis with anti-Rho-GDI antibody confirmed that the observed release of MMP-9 was not due to neutrophil lysis (S6A Fig). Degranulation was completely inhibited by WT-*Yptb* after stimulation with poly-RGD (Fig 6A) compared to uninfected controls, and was partially blocked by WT-*Yptb* after exposure to LPS+fMLP or IC (Fig 6B and 6C), indicating the combined actions of the





Yops are insufficient to halt degranulation under receptor stimulation (Fig 6B and 6C). Interestingly, degranulation was higher when cells were stimulated with either fMLP or IC and infected with $\Delta 5$ compared to stimulation of receptors in the absence of infection (S6B Fig), suggesting that simultaneous $\Delta 5$ binding via integrin receptor and stimulation of GPCR or FcyR enhances degranulation levels. YopH was sufficient to block tertiary granule release as $\Delta 5$ +pYopH-infected neutrophils released MMP-9 to the same level as WT-*Yptb* downstream of integrin, GPCR, and FcyR (Fig 6A-6C). Consistent with previous results [40, 41], significantly more degranulation was observed after infection with $\Delta yopH$ compared to WT-Yptb, but significantly less compared to infection with $\Delta 5$ (Fig 6A–6C). This demonstrates that YopH plays a role in reducing degranulation in murine neutrophils but is not the sole factor involved. Of note, SKAP2 was dispensable for MMP-9 release with approximately 6 ng/ml, 20 ng/ml and 25 ng/ml MMP-9 released after integrin, GPCR and FcyR stimulation, respectively (Fig 6D-6F). Supernatants of PMA-stimulated neutrophils showed 50% degranulation compared to triton X-100 treatment (S6C Fig). Combined, these results indicate that SKAP2 is dispensable for tertiary degranulation and suggests that SKAP2KO neutrophils may control growth of the $\Delta yopH$ mutant through degranulation.

Skap2KO neutrophils are defective for phagocytosis of Yptb

Finally, given that YopH participates in the ability of *Yersinia* to resist phagocytosis in isolated human neutrophils [29, 37, 39, 81, 82], we evaluated the ability of *Skap2KO* neutrophils to phagocytose WT-*Yptb*, $\Delta yopH$, or a strain defective for the type 3 secretion system, $\Delta yscF$. Using a gentamicin protection assay, we found that all three strains were phagocytosed at a





similar, low level by *Skap2KO* BM neutrophils and at significantly lower levels compared to that of WT BM neutrophils (Fig 7). This suggests that *Skap2KO* BM neutrophils are defective for invasin- β 1-mediated phagocytosis regardless of the presence of YopH or the T3SS. As expected, a significantly higher percentage of $\Delta yscF$ was phagocytosed in WT BM neutrophils than the WT-*Yptb* and $\Delta yopH$ mutant (Fig 7). The $\Delta yopH$ mutant was consistently phagocytosed by WT BM neutrophils more frequently than WT-*Yptb*, but this difference was not statistically significant suggesting that YopH has a modest role in preventing phagocytosis in WT BM neutrophils.

In summary, YopH inhibits ROS production, reduces degranulation and has a modest effect on phagocytosis of WT-*Yptb. Skap2KO* neutrophils have defects in ROS production and phagocytosis, but retain the ability to migrate to sites of *Yptb* infection and degranulate. Collectively, these results establish a role for SKAP2-mediated neutrophil signaling and its antimicrobial function towards *Yptb* and reveal how YopH acts to counter it.

Discussion

By exploring the interplay between *Yptb* effector YopH and the neutrophil adaptor protein, SKAP2, during tissue infection, we demonstrate the role YopH plays in protecting *Yptb* from neutrophil-mediated ROS production as well as the role SKAP2 plays in mediating neutrophil antimicrobial functions. The finding that inactivation of SKAP2 partially restores the growth of a $\Delta yopH$ mutant indicates that one or more SKAP2-controlled pathways must be inactivated



Fig 6. YopH reduces tertiary granule release, while SKAP2 is dispensable for tertiary granule release in neutrophils. Tertiary degranulation of (A-C) WT BM neutrophils that were simultaneously infected with WT-*Yptb*, *Yptb*- $\Delta 5+$ pBAD, *Yptb*- $\Delta 5+$ pYopH or *Yptb*- $\Delta yopH$ at a MOI of 20:1 or of (D-F) WT and *Skap2KO* BM neutrophils stimulated for 3 hrs by (A, D) plating on poly-RGD, (B, E) priming with LPS for 20 minutes and treating with 1 μ M fMLP, or (C, F) plating on IC. Cell free supernatants were analyzed for MMP-9 by ELISA with background subtraction of the unstimulated controls (FBS-plated controls). (A-C) Data for infections are expressed as fold increase relative to uninfected, unstimulated control samples from FBS coated plates. (D-F) Amount of MMP-9 (ng/ml) for WT and *Skap2KO* BM neutrophils with background subtraction of the unstimulated were used to determine the total granule content of WT and *Skap2KO* BM neutrophils. The data represent the means ± SEM from 3 independent experiments done in 2-3 replicates. Statistical significance was calculated using (A-C) one-way ANOVA followed by Tukey's Post-test or (D-F) Student t-test.



Fig 7. *SKAP2KO* neutrophils are defective in phagocytosis of a T3SS *Yptb* mutant. WT-BALB/c and *Skap2KO* BM neutrophils were infected with YPIII WT-*Yptb*, *ΔyopH*, and *ΔyscF* at MOI of 10:1 for 30 min. A gentamicin protection assay was used to determine the percentage of phagocytosed bacteria. Bars indicate mean of triplicate assays \pm SD. Individual dots show each replicate. Experiment shown is a representative of 3 independent experiments. Statistical significance was calculated using a Student's t-test to compare WT and *SKAP2KO* neutrophils and one-way ANOVA followed by Tukey's Post-test for comparisons between different infections of each neutrophil genotype.

by YopH for full virulence of *Yptb*. Thus, among the various proteins that have been previously identified as targets of YopH in neutrophils, macrophages, epithelial cells and T cells [29, 36, 44, 81, 83–87], SKAP2-controlled functions in neutrophils play a non-redundant role in host immune defenses in controlling *Yptb* infection. To our knowledge, this is the first identified signal-transduction pathway essential for YopH to target during *Yptb* infection in mammalian tissues.

What is YopH doing during infection? Our results indicate that YopH plays at least two important roles in blocking antimicrobial functions of neutrophils: inhibiting ROS production and reducing degranulation (Fig 8). A $\Delta yopH$ mutant survived significantly better in gp91^{phox-/-} mice than in WT mice, and YopH fully blocked ROS production downstream of integrin receptor, GPCR and FcyR stimulation in neutrophils. Furthermore, YopH interfered with the phosphorylation cascade triggered downstream of FcyR stimulation, including phosphorylation of Syk, SLP-76, PLC γ 2 and ERK1/2. Syk was not identified as a YopH target in a previous study; however, in that case, neutrophils were activated with bacteria alone [36] raising the possibility that Syk may be inactivated by YopH in vivo if it is activated through concurrent stimulation of several receptors at infectious loci. During tissue infection, other SKAP2-independent pathways may be triggered to generate ROS, which YopH can block, thus a $\Delta yopH$ mutant is susceptible to the ROS produced. In addition, in some cases NETosis depends on ROS production [65]. So our findings that YopH is critical to block ROS production support the idea that YopH may also block NETosis observed after infection of human neutrophils with a TT3S-minus strain but not WT Yptb strain [88]. We also found that degranulation was not dependent on SKAP2 after integrin and GPCR activation, receptors triggered by Yersinia [18, 19, 22–25]. Thus, degranulation could contribute to the partial elimination of the $\Delta yopH$ mutant observed in tissue infection of the Skap2KO mice (Fig 2A), because YopH was sufficient to reduce MMP-9 release from all three receptors tested (Fig 6A-6C). In addition, others have shown that YopH and YopE block degranulation-dependent release of lactoferrin, a



Fig 8. YopH dephosphorylates SKAP2-dependent and independent signaling proteins inhibiting ROS production and degranulation in neutrophils. (A) After integrin engagement, YopH inhibits signal propagation that is required for the activation of the NADPH oxidase complex and degranulation while SKAP2 is essential for ROS and phagocytosis and for maximal ERK1/2 phosphorylation, but not for degranulation or phosphorylation of Syk, SLP-76 and PLC₇2. (B) After GPCR stimulation of LPS-primed neutrophils with fMLP, YopH inhibits ROS and degranulation, where SKAP2 is required for ROS production, but not degranulation. (C) YopH inhibits ROS and degranulation after Fc₇R stimulation and reduces phosphorylation of Syk, SLP-76, PLC₇2 and ERK1/2, but SKAP2 is not required for ROS production, nor phosphorylation of these signaling proteins.

molecule in secondary granules [40, 41]. Interestingly, infecting neutrophils with a $\Delta 5$ mutant induced a significantly higher MMP-9 release than the activation of a single receptor with its corresponding ligand (S6B Fig). These results suggest concurrent stimulation of several receptors during infection can occur, which results in activation of multiple signaling pathways to effectively magnify the response. This may occur in a tissue infection milieu where neutrophils

are subjected to numerous innate signals, such as integrin stimulation from *Yptb* binding, and LPS and fMLP shed from *Yptb*. While it is tempting to speculate that YopH prevents phagocytosis in tissue infection as has been observed in isolated phagocytes [39], internalized $\Delta yopH$ was recovered in infected tissues at levels comparable to WT-*Yptb* [36], suggesting that YopH does not contribute significantly to anti-phagocytosis activities of *Yptb* in this model of *Yptb* tissue infection.

Precedence for more than one Yop blocking similar antimicrobial functions by targeting different proteins in receptor-mediated signaling cascades is abundant [5] and suggests that *Yptb* uses its arsenal of 6–7 Yops to block antimicrobial functions triggered by activation of different receptors and/or to alter antimicrobial responses of different cellular microdomains. Here, YopH was both necessary and sufficient to block ROS production downstream of two receptor classes; integrin and $Fc\gamma R$, however, its absence only partially crippled GPCR-mediated ROS production after infection with a $\Delta yopH$ mutant. These finding are consistent with the essential role of YopH in blocking integrin-mediated calcium flux, but not GPCR-mediated calcium release [38], and show that other Yops can compensate after activation of certain receptors. One Yop that could be acting with YopH to block GPCR-mediated ROS is YopO, as it has been reported to target Rho, Rac, and $G\alpha q$ -dependent signaling pathways of GPCR [89– <u>95</u>]. In contrast to the near complete restoration of growth of a $\Delta yopH$ mutant in $gp91^{phox-/-}$ mice reported here, a $\Delta yopE$ mutant was not restored for growth although a chimeric YopE that retained its Rac1 GAP but was deficient in other GAP activities was significantly restored for growth in gp91^{phox-/-} mice compared to C57BL/6J mice [67]. Thus, YopH appears to play a major role in protecting Yptb from ROS production during tissue infection, in part through its ability to block SKAP2-controlled pathways, while YopE may play a more prominent role at the phagocytic cup in inactivating Rac1 and RhoG rather than in inactivating Rac2 [67, 96-100].

Neutrophils kill bacteria using several different mechanisms including phagocytosis, ROS production, degranulation and NET production [101-103]. This raised the question of what anti-microbial functions are controlled by SKAP2, as Skap2KO mice partially restored $\Delta yopH$ growth. Together, our findings uncouple several key integrin- and GPCR-dependent antimicrobial responses from FcyR-dependent responses by virtue of their SKAP2-dependency for ROS production, and show that this adaptor is critical for some, but not all integrin- and GPCR-driven antimicrobial functions in neutrophils (Fig 8). SKAP2 may be part of a phospho-relay and/or could act a scaffolding function needed for full ERK1/2 phosphorylation [49]. ERK1/2 is required for phosphorylation of the $p47^{phox}$ and $p67^{phox}$ subunits of the NADPH oxidase complex in response to various stimuli in human neutrophils and therefore for ROS production [76–79]. The scaffolding function of SKAP2 may be essential to form stable signaling complexes downstream of integrin receptor and/or GPCR, but not FcyR. Hence, the defect of ROS in the integrin pathway may not be due to alteration of the phosphorylation state or activity of the proximal signaling proteins, but rather due to displacement of these proteins from the plasma membrane in the absence of SKAP2. In line with this hypothesis, the membrane localization of the SKAP2 binding partner, WASp, but not its phosphorylation, is abolished in *Skap2KO* neutrophils. This suggests that phosphorylation alone is not the only critical mediator of its activity, and possibly other SKAP2-associated-protein activities [49].

The fact that SKAP2 is critical for GPCR mediated ROS production was surprising, as the majority of the proteins known to interact with or predicted to be in the same signaling pathway as SKAP2 in neutrophils and other immune cells are dispensable downstream of GPCR; these include Syk, SLP-76, PRAM-1, and PLC $\gamma 2$ [51, 53, 54, 104], suggesting that SKAP2 signals through a different signaling pathway downstream of GPCR (Fig 8B). In addition, SKAP2 is not required for calcium flux in neutrophils following chemokine stimulation [49].

However, previous studies employed cytochalasin B to amplify the fMLP signal [51, 53, 54, 104], rather than LPS to prime neutrophils. Therefore, SKAP2 could be required for signaling pathways downstream of TLR4 receptor and/or downstream of GPCR in ROS production. Consistent with these findings, the proliferative response to LPS in *Skap2KO* B cells is strongly impaired, highlighting the critical role of SKAP2 in TLR4-mediated activation [70]. On the other hand, LPS-induced cytokine production was normal in *Skap2KO* dendritic cells, supporting the different functional requirement of SKAP2 in LPS activation in different immune cell types and functions [105]. Further, it is critical to evaluate these pathways as well as the role of SKAP2 in antimicrobial functions in human neutrophils to determine if these findings are applicable to humans as mouse and human neutrophils differ [55, 75, 103, 106].

While SKAP2 is necessary for limiting colonization of $\Delta yopH$ mutant, it does not appear to limit neutrophil migration to spleens in our infection models. This contrasts with the defect observed in a sterile inflammation model of hepatic necrosis where *Skap2KO* neutrophils were defective for migration [49]. It is noteworthy that the time courses of the two studies were different. In the sterile inflammation model, neutrophil recruitment was first observed after 30 min of tissue injury and infiltration reached maximum by 4 hours, whereas in our infection model no migration was observed for the first 24 hours in the spleen. Consistent with our results, neutrophils lacking SLP-76, which functions downstream of integrin receptors, also showed normal migration *in vivo* in response to *Staphylococcus aureus* infection, yet were defective in a sterile inflammation model of acute kidney injury [51, 107]. The difference in SKAP2-dependency in the context of microbial infection versus sterile inflammation may be due to multiple factors, including different chemokines released upon infection, a different adaptor protein compensating for the absence of SKAP2 in the infection model, or additional SKAP2-independent receptors activated during infection for tissue extravasation [51, 107].

In conclusion, *Yptb* employs YopH to target both SKAP2-dependent and independent signal-transduction pathways to cripple antimicrobial functions of murine neutrophils. *Skap2KO* neutrophils migrate efficiently to infected tissues, but only partially control the growth of a $\Delta yopH$ mutant. SKAP2 is required for ROS production after integrin and GPCR stimulation but not for degranulation, indicating that different antimicrobial functions can be triggered by activation of the same receptor but are not all dependent on SKAP2. However, since all these activities are blocked by YopH, future work is focused on uncovering the additional signaltransduction pathways that are essential YopH targets during tissue infection.

Materials and methods

Bacterial strains and growth conditions

Strains used in this work are listed in S1 Table. All mouse infections used *Yptb* IP2666 or derivatives while some neutrophil assays were performed with *Yptb* YPIIIpIB1 and derivatives as described in Figure Legends. The kanamycin resistant (Kan^R) IP2666 Δ yopH mutant was constructed by introducing a *yopH* deletion into IP2666-NdeI-Kan^R as previously described [30]. *Yptb* were grown overnight at 26°C in 2xYT media with aeration. For animal infections, overnight cultures were diluted 1:40 in 2xYT media incubated at 26°C with aeration for 8 hours, after which they were diluted 1:100 and incubated overnight with aeration and the OD₆₀₀ measured. The bacteria were diluted in PBS to achieve the required input dose for intravenous (I. V.) or intratracheal (I.T.) inoculation. For infection of neutrophils, *Yptb* were grown under Yop-inducing conditions as described [36]. Strains containing pBAD plasmids were grown in presence of 20 µg/ml chloramphenicol and 50 mM arabinose was added to the culture prior to the shift to 37°C and maintained in medium with antibiotics and inducer throughout the assays. OD₆₀₀ of each culture was measured and strains were diluted in KRP buffer (KrebsRinger-Phosphate: PBS—Corning; Cat# 21040CV + 1 mM CaCl₂ + 1.5 mM MgCl₂) with 5 mM glucose or 5 mM arabinose depending on bacterial strains used in infections. BM neutro-phils were infected at the indicated MOI.

Mouse infections

Female 8-12 week old BALB/c (Taconic labs), BALB/c-Skap2KO [48], C57BL/6J (Jackson labs), and C57BL/6J gp91^{phox-/-} (Jackson labs) mice were used for infections. Competition infections were performed using an equal mixture of unmarked and Kanamycin-resistant marked IP2666 strains. C57BL/6J and gp91^{phox-/-} were I.V. or I.T. infected with 1x10³ or 5x10³ colony forming units (CFU), respectively, with a 1:1 mixture of WT-Yptb IP2666 and YptbAyopH-Kan^R. Two days post infection spleens and lungs were isolated, weighed, homogenized and plated on L agar containing 0.5µg/mL irgasan. BALB/c and Skap2KO mice were inoculated I.V. with 1x10³ WT-*Yptb* IP2666 and Kan^R marked $\Delta yopH$ and mice were sacrificed three days post-infection. The competitive index (C.I.) (C.I. = $(CFU_{\Delta vopH}/CFU_{WT})_{output}/(CFU_{\Delta vopH}/CFU_{WT})_{input})$ was calculated as described [108] with the following modifications. In some cases, the CFU for each strain were determined by subtracting the number of *yopH*-Kan^R colonies from the total number of colonies recovered from L-irgasan plates to increase the limit of detection. For neutrophil (Ly6G⁺) cell depletions, mice were intraperitoneally injected with 100µl of 50µg/mL of 1A8 antibody (α -Ly6G⁺) (Fisher; Cat # bdb551459) 16 hours prior to and 24 hours post-infection. Depletion was confirmed by staining spleen homogenates with α -GR1⁺ and α -CD11b⁺ and analysis by FACS as previously described [21]. For determining the migration kinetics of neutrophils, 8 week-old female BALB/c and Skap2KO were either mock infected or IV infected with 1x10³ WT-Yptb-GFP strain (chloramphenicol, Cm^R). Spleens were collected at 24, 33, 48 hours post-infection for CFUs and immunohistochemistry.

Immunofluorescence and image analysis

Spleen tissues collected at 24, 33 and 48 hours were fixed in 4% paraformaldehyde for 4 hours at room temperature. Tissues were frozen-embedded in Sub Xero freezing media (Mercedes Medical; Cat # MER 5000) and cut into 10 µm sections using a cryostat microtome. Sections were permeabilized using 100% methanol followed by blocking with 2% BSA in PBS for 1 hour at room temperature. Ly6G⁺ cells were detected by staining sections with 1:100 rat antimouse monoclonal Ly6G (clone 1A8) antibody for 1 hour, washing 3 times with PBS, then incubating with 1:500 goat anti-rat Alexa Fluor 594 antibody (Invitrogen Cat# A-11007) and 1:1000 DAPI (Invitrogen) for 1 hour, and washing 3 times with PBS. Finally, samples were mounted on glass slides using ProLong Gold antifade (Invitrogen, Cat# P36930) and cured overnight in the dark at room temperature. Tissues were imaged using a 40X objective on a Nikon A1R confocal microscope (Nikon Instruments Inc.), with 0.5 µm z-steps using a filter set for DAPI (Ex/Em:350/470nm), GFP (Ex/Em:488/525nm) and mCherry (Ex/Em:561/ 595nm). Volocity software 6.3 was used for 3D reconstruction image quantification. For quantification of the microcolony size by volume, uninfected control images were used to set the threshold of the GFP intensity signal and objects smaller than 1µm³ were excluded. Volume was determined by the summation of the individual signals in the green channel after applying the threshold limits. Quantification of neutrophils was determined by determining the sum of the red fluorescent signal intensity (mCherry channel).

Neutrophil isolation

BM neutrophils were isolated as previously described [36]. Briefly, BM was harvested from tibia and femur bones in HBSS without Ca^{2+} or Mg^{2+} (HBSS-/-). Red blood cells were lysed

using sterile endotoxin free HyClone cold water (GE Heathcare; Cat# SH30529.FS) followed by addition of HBSS-/- to restore normal tonicity. Neutrophils were isolated using a three step, 55%, 65%, 75%, Percoll density gradient and centrifuged at 480 x g for 30 min without brakes. The cells at the 65–75% interface were collected, washed 2 times with HBSS-/-, and rested at room temperature for one hour. Purity of neutrophils ranged from 82–89% by FACS analysis with anti-CD11b, anti- Ly6C and anti-Ly6G antibodies (BioLegend). Neutrophils were then re-suspended in KRP neutrophil buffer to the desired concentration incubated for 20 min at room temperature and shifted to 37° C in 5% CO₂ for 10–15 min prior to initiation of the experiment.

Neutrophil ROS assays

For stimulating integrin receptors, wells were coated with 15µg/mL poly-RGD (Sigma; Cat# F5022) for 2 hours at 37°C, and then washed 2 times with PBS. To stimulate FcyR, wells were coated with 20µg/ml human serum albumin (Sigma; Cat # A9511) for 1 hour at 37°C, washed 2X with PBS, incubated with 10% FBS for 30 min, washed 2 times with PBS, then incubated with a 1:400 dilution of rabbit anti-human serum albumin (Sigma; Cat # A0433) in 10% FBS for 1 hour at 37°C, followed by washing 2 times with PBS. Control wells were coated with 10% FBS for 30 min, then washed 2 times with PBS. Assays for GPCR receptor were performed in the absence of Mg^{2+} to decrease the possibility of integrin dependent responses and surfaces were coated with 10% FBS to minimize adhesion to mimic cells in suspension. For stimulating GPCR, BM neutrophils were suspended in KRP without Mg²⁺ and primed with 10µg/mL LPS (Sigma; Cat# L4516) for 20 min then added to 10% FBS coated wells without centrifugation to maintain cells in suspension followed by stimulation using 1µM fMLP (Sigma; Cat # F3506). BM neutrophils suspended in KRP buffer at 37°C were loaded with 50 µM isoluminol (Sigma; Cat# A8264) and 15 U/ml of HRP (Sigma; Cat # P6782) as previously described [109] and 2 x10⁵ neutrophils/well were added to 96-well white high-bound plates (Thermo Fisher; Cat# 3922) coated with 10% FBS or different ligands and centrifuged at 190 x g for 1 min. For infections, bacteria resuspended in KRP buffer were added to wells at a MOI of 20:1 simultaneously with neutrophils when stimulating integrin and Fcy receptors. The bacteria and neutrophils were spun down on the ligand coated plates for 1 min to uniformly induce receptor binding. For GPCR stimulation, bacteria were added to neutrophils with LPS priming and kept in suspension. PMA (1µM) (Sigma; Cat# P1585) was used as a positive control. Chemiluminescence was measured using a BioTek Synergy HT microplate reader. ROS detection using cytochrome c (Sigma; Cat # C-7752) was performed as previously described [109–110] with the following modifications. BM neutrophils $(1X10^5 \text{ per } 50 \text{ µl})$ were added to 50 µl 200µM cytochrome c in 96 well high-binding Immulon 4HBX plates (Fisher; Cat# 3855) coated with ligands or FBS. Absorbance at 550nm and 490nm were recorded and the difference used to calculate the nanomoles of superoxide produced using an extinction coefficient of $2.11 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$ for cytochrome c. The total amount of ROS for each condition was corrected by subtracting its superoxidase dismutase (SOD) control [109].

Neutrophil degranulation assay

 2×10^5 BM neutrophils/well were plated on 96 well Immulon 4HBX plates coated with poly-RGD, IC or 10% FBS and incubated at 37°C, 5% CO₂. Infections were carried out at a MOI of 20:1 as described for ROS assays. After 3 hours, to determine the total amount of MMP-9, the control wells were treated with 0.01% triton X-100, without removing the supernatant, to lyse the cells and the supernatant collected. To determine the amount of released MMP-9, plates were placed on ice for 5 min to stop the reaction and then centrifuged for 10 min at 4°C at 190 x g. Supernatants were collected and stored at -80°C until analysis by ELISA using DuoSet (R&D Systems; Cat # DY6718) for mouse MMP-9, or by western blot using an antibody against Rho-GDI (Cell Signaling; Cat# 2564S). The remaining cells in the wells were collected and lysed for western blot analysis to determine the total amount of Rho-GDI using an antibody against Rho-GDI. For comparisons between WT and *Skap2KO* neutrophils the absolute amount of MMP-9 detected in supernatants was reported for the stimulated receptor after sub-tracting the unstimulated (FBS-coated wells) control values. The amount of MMP-9 released after infection with *Yptb* strains was reported as fold-change compared to the control of unstimulated, uninfected wells.

Western blot analysis

Cells were incubated at 37°C, 5% CO₂ for 5 min or 10 min for FcγR or integrin stimulation respectively. In some cases, BM neutrophils were infected with YPIII WT-*Yptb* or $\Delta yopH$ for 30 min and then spun down on the ligand coated surface. Cells were lysed in 1X Novex buffer and 2x10⁶ cell equivalents were resolved on a 4–12% NuPAGE gel (Invitrogen; Cat# NP0335BOX) in MOPS buffer. Proteins were transferred to Immobilon-FL filters and subjected to Western blot analysis with the following primary antibodies at a dilution of 1:500 anti-Syk (Cell Signaling; Cat# 2712), anti-SykpY352 (Cell Signaling Cat#2701), anti-SLP76 clone AS55 (Millipore; Cat# 05–1426), anti- SLP76pY128 (BD Pharmingen; Cat# 558367), anti-PLCγ2 (Cell Signaling; Cat# 3872), anti-PLCγ2pY1217 (Cell Signaling; Cat# 3871), anti-Erk1/2 pThr202/pTyr204 (Cell Signaling; Cat# 4370), anti-Erk1/2 (Cell Signaling; Cat# 9102), or at 1:1000 of anti-Rho-GDI (Cell Signaling; Cat# 2564S). Secondary LI-COR goat antimouse antibody IRDye 800 CW (Cat# 827–08364) and goat anti-rabbit IRDye 800 CW (Cat# 926–32211) were used at a dilution 1:20,000. ODYSSEY CLx LI-COR system was used to image the blots and IS Image Studio used for analysis and quantification of the bands.

Gentamicin protection assay

YPIII *WT-Yptb*, $\Delta yopH$ or $\Delta yscF$ strains were grown under Yop-inducing conditions and used to infect 1 x10⁴/well BM neutrophils at MOI of 10:1 in 96 well Immulon 4HBX plates coated with 10% FBS for 30 min at 37°C, 5% CO₂. Gentamicin was added to a final concentration of 100µg/ml for 60 min, and then cells were washed three times with PBS. BM neutrophils were centrifuged at 190 x g for 5 min between each wash. BM neutrophils were lysed with 100 µl 0.5% Triton-X-100 for 10 min followed by addition of 100µl 2xYT, and lysates were plated at different dilutions to determine the number of intracellular bacteria. Results are reported as the percent of the recovered CFU from gentamicin-treated cells divided by the non-gentamicin-treated control.

Statistical analyses

Statistical analyses were performed in GraphPad Prism software (version 7b). Specific tests used are indicated in the figure legends. Significant difference is indicated as p<0.05, p<0.01, p<0.001, p<0.001, p<0.001, p<0.001, p<0.001, p<0.001, p<0.001, p<0.0001, and ns = non-significant for all figures.

Ethics statement

This study was performed in accordance with the recommendations in the Guide for Care and Use of Laboratory Animals of the National Institutes of Health. The Tufts University Institutional Animal Care and Use Committee (IACUC) reviewed the study protocol under approval number B2015-35 and B2018-10.

Supporting information

S1 Fig. *Yptb* CFU in spleen of C57BL/6J or C57BL/6J $gp91^{phox-/-}$ mice and assessment of bone marrow neutrophil purity. (A) C57BL/6J or C57BL/6J $gp91^{phox-/-}$ mice were I.V. inoculated with a 1:1 mixture of IP2666 WT-*Yptb* and $\Delta yopH$ -Kan^R. Spleen were collected 2 days post-infection and CFU for WT-*Yptb* and $\Delta yopH$ -Kan^R was determined by plating on selective and non-selective plates. Statistical significance was calculated using one-way ANOVA with Sidak's multiple comparison test after log₁₀ transformation of data. (B) Bone marrow was harvested from the tibia and femur bones of mice and neutrophils were isolated using Percoll density gradient method. Cells at the 65–75% interface were collected and stained using with α -CD11b, α -Ly6G and α -Ly6C and analyzed by flow cytometry. Bone marrow isolated neutrophils were more than 80% pure. (TIFF)

S2 Fig. WT *Yptb*, *ΔyopM* and Δ 5+pYopH secrete equivalent amounts of YopH. Overnight cultures of IP2666 WT-*Yptb*, *Yptb*- Δ 5+pYopH and *Yptb*- Δ *yopM* were diluted 1:40 in low calcium media and grown for 2 hours at 26°C with aeration followed by addition of 50mM arabinose to induce YopH expression from the pBAD plasmid and shifted to 37°C for 2 hours with aeration. Proteins from culture supernatants were precipitated, resolved on a 4–20% gradient polyacrylamide gel and stained using coomassie blue dye. (TIFF)

S3 Fig. BALB/c and *Skap2KO* mice have equivalent *Yptb* CFU during infection. (A) WT-BALB/c and *Skap2KO* mice were intraperitoneally injected with 1A8 or an isotype control antibody and 16 hrs later inoculated I.V. with an equal mixture of 10^3 CFU IP2666 WT-*Yptb* and *ΔyopH*-Kan^R. Spleens were collected 3 days post-infection and plated for CFUs. Total CFU are shown. (B)WT-BALB/c and *Skap2KO* mice were I.V. infected with 1x10³ CFU *Yptb*-GFP, and sacrificed at 24, 33, or 48 hours post-infection. Spleens were collected, weighed, homogenized, and plated for CFU on selective plates. Each dot represents a mouse; horizontal bars represent the geometric mean. Significance was calculated using (A) two-way ANOVA followed by Tukey's Post-test and (B) Student's t-test. (TIFF)

S4 Fig. *Skap2KO* BM neutrophils fail to produce ROS following integrin and GPCR receptor stimulation but not after FcyR stimulation. Respiratory burst of WT-BALB/c and *Skap2KO* BM neutrophils ($1x10^5$ cells) was measured using the cytochrome c reduction test by (A-B) plating on a poly-RGD surface, (C-D) priming with 10µg/ml LPS for 20 min followed by stimulation with 1µM fMLP, or (E-F) plating on an IC coated surface. Absorbance at 550 nm and 490 nm was recorded and each condition was corrected by its superoxidase dismutase (SOD) control value. Total amount of ROS was estimated from measuring the AUC for each condition for the duration of the experiment. Data are shown as the means ± SD from triplicate measurements of one experiment, which are representative of at least 4 independent experiments. Statistical significance was calculated using Student t-test. (TIFF)

S5 Fig. *Skap2KO* mice are defective for SKAP2 and do not overexpress SKAP1. Lysates of BM neutrophils isolated from WT-BALB/c and *Skap2KO* were immunoblotted for SKAP2 and SKAP1. Anti-Rho-GDI was used as a loading control. No band for SKAP2 was detected in *Skap2KO* neutrophils confirming the absence of SKAP2. SKAP1 was detected at very low levels in both genotypes. (TIFF)

S6 Fig. Release of neutrophil granules is not due to cell lysis and degranulation induced after infection and receptor stimulation is significantly higher than stimulation of single **receptors alone.** (A) Supernatants and cell lysates $(2 \times 10^5 \text{ cells})$ from infected and stimulated samples were immunoblotted with anti-Rho-GDI as an indicator of cell lysis. A positive control for degranulation was treatment with PMA, and the negative control was cells added to FBS-coated wells. Rho-GDI was detected in the supernatants of cells treated with triton X-100. (B) BM neutrophils (2x10⁵) were stimulated with poly-RGD, fMLP or IC to stimulate integrin, GPCR, or FcyR respectively and left uninfected or simultaneously infected with $Yptb\Delta 5$ at a MOI of 20:1 for 3 hrs. Cell free supernatants from equivalent numbers were analyzed by ELISA for MMP-9 release. Data presented are expressed as fold increase relative to control uninfected and unstimulated samples. The data represent the means \pm SEM from 3 independent experiments done in 2-3 replicates. (C) WT-BALB/c and Skap2KO BM neutrophils were stimulated for 3 hrs by 1µM PMA. Cell-free supernatants were analyzed for MMP-9 by ELISA. The data represent the means \pm SEM from 3 independent experiments done in triplicate. Statistical significance was calculated using Student t-test. (TIFF)

S1 Table. List of strains. (DOCX)

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