Accum	ulation of fluorophore-labeled SNX fusio at internalized borreliae (Figure 1B)	n constructs		
Overexpression of	Accumulation at internalized borr	eliae (Mean ± SEM)		
SNX1-GFP	48.33% ± 4.41%			
GFP-SNX3	73.33% ± 10.14	%		
SNX4-GFP	11.67% ± 2.89%	6		
GFP-SNX8	20.00% ± 11.55	%		
mCherry-SNX9	3.33% ± 1.67%			
GFP-SNX12	5.00% ± 2.89%			
SNX17-GFP	16.67% ± 1.67%	6		
SNX27-GFP	1.67% ± 1.67%			
SNX31-GFP	11.67% ± 4.419	6		
Effect of double k	nockdown of SNX1 and SNX3 on the mor	phology of internalized		
ciPNA target	Dorrellae (Figure 2B)	Quantification		
Sirina target		(Mean ± SEM)		
Non-specific control		70.40% ± 6.17%		
SNX1 #1	compacted	60.13% ± 10.35%		
SNX3 #1		27.67% ± 5.25%		
SNX1 #1 + SNX3 #1		37.20% ± 6.18%		
Fluorescence	intensity of DQ-BSA/borreliae phagosom treatment (Figure 21)	es after SNX3 siRNA		
siRNA target	Quantification (Mean	± SEM)		
Non-specific control	1311 a.u. ± 51.96	a.u.		
SNX3 #1	1058 a.u. ± 51.90	a.u.		
SNX3 #2	1135 a.u. ± 76.17	a.u.		
Flu	uorescence intensity of DQ-BSA after SNX treatment (Figure 2J)	(3 siRNA		
siRNA target	Quantification (Mean	± SEM)		
Non-specific control	1525 a.u. ± 17.18	a.u.		
SNX3 #1	1587 a.u. ± 23.89 a.u.			
SNX3 #2	SNX3 #2 1470 a.u. ± 17.11 a.u.			

## Table S1. Values for specified parameters following various treatments

survival assa	y after SNX3 knock	down - Bacteria num	ber [10⁴/ml] <mark>(Fi</mark> g	gure 2K)	
		days			
siRNA target	7	8	9	10	
Non-specific control	2.21 ± 0.20	3.27 ± 1.53	11.93 ± 5.53	19.23 ± 6.69	
SNX3	2.01 ± 0.89	5.38 ± 3.24	18.04 ± 7.53	44.40 ± 13.41	
SNX1	expression level a	after SNX3 knockdo	wn <mark>(Figure S3</mark> B	)	
siRNA target	Quanti protein lev	fication SNX1 vel (Mean ± SEM)	Corres protein le	ponding SNX3 vel (Mean ± SEM)	
Non-specific control	10	00% ± 0%	1	00% ± 0%	
SNX3 #1	8.77	7% ± 3.12%	79.3	37% ± 4.38%	
SNX3 #2	26.2	3% ± 7.61%	95.4	17% ± 7.06%	
	·				
SNX3	expression level a	after SNX1 knockdo	wn (Figure S3D	)	
siRNA target	Quanti protein lev	ification SNX3Corresponding SNX1vel (Mean ± SEM)protein level (Mean ± SEM)			
Non-specific control	100% ± 0%		00% ± 0% 100% ± 0%		
SNX1 #1	5.23	93.30% ± 8.14%		30% ± 8.14%	
SNX1 #2	8.39	39% ± 3.51%         99.73% ± 11.14%		3% ± 11.14%	
Сог	mpaction after Wo	ortmannin treatme	nt (Figure 3Q)		
Wortmannin concentration	Morphology of	internalized borrelia	e Qua (M	antification ean ± SEM)	
control		mnacted	49.9	90% ± 9.27%	
1 µM		inpacted	21.6	53% ± 2.15%	
SNX3 accumulation	n at <i>Borrelia</i> phag	osomes after Wortr	mannin treatm	ent (Figure 3T)	
Wortmannin co	ncentration	Quanti	fication (Mean ±	SEM)	
contro	bl		62.43% ± 5.41%		
1 μΝ	1		2.97% ± 1.63%		
	(			• • • • • • • • • • • • •	
Effect of SNX3 res	cue after knockdo	(Figure 4D)	logy of internal	ized borreliae	
siRNA target	overexpression of	Morphology of internalized borrel	Qui iae (M	antification ean ± SEM)	
	GFP control		71.2	27% ± 4.78%	
Non-specific control	siRNA insensitive GFP-SNX3	RNA insensitive GFP-SNX3 compacted		13% ± 4.83%	
	siRNA insensitive GFP-SNX3-Y71A		47.7	47.70% ± 7.39%	

	GFP control		34.70% ± 6.35%
SNX3 #1	siRNA insensitive GFP-SNX3	compacted	64.63% ± 2.81%
	siRNA insensitive GFP-SNX3-Y71A		31.67% ± 3.35%
	sumulation of GED	labolod SNX deletion c	onstructs
Att	at internaliz	ed borreliae (Figure 5B)	
Overexpression of	Accumu	lation at internalized born	reliae (Mean ± SEM)
GFP-SNX3-FL		61.67% ± 10.93	%
GFP-SNX3-ΔN		41.67% ± 4.419	6
GFP-SNX3-PX+C		38.33% ± 3.33%	6
GFP-SNX3-ΔC		11.67% ± 7.27%	6
GFP-SNX3-N+α		0%	
GFP-SNX3-C		0%	
Effect of rescue by	different SNX3 co internalized	nstructs after knockdov d borreliae (Figure 5S1)	wn on the morphology of
	Morphology of	overexpression	Quantification
siRNA target	internalized borreliae	construct	(Mean ± SEM)
	compacted	GFP-C1	67.50% ± 3.64%
		GFP-SNX3-FL	58.08% ± 0.78%
		GFP-SNX3-ΔN	64.83% ± 7.61%
Non-specific control		GFP-SNX3-PX+C	62.50% ± 7.51%
		GFP-SNX3-ΔC	37.28% ± 6.12%
		GFP-SNX3-N+α	41.9% ± 4.33%
		GFP-SNX3-C	42.85% ± 4.01%
		GFP-C1	34.03% ± 2.85%
		GFP-SNX3-FL	68.23% ± 1.76%
		GFP-SNX3-ΔN	55.23% ± 6.65%
SNX3	compacted	GFP-SNX3-PX+C	56.48% ± 6.98%
		GFP-SNX3-ΔC	41.45% ± 6.55%
		GFP-SNX3-N+α	36.48% ± 5.70%
		GFP-SNX3-C	40.65% ± 3.72%
			wholes of intervalies d
Effect of double k	borr	eliae (Figure 6I)	phology of internalized
siRNA target	Morphology of	internalized borreliae	Quantification (Mean ± SEM)
Non-specific control	compacted		70.20% ± 5.06%

treatment (Figure 7G)				
Fluorescence intensity of DQ-BSA/borreliae phagosomes after Gal9 siRNA				
SNX3 #1 + Gal9 #1		39.23% ± 9.37%		
Gal9 #1		39.03% ± 2.84%		
SNX3 #1		35.67% ± 2.04%		

siRNA target	(Mean ± SEM)
Non-specific control	1311 a.u. ± 51.96 a.u.
Gal9 #1	1074 a.u. ± 60.87 a.u.
Gal9 #2	1048 a.u. ± 57.07 a.u.

Fluorescence intensity of DQ-BSA after Gal9 siRNA treatment (Figure 7H)				
siRNA target	Quantification (Mean ± SEM)			
Non-specific control	1525 a.u. ± 17.18 a.u.			
Gal9 #1	1591 a.u. ± 21.37 a.u.			
Gal9 #2	1590 a.u. ± 33.30 a.u.			

Effect of rescue by different Gal9 constructs after knockdown on the morphology of					
siRNA target borreliae (Figure 8B) borreliae (Figure 8B) Overexpression construct (Mean ± SEM)					
		GFP-C1	54.32% ± 2.80%		
		GFP-Gal9-FL	54.69% ± 3.12%		
		GFP-Gal9-∆CRD2	44.40% ± 5.02%		
Non-specific control	Compacted	GFP-Gal9-∆N	52.19% ± 5.54%		
		GFP-Gal9-linker+CRD2	53.28% ± 5.03%		
		GFP-Gal9-N+CRD1	41.13% ± 8.34%		
		GFP-Gal9-CRD1+linker	51.57% ± 8.91%		
	Compacted	GFP-C1	36.56% ± 1.62%		
		GFP-Gal9-FL	56.17% ± 3.08%		
		GFP-Gal9-∆CRD2	43.84% ± 1.30%		
Gal9		GFP-Gal9-∆N	56.57% ± 5.00%		
		GFP-Gal9-linker+CRD2	35.24% ± 3.66%		
		GFP-Gal9-N+CRD1	35.78% ± 4.64%		
		GFP-Gal9-CRD1+linker	41.89% ± 4.60%		

Effect of rescue by different Gal9 CRD mutants after knockdown on the morphology of internalized borreliae (Figure 8D)				
siRNA target	Morphology of internalized borreliae	Overexpression construct	Quantification (Mean ± SEM)	
Non-specific control	Compacted	GFP-C1	58.52% ± 3.98%	
		GFP-C1	40.46% ± 0.46%	
		GFP-Gal9-FL	59.32% ± 4.32%	
Gal9	Compacted	GFP-Gal9-A46V	51.08% ± 3.46%	
Gais	Compacted	GFP-Gal9-N137A	53.36% ± 5.74%	
		GFP-Gal9-R252A	50.00% ± 4.55%	
		GFP-Gal9-A46V+N137A	57.81% ± 1.29%	
Ef	fect of SNX3 knoc cence intensity at	kdown on Rab5a and ga borreliae phagosomes	llectin-9 (Figure 9C+D)	
Protein	Protein siRNA target Fluorescence intensity at borreliae phagosomes (Mean + SEM)			
Dahfa	Control		38078 a.u. ± 5603 a.u.	
Kabba	SNX3		17497 a.u. ± 2468 a.u.	
Galectin-9		Control	2837 a.u. ± 291.2 a.u.	
		SNX3	1685 a.u. ± 144.5 a.u.	
Effect of SNX3 knockdown on phagosomes size (Figure 9E)				
siRNA target	Volur	ne of borreliae phagosom	es (Mean ± SEM)	
control	2.80 ± 0.31 μm³			
SNX3	2.65 ± 0.25 μm³			

siRNAs			
Name/target	Sequence (5′ → 3′)		
SNX3 #1	GCGUCAGCUUCCUUUUAGA-(dTdT)		
SNX3 #2	CUCAUAUGCUCAGUUUUGU-(dTdT)		
SNX1 #1	AAGAACAAGACCAAGAGCCAC-(dTdT)		
SNX1 #2	GAACAAGACCAAGAGCCAC-(dTdT)		
Gal9 #1	GGACUUCAGAUCACUGU-(dTdT)		
Gal9 #2	GGAAGACACACAUGCCUUUCC-(dTdT)		

## Table S2. siRNAs, oligonucleotides, plasmids, and antibodies used in this study

	Oligonucleotides					
#	Name	Sequence (5' $\rightarrow$ 3')	Restriction site			
P01	SNX3_fw	AAAAAAGGTACCCCGCGGAGACCGTGGCTGACACCC	Kpnl			
P02	SNX3_ΔC_rev	TTTTTTGGATCCTCAATCTATTATTTCATCTTGTAAAAAC	BamHI			
P03	SNX3_N+a_rev	TTTTTTTTGGATCCTCAGCTGGGGGGGTCCGTAGG	BamHI			
P04	SNX3_ΔN_fw	AAAAAAGGTACCCCAACCTGAATGACGCCTACGGACCC	Kpnl			
P05	SNX3_rev	TTTTTTGGATCCTCAGGCATGTCTTATTTTAGATGG	BamHI			
P06	SNX3_PX+C_fw	AAAAAAGGTACCCCAACTTCCTCGAGATCGATGTGAGC	Kpnl			
P07	SNX3_C_fw	AAAAAAGGTACCCCAAAAGCTATACTCCATCT	Kpnl			
P08	SNX3_C_rev	TTTTTTGGATCCTCAGGCATGTCTTATTTT	BamHI			
P09	Gal9_fw	AAAAAACTCGAGCCGCCTTCAGCGGTTCCCAGGCTCCCTAC	Xhol			
P10	Gal9_rev	TTTTTTGGTACCTCATGTCTGCACATGGGTCAGCTGGATGTC	Kpnl			
P11	SNX3_PX_fw	GAAAGAATCTACTGTTAGAAGAAGAGCCAGTGACTTTG	-			
P12	SNX3_PX_rev	CAAAGTCACTGGCTCTTCTTCTAACAGTAGATTCTTTC	-			
P13	SNX3_si- insens_fw	GAAAGCGTTTTTGCGGCAGCTTCCATTCAGAGGAGATG	-			
P14	SNX3_si- insens_rev	CATCTCCTCTGAATGGAAGCTGCCGCAAAAACGCTTTC	-			
P15	SNX3_GST_fw	AAAAAAGGATCCGCGGAGACCGTGGCTGACACCC	BamHI			
P16	SNX3_GST_rev	AAAAAAGAATTCTCAGGCATGTCTTATTTTAGATGG	EcoRI			
P17	KIF5A_stop_fw	CGGCATGGACGAGCTGTACAAGTGATCCATGAATGGAGC	-			
P18	KIF5A_stop_rev	GCTCCATTCATGGATCACTTGTACAGCTCGTCCATGCCG	-			

P19	Gal9_CRD1_fw	AAAAAACTCGAGCCTTTTCTGGGACTATTCAAGGAGG	Xhol
P20	Gal9_CRD1_rev	TTTTTTGGTACCTCAGAAGCTGATGTAGGACAGCTGC	Kpnl
P21	Gal9_linker_fw	AAAAAACTCGAGCCCAGAACCCCCGCACAG	Xhol
P22	Gal9_linker_rev	TTTTTTGGTACCTCAAGGCATCGGATAGGCGGG	Kpnl
P23	Gal9_si-insens_fw	CAGGACGGACTACAAATAACAGTCAATGGGACC	-
P24	Gal9_si- insens_rev	GGTCCCATTGACTGTTATTTGTAGTCCGTCCTG	-
P25	Gal9_A46V_fw	GGAACCAGGTTTGTTGTGAACTTTCAGACTGG	-
P26	Gal9_A46V_rev	AAAGTTCACAACAAACCTGGTTCCACTGGCAGC	-
P27	Gal9_N137A_fw	ATCTCCGTCGCTGGCTCTGTGCAGCTGTCCTA	-
P28	Gal9_N137A_rev	TGCACAGAGCCAGCGACGGAGATGGTGTCCAC	-
P29	Gal9_R252A_fw	CAGTGCTCAGGCGTTCCACATCAACCTGTGCT	-
P30	Gal9_R252A_rev	TTGATGTGGAACGCCTGAGCACTGGGCAGGAC	-

Antibodies and staining reagents					
Antibody	Company	Species	Dilution IF	Dilution WB	
anti-SNX3 (C16)	Santa Cruz	goat polyclonal	1/100	-	
anti-SNX3	Proteintech	rabbit polyclonal	1/200	1/200	
anti-SNX1	BD Biosciences	mouse monoclonal	1/100	1/1000	
anti-Gal9	Proteintech	rabbit polyclonal	1/200	1/500	
anti-Borrelia burgdorferi	Antibodies online	rabbit polyclonal	1/1000	1/1000	
anti- <i>Borrelia burgdorferi</i> Bss42	Novus Biologicals	mouse monoclonal	1/1000	1/1000	
anti-GFP	Novus Biologicals	rabbit polyclonal	-	1/5000	
anti-pan-actin	Merck Millipore	mouse monoclonal	-	1/5000	
anti-mouse-HRP	GE Healthcare	sheep polyclonal	-	1/5000	
anti-rabbit-HRP	GE Healthcare	donkey polyclonal	-	1/5000	
anti-goat-HRP	abcam	rabbit polyclonal	-	1/5000	
anti-rabbit- AlexaFluor488/568/647	Thermo Scientific	donkey or goat	1/200	-	
anti-mouse- AlexaFluor488/568/647	Thermo Scientific	donkey or goat	1/200	-	
anti-goat- AlexaFluor488/568/647	Thermo Scientific	donkey	1/200	-	

Phalloidin AlexaEluor488/568/647	Thermo Scientific		-	1/50	-		
Plasmids							
Name	Provided by/		Reference				
GFP-SNX3	S. Grinstein		Braun et al, Cell Microbiol. 2010 Sep 1:12(9):1352-67				
RFP-SNX3	J. Gruenberg		Pons et al, PLoS Biol. 2008 Sep 2;6(9):e214				
Rab5a-RFP	S. Grinstein						
SNX1-GFP	S. Grinstein						
SNX4-GFP	L. Binkle						
GFP-SNX8	K. Sandvig		Dyve et al, Biochem Biophys Res Commun. 2009 Dec 4;390(1):109-14				
mCherry-SNX9	V. Haucke		Posor et al, Nature. 2013 Jul 11;499(7457):233-7				
GFP-SNX12	J. Gruenberg		Pons et al, PLoS One. 2012;7(6):e38949				
SNX17-GFP	R. Boettcher		Böttcher et al, Nat Cell Biol. 2012 May 6;14(6):584-92				
SNX27-GFP	R. Boettcher		Tseng et al, J Mol Biol. 2014 Sep 9;426(18):3180-3194				
SNX31-GFP	R. Boettcher		Tseng et al, J Mol Biol. 2014 Sep 9;426(18):3180-3194				
PX-p40phox-GFP	Addgene #19010		Derived from Addgene, desposited by M. Yaffe				
PHD-Ing2x3-GFP	Addgene #21589		Derived from Yuan	Addgene, despo	sited by J.		
MLNx2-GFP	T. Balla		Hammond et al, PLoS One. 2015 Oct 13;10(10):e0139957				
P4M-SidM-GFP	T. Balla		Hammond et al, J Cell Biol. 2014 Apr 14;205(1):113-26				
TAPP1-PH-GFP	T. Balla		Kimber et al, Biochem J. 2002 Feb 1;361(Pt 3):525-36				
PH-PLCδ1-GFP	T. Balla		Várnai & Balla, J Cell Biol. 1998 Oct 19;143(2):501-10.				
PH-AKT-GFP	T. Balla		Sason et al, N Jan;20(1):544	1ol Biol Cell. 2009 -55	)		
EGFP-C1	Clontech						
EGFP-C2	Clontech						
pGEX-2T	GE Healthcare						
GFP-SNX3 siRNA- insensitive			This work				
GFP-SNX3-Y71A			This work				
GFP-SNX3-Y71A siRNA- insensitive			This work				
GST-SNX3			This work				

GST-SNX3-Y71A		This work
GFP-SNX3-ΔN		This work
GFP-SNX3-PX+C		This work
GFP-SNX3-N+α		This work
GFP-SNX3-ΔC		This work
GFP-SNX3-C		This work
GFP-SNX3-ΔN siRNA-		This work
GFP-SNX3-PX+C siRNA-		This work
insensitive		
insensitive		This work
GFP-Gal9		This work
KIF5A-GFP-CIBN	Addgene #102252	Derived from Addgene, deposited by B. Cui
KIF5A-GFP		This work
Myo1e-GFP	M. Krendel	Ouderkirk & Krendel, Exp Cell Res. 2014 Apr 1;322(2):265-76
GFP-Rab35	A. Echard	Klinkert & Echard, Traffic. 2016 Oct;17(10):1063-77
GFP-Gal9-∆CRD2		This work
GFP-Gal9-∆N		This work
GFP-Gal9-linker+CRD2		This work
GFP-Gal9-N+CRD1		This work
GFP-Gal9-CRD1+linker		This work
GFP-Gal9-ACRD2		This work
siRNA-insensitive		
GFP-Gal9-AN		This work
GEP-Gal9-N+CRD1		
siRNA-insensitive		This work
GFP-Gal9-CRD1+linker		
siRNA-insensitive		This work
GFP-Gal9-A46V		This work
siRNA-insensitive		I NIS WORK
GFP-Gal9-N137A		This work
siRNA-insensitive		
GFP-Gal9-R252A		This work
siRNA-insensitive		
GFP-Gal9-A46V+N173A		This work
siRNA-insensitive		
flotillin-1-mCherry	A. Echard	Baumann et al., J Biol Chem 2012 Nov;287(47): 39664-72

flotillin-2-mCherry	A. Echard	Baumann et al., J Biol Chem 2012 Nov;287(47): 39664-72
GFP-Rab6a	J. Kremerskothen	Rzomb et al., Infect Immun. 2003
		Oct;/1(10):5855-70
GFP-Rab8a	D. Sheff	
		May;19(5):2059-68
GFP-Rab10	K. Simons	Schucke et el., Traffic 2007 Jan;8(1): Epub
		2006
GFP-Rab18	Addgene #49550	Derived from Addgene, deposited by M.
		Scidmore
GFP-Rab21a	A.T. Jones	Simpson et al., J Cell Sci 2004
		Sept;117:6297-6311
GFP-SNAP23	M. Coppolino	Williams et al., Mol Biol Cell 2014
		Jul;25(13):2061-70