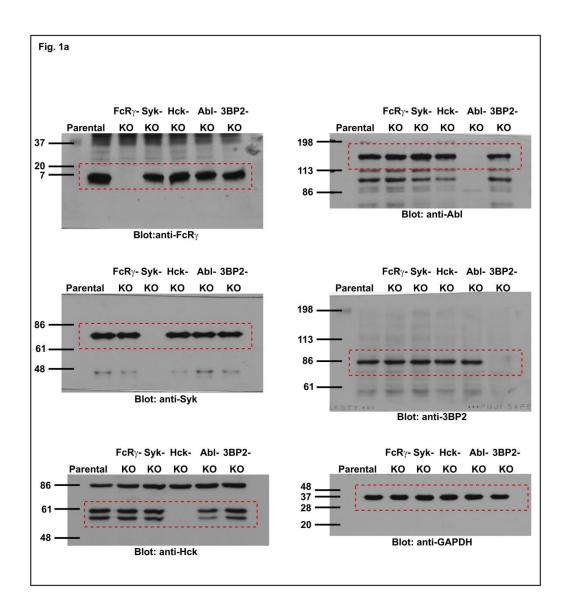
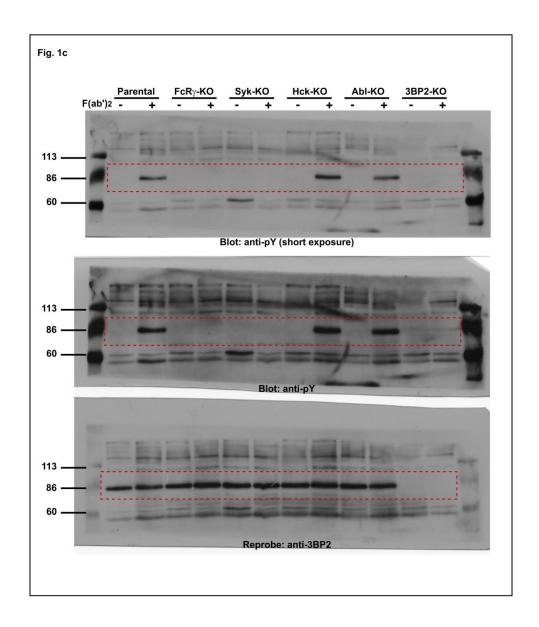
Supplementary information

The title of the manuscript: Syk-dependent tyrosine phosphorylation of 3BP2 is required for optimal FcR γ -mediated phagocytosis and chemokine expression in U937 cells

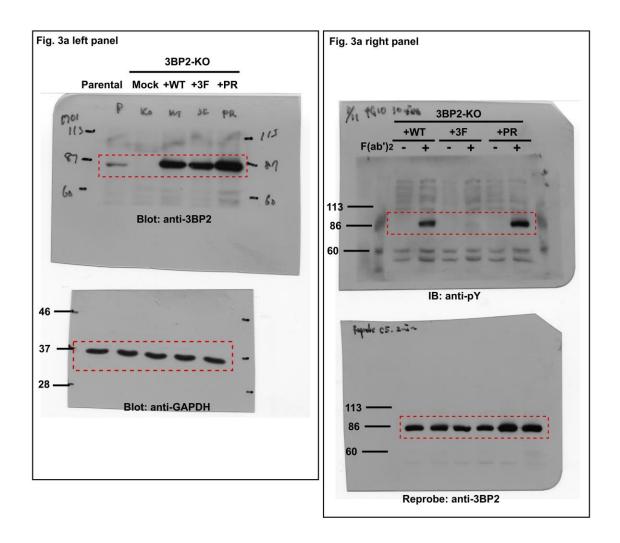
The author list: Kazuyasu Chihara, Yuji Kato, Hatsumi Yoshiki, Kenji Takeuchi, Shigeharu Fujieda, Kiyonao Sada



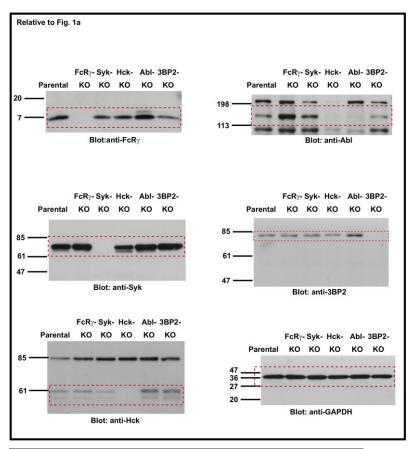
Full-length images of immunoblots presented in the main figures

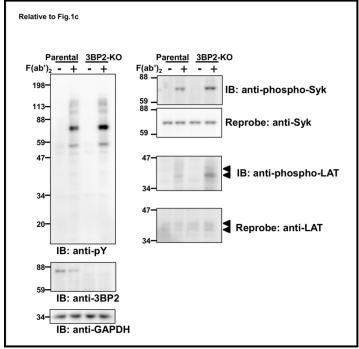


Full-length images of immunoblots presented in the main figures

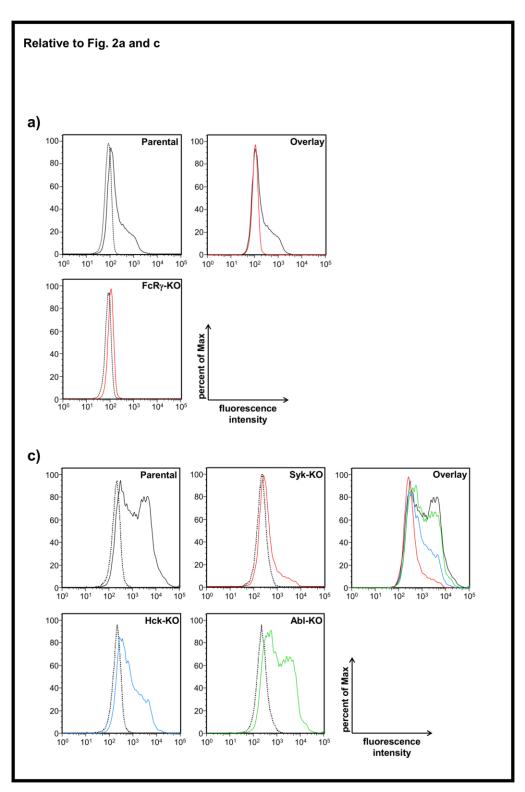


Full-length images of immunoblots presented in the main figures

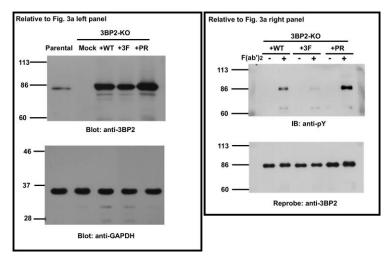


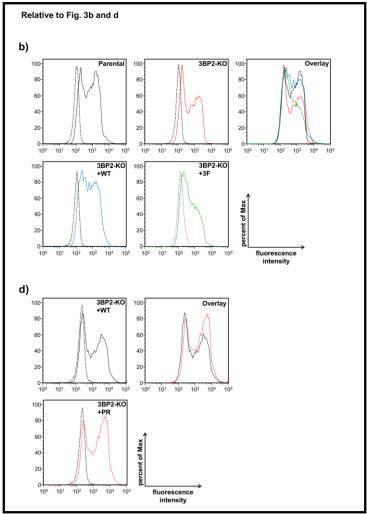


Similar results obtained from the other clones



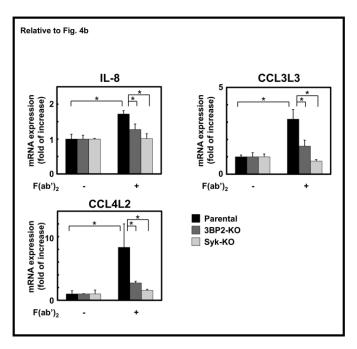
Supplementary Figure S2Similar results obtained from the other clones

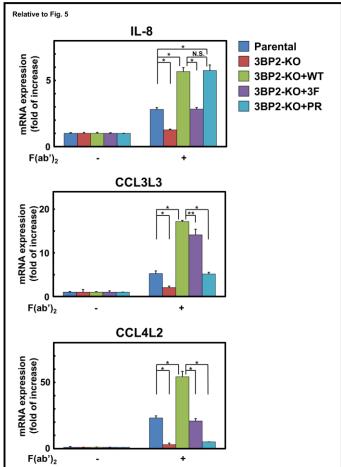




Supplementary Figure S2

Similar results obtained from the other clones



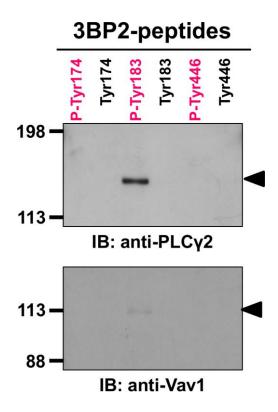


Supplementary Figure S2

Similar results obtained from the other clones

Supplementary Materials

Peptide binding experiment was performed as described (Shukla *et al. J. Biol. Chem.*, 2009). In brief, biotinylated peptides corresponding to residues between 166 to 182, 177 to 193 or 438 to 453 of mouse 3BP2 were synthesized (3BP2-peptide Tyr174, 183 and 446 respectively). Peptides containing phosphorylated tyrosine residue corresponding to Tyr174, 183 or 446 of mouse 3BP2 were also synthesized (3BP2 peptide P-Tyr174, 183 and 446 respectively). U937 cell lysates were reacted with these peptides prebound to streptavidin beads. Interaction of the peptides with PLC γ 2 and Vav1 were analyzed by immunoblotting. Anti-PLC γ 2 and anti-Vav1 antibodies were obtained from Santa Cruz Biotechnology.



Supplementary Figure S3. Interaction of 3BP2 with PLCy2 and Vav1.

U937 cell lysates were reacted with the indicated peptides prebound to streptavidin beads. Interaction of the peptides with $PLC\gamma2$ and Vav1 were analyzed by immunoblotting. Molecular size markers are indicated on the left in kDa. Data are representative of three independent experiments. Arrowheads indicate the position of $PLC\gamma2$ (upper panel) or Vav1 (lower panel).

Supplementary Table S1

Oligo DNA pairs to generate the single-guide RNA for the indicated genes

FcRγ: 5'-CACCGAGTAAGAGCAAGACCACTGC-3'

3'-CTCATTCTCGTTCTGGTGACGCAAA-5'

Syk: 5'-CACCGTTTCGGCAACATCACCCGGG-3'

3'-CAAAGCCGTTGTAGTGGGCCCCAAA-5'

Hck: 5'-CACCGTGGGATCCGGCACGTACAC-3'

3'-CACCCTAGGCCGTGCATGTGCAAA-5'

Abl: 5'-CACCGTCTGAGTGAAGCCGCTCGT-3'

3'-CAGACTCACTTCGGCGAGCACAAA-5'

3BP2: 5'-CACCGGGTACCGCCCTTCTTGTGC-3'

3'-CCCATGGCGGGAAGAACACGCAAA-5'

Supplementary Table S2

Calculated fold changes of FcyRI-induced genes in the indicated cell lines

Affymetrix Human	Cell type→	Parental		3BP2-KO		Syk-KO	
Gene 2.0 ST array	F(ab') ₂ →	-	+	-	+	-	+
Probe set ID	Gene Symbol	Log2 (fold change)		Log2 (fold change)		Log2 (fold change)	
16934764	ELFN2	-0.801667	0.248333	0.308333	-0.081666	0.298333	0.028333
17049237	STAG3	-0.926667	0.213333	0.083333	0.143333	-0.076667	0.563334
16723653	CD44	0.110000	1.390000	0.250000	-0.020000	-1.700000	-0.030000
16989736	EGR1	-0.683333	1.006666	-0.463333	0.686667	-0.313333	-0.233333
17075395	EGR3	-0.281667	1.338333	-0.531667	0.498333	-0.401667	-0.621667
16833426	CCL4L2	-0.298333	0.741667	-0.308333	0.201667	-0.098333	-0.238333
16761201	CD69	-0.551667	0.718333	-0.121666	0.338334	-0.451667	0.068333
16689869	F3	0.131667	1.461667	-0.368333	0.291667	-0.948333	-0.568334
17035418	TNF	0.438333	1.798334	-0.291667	-0.011667	-0.961667	-0.971667
16967771	IL8	-0.040000	2.700000	-0.330000	1.340000	-1.660000	-2.010000
17031687	IER3	0.335000	1.395000	0.085000	0.635000	-1.115000	-1.335000
16843602	CCL3L3	-0.233333	1.226667	-0.093333	0.076667	-0.303333	-0.673333
16901986	IL1B	-0.113333	0.936667	-0.293333	-0.203333	-0.123333	-0.203333