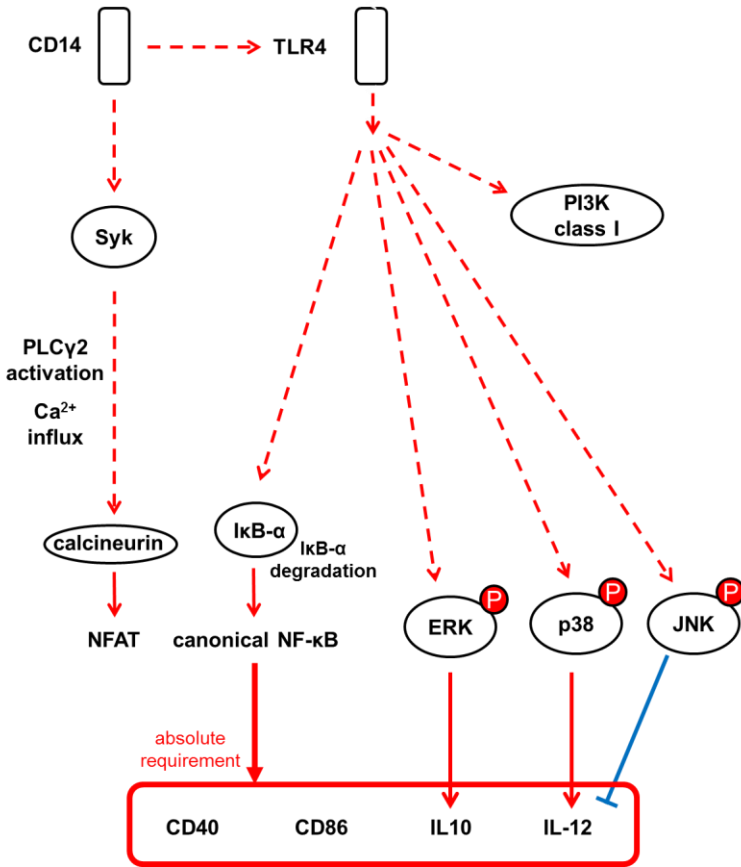


(a) Overview of relevant pathways



(b) PI3K-Akt-GSK3 pathway

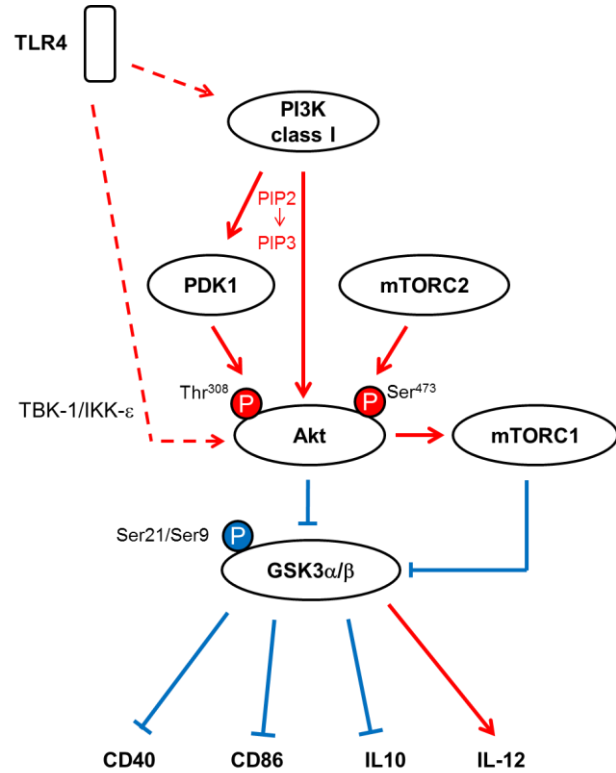


FIG S1. Summary of the known participation of the NF-κB, MAP kinases, Ca²⁺-calcineurin-NFAT and PI3K-Akt-GSK3 pathway in the activation of GM-CSF-BMDCs and other myeloid cell models in response to LPS. (See full legend on the next page).

FIG S1. Summary of the known participation of the NF- κ B, MAP kinases, Ca²⁺-calcineurin-NFAT and PI3K-Akt-GSK3 pathway in the activation of GM-CSF-BMDCs and other myeloid cell models in response to LPS.

For clarity, a general view of the pathways is presented in (a) and the PI3K-Akt-GSK3 pathway is detailed in (b). Only major points relevant to the present paper are shown. Red and blue lines represent activating and inhibitory effects respectively. Similarly, red and blue circles represent activating and inhibitory phosphorylation events respectively, and only phosphorylation events analyzed in the paper are represented.

LPS is mainly sensed by TLR4, in cooperation with CD14. TLR4 activation leads, through a complex series of steps, to the largely parallel activation (with certain early steps in common) of each of the MAP kinases (p38, JNK and ERK) and of the canonical NF- κ B pathway. The central step in canonical NF- κ B activation is the proteolytic degradation of I κ B- α , which releases active NF- κ B subunits for their nuclear translocation. Canonical NF- κ B activation is required for up-regulation of the co-stimulatory molecules CD86 and CD40 and for expression of the cytokines IL-10 and IL-12. The p38 MAP kinase is needed for full IL-12 expression; contradictory data are available about its effects on CD86, CD40 and IL-10. JNK downregulates IL-12 expression whereas its effects on CD86, CD40 and IL-10 are not well established. ERK is dispensable for CD86 expression, whereas its effects on CD40 expression are not established; ERK is needed for IL-10 expression and dispensable for IL-12 expression.

Also in response to LPS, CD14 initiates, independently of TLR4 and dependent on Syk and phospholipase C γ 2 (PLC γ 2), Ca²⁺ influx leading to calcineurin activation and therefore the activating dephosphorylation of NFAT. Information on the role of the NFAT pathway on expression of CD86, CD40, IL-10 and IL-12 in LPS-stimulated DCs is lacking or contradictory.

TLR4 activation additionally leads to activation of PI3K, through complex steps that are omitted for simplicity. PI3K catalyzes the conversion of phosphatidylinositol(4,5) biphosphate (PIP2) into phosphatidylinositol(3,4,5)trisphosphate (PIP3). PIP3 promotes the recruitment of PDK1 and Akt to the plasma membrane, facilitating the activation by phosphorylation of both these kinases. Active PDK1 and the mTORC2 complex doubly phosphorylate Akt, causing its full activation. The kinases TBK-1 and IKK- ϵ are also required for LPS-driven Akt phosphorylation in BMDCs, but the exact mechanism by which they act is unclear. Akt directly phosphorylates GSK3, inactivating this kinase. Akt additionally, and indirectly, activates the mTORC1 complex, which also leads to GSK3 phosphorylation through the p70S6K kinase. Active GSK3 has negative effects on CD86 and CD40 and IL-10 expression, and a positive effect on IL-12 expression. Thus, activation of the PI3K pathway enhances the expression of CD40, CD86 and IL-10, through a “double-negative” (inhibition-inhibition) mechanism; in contrast, activation of the pathway inhibits expression of IL-12.

The diagram is based on references (1-22); see “References in Supplementary Materials”, at the end of this document.

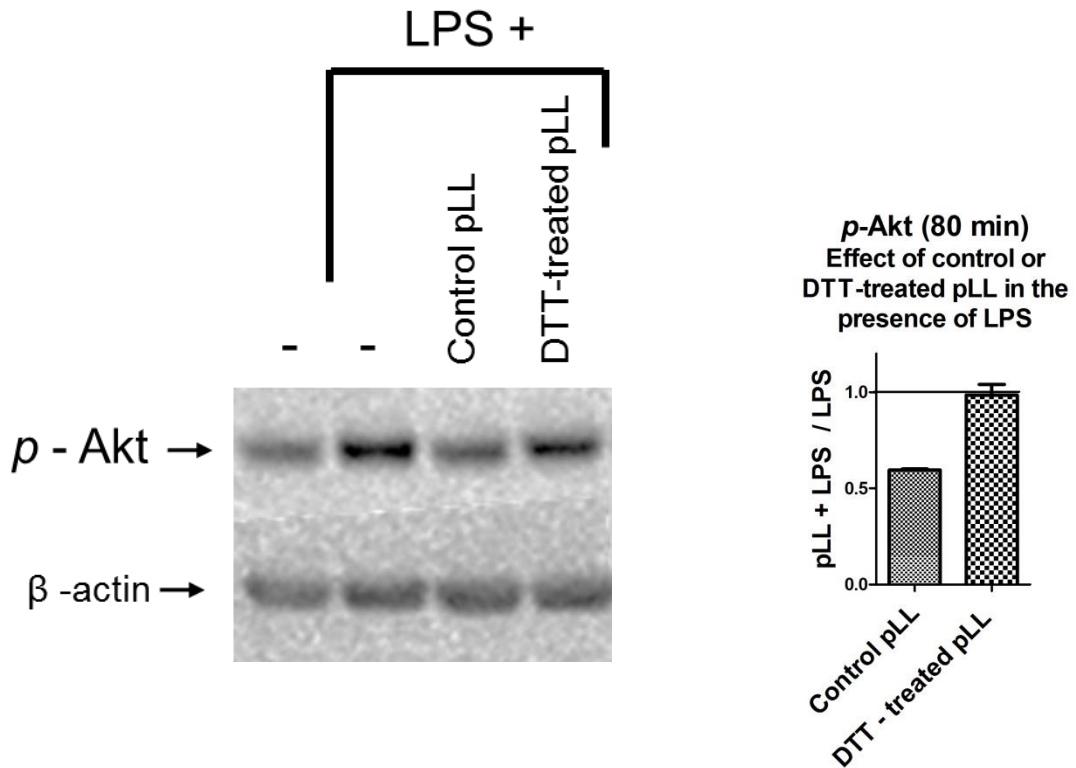


FIG S2 (related to Fig. 1). Reduction of disulphides in pLL abrogates its capacity to inhibit Akt phosphorylation in GMCSF-BMDCs. GMCSF-BMDCs were exposed to medium only, LPS, or LPS together with either pLL or pLL previously treated with DTT for disulfide reduction. After 80 min, cell lysates were prepared, and subsequently analyzed for phosphorylated Akt. The result shown is representative of 2 independent experiments. The graphs show the quotients of p-Akt values (normalized over loading controls) for cells treated with pLL (either control or DTT-treated) plus LPS over cells exposed to LPS only; values shown are mean and range of the 2 independent experiments.

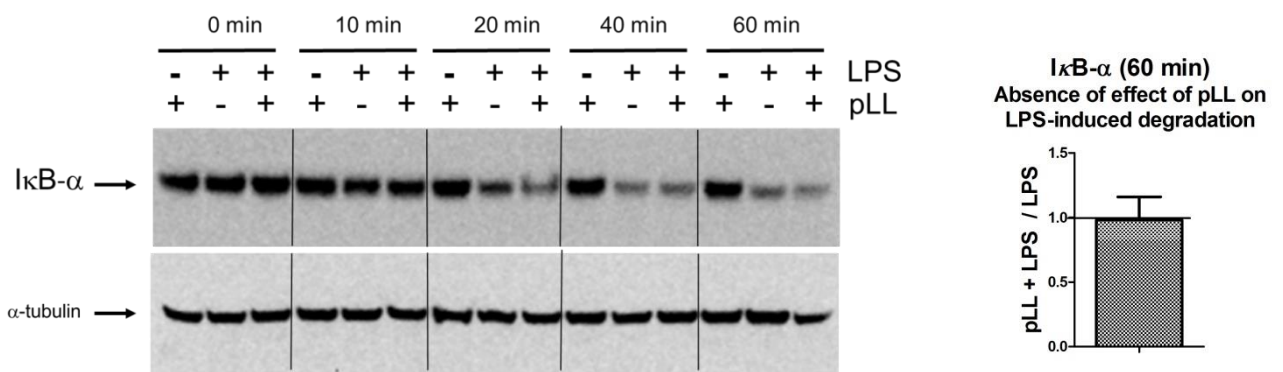


FIG S3. The effects of pLL on GMCSF-BMDCs are not associated with gross alteration in the activation of the canonical NF- κ B pathway. GMCSF-BMDCs were exposed to pLL, LPS, or both stimuli together, for the indicated times. Cell lysates were analyzed for total I κ B- α . The Western blot shown is representative of 2 independent experiments. The graph shows the quotients of the I κ B- α amounts (normalized over loading controls) for cells treated with pLL plus LPS over cells exposed to LPS only, for the 60-min timepoint; values are the mean and range of the 2 independent experiments.

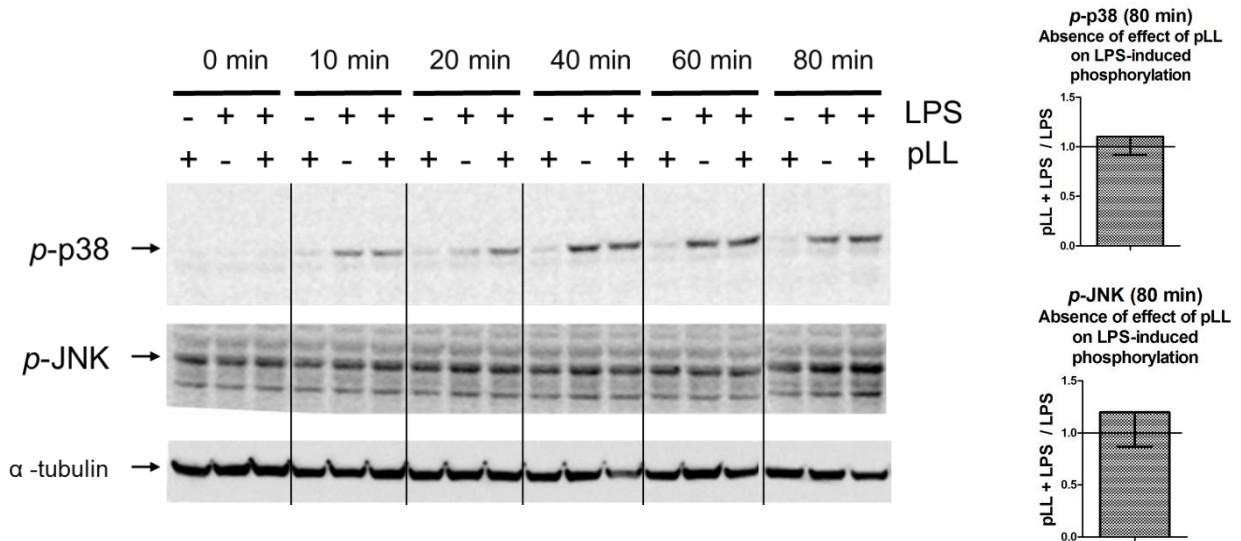


FIG S4. The effects of pLL on GMCSF-BMDCs are not associated with gross alterations in the activation of the p38 or JNK MAP kinases. GMCSF-BMDCs were exposed to pLL, LPS, or both stimuli together, for the indicated times. Cell lysates were analyzed for phosphorylated p38 and JNK MAP kinases. Results shown are representative of 3 independent experiments. The graphs show the quotients of *p*-p38 or *p*-JNK values respectively (normalized over loading controls) for cells treated with pLL plus LPS over cells treated with LPS only, for the 80-min timepoint; values shown are the mean and SD of the 3 independent experiments.

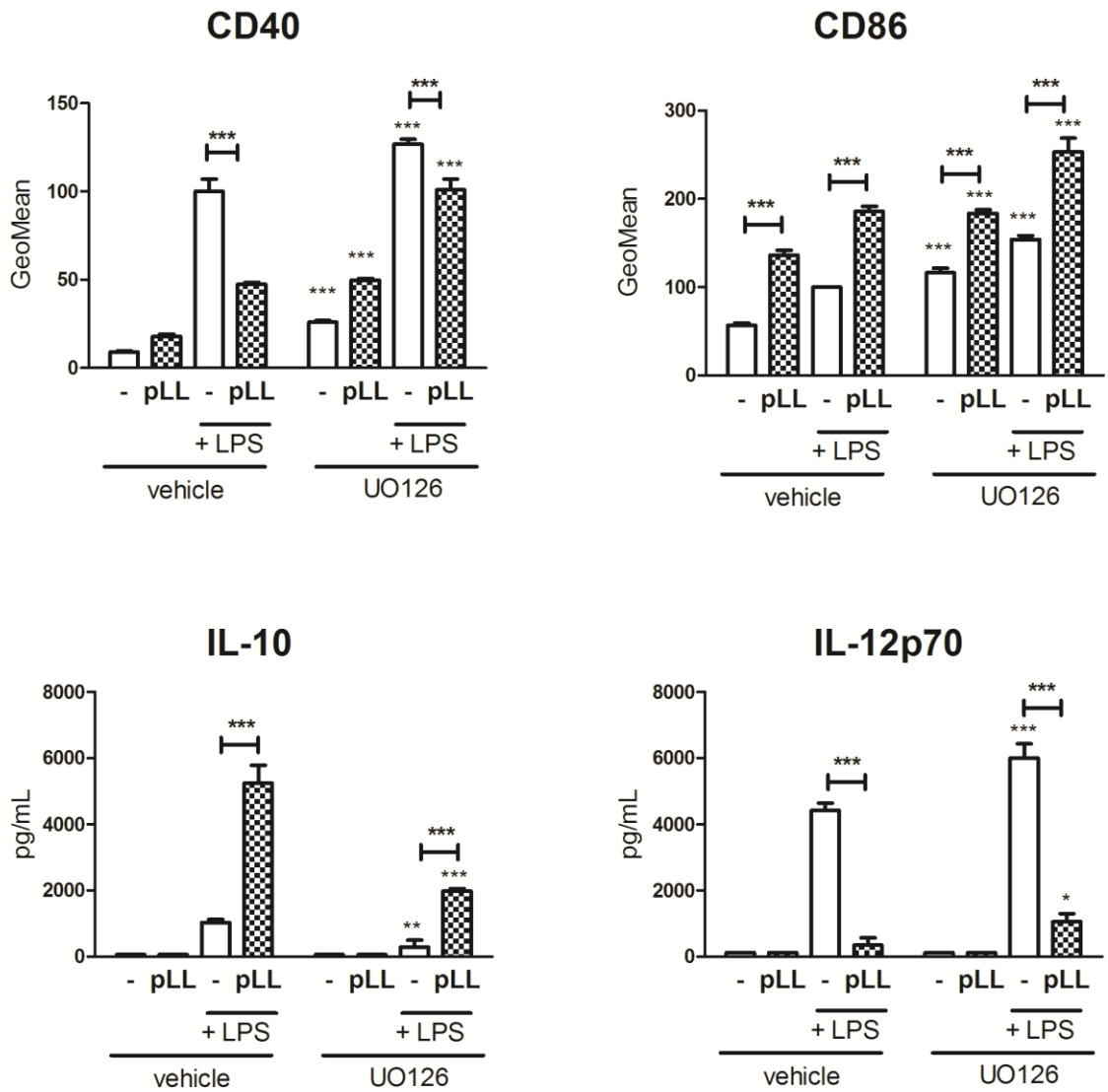


FIG S5. The effects of pLL on GMCSF-BMDCs are not abrogated by inhibition of ERK. GMCSF-BMDCs were exposed to pLL, LPS, or both stimuli together, for 18 h, in the absence or presence of the ERK activation inhibitor UO126. Cell surface expression of CD40 and CD86 and levels of IL-10 and IL-12p70 in supernatants were measured. All values plotted correspond to means \pm SD of triplicate wells. Results shown are representative of 2 independent experiments.

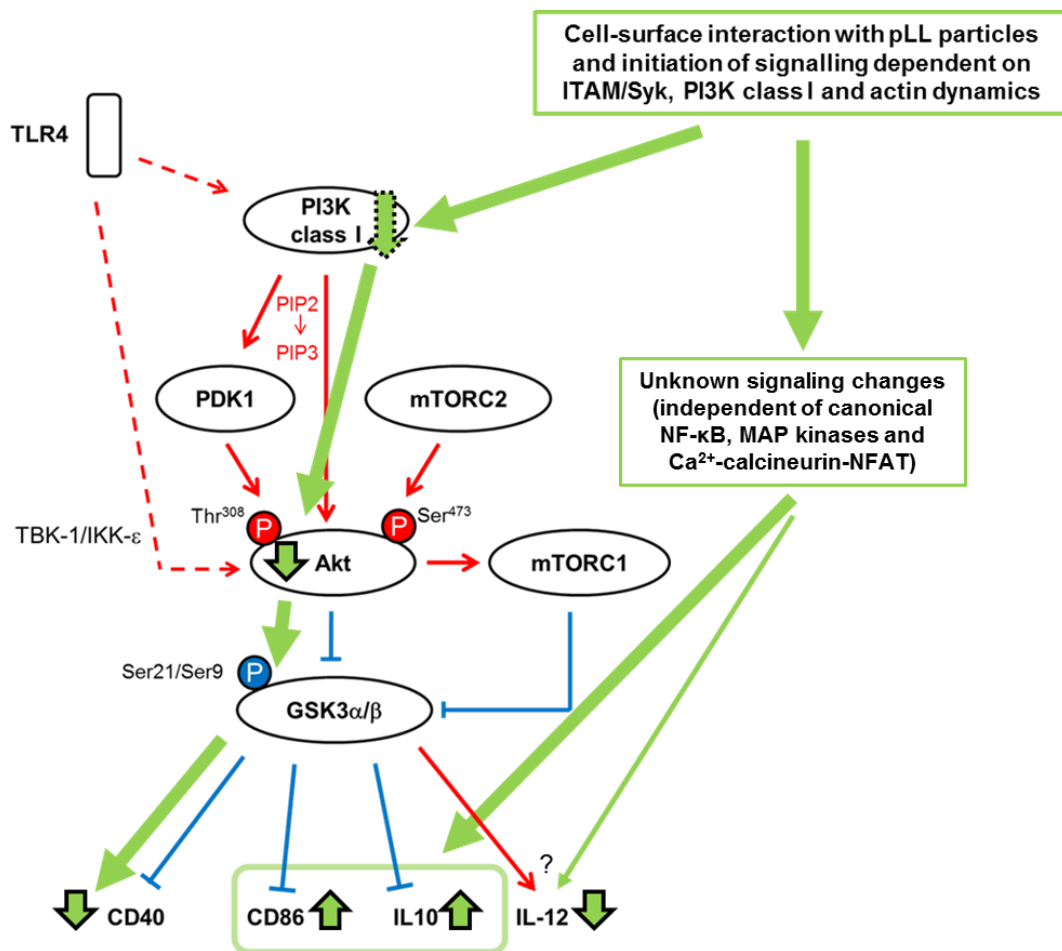


FIG S6. Summary of proposed mechanisms underlying the effects of pLL on GMCSF-BMDCs. The proposed changes brought about by pLL are shown in green, plotted onto the diagram shown in Fig. S1b (which summarizes relevant changes in the PI3K-Akt-GSK3 pathway induced by LPS). Mechanistic links are represented by long arrows without borders, and changes in kinase activity or in protein expression are shown by short arrows with black borders.

The cell-surface interaction and signaling envisaged is similar to membrane affinity-triggered signaling (MATS). MATS requires actin remodeling, PI3K and Syk, although the precise mechanistic relationships between these players have not been elucidated (23-25); the role of Syk in MATS would be independent from its role in LPS-stimulated, CD14-dependent signaling shown in Fig. S1a. We propose that in the context of a MATS-like interaction with pLL particles, receptor-activated PI3K class I activity is diminished, possibly as a consequence of decreased availability of its PIP2 substrate (24, 26).

Diminished PI3K activity in the presence of pLL blunts Akt activation, and hence enhances GSK3 activity, which in turn blunts CD40 up-regulation. Parallel, unidentified signaling changes downstream of the MATS-like interaction, independent of the NF-κB, MAP kinases and Ca²⁺-calcineurin-NFAT pathways, are responsible for the effects of pLL on CD86 and IL-10; these unidentified changes would overwhelm the opposite effects on CD86 and IL-10 brought about by the enhanced GSK3 activity. The effects of pLL on IL-12 expression would be brought about by Akt-dependent and Akt-independent signaling changes interacting in complex ways that we do not have a model for, as indicated by the question mark in the diagram.

Although not represented in the diagram, the MATS-like interaction with pLL and the blunting of receptor-activated PI3K activity would also arise in the absence of LPS stimulation, explaining that pLL also inhibits Akt activation in response to IL-4 or GM-CSF (27).

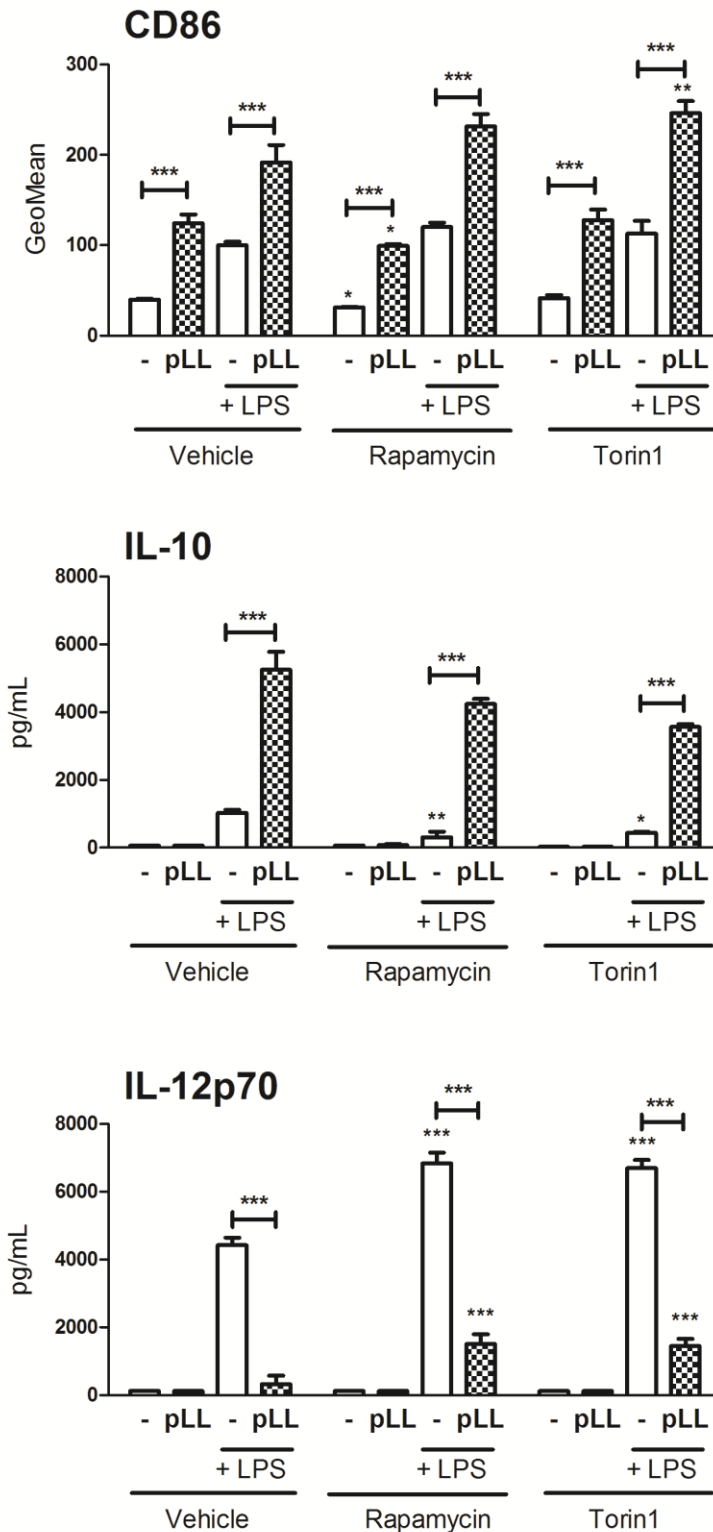


FIG S7 (related to Figs. 2 and 3). Effects of mTORC1 inhibitors on CD86, IL-10 and IL-12 expression in GMCSF-BMDCs, in the absence and presence of pLL. GMCSF-BMDCs were exposed to pLL, LPS, or both stimuli together, for 18 h, in the absence or presence of the mTORC1 inhibitors rapamycin or torin1. Cell surface expression of CD86, and IL-10 and IL-12p70 levels in supernatants were measured. All data shown correspond to means \pm SD of triplicate wells. Results shown are representative of 3 independent experiments.

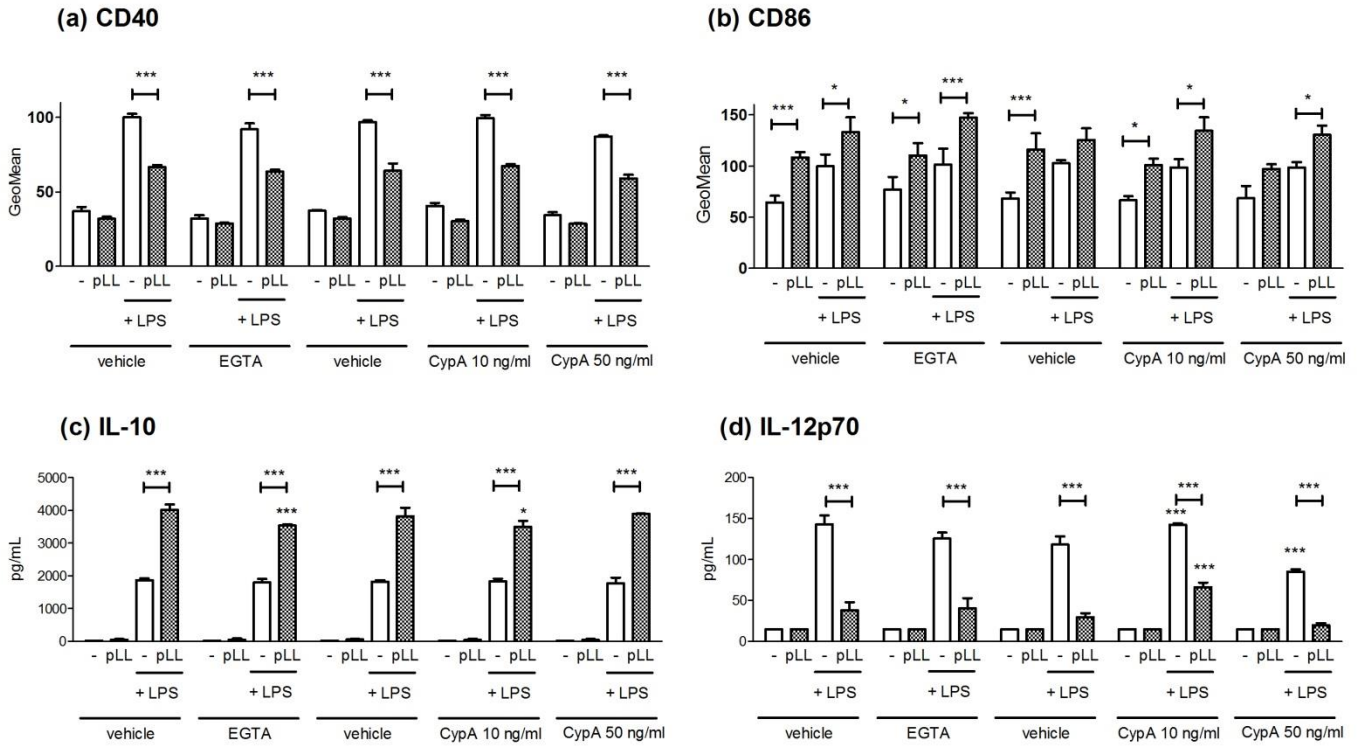


FIG S8. The effects of pLL on GMCSF-BMDCs are not abrogated by inhibitors of the calcium-calcineurin-NFAT pathway. GMCSF-BMDCs were exposed to pLL, LPS, or both stimuli together, for 18 h, in the absence or presence of the extracellular calcium sequestering agent EGTA or the calcineurin inhibitor cyclosporine A at two different concentrations. Cell surface expression of CD40 and CD86 and levels of IL-10 and IL-12p70 in supernatants were measured. All values plotted correspond to means \pm SD of triplicate wells. Results shown are representative of 2 or 3 independent experiments. The small effects of EGTA and cyclosporine A (10 ng/mL) on IL-10 and IL-12p70 production were not reproducible across experiments.

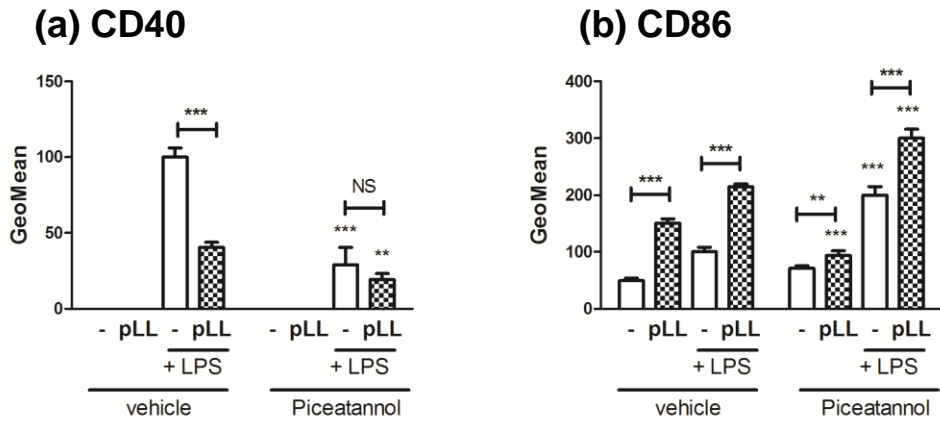
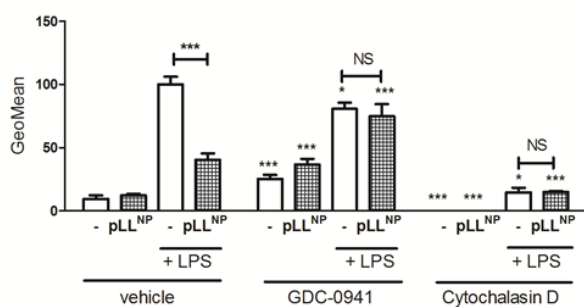
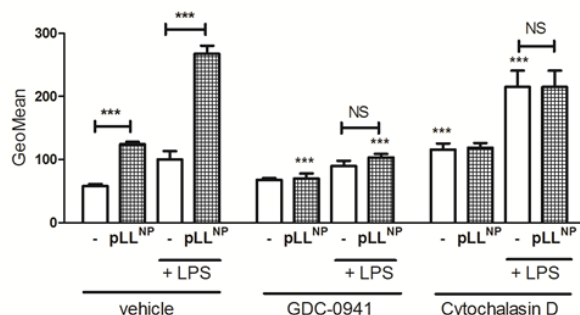


FIG S9. Syk may be involved in the effects of pLL on the phenotype of GMCSF-BMDCs. GMCSF-BMDCs were exposed to pLL, LPS, or both stimuli together, for 18 h, in the absence or presence of the Syk inhibitor piceatannol. Cell surface expression of CD40 (a) and CD86 was measured. All values plotted correspond to means \pm SD of triplicate wells. Results shown are representative of three independent experiments.

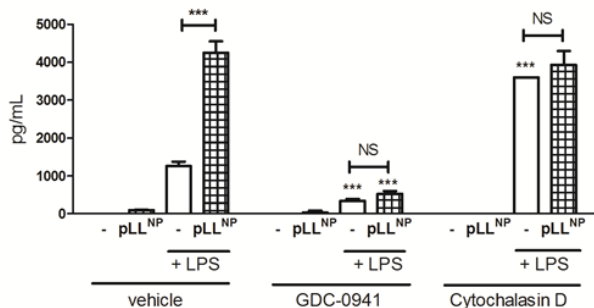
(a) CD40



(b) CD86



(c) IL-10



(d) IL-12p70

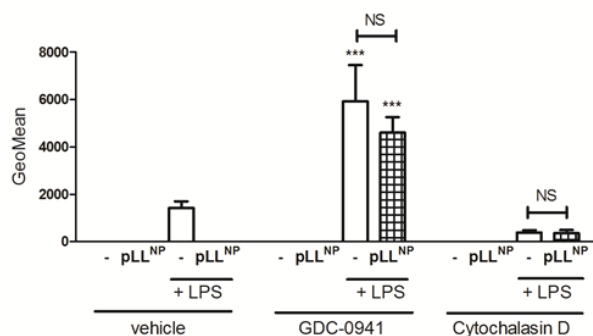


FIG S10 (related to Fig. 7). The phenotypic effects of non-phagocytosable pLL on GMCSF-BMDCs require actin dynamics and functional PI3Ks class I. GMCSF-BMDCs were exposed to pLL selected for non-phagocytosable particle size range (“pLL^{NP}”), LPS, or both stimuli together, for 18 h in the absence and presence of PI3K class I inhibitor (GDC-0941) or actin dynamics inhibitor (cytochalasin D), and cell surface expression of CD40 (a) and CD86 (b), and IL-10 (c) and IL-12p70 (d) levels in supernatants were measured. All data shown correspond to means +/- SD of triplicate wells. Results shown are representative of 2 independent experiments.

CD40

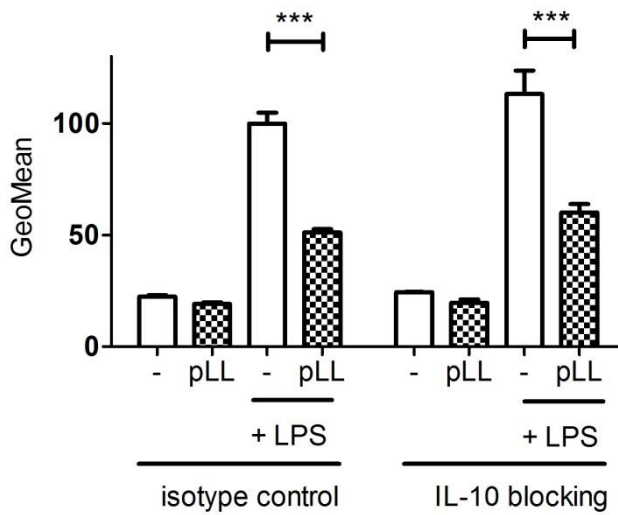


FIG S11. Paracrine IL-10 does not contribute to blunted CD40 up-regulation in GM-CSF-BMDCs exposed to pLL and co-stimulated with LPS. GM-CSF-BMDCs were exposed to pLL, LPS, or both stimuli together, for 18 h, in presence of either control antibody or an antibody blocking mouse IL-10 (both at 1 $\mu\text{g}/\text{mL}$), and cell surface expression of CD40 was measured. Data shown correspond to means \pm SD of triplicate wells. Results shown are representative of 3 independent experiments.

Respiration (pmol/min.µg protein)	Medium only	pLL	LPS	pLL + LPS
Basal	20 ± 2	24 ± 4	4.0 ± 0.8	6 ± 2
ATP independent	3.1 ± 0.3	3.9 ± 0.6	1.9 ± 0.5	3 ± 2
ATP dependent	17 ± 2	20 ± 3	2.1 ± 0.3	2.4 ± 0.4
Maximum	28 ± 2	32 ± 3	0.8 ± 0.4	2 ± 1
Non-mitochondrial	5 ± 3	10 ± 1	17 ± 1	16 ± 2

TABLE S1 Mitochondrial respiratory parameters of GMCSF-BMDCs stimulated with pLL and/or LPS. Respiratory parameters were calculated from the data shown in Fig. 4 as described in the Methods section. Results are presented as mean ± SD of triplicate wells.

Target	Supplier	Catalog number	Dilution
I κ B- α	Cell Signaling Technology	9242	1:1000
phospho-p38 (Thr ¹⁸⁰ /Tyr ¹⁸²)	Cell Signaling Technology	4511	1:1000
phospho-JNK/SAPK (Thr ¹⁸³ /Tyr ¹⁸⁵)	Cell Signaling Technology	4668	1:1000
phospho-ERK 1/2 (Thr ²⁰² /Tyr ²⁰⁴)	Cell Signaling Technology	4370	1:1000
phospho-Akt (Ser ⁴⁷³)	Cell Signaling Technology	4060	1:1000
Akt	Cell Signaling Technology	9272	1:1000
phospho-GSK3- α/β (Ser ²¹ /Ser ⁹)	Cell Signaling Technology	8566	1:1000
GSK3- β	Cell Signaling Technology	12456	1:1000
α -tubulin	Santa Cruz Biotechnology	sc-8035	1:250
β -actin	Cell Signaling Technology	4967	1:1000

Table S2. Antibodies used in Western blotting. The anti- α -tubulin primary antibody is a mouse IgM, and it was detected using a corresponding secondary antibody from Invitrogen (M31507). All the other primary antibodies are rabbit IgG, and were detected using corresponding secondary antibodies from Cell Signaling Technologies (7074) or Calbiochem (401353).

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