



FIG S3. CAR, SREC1, SR-A and MARCO expression in BMM and MPI cells. (A) Comparison of mRNA expression (normalized intensity values; NI) in BMM and MPI cells using microarray data published in (1).

(B) mRNA expression of indicated genes validated by qRT-PCR. Lung and B cell total RNA was used as positive and negative control, respectively. n. d.: not detectable

(C) FACS analysis of SR-A and MARCO expression in BMM and MPI cells. Open histograms: specific antibody, grey filled histograms: isotype control