

Figure S1. Family-wide transcriptomic analysis of porcine cytokines/chemokines and their receptor-related genes. Using PRRSV-infected macrophages at different activation statuses, the expression of most cytokines/chemokines, and their receptors at the transcriptomic level was revealed. In addition to inflammatory cytokines analyzed, including IL-1 β , IL-6, IL-8, and IL-12, multiple status-specific identifiers could be novel signature genes. (A) Differential expression of cytokines and their receptor-related genes. (B) Differential expression of chemokines and their receptor-related genes. (C) Differential expression of porcine Toll-like receptors genes (TLRs), which act as major innate immune receptors in regulation of cytokine/chemokine production corresponding to different activation statuses in monocytic cells. Transcripts in heatmaps were labeled with the official gene symbols from <http://www.ncbi.nlm.nih.gov/gene>. The color scale under each heatmap illustrates the midpoint and range of RPKM values of listed transcripts.

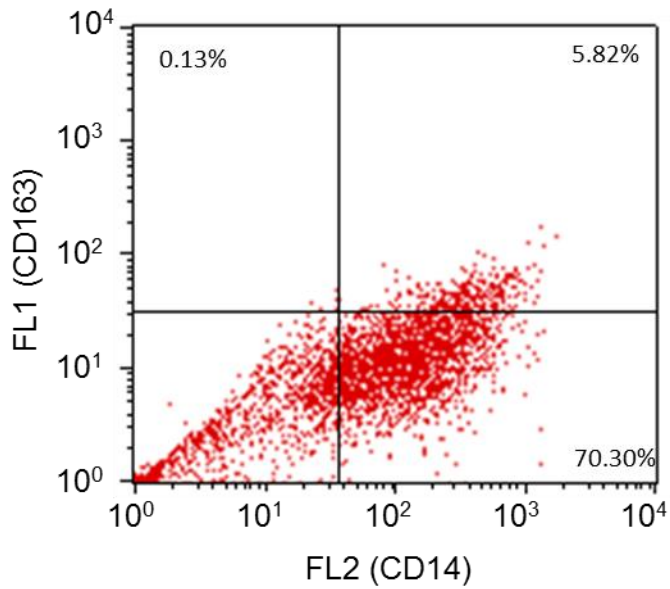


Figure S2. Two-color immunofluorescence facilitated flow cytometric analysis of purified monocytes with mAbs to CD14 and CD163. Monocytes used in this study are predominant CD14⁺ CD163^{low} (>90%) and few CD14⁺CD163^{high} (5.8%).

Supplemental Table S1: PCR primers used for RT-PCR assay of porcine GBPs and PRRSV detection

Gene	Primer sequence (5'-3')	Product size (bp)	GenBank RefSeq RNA Accession
<i>GBP1</i>	Forward: CTGAGTCCGGCACGAGTTTG Reverse: CACCAGTAAGCATCGCGCAT	167	NM_001128473.1
<i>GBP2</i>	Forward: ACAGGGATTCAAGAAGGAGA Reverse: TAGTTGAGTGGCAATGGTTT	191	NM_001128474.1
<i>GBP4</i>	Forward: AGGAAACGATGCTGGAGCACAA Reverse: ATGCAATGCTAGCCACGTCAAG	177	XM_001927929.3
<i>GBP5</i>	Forward: TATGACCAGCTGATGGGCCAGAAT Reverse: TCTGTTTGGCTTCTAGCAGGGTCT	175	XM_003125920.3
<i>GBP6</i>	Forward: TGCTGCAGAATGAAGAGGCA Reverse: TTTCTGGGCACTTGGCAAT	173	XM_003355299.1
<i>GBP1L</i>	Forward: ACCAGCTGACAGAGAAGCTGGAAA Reverse: TCTCTGCATTCATCTCCTGGGACT	174	XM_003481515.1
<i>GBP2L</i>	Forward: CAGAAACAACCTCCTTGAGGAGGGA Reverse: GGCGATGGTTTGGCATCTTGAAGT	199	XM_003125922.3
<i>GBP_{hp}</i>	Forward: GCTCAGTATGTGTGGGGCTT Reverse: TGATACGGTGAGTTCTGACACA	167	XM_003125925.1
<i>PRRSV-N</i>	Forward: ATGGCCAGCCAGTCAATCA Reverse: ACGGTCGCCCTAATTGAATAGGTG	436	AF494042
<i>GAPDH</i>	Forward: TGGYATCGTGGAAGGRCTCAT Reverse: RTGGGWGTYGCTGTTGAAGTC	370	NM_001206359.1