## Molecular Interaction Map of Macrophage

MI-MP-2004D Ver

Kanae Oda (1,2,3), Yukiko Matsuoka (4), Hiroaki Kitano (1,2,5)

(1) The Systems Biology Institute, (2) Department of Fundamental Science and Technology, Keio University,
(3) Department Molecular Epidemiology, Medical Research Institute, Tokyo Medical and Dental University,
(4) ERATO-SORST Kitano Symbiotic Systems Project, Japan Science and Technology Agency,
(5) Sony Computer Science Laboratories, Inc.

## Ver.2.0 update

Toll-like Receptor (TLR) pathways added to Macrophage Map Version 1.0 (Oda, et al., AfCS Report...)

Toll-like receptors (TLRs) signalings via myeloid differentiation primary response gene 88 (MyD88) -dependent and MyD88-independent pathways are added.

As input signals, we selected triacyl lipopeptide for TLR1/TLR2 complex, double-stranded RNA (ddRNA) for TLR3, flagellin for TLR5, lipoprotein for TLR2/TLR6 complex, single-stranded RNA (ssRNA) for TLR7, nonmethylated CpG DNA for TLR9.

