ABSTRACT

The invention of the application provides gene groups individually comprising 100 genes expressed most in human monocytes, GM-macrophages and M-macrophages, respectively. The cDNAs of the individual genes belonging to these gene groups contain nucleotide sequences of SQ ID Nos. 1 to 100, 101 to 200, and 201 to 300 consecutive to the nucleotide sequence CATG closest to the polyA region. Additionally, the invention of the application provides the cDNA groups of the genes belonging to the gene groups, the novel gene groups contained in these gene groups, and the oligonucleotide groups identifying these genes and cDNAs.