

SUPPLEMENTARY TABLE

Supplementary Table 2. Analysis of macrophage markers genes by univariate COX.

Gene	HR	HR.95L	HR.95H	P-value
CYP19A1	4.522	2.342	8.733	7.000*10 ⁻⁶
LIMD2	0.912	0.871	0.955	8.030*10 ⁻⁵
MYO9B	0.944	0.914	0.975	5.717*10 ⁻⁴
DOT1L	0.885	0.822	0.953	1.202*10 ⁻³
CD52	0.986	0.977	0.995	1.462*10 ⁻³
CCL1	1.407	1.128	1.754	2.407*10 ⁻³
CD48	0.938	0.900	0.978	2.535*10 ⁻³
BCAP31	1.003	1.001	1.005	6.353*10 ⁻³
NCKAP1L	0.914	0.856	0.976	6.977*10 ⁻³
VPS35	1.037	1.010	1.066	7.119*10 ⁻³
MMP19	0.938	0.894	0.984	9.268*10 ⁻³
CLCN7	0.933	0.886	0.983	9.412*10 ⁻³
FDX1	1.078	1.018	1.140	9.683*10 ⁻³
UTP3	1.028	1.006	1.051	1.257*10 ⁻²
TMEM9B	1.030	1.006	1.054	1.369*10 ⁻²
IL10	0.622	0.420	0.921	1.761*10 ⁻²
FANCE	0.975	0.954	0.996	2.078*10 ⁻²
ANXA2	1.002	1.000	1.003	2.096*10 ⁻²
S1PR2	0.900	0.822	0.984	2.128*10 ⁻²
SDCBP	1.014	1.002	1.026	2.437*10 ⁻²
MYH11	1.013	1.002	1.025	2.625*10 ⁻²
SLC6A7	0.013	0.000	0.606	2.669*10 ⁻²
AGPS	1.035	1.004	1.068	2.672*10 ⁻²
DNAJC13	0.958	0.922	0.995	2.702*10 ⁻²
CD84	0.881	0.788	0.986	2.781*10 ⁻²
STIP1	1.006	1.001	1.011	3.085*10 ⁻²
MYOF	1.009	1.001	1.017	3.282*10 ⁻²
IL12B	0.208	0.049	0.886	3.365*10 ⁻²
NAGPA	0.863	0.752	0.990	3.483*10 ⁻²
DNASE1L3	1.018	1.001	1.035	3.803*10 ⁻²
OSBPL11	0.958	0.919	0.998	4.025*10 ⁻²
SCAMP2	1.016	1.001	1.032	4.050*10 ⁻²
ADAMDEC1	0.966	0.934	0.999	4.106*10 ⁻²
DLAT	1.036	1.001	1.073	4.375*10 ⁻²
UBXN6	0.981	0.963	0.999	4.409*10 ⁻²
GP1BA	0.749	0.561	1.000	4.981*10 ⁻²