Supplementary Table S3. Protein bands intensity signal obtained by Western-blot analysis.

	CTRL group	MM group	PM group	p-value
APOA1	1358 ± 116	139 ± 52	78 ± 44	0.00008 (MM vs CTRL) 0.00006 (PM vs CTRL)
TTHY	886 ± 191	4149 ± 1112	3685 ± 1028	0.0074 (MM vs CTRL) 0.0097 (PM vs CTRL)
S10A8	697 ± 233	1917 ± 151	1304 ± 330	0.0016 (MM vs CTRL) 0.0429 (MM vs PM)
UROM	717 ± 64	893 ± 100	1170 ± 99	0.0027 (PM vs CTRL) 0.0268 (PM vs MM)
AMBP	404 ± 157	836 ± 237	1694 ± 92	0.00025 (PM vs CTRL) 0.00426 (PM vs MM)
GELS	592 ± 75	727 ± 159	1514 ± 172	0.0011 (PM vs CTRL) 0.0044 (PM vs MM)
PTGDS	4485 ± 1235	5570 ± 1216	9163 ± 1183	0.0091 (PM vs CTRL) 0.0214 (PM vs MM)

Reported data are indicative of protein intensity signal derived from band optical density and area detected in each group. Data are expressed as mean of three independent experiments for each protein ± standard deviation. P-values derived from Student's *t*-test ($p<0.05$ is considered as statistically significant).