

Table 1. Results of indentification of protein spots changed by doxycycline treatment

PROTEIN SPOT (SSP)	PROTEIN SCORE	PROTEIN SCORE C.I. %	TOTAL ION SCORE	TOTAL ION C.I. %	PROTEIN IDENTIFICATION
1201	122	100	111	100	sRAGE
1202	621	100	537	100	Peroxiredoxin II
2202	436	100	400	100	Albumin, isoform CRA_b
2403	249	100	135	100	Unnamed protein
3202	511	100	452	100	Unnamed protein
3204	170	100	164	100	Albumin, isoform CRA_a
3401	1170	100	918	100	Apolipoprotein A-I
5402	883	100	701	100	Albumin
7704	1150	100	918	100	Serum Albumin

Table 2. Correlations between levels of identified protein

PROTEINS	R=	p=
sRAGE and Albumin	0.87	0.001
sRAGE and Albumin, isoform CRA_a	0.96	0.0001
sRAGE and Apolipoprotein A-I	0.72	0.029
Peroxiredoxin II and Serum albumin	0.86	0.005
Albumin and Albumin, isoform CRA_a	0.85	0.002
Albumin and Unnamed protein (SSP 2403)	0.73	0.022
Apolipoprotein A-I and Albumin, isoform CRA_a	0.67	0.049