Supplementary Table S6. Logistic regression relating *SERPINA3* expression to tumor-associated immune-cell signatures

	Term	p-value	FDR	Estimate
G1	B cells (G1)	0.000854481	0.001424136	−0.782687075
G2	Plasma cells (G2)	4.60135758176704e-07	2.30067879088352e-06	0.641045432
G3	Monocytes/Macrophages (G3)	0.003177761	0.004539659	0.843981651
G4	Dendritic cells (G4)	0.000752671	0.001424136	−1.086410369
G5	Lymphocytes (G5)	1.39614103583034e-05	4.65380345276779e-05	0.627539321
G6	Exhausted CD8 T cells (G6)	0.007863757	0.009829696	0.922081598
G7	Regulatory T cells (G7)	0.126559287	0.134187324	0.587426651
G8	Cytotoxic lymphocytes (G8)	6.61755751773632e-13	6.61755751773632e-12	−2.368179715
G9	Exhausted/HS CD8 T cells (G9)	0.134187324	0.134187324	0.539359275
G10	Memory T cells (G10)	0.000294993	0.000737483	0.908751447

SERPINA3: serpin family A member 3; FDR: false discovery rate.