

S6 table. Gene ontology analysis of albumin-depleted proteins with p -value ≤ 0.05 (Holm-Bonferroni corrected)

Cellular component Bonferroni method $p \leq 0.05$	
Exosomes	YWHAG; ACTB; SERPINA1; AHSBG; A2M; APOA1; APOA2; APOD; APOE; APOL1; C4BPA; C4BPB; CALM1; CALM2; CALM3; CD5L; CP; CLU; CFL1; C1QB; C1QC; C1R; C3; C4A; C4B; CFH; DCD; ECM1; FGA; FGB; FGG; FCN2; FCN3; LGALS3BP; GSN; GAPDH; HP; HPR; HBA1; HBA2; HBB; HBD; HPX; JCHAIN; ITGA2B; ITIH4; JUP; KRT10; KRT12; KRT13; KRT14; KRT16; KRT17; KRT18; KRT19; KRT20; KRT9; KRT77; KRT2; KRT3; KRT4; KRT5; KRT6A; KRT6B; KRT7; KRT75; KRT8; MASP1; MYL6; MYL12B; POTEF; S100A9; TF; SH3BGR3; TTR; TPM3; TPM4; PROS1; VTN; AZGP1;
Cytoplasm	YWHAG; ACTB; SERPINA1; AHSBG; APOA1; APOA2; APOE; APOH; CALM1; CALM2; CALM3; CASP14; CIT; CLU; CFL1; C1QA; C1QC; C3; DCD; KHSRP; FGA; FGB; FGG; GSN; GFAP; GAPDH; HP; HBA1; HBA2; HBB; HBD; ITGA2B; JUP; KRT36; KRT10; KRT12; KRT13; KRT14; KRT16;
Extracellular	ACTB; ORM1; ORM2; SERPINA1; AHSBG; A2M; APOA1; APOA2; APOC3; APOD; APOE; APOL1; APOH; C4BPA; C4BPB; CASP14; CD5L; CP; CLU; CFL1; C1QA; C1QB; C1QC; C1R; C1S; C3; C4A; CFH; DCD; ECM1; FGA; FGB; FGG; FCN2; FCN3; LGALS3BP; GSN; GAPDH; HP; HPR; HBA1; HBB; HBD; HPX; JCHAIN; ITIH4; JUP; KRT17; KRT18; KRT9; KRT86; KRT2; KRT8; MASP1; OIT3; PPBP; CFP; S100A9; F2; TF; TTR; TPM3; TPM4; WDR48; GC; PROS1; VTN; AZGP1;
Extracellular region	ORM1; SERPINA1; AHSBG; A2M; APOA1; APOA2; APOC3; APOD; APOE; C4BPA; C4BPB; CALM1; CALM2; CALM3; CD5L; CP; C1QA; C1QB; C1QC; C1R; C1S; C3; C4A; DCD; FGA; FGB; FGG; FCN3; GSN; HP; HPX; JCHAIN; ITIH4; MASP1; PPBP; CFP; F2; TF; TTR; GC; PROS1; VTN; AZGP1;
Lysosome	YWHAG; ACTB; ORM1; SERPINA1; AHSBG; A2M; APOA1; APOA2; APOD; APOH; C4BPA; CALM1; CALM2; CALM3; CASP14; CP; CLU; CFL1; C4A; C4B; CFH; DCD; FGA; LGALS3BP; GSN; GAPDH; HP; HBA1; HBA2; HBB; HBD; HPX; ITGA2B; ITIH4; JUP; S100A9; TF; TTR; TPM3; TPM4; WDR48; VTN; AZGP1;
Cytoskeleton	YWHAG; ACTB; APOA1; APOE; CLU; CFL1; DCD; FGA; FGB; FGG; GSN; GAPDH; ITGA2B; JUP; KRT10; KRT12; KRT13; KRT14; KRT16; KRT18; KRT9; KRT77; KRT2; KRT5; KRT8; MYL6; MYL12A; PPBP; TTR; TPM3; TPM4;
Extracellular space	ORM1; ORM2; SERPINA1; AHSBG; APOA1; APOC3; APOD; APOE; APOL1; APOH; CD5L; CP; CLU; CFH; FGA; FGB; FGG; LGALS3BP; HPX; MASP1; CFP; F2; VTN;
Intermediate filament	GFAP; KRT10; KRT14; KRT16; KRT17; KRT18; KRT19; KRT20; KRT86; KRT2; KRT3; KRT4; KRT5; KRT6A; KRT7; KRT8;
Very-low-density lipoprotein particle	APOA1; APOA2; APOC3; APOE; APOL1; APOH;
Platelet alpha granule lumen	SERPINA1; A2M; FGA; FGB; FGG; PPBP;
Spherical high-density lipoprotein particle	APOA1; APOA2; APOC3; CLU; HPR;
High-density lipoprotein particle	APOA1; APOA2; APOE; APOL1; APOH;
Keratin filament	KRT14; KRT18; KRT3; KRT5;
Chylomicron	APOA2; APOC3; APOE; APOH;
Hemoglobin complex	HBA1; HBA2; HBB; HBD;
Fibrinogen complex	FGA; FGB; FGG;
Platelet alpha granule	FGA; FGB; FGG;
Complement component C1 complex	C1QA; C1QB;
Molecular function Bonferroni method $p \leq 0.05$	
Structural constituent of cytoskeleton	ACTB; GSN; KRT14; KRT16; KRT17; KRT18; KRT19; KRT85; KRT86; KRT3; KRT4; KRT7; MYL6; MYL12B; TPM4;
Transporter activity	APOA1; APOA2; APOC3; APOD; APOE; APOH; HP; HBA1; HBA2; HBB; HBD; HPX; TF; TTR; GC;
Structural molecule activity	GFAP; KRT36; KRT10; KRT12; KRT13; KRT9; KRT77; KRT5; KRT6A; KRT6B; KRT75; KRT8;
Complement activity	C4BPA; C4BPB; CLU; C1QA; C1QB; C1QC; C1R; C1S; C3; C4A; CFP;
Biological process Bonferroni method $p \leq 0.05$	
Cell growth and/or maintenance	ACTB; CFL1; ECM1; GSN; GFAP; KRT36; KRT10; KRT12; KRT13; KRT14; KRT16; KRT17; KRT18; KRT19; KRT9; KRT85; KRT86; KRT77; KRT2; KRT3; KRT4; KRT5; KRT6A; KRT6B; KRT7; KRT75; KRT8; MYL6; MYL12B; TMSB4X; TPM3; TPM4; VTN;
Immune response	ORM1; ORM2; C4BPA; C4BPB; CD5L; CLU; C1QA; C1QB; C1QC; C1R; C1S; C3; C4A; CFH; DCD; FCN2; FCN3; LGALS3BP; HP; HPR; JCHAIN; CFP; AZGP1;