

Table 1. MS analysis of AP-1A-coated liposomes

Protein	NCBI GI number	Predicted Mr (Da)	LC-MS/MS analysis		MALDI-TOF/TOF analysis			
			No. of sequenced peptides	Mascot score	Matched/total peptides	Sequence coverage	Peak number	
Coat components								
Clathrin heavy chain	66773801	191,557	33	317	47/75	38%	2	
Clathrin light chain B	62510439	25,172	3					
Clathrin light chain A	2493731	25,557	3					
AP-1 β 1	21541948	104,540	26	215	28/52	36%	5	
AP-1 γ 1	113349	91,202	21	203	27/47	43%	6	
AP-1 μ 1	543817	48,512	18	161	22/60	53%	8	
AP-1 α 1B	21541960	18,917	4					
AP-1 α 1A	48428720	18,721	5					
AP-1 γ -subunit binding protein 1 (γ -synergin)	34996507	122,005		100	16/37	18%		
ARF-1/ARF-3	51316986 47117658	20,553 20,457	9	75	9/39	56%	11	
ARF-GEF 2 (brefeldin A-inhibited)	63492672	181,465	8	78	25/81	22%	1	
G protein-coupled receptor kinase-interactor 1 (ARF-GAP Git1)	58864889	84,195	11	94	14/48	32%	7	
G protein-coupled receptor kinase-interactor 2 (ARF-GAP Git2)	18203126	78,747	9					
Arfaptin 1	63501125	41,518	3					
Arfaptin 2	67460562	37,773	5					
Actin nucleation								
Nck-associated protein 1 (HEM-2)	26986194	128,784	25	69	12/29	14%	4	
CYFIP2	19526988	145,589		388*	46/86	41%	3	
Abi-1	50400517	52,255	5					
Abi-2	50400259	49,356	7					
SH3 adapter protein SPIN90 (Nck-interacting protein with SH3 domain)	57015413	78,523	6					
WASP-family protein member 1	16877274	61,471	6					
WASP-family protein member 3	20071942	55,170	4					
ARP 2/3 complex subunit 1A	59797974	41,600	4					
ARP 2/3 complex subunit 2	23621467	34,357	4					
ARP 2/3 complex subunit 4	38372626	19,523	5					
ARP 2/3 complex subunit 3	62899893	20,525	2					
p21-Rac1	51702788	21,436	4	108	5/51	31%	10	
Cdc42 homolog	46397379	21,297	3					
SLIT-ROBO Rho-GAP3 (WAVE-associated Rac-GAP)	48428625	124,341	32	85	18/60	21%		
Rho-GEF 7 (PAK-interacting exchange factor β)	18202873	79,782	13	178	13/20	16%		
PAK 3	47117898	62,358	11					
Lipid modification								
PI-4-Kinase III β	51262078	89,887		78†	7/30	14%		
Membrane fusion								
Rab11B	1172815	24,474	11	86	9/47	54%	9	
Rab14	46577103	23,882	10	165	15/47	74%	9	
Rab4	15986733	23,539	4	131‡	6/50	38%		

* Combined mascot score: 2 sequenced peptides ([M+H]⁺ 1420.83¹⁰⁵³YAPLHLVPLIER¹⁰⁶⁴ and 1824.00⁴⁷⁷NTIYAALQDFAQVTLR⁴⁹²).

†Combined mascot score: 1 sequenced peptide ([M+H]⁺ 1187.63⁴⁸²EPVFIAAGDIR⁴⁹²).

‡Combined mascot score: 1 sequenced peptide ([M+H]⁺ 1406.75⁸⁰GAAGALLVYDITSR⁹³).