	Accession	Identified interaction	CenpA.S..Parental S exp1	CenpA.S..H3.S exp1	CenpA.S..CenpA.RdCycl exp1	CenpA.S.mg1..Trex.S.mg1 exp2	CenpA.S.mg1..H3.S.mg1 exp2	CenpA.S...CenpA.RdCycl	CenpA.S.mg1..CenpA.G1S.mg1 exp2	CenpaA.G1S.mg1..CenpA.G1S exp2
1	P49450	Histone H3-like centromeric protein A OS=Homo sapiens GN=CENPA PE=1 SV=1 - [CENPA_HUMAN]	100	100	0.576	71.45		0.479	1.698	
2	Q8NCD3	Holliday junction recognition protein OS=Homo sapiens GN=HJURP PE=1 SV=2 - [HJURP_HUMAN]	100	100	0.882	100	100	0.668	1.115	
3	P62987	Ubiquitin-60S ribosomal protein L40 OS=Homo sapiens GN=UBA52 PE=1 SV=2 - [RL40_HUMAN]	0.734	77.757						
4	P49411	Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2 - [EFTU_HUMAN]		50.223	100					
5	Q6PF15	Kelch-like protein 35 OS=Homo sapiens GN=KLHL35 PE=2 SV=2 - [KLH35_HUMAN]		34.018						
6	E9PKH2	Uncharacterized protein OS=Homo sapiens GN=SERPINH1 PE=4 SV=1 - [E9PKH2_HUMAN]		23.11						
7	Q03188	Centromere protein C 1 OS=Homo sapiens GN=CENPC1 PE=1 SV=2 - [CENPC_HUMAN]	1.231	8.529		100	96.004		0.856	
8	P62263	40S ribosomal protein S14 OS=Homo sapiens GN=RPS14 PE=1 SV=3 - [RS14_HUMAN]	45.577	7.385	0.693					
9	P10412	Histone H1.4 OS=Homo sapiens GN=HIST1H1E PE=1 SV=2 - [H14_HUMAN]	0.82	3.556	0.641					
10	B5ME40	Uncharacterized protein OS=Homo sapiens GN=SREK1IP1 PE=4 SV=2 - [B5ME40_HUMAN]	0.598	3.541	0.836					
11	P84243	Histone H3.3 OS=Homo sapiens GN=H3F3A PE=1 SV=2 - [H33_HUMAN]	1.087	2.642	1.642					
12	Q71DI3	Histone H3.2 OS=Homo sapiens GN=HIST2H3A PE=1 SV=3 - [H32_HUMAN]	1.009	1.461	1.9					
13	P16401	Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3 - [H15_HUMAN]	0.888	1.38	0.738	1.063	1.832	2.21	1.013	0.922
14	P23284	Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PPIB PE=1 SV=2 - [PPIB_HUMAN]	0.997	1.33	4.052					
15	P16403	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2 - [H12_HUMAN]	7.076	1.311	0.51	0.86	1.431	3.669	1.23	5.126
16	P42766	60S ribosomal protein L35 OS=Homo sapiens GN=RPL35 PE=1 SV=2 - [RL35_HUMAN]	2.628	1.114	3.059		0.916	0.974	1.169	1.813
17	Q969Q0	60S ribosomal protein L36a-like OS=Homo sapiens GN=RPL36AL PE=1 SV=3 - [RL36L_HUMAN]	0.822	1.049	3.741					
18	P49207	60S ribosomal protein L34 OS=Homo sapiens GN=RPL34 PE=1 SV=3 - [RL34_HUMAN]	0.732	0.997	0.821	2.869		66.907		
19	P62861	40S ribosomal protein S30 OS=Homo sapiens GN=FAU PE=1 SV=1 - [RS30_HUMAN]	0.686	0.993	0.851	1.05		2.803	1.081	
20	P26373	60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4 - [RL13_HUMAN]	0.663	0.975	0.89	34.827		0.865	1.616	2.276
21	Q2VPJ6	HSP90AA1 protein (Fragment) OS=Homo sapiens GN=HSP90AA1 PE=2 SV=1 - [Q2VPJ6_HUMAN]	0.951	0.944	1.082		50.07	1.259		
22	P06748	Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2 - [NPM_HUMAN]	5.772	0.907	0.917	97.695	0.284	0.798		
23	Q9NWT1	p21-activated protein kinase-interacting protein 1 OS=Homo sapiens GN=PAK1IP1 PE=1 SV=2 - [PK1IP_HUMAN]	0.748	0.879	2.134					
24	B5BU25	U2 small nuclear RNA auxiliary factor 2 isoform b OS=Homo sapiens GN=U2AF2 PE=2 SV=1 - [B5BU25_HUMAN]	0.82	0.845	1.774					
25	Q6DCA8	BCLAF1 protein (Fragment) OS=Homo sapiens GN=BCLAF1 PE=2 SV=1 - [Q6DCA8_HUMAN]		0.646			83.348			
26	Q13823	Nucleolar GTP-binding protein 2 OS=Homo sapiens GN=GNL2 PE=1 SV=1 - [NOG2_HUMAN]	0.832	0.595	0.625	97.332	0.017			
27	P11388	DNA topoisomerase 2-alpha OS=Homo sapiens GN=TOP2A PE=1 SV=3 - [TOP2A_HUMAN]	0.69	0.518	1.528	100	0.021	0.418	4.31	
28	Q16777	Histone H2A type 2-C OS=Homo sapiens GN=HIST2H2AC PE=1 SV=4 - [H2A2C_HUMAN]	0.358	0.362	0.597	100			23.512	
29	P62805	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2 - [H4_HUMAN]	0.832	0.201	0.843	16.882	0.092	0.658	1.876	0.818
30	B4DZC0	cDNA FLJ51771, highly similar to SWI/SNF-related matrix-associatedactin-dependent regulator of chromatin subfamily A member5 (EC 3.6.1.-) OS=Homo sapiens PE=2 SV=1 - [B4DZC0_HUMAN]		0.147		100	0.017			
31	P46013	Antigen KI-67 OS=Homo sapiens GN=MKI67 PE=1 SV=2 - [KI67_HUMAN]	0.864	0.118	0.732	100	0.016	0.317	39.449	
32	Q86T24	Transcriptional regulator Kaiso OS=Homo sapiens GN=ZBTB33 PE=1 SV=2 - [KAISO_HUMAN]		0.08		100	0.095	1.659	2.726	
33	Q09666	Neuroblast differentiation-associated protein AHNAK OS=Homo sapiens GN=AHNAK PE=1 SV=2 - [AHNK_HUMAN]		0.028		100	0.01			
34	P12270	Nucleoprotein TPR OS=Homo sapiens GN=TPR PE=1 SV=3 - [TPR_HUMAN]		0.011		100	0.011			
35	Q9NVP2	Histone chaperone ASF1B OS=Homo sapiens GN=ASF1B PE=1 SV=1 - [ASF1B_HUMAN]		0.01		100	0.01		1.904	
36	A6NDY5	Uncharacterized protein OS=Homo sapiens GN=SLC39A4 PE=4 SV=2 - [A6NDY5_HUMAN]	100		1.013					
37	A8K4D5	Kynureninase OS=Homo sapiens PE=2 SV=1 - [A8K4D5_HUMAN]					99.75			
38	A8K9C4	Elongation factor 1-alpha OS=Homo sapiens PE=2 SV=1 - [A8K9C4_HUMAN]					5.182			
39	A8K9J7	Histone H2B OS=Homo sapiens PE=2 SV=1 - [A8K9J7_HUMAN]				1.85	0.44	0.506	1.089	0.83
40	A8MPX7	Chromosome X open reading frame 56, isoform CRA_b OS=Homo sapiens GN=CXorf56 PE=2 SV=1 - [A8MPX7_HUMAN]				100	0.01			
41	B2R5U1	cDNA, FLJ92620, highly similar to Homo sapiens staphylococcal nuclease domain containing 1 (SND1),mRNA OS=Homo sapiens PE=2 SV=1 - [B2R5U1_HUMAN]					74.434			
42	B3GQS7	Mitochondrial heat shock 60kD protein 1 variant 1 OS=Homo sapiens GN=HSPD1 PE=2 SV=1 - [B3GQS7_HUMAN]			1.023		95.912			
43	B3KM80	Nucleolin, isoform CRA_c OS=Homo sapiens GN=NCL PE=2 SV=1 - [B3KM80_HUMAN]				78.657	0.191			
44	B3KM90	cDNA FLJ10529 fis, clone NT2RP2000965, highly similar to Targeting protein for Xklp2 OS=Homo sapiens PE=2 SV=1 - [B3KM90_HUMAN]					0.01		6.926	
45	B3KNZ4	cDNA FLJ30804 fis, clone FEBRA2001267, highly similar to Eukaryotic translation initiation factor 3 subunit 8 OS=Homo sapiens PE=2 SV=1 - [B3KNZ4_HUMAN]					83.781			
46	B3KVC6	cDNA FLJ16395 fis, clone TRACH2024139, highly similar to DNA polymerase eta (EC 2.7.7.7) (Fragment) OS=Homo sapiens PE=2 SV=1 - [B3KVC6_HUMAN]				100	0.014			
47	B4DRA2	cDNA FLJ57828, highly similar to Treacle protein (Fragment) OS=Homo sapiens PE=2 SV=1 - [B4DRA2_HUMAN]							7.536	
48	B4DRX3	60S ribosomal protein L6 OS=Homo sapiens PE=2 SV=1 - [B4DRX3_HUMAN]					2.62	0.68	0.881	1.216
49	B4E102	cDNA FLJ57766, moderately similar to Eukaryotic initiation factor 4A-I (EC 3.6.1.-) OS=Homo sapiens PE=2 SV=1 - [B4E102_HUMAN]					100			
50	B7Z4B4	cDNA FLJ59338, highly similar to Retinoic acid receptor gamma-2 OS=Homo sapiens PE=2 SV=1 - [B7Z4B4_HUMAN]				100			100	100
51	B7Z5E7	cDNA FLJ51046, highly similar to 60 kDa heat shock protein, mitochondrial OS=Homo sapiens PE=2 SV=1 - [B7Z5E7_HUMAN]				100		50.005	100	
52	E7ETG7	Uncharacterized protein OS=Homo sapiens GN=MCCC1 PE=4 SV=1 - [E7ETG7_HUMAN]					2.046			
53	E9PP36	Uncharacterized protein OS=Homo sapiens GN=RPL8 PE=4 SV=1 - [E9PP36_HUMAN]				13.677	0.078		0.073	
54	O95138	Nuclear autoantigenic sperm protein autosomal variant (Fragment) OS=Homo sapiens PE=2 SV=1 - [O95138_HUMAN]			4.641					
55	P07195	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2 - [LDHB_HUMAN]					8.713			
56	P0CG48	Polyubiquitin-C OS=Homo sapiens GN=UBC PE=1 SV=2 - [UBC_HUMAN]						0.633	1.218	28.772
57	P61956	Small ubiquitin-related modifier 2 OS=Homo sapiens GN=SUMO2 PE=1 SV=2 - [SUMO2_HUMAN]				100	0.036		1.331	
58	Q14568	Putative heat shock protein HSP 90-alpha A2 OS=Homo sapiens GN=HSP90AA2 PE=1 SV=2 - [HS902_HUMAN]							32.729	
59	Q14980	Nuclear mitotic apparatus protein 1 OS=Homo sapiens GN=NUMA1 PE=1 SV=2 - [NUMA1_HUMAN]				100	0.012	0.242	2.151	
60	Q16380	Metastasis-associated gene protein (Fragment) OS=Homo sapiens GN=metastasis-associated gene PE=2 SV=1 - [Q16380_HUMAN]				49.159		1.438		
61	Q5T3N0	Annexin A1 (Fragment) OS=Homo sapiens GN=ANXA1 PE=2 SV=1 - [Q5T3N0_HUMAN]					32.767			
62	Q5T624	Nuclear autoantigenic sperm protein (Histone-binding) OS=Homo sapiens GN=NASP PE=2 SV=1 - [Q5T624_HUMAN]							36.741	
63	Q6P0N0	Mis18-binding protein 1 OS=Homo sapiens GN=MIS18BP1 PE=1 SV=1 - [M18BP_HUMAN]				100			1.703	
64	Q8IUB0	Elongation factor 1-alpha OS=Homo sapiens PE=2 SV=1 - [Q8IUB0_HUMAN]				4.204			1.862	
65	Q96C32	UBC protein OS=Homo sapiens PE=2 SV=1 - [Q96C32_HUMAN]				3.468	0.149			
66	Q96KK5	Histone H2A type 1-H OS=Homo sapiens GN=HIST1H2AH PE=1 SV=3 - [H2A1H_HUMAN]	0.319			16.804	1.065	2.2	2.485	51.04
67	Q9BSL9	ATPIF1 protein OS=Homo sapiens PE=2 SV=1 - [Q9BSL9_HUMAN]					62.364			
68	Q9P2E9	Ribosome-binding protein 1 OS=Homo sapiens GN=RRBP1 PE=1 SV=4 - [RRBP1_HUMAN]				35.818			3.904	