

ID	Count	Intersection	Fisher's p-value	Net	ID	Gene Count	Genes	Top Ontology	PValue	Bonferroni	
1	8	CDK1, CDK2, MAD2L1, NUSAP1, RRM2, TOP2A, TPX2, UBE2T	0.00E+00	Single-Cell	15	26	ABHD4, ALG9 , AURKB , BIRC5 , BUB1 , CCNB2 , CDC6 , CDK1, CDK2, CENPF , DHFR, DSN1 , DTL , FANCI , HMGB2, KIF4A , KIF15 , MAD2L1, MLF1IP , NCAPG2 , NUSAP1, PBK , RRM2, TOP2A, TPX2, UBE2T	BP	nuclear division	1.248E-19	4.99046E-17
								CC	spindle	1.06759E-15	1.01E-13
								MF	ATP binding	6.26055E-05	6.43E-03
				Bulk	1	34	ABHD8 , APOC1 , CCNB1, CDC20 , CDK1, CDK2, CKS1B, CTPS , DNMT1, FOXM1 , KPNA2, LMNB1 , LMNB2 , MAD2L1, MCM2 , MCM3 , MCM4, MCM6 , NUSAP1, PRC1 , PTTG1, RACGAP1, RCC1 , RNASEH2A, RRM2, SMC4, TOP2A, TPX2, TYMS , UBE2C , UBE2T, UHRF1, ZWINT	BP	cell cycle	3.06719E-19	1.58E-16
								CC	nucleus	7.72832E-10	8.11E-08
								MF	ATP binding	0.000136382	1.90E-02
2	10	RPL10, RPL17, RPL24, RPL27, RPL31, RPL34, RPL37A, RPS15A, RPS27, RPS27A	0.00E+00	Single-Cell	3	49	ACIN1, C12orf29 , EEF1A1, GNB2L1, NPM1, RPL3, FAU, RPL7, RPL8, RPL9P9 , RPL10, RPL12, RPL15, RPL17, RPL19, RPL21, RPL23, RPL23A, RPL24, RPL26, RPL27, RPL27A, RPL28, RPL29, RPL30, RPL31, RPL32, RPL34, RPL35, RPL35A, RPL36, RPL37A, RPLP1, RPLP2, RPS5, RPS6, RPS14, RPS15, RPS15A, RPS16, RPS18, RPS19, RPS21, RPS25, RPS27, RPS27A, RPS28, SNORA66 , SNORD73A	BP	translational elongation	7.81991E-90	3.09E-87
								CC	ribosome	5.27654E-75	2.74E-73
								MF	structural constituent of ribosome	5.19883E-73	3.22E-71
				Bulk	4	21	ABCE1, ADD3, GASS , RPL10, RPL17, RPL24, RPL27, RPL31, RPL34, RPL37A, RPL39 , RPL5 , RPL9, RPS13, GASS , ia, RPS27, RPS27A, RPS3A , RPL39 , RPL5 , ;	BP	translational elongation	3.54033E-32	8.46E-30
								CC	ribosome	7.42534E-27	3.56E-25
								MF	structural constituent of ribosome	3.34645E-28	1.51E-26
3	2	CALR, HSPA5	1.00E-04	Single-Cell	4	9	AASDHPPT, ABCB6, CALR, HSPA5, DNAJB11, HYOU1, MIR3652 , PDIA6, PPIB	BP	protein folding	6.5377E-05	1.53E-02
								CC	endoplasmic reticulum lumen	5.90351E-11	4.19E-09
								MF	unfolded protein binding	1.46063E-05	1.11E-03
				Bulk	40	11	AATF, ABCE1, ANXA2, CALR, CHPF2 , HSPA5, MANF , PDIA4, PLOD3, TUBA1C	BP	response to biotic stimulus	0.018540391	9.96E-01
								CC	endoplasmic reticulum lumen	0.000878763	7.12E-02
								MF	enzyme inhibitor activity	0.010497711	6.77E-01
4	4	RPS5, RPS15, RPS16, RPS19	0.00E+00	Single-Cell	3	49	ACIN1, C12orf29 , EEF1A1, GNB2L1, NPM1, RPL3, FAU, RPL7, RPL8, RPL9P9 , RPL10, RPL12, RPL15, RPL17, RPL19, RPL21, RPL23, RPL23A, RPL24, RPL26, RPL27, RPL27A, RPL28, RPL29, RPL30, RPL31, RPL32, RPL34, RPL35, RPL35A, RPL36, RPL37A, RPLP1, RPLP2, RPS5, RPS6, RPS14, RPS15, RPS15A, RPS16, RPS18, RPS19, RPS21, RPS25, RPS27, RPS27A, RPS28, SNORA66 , SNORD73A	BP	translational elongation	7.81991E-90	3.09E-87
								CC	ribosome	5.27654E-75	2.74E-73
								MF	structural constituent of ribosome	5.19883E-73	3.22E-71
				Bulk	5	10	AASS, ABCB7 , RPL13AP5 , RPL18 , RPS15, RPS16, RPS19, RPS5, RPS9	BP	NONE	-	-
								CC	NONE	-	-
								MF	NONE	-	-
7	3	AASS, HLA-B, HLA-A, HLA-C	0.00E+00	Single-Cell	18	7	AARSD1, AASS, CD74, HLA-B, HLA-A, HLA-C, IFI6	BP	antigen processing and presentation of peptide antigen	3.78032E-05	9.07E-03
								CC	MHC class I protein complex	0.008770777	3.39E-01
								MF	MHC class I receptor activity	0.005601304	1.78E-01
				Bulk	7	10	AASS, ABCD3, B2M, HLA-B, HLA-A, HLA-C, PSMB8 , PSMB9 , TAP1	BP	antigen processing and presentation	3.8373E-08	6.33E-06
								CC	MHC class I protein complex	6.23979E-05	3.55E-03
								MF	MHC class I receptor activity	0.007833546	3.61E-01