Gene	Feature	I	I:Pval	1	l:Pval	K	K:Pval
Cyclophilin	exonNum	1.000e+00	1.00e+00	9.999e-01	NA	NA	NA
EF1a	exonNum	-1.437e-01	4.78e-01	6.611e-05	1.000e+00	5.419e-04	6.076 e-01
Hsp90	exonNum	-1.963e-01	7.14e-01	6.611e-05	1.000e+00	3.345 e-03	2.142e-01
IPS1	exonNum	-1.859e-03	1.68e-01	9.999e-01	8.885 e-02	5.294 e-01	6.006e-03
PHO2	exonNum	-7.807e-02	3.35e-01	9.999e-01	1.702e-01	4.454 e - 04	7.397e-01
PHR1	exonNum	-1.673e-01	5.62e-01	9.999e-01	5.313e-02	4.395e-04	7.287e-01
SIZ1	exonNum	-2.869e-01	8.61e-01	6.611 e-05	1.000e+00	3.075e-04	8.128e-01
Cyclophilin	ORFLen	1.655e-02	1.32e-01	2.086e-01	8.010e-01	6.377e-01	3.103 e-02
EF1a	ORFLen	4.140e-02	1.52e-01	2.209 e-01	7.115e-01	4.758e-01	5.305 e-02
Hsp90	ORFLen	6.178e-02	1.30e-01	6.611 e-05	1.000e+00	4.597e-02	1.512e-01
IPS1	ORFLen	-3.980e-02	3.13e-01	9.999e-01	2.980e-02	5.495 e-04	6.036 e-01
PHO2	ORFLen	2.489e-02	2.00e-01	9.999e-01	4.739e-02	2.327e-02	2.372 e-01
PHR1	ORFLen	-1.127e-01	3.94e-01	9.999e-01	3.381e-02	5.143e-03	1.371e-01
SIZ1	ORFLen	-3.078e-02	2.68e-01	6.611 e-05	1.000e+00	3.235 e-02	1.902e-01
Cyclophilin	mRNAlen	-2.543e-01	7.19e-01	6.611 e-05	1.000e+00	4.716e-04	6.827 e-01
EF1a	mRNAlen	-6.216e-02	2.86e-01	9.999e-01	4.975e-02	3.908e-04	7.808e-01
Hsp90	mRNAlen	-1.343e-01	4.81e-01	6.611 e-05	1.000e+00	7.682e-04	6.206 e-01
IPS1	mRNAlen	-1.394e-01	4.66e-01	9.999e-01	8.567e-02	2.854e-04	8.789 e-01
PHO2	mRNAlen	-1.511e-01	4.69e-01	9.999e-01	1.255e-01	5.059e-04	6.637 e-01
PHR1	mRNAlen	-3.971e-02	2.65e-01	9.999e-01	3.684 e-02	3.713e-04	7.518e-01
SIZ1	mRNAlen	-2.190e-01	7.09e-01	9.999e-01	3.750e-02	1.401e-03	3.874 e- 01
Cyclophilin	GC	2.418e-01	4.90e-02	9.999e-01	3.349e-02	7.404e-04	5.195 e-01
EF1a	GC	-1.541e-01	5.14e-01	9.999e-01	3.383e-02	2.906e-04	8.458 e-01
Hsp90	GC	2.598e-02	2.07e-01	9.999e-01	3.252 e-02	2.515e-04	8.999e-01
IPS1	GC	-1.887e-04	1.91e-01	9.999e-01	5.497e-02	6.367 e-04	6.727 e-01
PHO2	GC	-1.491e-01	5.02e-01	9.999e-01	1.067e-01	1.224 e-02	2.603 e-02
PHR1	GC	-1.927e-01	5.88e-01	9.999e-01	2.984e-02	2.505 e-04	9.199 e-01
SIZ1	GC	-1.445e-01	4.61e-01	9.999e-01	2.955e-02	2.179e-04	9.369 e-01