

Table S1 The risk score in the model of screened RNAs and their regression coefficients

Gene	Gene ID	HR	L95CI	H95CI	P value	Risk coefficients	Adjusted coefficients for GEO data
<i>ENSG00000167676</i>	<i>PLIN4</i>	0.970813	0.95365	0.988284	0.001134	-0.046041241	-1.078
<i>ENSG00000135333</i>	<i>EPHA7</i>	1.148573	1.114116	1.184095	4.94E-19	0.057248905	0.025
<i>ENSG00000106105</i>	<i>GARS</i>	1.010443	1.005553	1.015357	2.70E-05	0.003292712	0
<i>ENSG00000154277</i>	<i>UCHL1</i>	0.996983	0.995261	0.998708	0.000614	0.000725265	0.168
<i>ENSG00000142178</i>	<i>SIK1</i>	0.937606	0.889847	0.987928	0.015722	-0.062224827	0.318
<i>ENSG00000154920</i>	<i>EME1</i>	1.093537	1.045657	1.143609	9.06E-05	0.111282013	0.077
<i>ENSG00000165795</i>	<i>NDRG2</i>	1.00478	1.001924	1.007644	0.001026	0.016181747	0.487
<i>ENSG00000110651</i>	<i>CD81</i>	1.003416	1.001144	1.005693	0.003193	0.007979442	1.928
<i>ENSG00000156345</i>	<i>CDK20</i>	0.67721	0.604511	0.758653	1.73E-11	-0.261351744	1.163
<i>ENSG00000170899</i>	<i>GSTA4</i>	1.011585	1.005976	1.017225	4.91E-05	0.021256979	0.047
<i>ENSG00000123415</i>	<i>SMUG1</i>	1.039328	1.011122	1.068322	0.005999	0.001926979	-1.146
<i>ENSG00000102010</i>	<i>BMX</i>	1.027172	1.015871	1.038599	2.04E-06	0.001172657	-8.789
<i>ENSG00000172159</i>	<i>FRMD3</i>	0.854756	0.730807	0.999727	0.049602	-0.201949924	0.196
<i>ENSG00000131016</i>	<i>AKAP12</i>	1.022967	1.013535	1.032487	1.55E-06	0.014111537	0.335
<i>ENSG00000162881</i>	<i>OXER1</i>	0.816902	0.742772	0.89843	3.09E-05	-0.336559988	-383.383
<i>ENSG00000141738</i>	<i>GRB7</i>	1.002453	1.001527	1.00338	2.02E-07	0.003410404	-0.108
<i>ENSG00000237187</i>	<i>NR2F1-AS1</i>	1.401759	1.268647	1.548837	3.26E-11	0.229409557	0.017
<i>ENSG00000065833</i>	<i>ME1</i>	0.976762	0.968515	0.985079	5.47E-08	-0.012850624	-0.328
<i>ENSG00000156298</i>	<i>TSPAN7</i>	1.012817	1.009808	1.015836	4.92E-17	0.003315713	-0.367
<i>ENSG00000235501</i>	<i>RP4-639F20.1</i>	1.047363	1.029268	1.065775	1.95E-07	0.041037196	0
<i>ENSG00000185803</i>	<i>SLC52A2</i>	1.005158	1.000816	1.009519	0.019841	0.005146729	-0.885
<i>ENSG00000179750</i>	<i>APOBEC3B</i>	1.009378	1.004865	1.01391	4.45E-05	0.009665938	-0.055
<i>ENSG00000065413</i>	<i>ANKRD44</i>	0.659954	0.549185	0.793065	9.28E-06	-0.315784717	-1.585
<i>ENSG00000185070</i>	<i>FLRT2</i>	1.133439	1.05655	1.215925	0.000475	0.080010429	0
<i>ENSG00000273338</i>	<i>RP11-386I14.4</i>	0.881374	0.803079	0.967301	0.007805	-0.08995099	0
<i>ENSG00000112312</i>	<i>GMNN</i>	0.988447	0.978017	0.998988	0.031795	-0.019084293	-0.049
<i>ENSG00000211445</i>	<i>GPX3</i>	1.000643	1.000228	1.001059	0.002371	0.000273901	0.388
<i>ENSG00000108551</i>	<i>RASD1</i>	1.003579	1.001167	1.005997	0.003614	0.002603366	0.133
<i>ENSG00000102804</i>	<i>TSC22D1</i>	1.007649	1.003577	1.011737	0.000225	0.00741084	0.308
<i>ENSG00000125877</i>	<i>ITPA</i>	1.010399	1.004943	1.015884	0.00018	0.00631014	0.067

HR, hazard ratio; L95CI, lower value of 95% confidence interval; H95CI, upper value of 95% confidence interval; GEO, Gene Expression Omnibus.