

Table S1 Univariate Cox regression for each autophagy-related lncRNAs

Gene	KM	B	SE	HR	HR.95L	HR.95H	P value
AL391244.3	5.45E-07	0.524429	0.080766	1.689494	1.442144	1.979268	8.40E-11
LINC01843	6.20E-07	-0.2271	0.045867	0.796843	0.728335	0.871795	7.37E-07
AC011462.4	2.04E-07	0.180471	0.033516	1.197782	1.121629	1.279105	7.26E-08
AC103706.1	5.31E-07	0.427881	0.051977	1.534003	1.385425	1.698515	1.84E-16
SNHG15	4.21E-08	0.074191	0.014596	1.077012	1.046637	1.108269	3.72E-07
AC121338.2	1.21E-07	-1.08526	0.190731	0.337814	0.232449	0.49094	1.27E-08
AL590094.1	3.38E-08	0.414275	0.068122	1.513273	1.324133	1.729431	1.19E-09
AP003352.1	6.64E-09	0.284143	0.052247	1.328622	1.199302	1.471887	5.38E-08
EPB41L4A-DT	1.64E-09	-0.53498	0.083571	0.585679	0.497191	0.689915	1.54E-10
AC026356.2	7.56E-07	0.346607	0.065167	1.41426	1.244685	1.606939	1.04E-07
SNHG17	4.15E-12	0.142409	0.022011	1.153049	1.104364	1.20388	9.80E-11
LINC00460	4.38E-07	0.113946	0.01529	1.120692	1.087606	1.154785	9.16E-14
HOTAIRM1	4.67E-08	0.1151	0.016709	1.121986	1.085837	1.159339	5.64E-12
AC084876.1	5.01E-07	0.490199	0.073549	1.632642	1.413467	1.885802	2.65E-11
AC027796.4	7.68E-07	0.268466	0.046421	1.307957	1.194208	1.432541	7.32E-09
MELTF-AS1	6.41E-09	0.155745	0.019183	1.168529	1.12541	1.213299	4.70E-16
AC010973.2	5.44E-08	0.389931	0.064532	1.47688	1.301415	1.676002	1.52E-09