

**Supplementary Table 1. Pleiotropy and heterogeneity analyses of Two-Sample MR between cathepsins family and different histological typ**

exposure	outcome	MR-Egger intercept		MR-PRESSO global	MR-IWW			MR-Egger		
		Egger intercept	p_value	p_value	Q	Q_df	Q_pval	Q	Q_df	Q_pval
cathepsin B	Overall lung cancer	-0.002	0.619	0.841	10.268	15	0.803	10.244	14	0.744
	Adenocarcinoma	-0.019	0.278	0.595	12.638	14	0.555	13.915	15	0.532
	Squamous carcinoma	0.004	0.835	0.645	11.318	13	0.584	11.273	12	0.506
	SCLC	0.059	0.261	0.084	20.612	11	0.038	18.054	10	0.054
cathepsin E	Overall lung cancer	-0.004	0.751	0.820	5.254	9	0.812	5.146	8	0.742
	Adenocarcinoma	-0.016	0.376	0.586	7.264	9	0.610	6.386	8	0.604
	Squamous carcinoma	-0.012	0.658	0.473	7.815	8	0.452	7.583	7	0.371
	SCLC	-0.037	0.299	0.299	12.465	9	0.188	10.797	8	0.213
cathepsin F	Overall lung cancer	-0.014	0.320	0.515	9.209	10	0.512	8.100	9	0.524
	Adenocarcinoma	-0.026	0.284	0.173	15.529	10	0.114	13.573	9	0.138
	Squamous carcinoma	-0.005	0.816	0.701	7.676	10	0.660	7.619	9	0.573
	SCLC	-0.041	0.264	0.667	7.502	10	0.677	6.086	9	0.731
cathepsin G	Overall lung cancer	-0.001	0.967	0.309	12.134	10	0.276	12.131	9	0.206
	Adenocarcinoma	-0.019	0.366	0.750	6.742	10	0.750	5.837	9	0.756
	Squamous carcinoma	0.003	0.908	0.399	9.894	9	0.359	9.876	8	0.274
	SCLC	0.021	0.674	0.074	17.291	10	0.068	16.936	9	0.050
cathepsin H	Overall lung cancer	-0.004	0.702	0.270	15.432	9	0.080	15.134	8	0.057
	Adenocarcinoma	-0.007	0.603	0.433	10.420	9	0.318	10.052	8	0.261
	Squamous carcinoma	0.010	0.538	0.347	13.054	10	0.221	12.485	9	0.187
	SCLC	0.021	0.421	0.430	7.676	8	0.466	6.945	7	0.435
cathepsin L2	Overall lung cancer	-0.022	0.169	0.156	15.009	10	0.132	12.026	9	0.212
	Adenocarcinoma	-0.010	0.630	0.452	9.263	9	0.413	8.982	8	0.344
	Squamous carcinoma	0.003	0.931	0.072	17.830	10	0.058	17.814	9	0.037
	SCLC	-0.015	0.663	0.884	3.349	8	0.911	3.143	7	0.871
cathepsin O	Overall lung cancer	0.016	0.227	0.799	6.806	10	0.744	5.127	9	0.823
	Adenocarcinoma	0.023	0.215	0.871	5.882	10	0.825	4.105	9	0.904
	Squamous carcinoma	-0.001	0.975	0.855	6.322	10	0.788	6.321	9	0.707
	SCLC	0.013	0.804	0.212	8.970	7	0.255	8.871	6	0.181
cathepsin S	Overall lung cancer	0.019	0.022	0.074	30.317	19	0.048	22.495	18	0.211
	Adenocarcinoma	0.018	0.066	0.211	24.987	20	0.202	20.805	19	0.348
	Squamous carcinoma	0.023	0.049	0.134	29.214	22	0.139	24.204	21	0.283
	SCLC	0.025	0.141	0.777	14.462	19	0.756	12.087	18	0.843
cathepsin Z	Overall lung cancer	0.025	0.141	0.757	14.462	19	0.756	12.087	18	0.843
	Adenocarcinoma	0.025	0.141	0.763	14.462	19	0.756	12.087	18	0.843
	Squamous carcinoma	0.025	0.141	0.766	14.462	19	0.756	12.087	18	0.843
	SCLC	0.025	0.141	0.760	14.462	19	0.756	12.087	18	0.843

Supplementary Table 2. The results of reverse MR analysis of different histological of lung cancer and cathepsins family

exposure	outcome	nsnp	Inverse variance weighted			MR-Egger			Weighted median			MR-Egger intercept		MR-PRESSO global		MR-IVW			MR-Egger		
			p_value	OR	95% OR	p_value	OR	95% OR	p_value	OR	95% OR	Egger intercept	p_value	p_value	Q	Q_df	Q_pval	Q	Q_df	Q_pval	
Overall lung cancer	Cathepsin B	15	0.352	1.074	0.924-1.278	0.312	1.255	0.822-1.916	0.373	1.101	0.891-1.361	-0.018	0.454	0.644	11.340	14	0.659	10.743	13	0.632	
	Cathepsin E	15	0.502	1.058	0.897-1.247	0.600	1.141	0.705-1.846	0.554	1.070	0.860-1.337	-0.009	0.747	0.262	16.959	14	0.258	16.819	13	0.208	
	Cathepsin F	15	0.504	0.948	0.811-1.108	0.554	0.868	0.551-1.369	0.367	0.907	0.731-1.122	0.010	0.692	0.360	15.243	14	0.362	15.053	13	0.304	
	Cathepsin G	15	0.747	1.025	0.880-1.194	0.427	1.201	0.775-1.863	0.710	1.039	0.848-1.272	-0.018	0.463	0.443	14.572	14	0.408	13.957	13	0.377	
	Cathepsin H	15	0.447	1.107	0.852-1.439	0.789	1.114	0.515-2.408	0.953	1.006	0.813-1.245	-0.001	0.988	0.063	43.145	14	0.000	43.144	13	0.000	
	Cathepsin L2	15	0.557	0.955	0.819-1.113	0.233	0.760	0.494-1.169	0.873	0.983	0.802-1.210	0.026	0.286	0.415	14.744	14	0.396	13.462	13	0.413	
	Cathepsin O	15	0.226	1.110	0.937-1.316	0.326	1.291	0.790-2.108	0.460	1.082	0.877-1.336	-0.017	0.532	0.236	18.003	14	0.207	17.449	13	0.180	
	Cathepsin S	15	0.325	1.078	0.928-1.252	0.520	0.867	0.568-1.324	0.422	1.082	0.892-1.313	0.025	0.301	0.981	5.112	14	0.984	3.950	13	0.992	
Cathepsin Z	15	0.191	1.117	0.947-1.317	0.365	0.807	0.516-1.263	0.284	1.121	0.909-1.383	0.037	0.152	0.255	17.102	14	0.251	14.521	13	0.338		
Adenocarcinoma	Cathepsin B	14	0.763	0.979	0.854-1.123	0.536	1.169	0.724-1.888	0.322	0.914	0.765-1.092	-0.024	0.464	0.243	16.042	13	0.247	15.312	12	0.225	
	Cathepsin E	14	0.255	1.074	0.950-1.215	0.496	1.164	0.761-1.780	0.150	1.131	0.956-1.338	-0.011	0.705	0.617	10.781	13	0.629	10.631	12	0.561	
	Cathepsin F	14	0.495	1.049	0.914-1.205	0.967	1.011	0.616-1.659	0.967	1.076	0.908-1.275	0.005	0.880	0.250	16.352	13	0.231	16.319	12	0.177	
	Cathepsin G	14	0.595	1.034	0.914-1.170	0.158	1.385	0.906-2.118	0.281	1.089	0.933-1.270	-0.040	0.183	0.712	9.705	13	0.718	7.712	12	0.807	
	Cathepsin H	14	0.598	1.062	0.848-1.331	0.476	0.746	0.342-1.627	0.800	0.977	0.817-1.169	0.048	0.372	0.142	43.402	13	0.000	40.497	12	0.000	
	Cathepsin L2	14	0.184	0.920	0.813-1.040	0.131	0.704	0.461-1.076	0.579	0.954	0.806-1.128	0.037	0.221	0.847	7.529	13	0.873	5.863	12	0.923	
	Cathepsin O	14	0.731	0.979	0.865-1.107	0.117	0.693	0.454-1.060	0.970	1.003	0.843-1.195	0.047	0.122	0.417	12.171	13	0.514	9.406	12	0.668	
	Cathepsin S	14	0.466	1.051	0.920-1.200	0.691	1.103	0.687-1.773	0.487	1.059	0.901-1.245	-0.007	0.836	0.327	12.171	13	0.514	9.406	12	0.668	
Cathepsin Z	14	0.328	1.078	0.928-1.251	0.260	0.745	0.456-1.215	0.546	1.057	0.884-1.263	0.051	0.148	0.125	12.171	13	0.514	9.406	12	0.668		
Squamous carcinoma	Cathepsin B	6	0.033	1.189	1.014-1.395	0.438	1.264	0.741-2.156	0.038	1.224	1.011-1.481	-0.011	0.826	0.804	2.523	5	0.773	2.468	4	0.650	
	Cathepsin E	6	0.763	1.032	0.843-1.263	0.325	0.694	0.366-1.315	0.640	1.055	0.842-1.323	0.074	0.272	0.172	8.065	5	0.153	5.738	4	0.220	
	Cathepsin F	6	0.437	0.917	0.737-1.141	0.279	0.630	0.306-1.300	0.037	0.789	0.631-0.986	0.070	0.347	0.118	9.431	5	0.093	7.352	4	0.118	
	Cathepsin G	6	0.870	1.015	0.846-1.219	0.470	1.296	0.685-2.450	0.967	0.996	0.813-1.220	-0.045	0.476	0.286	6.590	5	0.253	5.708	4	0.222	
	Cathepsin H	6	0.758	0.975	0.832-1.144	0.979	0.992	0.582-1.693	0.601	0.950	0.782-1.153	-0.003	0.950	0.982	0.651	5	0.986	0.647	4	0.958	
	Cathepsin L2	6	0.146	0.882	0.744-1.045	0.136	0.602	0.353-1.027	0.635	0.950	0.767-1.176	0.071	0.216	0.350	5.664	5	0.340	3.511	4	0.476	
	Cathepsin O	6	0.964	1.004	0.842-1.197	0.566	0.820	0.439-1.530	0.979	1.003	0.805-1.250	0.038	0.541	0.307	6.072	5	0.299	5.464	4	0.243	
	Cathepsin S	6	0.678	0.967	0.824-1.134	0.640	1.147	0.673-1.956	0.916	1.010	0.834-1.223	-0.032	0.546	0.585	3.935	5	0.559	3.502	4	0.478	
Cathepsin Z	6	0.798	0.970	0.770-1.222	0.908	0.947	0.399-2.250	0.512	0.926	0.736-1.165	0.004	0.957	0.082	10.524	5	0.062	10.515	4	0.033		
SCLC	Cathepsin B	26	0.641	1.017	0.948-1.091	0.156	1.115	0.964-1.291	0.965	1.002	0.914-1.099	-0.024	0.173	0.293	28.358	25	0.292	26.203	24	0.343	
	Cathepsin E	26	0.841	1.007	0.942-1.075	0.516	1.048	0.911-1.205	0.345	1.049	0.950-1.157	-0.010	0.528	0.581	23.060	25	0.574	22.650	24	0.541	
	Cathepsin F	26	0.509	1.024	0.954-1.100	0.935	1.006	0.863-1.173	0.457	1.037	0.942-1.141	0.004	0.802	0.254	28.986	25	0.264	28.909	24	0.224	
	Cathepsin G	26	0.897	1.004	0.940-1.073	0.778	1.021	0.887-1.174	0.958	0.998	0.909-1.095	-0.004	0.802	0.874	17.361	25	0.868	17.297	24	0.836	
	Cathepsin H	26	0.442	0.973	0.906-1.044	0.761	0.976	0.838-1.138	0.645	0.977	0.886-1.078	-0.001	0.957	0.259	28.798	25	0.272	28.795	24	0.228	
	Cathepsin L2	26	0.148	1.050	0.983-1.122	0.517	1.048	0.911-1.205	0.470	1.033	0.946-1.127	0.000	0.977	0.920	15.533	25	0.928	15.532	24	0.904	
	Cathepsin O	26	0.367	1.031	0.965-1.101	0.890	1.010	0.878-1.162	0.408	1.038	0.950-1.136	0.005	0.749	0.919	16.098	25	0.912	15.993	24	0.888	
	Cathepsin S	26	0.464	0.972	0.901-1.049	0.253	0.908	0.772-1.067	0.608	0.976	0.891-1.070	0.018	0.358	0.116	33.419	25	0.121	32.238	24	0.121	
Cathepsin Z	26	0.798	0.970	0.770-1.223	0.908	0.947	0.399-2.250	0.512	0.926	0.736-1.165	0.004	0.957	0.082	10.524	5	0.062	10.515	4	0.033		

Supplementary Table 3. The results of multivariable MR analysis between cathepsins family and lung cancer risk.

outcome	exposure	SNPs	IVW			MR-egger			MR-IVW			MR-Egger			MR-Egger intercept	
			pvalue	OR	95% OR	pvalue	OR	95% OR	Q	Q_df	Q_pval	Q	Q_df	Q_pval	Egger intercept	p_value
overall lung cancer	cathepsin B	88	0.994	1.000	0.957 - 1.045	0.543	1.016	0.966 - 1.068								
	cathepsin E		0.283	1.037	0.970 - 1.109	0.118	1.057	0.986 - 1.133								
	cathepsin F		0.431	1.023	0.966 - 1.084	0.696	1.012	0.954 - 1.073								
	cathepsin G		0.219	1.039	0.977 - 1.105	0.298	1.033	0.972 - 1.098								
	cathepsin H		0.000	1.070	1.033 - 1.109	0.000	1.074	1.038 - 1.113	89.810	79.000	0.191	85.904	78.000	0.253	-0.003	0.219
	cathepsin O		0.419	0.967	0.892 - 1.049	0.231	0.951	0.877 - 1.032								
	cathepsin S		0.725	1.007	0.968 - 1.048	0.760	1.006	0.967 - 1.047								
	cathepsin L2		0.924	1.003	0.935 - 1.077	0.623	1.018	0.948 - 1.094								
	cathepsin Z		0.333	0.976	0.928 - 1.025	0.478	0.982	0.935 - 1.032								
Adenocarcinoma	cathepsin B	88	0.982	1.001	0.944 - 1.060	0.402	1.029	0.963 - 1.099								
	cathepsin E		0.488	1.032	0.944 - 1.127	0.267	1.053	0.961 - 1.155								
	cathepsin F		0.680	1.016	0.941 - 1.097	0.999	1.000	0.926 - 1.080								
	cathepsin G		0.152	1.061	0.978 - 1.151	0.189	1.055	0.974 - 1.144								
	cathepsin H		0.000	1.094	1.046 - 1.143	0.000	1.098	1.050 - 1.147	82.350	79.000	0.376	78.705	78.000	0.456	-0.005	0.106
	cathepsin O		0.159	0.926	0.832 - 1.031	0.077	0.908	0.815 - 1.011								
	cathepsin S		0.359	1.025	0.973 - 1.079	0.352	1.024	0.974 - 1.078								
	cathepsin L2		0.513	1.033	0.937 - 1.138	0.317	1.051	0.954 - 1.158								
	cathepsin Z		0.796	0.991	0.927 - 1.060	0.980	1.001	0.937 - 1.070								
Squamous carcinoma	cathepsin B	89	0.442	1.028	0.958 - 1.103	0.341	1.041	0.958 - 1.131								
	cathepsin E		0.941	0.996	0.888 - 1.116	0.916	1.007	0.891 - 1.137								
	cathepsin F		0.340	1.046	0.954 - 1.147	0.439	1.038	0.944 - 1.142								
	cathepsin G		0.626	1.026	0.926 - 1.136	0.674	1.022	0.922 - 1.134								
	cathepsin H		0.175	1.037	0.984 - 1.094	0.157	1.039	0.985 - 1.096	90.022	80.000	0.208	89.465	79.000	0.197	-0.002	0.557
	cathepsin O		0.405	0.947	0.833 - 1.076	0.339	0.938	0.823 - 1.070								
	cathepsin S		0.459	1.023	0.963 - 1.087	0.471	1.023	0.962 - 1.087								
	cathepsin L2		0.830	1.012	0.905 - 1.133	0.712	1.022	0.910 - 1.148								
	cathepsin Z		0.561	0.976	0.901 - 1.058	0.636	0.980	0.903 - 1.064								
SCLC	cathepsin B	83	0.180	0.924	0.824 - 1.037	0.069	0.888	0.782 - 1.009								
	cathepsin E		0.143	1.131	0.959 - 1.332	0.030	1.200	1.018 - 1.415								
	cathepsin F		0.580	1.042	0.900 - 1.207	0.562	1.044	0.903 - 1.206								
	cathepsin G		0.422	1.066	0.912 - 1.247	0.529	1.049	0.903 - 1.219								
	cathepsin H		0.154	1.063	0.978 - 1.155	0.134	1.063	0.981 - 1.152	82.265	74.000	0.239	74.103	73.000	0.442	0.007	0.266
	cathepsin O		0.365	1.102	0.893 - 1.360	0.434	1.085	0.884 - 1.333								
	cathepsin S		0.422	1.041	0.943 - 1.149	0.395	1.042	0.948 - 1.145								
	cathepsin L2		0.990	1.001	0.837 - 1.197	0.937	1.007	0.844 - 1.202								
	cathepsin Z		0.109	0.900	0.791 - 1.024	0.097	0.899	0.794 - 1.019								