

Table S5 the detail data of nomogram and calibration

ID	futime	fustat	NIFK-AS1	AC026355.2	SEPSECS-AAL360270.1	AC010999.2	ABCA9-AS1
TCGA-62-A	3.4136986	0	1.6458861	2.0037821	1.5109113	0.2747686	0.6912661
TCGA-L9-A	0.1589041	1	1.7285301	1.1073531	0.8864331	1.7777038	0.1142337
TCGA-78-8	9.2082192	1	2.2767051	2.1504297	0.8904856	0.0635029	0.5479433
TCGA-62-A	1.6273973	1	1.6230709	0.5665716	1.2525973	1.0546401	0
TCGA-38-4	3.1424658	1	1.8482373	0.9387751	0.6263924	0.6300326	0.0257376
TCGA-86-A	1.1369863	0	1.926417	2.3189242	0.3170725	0.8433407	0
TCGA-97-8	1.3671233	0	1.8217102	2.0355535	0.230449	0.8094144	0.4565959
TCGA-50-6	0.5178082	1	1.5531147	0.4722798	0.5809173	1.4765891	0
TCGA-78-7	4.1863014	1	2.7582819	0.8914192	1.0702519	0.3674829	0.2055176
TCGA-97-A	1.6821918	0	1.6142858	0.7426961	1.2848102	1.1379625	0
TCGA-78-7	2.2109589	1	2.2157097	0.9280481	1.8459918	2.5246404	0.3743996
TCGA-50-8	3.0821918	0	1.9480027	1.4671751	1.400374	0.7987556	0.2424501
TCGA-69-8	0.3534247	0	1.8317962	0.8284691	1.3331375	1.8219143	0.2675957
TCGA-86-A	0.3178082	1	1.8528378	0.2757223	1.6521423	0.0736832	0.2719037
TCGA-78-8	19.347945	0	1.8997943	3.4568456	0.651775	2.2863489	0.1823109
TCGA-97-8	0.4493151	1	2.4749551	1.2444	1.4936473	0.795102	0.3605328
TCGA-44-7	2.4356164	0	1.6638455	0.6084308	0.751378	0.5211513	0
TCGA-78-7	0.490411	1	2.2463169	1.7959331	1.1412366	0.9758503	0
TCGA-44-4	2.2136986	1	2.2070338	6.2884084	2.1200699	2.3212354	1.073786
TCGA-71-6	0.7013699	0	1.3074285	3.4390247	0.988921	1.7126398	0.1054099
TCGA-05-5	0.169863	0	1.4943644	1.13888	0.7441611	1.1352065	0
TCGA-05-4	2.0849315	0	2.3566509	0.7472585	0.902035	1.4537019	0.0932897
TCGA-78-7	13.591781	1	1.5716283	1.3148705	1.6388616	1.8580208	0.4850146
TCGA-73-4	1.2794521	0	1.4670186	0.2979544	0.3188085	1.5837598	0.154324
TCGA-38-4	8.1452055	0	2.5611061	0.6304055	0.8901743	0.5254676	0.1633699
TCGA-67-3	1.169863	0	2.298629	3.801376	1.4158668	0.9159415	0
TCGA-97-A	1.709589	0	1.8169683	2.5622302	1.5087836	0.6410836	0.1384213
TCGA-78-7	2.1671233	0	2.3810341	1.2489598	1.5018722	0.4018489	0.1172961
TCGA-67-3	1.6712329	0	2.2606881	2.8660178	0.6911768	1.8722378	0.2077679
TCGA-49-4	1.0547945	1	2.3033716	0.9974731	1.3241769	0.8426973	0
TCGA-55-8	1.6383562	1	1.8163539	1.4661833	0.6991962	1.1653653	0.1949696
TCGA-99-A	1.8027397	0	2.1718793	1.3562002	1.6678467	0.5840966	0
TCGA-49-4	6.3506849	1	1.9117683	2.9064866	1.073409	0.5083272	0.0809305
TCGA-44-2	0.2657534	1	2.2703495	2.9008443	1.4187477	1.5097634	0.9962682
TCGA-62-8	3.3315068	0	2.1639173	1.8326875	1.4057201	1.8278597	0
TCGA-05-4	0.0027397	0	1.6275601	0.7598376	0.9165531	1.4690524	0.2273714
TCGA-55-6	0.3232877	1	1.7196654	0.7055799	0.5672534	1.7233401	0.2715452
TCGA-73-4	6.890411	0	2.0787806	1.0085586	1.5945008	1.0164963	0.2471984
TCGA-44-A	0.9616438	0	1.5695883	1.5507529	0.8443053	0.9710745	0.1528973
TCGA-NJ-A	6.1945205	0	1.7037652	1.381228	1.3090597	0.4500631	0.1784916
TCGA-05-4	2	0	1.833133	0.4839834	0.78668	1.0641931	0
TCGA-NJ-A	1.690411	0	2.4687397	2.5512453	1.4946204	0.5693939	0.3096419

TCGA-44-6	1.8027397	0	1.9414444	1.8648101	0.7482036	0.6209617	0	0.1909309
TCGA-50-8	2.2712329	0	2.6739182	1.027791	1.8027722	0.4577519	0	0.032383
TCGA-78-7	0.7068493	1	1.2649567	0.4684269	0.9544197	0.7263953	0.1478277	0
TCGA-05-4	2.1671233	0	1.9730234	1.3443172	1.1956626	1.3879139	0	0
TCGA-49-4	0.7342466	1	1.7248685	0.9985566	0.7623063	1.2379308	0.1578198	0.1272371
TCGA-78-7	1.7150685	1	1.772308	0.5916794	0.4422269	1.3116775	0.1696685	0.3769571
TCGA-55-7	1.8356164	0	1.8586176	0.5526722	0.4806787	1.0206978	0.2594232	0.1774714
TCGA-55-6	3.3945205	0	1.6006979	0.3797318	1.1434587	0.6909981	0.3070787	0.0336521
TCGA-78-7	3.2794521	1	1.572065	0.3660284	1.0361868	1.8428582	0.0990932	0
TCGA-99-8	3.0630137	0	1.6382129	0.8234233	0.6979517	0.6388616	0.186247	0
TCGA-05-4	1.5835616	0	1.3711127	0.3677065	0.7355222	0.8436623	0	0
TCGA-78-7	7.3452055	1	2.9411814	3.1706622	1.7710827	1.285343	0.1127668	1.1044709
TCGA-78-7	1.6054795	1	2.3720606	0.3995539	2.0441843	1.3717819	0.1437852	0.2735756
TCGA-97-A	1.7863014	0	2.1608557	1.429857	1.3278598	0.8304987	0	0.0364682
TCGA-64-1	6.8164384	0	1.629986	1.192131	0.8076022	1.7601358	0.116897	0.047678
TCGA-64-1	3.0849315	0	2.2335813	2.8055814	0.9096579	1.0976782	0	0
TCGA-86-8	2.569863	0	1.8432201	1.3234277	1.2047042	0.5835191	0	0.1777265
TCGA-78-7	2.2630137	1	1.2555007	1.7439458	0.9963887	1.5870769	0	0
TCGA-99-8	2.9041096	0	2.3296977	0.1096949	0.574925	1.7646434	0.2086421	0
TCGA-L9-A	1.8191781	0	1.5395807	1.1926992	0.398022	0.9309265	0	0
TCGA-55-8	1.4821918	0	1.6663476	1.0278618	1.4797997	2.1805935	0.4234168	0.3725064
TCGA-86-A	2.4547945	0	1.5999366	1.8958842	1.4172179	1.2582174	0	0.0832474
TCGA-44-3	2.8109589	1	1.9068136	1.8295655	0.9040409	0.875151	0	0.0662614
TCGA-69-7	0.630137	0	2.2128494	0.5965536	0.9237957	1.1021878	0.1164978	0.2026371
TCGA-35-4	0.6164384	0	1.3888508	0.7017708	0.4311415	1.0234683	0.3002995	0
TCGA-35-4	0.4986301	0	1.9149854	1.0100637	0.4955928	0.7494914	0.3780676	0
TCGA-55-6	2.7260274	1	1.3021728	2.3938555	0.7968469	1.5688104	0	0
TCGA-86-8	2.6	0	2.6038578	1.6152279	1.96857	1.5786972	0.3392514	0
TCGA-05-4	0.830137	1	1.8861209	1.1050746	1.2975436	1.0222613	0	0.9958824
TCGA-05-4	1	0	1.710305	1.5953604	1.5354558	0.2971327	0.5682267	0.8038057
TCGA-05-4	1.2465753	0	1.3309599	0.3640121	0.6059684	1.2367681	0	0
TCGA-MP-	6.1589041	0	2.3208601	4.3677345	2.7864709	1.4128357	0.1351408	0.0554733
TCGA-97-8	1.509589	0	1.5070081	2.325041	0.6121166	0.151729	0	0.0250288
TCGA-97-A	1.7232877	0	1.7589001	1.0513025	0.7054029	0.4731115	0.1322478	0.0272956
TCGA-05-4	2.4986301	0	1.8126214	0.6050202	0.6574571	1.3938555	0.3618806	0.0483757
TCGA-38-4	3.7178082	1	1.6224619	1.6856711	0.6531518	0.974823	0.628447	0
TCGA-78-7	19.857534	0	2.5607883	1.5471043	1.4518573	0.3640121	0.2501127	0
TCGA-MP-	2.6027397	1	2.7832471	2.0251351	1.4768483	2.1369788	0.5029416	0.3450557
TCGA-50-5	1.1890411	1	3.0737346	0.2691529	1.2688535	1.5046204	0.4896457	1.061638
TCGA-MP-	4.1123288	1	1.7973866	2.8881685	0.6768992	0.7228157	0	0.0721744
TCGA-78-7	1.6246575	1	1.7887275	1.1081564	1.0770379	0.655535	0	0.5676428
TCGA-73-4	1.9479452	1	1.6257381	1.423363	0.5981746	0.4166236	0.287945	0.015783
TCGA-69-7	1.1260274	0	2.6375407	1.3329084	2.2825288	0.8304176	0.6925165	0.6428402

TCGA-73-7	3.2575342	0	2.5056369	3.0379796	1.0243906	0.8342258	0.3810618	0.0216219
TCGA-78-7	0.8794521	1	1.5897156	0.5045695	1.0193461	1.6248028	0	0
TCGA-05-4	0.3315068	1	1.9805006	0.8274939	0.6782522	0.9194545	0	0.0538064
TCGA-38-4	2.939726	1	2.910771	0.0973411	1.3837181	0.4670708	0	0.0581086
TCGA-05-4	0.7506849	1	1.21456	0.4239546	0.6628435	1.0345679	0.1534162	0.0319597
TCGA-55-8	2.4767123	0	2.1519562	1.70982	1.3631711	0.9670948	0.3879139	0.3350258
TCGA-55-7	2.4356164	0	1.6830227	4.9093363	0.8902521	0.8127036	0.5017194	0.2618317
TCGA-55-6	0.6493151	1	1.5132886	2.5776101	0.5668639	1.5134402	0.2053925	0
TCGA-62-A	2.4931507	0	2.0760802	3.3309742	1.1025238	1.3238888	1.0579008	0.1476974
TCGA-62-8	1.2164384	1	1.7126398	1.9564656	0.8410875	1.7650679	0.1493894	0.0614306
TCGA-MP-	1.1753425	1	1.7083638	2.239123	0.6586457	0.7759355	0	0.078883
TCGA-44-8	0.7123288	0	2.9218362	1.1307332	1.1749177	1.0497702	0.2391536	0
TCGA-73-4	2.1917808	0	1.8062828	1.4118601	0.7774513	0.9150237	0.1082233	0.0221902
TCGA-78-7	0.6684932	1	2.1740868	1.4138107	1.1765143	1.4826934	0.108491	0
TCGA-55-7	2.3890411	0	1.851759	2.0312187	0.9375704	1.411155	0	0
TCGA-62-8	0.3808219	1	1.7466567	0.6686638	0.9776096	1.2503553	0.3013535	0
TCGA-86-8	1.1589041	0	2.3610946	0.2744108	1.0710762	1.4979955	0.2705887	0
TCGA-50-5	1.260274	1	1.4259963	0.7796379	0.8061178	1.842295	0	0
TCGA-38-7	2.1917808	1	1.5662307	1.0613615	0.5462648	1.0308304	0	0
TCGA-05-4	0.0027397	1	1.9119216	2.213565	0.5247658	1.5071096	0.2492633	1.5630605
TCGA-49-4	4.6575342	0	1.5301706	0.7564256	1.0061185	0.5988416	0	0.1074201
TCGA-62-A	4.5287671	1	1.6932305	1.2653769	0.9901562	0.5101518	0.2600257	0.0281448
TCGA-44-3	3.2410959	0	2.3708896	5.0359406	1.7080989	2.587509	1.7122876	0.597698
TCGA-05-4	0.2493151	1	1.7066853	0.3514022	0.7063758	2.4935961	0	0
TCGA-44-6	1.630137	0	1.7376435	1.7195339	0.9919707	0.5797594	0	0
TCGA-78-7	1.8246575	1	1.8387905	0.504061	1.3343394	2.35834	0.1856129	0.2529
TCGA-69-7	1.890411	0	1.8263146	1.6436253	0.6030269	1.0174231	0	0
TCGA-95-7	1.3068493	0	1.6869711	0.3842709	1.2785201	1.403595	0.2635152	0
TCGA-86-7	2.8657534	1	1.6314768	0.9668734	0.8694766	2.0313599	0.0938306	0
TCGA-49-4	3.8931507	1	2.4113449	1.761966	0.6310577	0.6293799	0	0.0182069
TCGA-91-6	0.2164384	0	2.5512453	2.273307	0.175301	1.3050366	0	0.044604
TCGA-50-6	3.5287671	1	1.8273313	0.8554316	0.5378427	1.1428708	0	0
TCGA-50-5	1.309589	1	1.7035437	2.7525131	0.8880711	0.9997114	0	0
TCGA-91-6	1.0191781	0	2.1673258	0.2873541	0.8494389	0.7763567	0	0
TCGA-49-4	2.4547945	1	2.4302317	1.0308304	0.8823692	0.4013028	0.1646576	0.0343566
TCGA-35-3	0.0383562	0	1.9443337	4.7970234	0.8890062	1.2120712	0.4205094	0
TCGA-97-A	1.5561644	0	2.082975	2.3025238	1.7816115	0.4072984	0.1751733	0.1072861
TCGA-53-7	1.1616438	0	1.8211387	1.0619144	0.8777442	0.5112656	0.5450788	0
TCGA-78-8	6.4657534	0	1.9972563	3.0222258	1.0434145	0.683921	0.4534386	0
TCGA-86-8	0.339726	1	2.0259147	1.7756827	0.5383395	1.1716552	0	0.0771062
TCGA-49-6	1.430137	0	1.4276062	0.5605193	0.5550308	1.7154541	0	0
TCGA-MP-	2.030137	0	2.146753	2.7413375	1.0941009	0.4938522	0.516822	0
TCGA-99-7	2.0465753	0	2.1324452	0.581303	2.153935	1.0611541	0.3709454	0.0278618

TCGA-55-1	3.2273973	0	2.3415019	0.4325319	1.6915788	1.3757345	0.4269624	0.2143735
TCGA-95-7	1.0328767	0	1.762774	0.7567671	0.2584586	1.4815573	0.2028878	0.0427844
TCGA-05-4	3.9205479	0	1.4616333	1.6666657	0.514905	1.2140626	0.1433934	0.0149265
TCGA-78-7	0.7972603	1	2.0398753	1.4268551	0.7409707	0.6864333	0.1152996	0.0926134
TCGA-50-5	6.5561644	1	1.4302317	1.2467121	1.3502709	0.9871753	0	0
TCGA-78-7	1.909589	1	1.7881428	1.5262692	1.3467584	1.5639879	0	0.047678
TCGA-62-8	4.1041096	1	2.1884326	0.7773671	1.4190534	0.8042188	0.1897929	0.1164978
TCGA-78-8	3.3123288	1	1.336569	0.4881036	0	0.1377658	0.3383388	0
TCGA-44-3	2.8383562	0	2.3632739	5.1001984	1.6948208	2.3353975	1.4157226	0.8759897
TCGA-55-A	1.4958904	0	2.0604279	1.6917575	1.1669082	2.1144337	0.2740529	0
TCGA-78-7	2.6	1	2.3312753	1.8125393	1.6075315	1.417974	0.0806577	0
TCGA-55-A	0.0767123	0	1.4540179	1.1560082	0.5089357	1.4678533	0.1819294	0.0381552
TCGA-50-5	3.3835616	1	2.0451284	0.3230818	1.2030132	1.1693478	0.1681285	0.1028597
TCGA-05-5	0.4136986	0	1.857583	3.0980152	1.5496199	0.7616257	0	0.0458623
TCGA-62-A	1.1342466	1	2.1198541	0.5751187	0.7311832	1.3698294	0.1642714	0.0342157
TCGA-97-A	1.6465753	0	2.6322449	0.951364	1.4489538	1.0324535	0.1570437	0.2428159
TCGA-95-7	1.5561644	0	1.8538759	2.2292187	0.6004601	1.0906502	0.097476	0.0199154
TCGA-55-7	2.8493151	0	2.0620526	1.0500489	0.6125884	1.6870607	0.4345615	0.2752455
TCGA-95-8	0.230137	0	1.8360049	1.6260185	0.8212204	0.6474527	0.458067	0
TCGA-55-1	1.9205479	1	2.6495005	0.9327415	1.1957885	1.9589911	0.3509498	0.2701103
TCGA-44-2	3.5643836	0	2.7483539	5.2143152	2.8668486	1.6344306	1.7849855	0.9145263
TCGA-78-7	3.3287671	1	1.5189883	2.1528	0.3395935	1.2313095	0	0
TCGA-64-5	0.169863	1	1.357552	0.198997	0.5443865	1.1228052	0.3688242	0
TCGA-78-7	9.9589041	0	1.4807304	0.5103543	0.9587313	0.4812989	0.2738142	0
TCGA-MP-	0.4410959	1	1.1471113	3.0422241	0.705049	1.2708877	0	0.031254
TCGA-44-2	3.1753425	0	1.7868891	0.6673926	0.8511194	0.6837414	0.220949	0.0236102
TCGA-73-4	4.3835616	1	1.4639363	0.2000021	0.2635152	1.6703418	0.1971105	0
TCGA-44-7	2.4164384	0	3.3186928	0.3729521	0.9368923	1.1997509	0.133432	0.0275787
TCGA-55-8	1.6410959	0	1.763794	0.3828332	0.8302553	0.7270927	0.2624332	0.0563061
TCGA-55-8	1.6630137	1	2.0860364	1.4549655	1.278639	0.8745216	0.1787465	0.0374525
TCGA-55-8	1.4986301	0	2.0705954	0.2488991	0.7457106	0.3922075	0.3188085	0.0176369
TCGA-44-A	0.7863014	0	1.8415706	2.0559938	0.915865	1.3255591	0.175812	0
TCGA-64-5	3.5753425	0	2.7036766	1.1768334	1.6325474	0.6770797	1.0901082	0
TCGA-86-7	1.6575342	0	1.9698596	2.2993326	1.9301696	0.7359554	0.2333973	0.1890338
TCGA-62-8	7.3863014	0	1.83289	5.1265393	1.7534334	2.9390761	0.1548424	0.0322419
TCGA-95-7	0.2383562	1	1.6429788	0.4371212	0.6636634	1.1154994	0.231801	0.0492125
TCGA-69-7	0.5534247	0	1.3072536	2.622743	0.6971511	0.3170725	0.1480881	0.0308304
TCGA-55-7	2.7260274	1	1.8181961	1.2189051	1.0341453	1.1551663	0	0
TCGA-MP-	0.4575342	1	1.859254	2.0840302	0.9795132	1.1005068	0.5990321	0.1674864
TCGA-95-A	1.5150685	0	1.3953919	2.2516888	0.782073	0.4138107	0.1628545	0
TCGA-86-7	3.169863	0	2.499476	0.432425	0.4978423	1.0310422	0	0.0957219
TCGA-50-5	1.709589	1	1.212009	0.8919635	0.3524196	0.1733833	0	0
TCGA-64-5	7.3315068	0	2.5934016	0.8951475	0.921284	1.0061903	0.4466797	0.3116194

TCGA-4B-A	0.8219178	1	1.5191896	0.9046575	0.9159415	0.4592218	0.3036931	0.8678174
TCGA-83-5	2.2575342	0	1.6560844	1.3683214	0.7417474	1.8250529	0.1632411	0.0673634
TCGA-44-A	1.1369863	0	1.4833128	2.7146192	1.3902274	1.5402257	0	0.5581697
TCGA-L4-A	1.5835616	0	1.5265196	1.6076262	1.1286233	1.877705	0	0
TCGA-97-8	1.5561644	0	1.963844	1.1071522	0.8646517	0.5005981	0.088413	0
TCGA-78-7	0.4739726	1	1.3972553	2.6843475	0.9772433	0.9355352	0.3662523	1.6730136
TCGA-50-5	0.7726027	1	1.8611614	0.5401265	0.9622899	1.165301	0.291132	0.1231364
TCGA-64-5	4.2712329	0	1.3827779	2.359999	0.605115	1.2185952	0	0
TCGA-NJ-A	0.1369863	0	1.7774513	0.70982	0.5062973	2.5112909	0	0.032383
TCGA-44-8	0.7808219	0	2.1713349	0.4645637	0.6205864	0.7731524	0.3915478	0.0354832
TCGA-44-2	3.6273973	0	2.1659119	4.3316049	0.8448677	0.580435	0.088413	0
TCGA-86-8	2.3616438	0	2.2454044	3.1489666	1.1796384	1.1615008	0	0.3765127
TCGA-MP-	0.9205479	1	2.3037515	2.3048906	1.9012241	2.3558902	0	1.3708338
TCGA-05-4	0.0027397	0	1.9289198	1.6444101	1.2591218	0.8621141	0.2514464	0
TCGA-05-4	1.6630137	0	2.0128547	3.7405175	1.0194884	1.23444	0.5058909	0
TCGA-73-4	0.3589041	0	1.6999957	1.0243906	0.3171883	1.4544918	0.3779566	0.0170667
TCGA-44-3	3.1863014	0	1.6972845	3.0439569	0.85591	0.4811956	0.5212518	0.1200865
TCGA-73-A	0.9315068	1	2.0909889	0.9492735	1.0025945	2.1239641	0.5443865	0
TCGA-97-A	1.7369863	0	1.8467942	0.4005379	1.1140338	0.9279723	0.1441769	0.1973622
TCGA-62-A	5.6630137	0	2.051024	2.5852029	1.9626601	1.0627434	0.3176514	0
TCGA-35-5	0.7232877	0	1.866908	1.3654126	0.7573647	0.1350094	0	0
TCGA-MP-	7.169863	1	1.4068087	1.0800436	0.6156988	0.812375	0.2624332	0
TCGA-05-4	2.0849315	0	1.8956904	1.503654	0.7438165	1.3777901	0	0
TCGA-05-4	2.5013699	0	2.2174169	2.8187892	0.9396027	1.038998	0	0.1998765
TCGA-86-8	2.0273973	0	1.4115889	2.750778	1.2854614	3.4628507	0	0.0729976
TCGA-55-6	2.0821918	1	1.3761792	0.3959402	0.4898512	1.0433445	0	0
TCGA-69-7	1.1342466	0	2.334082	0.8539158	1.6372856	1.0033144	0.2128806	0.0886844
TCGA-55-8	1.0273973	1	3.0109588	0.1988713	1.5714341	0.8257042	0.7890197	0
TCGA-53-7	2.8575342	1	2.1445359	0.1914363	1.6068686	2.436535	0.1886541	0.2570106
TCGA-62-A	4.7260274	1	1.8856134	0.1601457	1.2290957	1.7755563	0.4294285	0
TCGA-78-7	2.6739726	1	2.1075874	4.1559597	1.7538184	1.3726736	0.5462648	1.0920044
TCGA-NJ-A	3.9232877	0	1.4764335	2.5258434	1.2750667	2.4916479	0.2906603	0
TCGA-50-5	0.6849315	1	2.0199509	1.4294821	1.0649518	1.4729556	0.162339	1.0975434
TCGA-55-6	5.8547945	0	1.2973088	2.9288061	0.5768607	1.0032424	0.1319845	0
TCGA-44-2	3.5068493	0	2.2156682	5.6207438	2.1087475	1.7441898	1.2164859	1.587941
TCGA-44-2	3.9150685	0	2.0842117	4.7293352	1.469313	2.3526173	1.2053091	0.9030126
TCGA-64-5	2.3726027	0	1.4120227	1.6689814	0.458277	0.9900109	0	0
TCGA-95-7	1.3041096	0	2.8283472	3.1076544	1.7055799	0.6995516	0.8914192	0
TCGA-86-6	0.490411	0	1.5068558	0	1.1868808	0.9344033	0.0802483	0.0163537
TCGA-78-7	4.4438356	1	1.9583601	1.2559841	1.0924104	1.2520523	0.1003049	0.0204844
TCGA-05-4	1.2520548	1	1.8428984	1.063641	0.9851361	0.9177754	0.1625968	0
TCGA-05-5	2.4164384	0	1.7055356	0.9689386	0.5648171	0.6586457	0	0.0382957
TCGA-50-5	4.0383562	0	1.5433969	0.5576797	1.0719686	0.7359554	0.2077679	0

TCGA-05-5	0.7534247	1	1.4339743	0.7923556	0.6927843	0.7895206	0.2390314	0
TCGA-86-A	2.0191781	1	1.5334637	2.7914807	0.6244285	0.5070081	0.2730981	0
TCGA-MP-	0.2027397	1	1.5061449	2.8489183	0.7775354	0.9979066	0.3272275	0.0714881
TCGA-64-1	3.1972603	1	2.1686099	0.9787082	1.6993295	1.2125693	0	0
TCGA-05-4	4.1726027	0	2.0243197	1.2329063	0.6994628	0.8607642	0	0
TCGA-50-5	8.4767123	0	1.4864055	1.8676593	0.6057788	0.7524061	0	0
TCGA-86-8	0.0520548	1	1.5710943	0.6042611	1.1009776	1.7167715	0	0.3083609
TCGA-50-5	3.9506849	0	1.5977219	1.7270709	1.1881477	0.9951227	0.0462815	0.0370307
TCGA-86-8	2.2082192	0	1.8211796	4.7771619	1.3028747	1.3726178	0.1928254	0.1557492
TCGA-50-5	5.060274	0	2.593306	0.3846025	1.1471113	1.2440346	0	0
TCGA-69-A	1.6191781	0	1.9712216	2.3466449	0.8356816	0.898634	0	0.1378969
TCGA-50-6	2.1287671	1	2.0330529	0.4460444	1.4888235	0.8085086	0.3078948	0.3612069
TCGA-49-4	2.9616438	1	1.2514464	1.0738202	0.0858325	0.5706572	0	0
TCGA-44-2	2.0849315	1	1.8466203	3.0655838	1.2696516	1.3267866	0.3329275	0.4730769
TCGA-50-6	0.060274	1	2.1492594	0.5014137	0.7891867	1.3315906	0	0
TCGA-50-6	0.3260274	1	1.9972924	0.2617113	1.2913678	0.9191494	0.2580968	0.108491
TCGA-49-4	2.3424658	1	1.587221	0.7411434	1.2371354	0.8760945	0.2207014	0.0920721
TCGA-73-4	0.769863	1	0.979879	3.6998847	0.8316341	0.6104166	0.3538878	0
TCGA-NJ-A	5.9205479	0	1.4122396	2.0003606	0.955164	0.6660294	0.276318	0
TCGA-44-7	1.969863	0	1.3741214	1.4591168	0.7418337	0.4718638	0.0892269	0
TCGA-64-1	1.7205479	1	1.7593689	3.7171445	0.6848187	0.2571313	0	0.0545012
TCGA-55-6	5.7780822	0	1.9192638	1.874797	0.5649147	0.7816535	0.2855798	0
TCGA-86-7	1.7123288	1	2.0197731	1.2372578	1.0174944	1.1033298	0.3418719	0
TCGA-86-8	1.9013699	1	1.8934787	1.8551127	1.0980152	1.2956055	0	0.1411712
TCGA-MP-	4.8712329	1	1.724301	1.9676113	0.9234915	0.7642612	0.2687937	0.0291348
TCGA-86-8	2.2986301	0	1.9595476	1.5229597	1.1779178	0.7650255	0	0.0739573
TCGA-67-3	1.569863	0	2.0897354	2.092681	0.380397	0.7625614	0.1268407	0.1264443
TCGA-86-8	1.7890411	0	2.408413	0.2832106	2.0554733	0.6723801	0.8910303	0.328262
TCGA-78-7	8.6821918	1	2.3099038	1.0158543	1.187261	0.7990044	0.1263121	0.3694944
TCGA- MN	2.2657534	0	1.5253172	1.0287106	0.6655747	1.6939441	0.4568061	0
TCGA-49-4	2.7369863	1	0.9830209	0.2913678	0.1642714	1.1896664	0	0
TCGA-62-A	3.9835616	1	2.3156822	0.374177	2.1478928	2.315885	0.7900213	0.8357624
TCGA-86-8	0.0657534	0	1.8671848	2.2276179	1.1641426	2.460297	0.1095612	0.0224743
TCGA-86-7	2.5945205	0	1.8113477	0.514501	1.0684645	1.4644069	0	0
TCGA-67-3	1.6712329	0	1.7749662	3.2508252	0.988412	0.3246954	0	0.1104969
TCGA-38-4	2.3671233	1	1.2105759	0.1792562	0.5582676	0.729879	0.1767057	0
TCGA-55-6	4.4712329	1	2.5541959	3.4506306	1.2808376	0.5354558	0.2895984	0.8813514
TCGA-86-8	0.9671233	0	1.7137398	3.8957389	1.142152	1.9626231	0.3719491	0
TCGA-05-4	3.0849315	0	1.5489791	0.5832302	0.5111644	0.8690818	0.2185952	0
TCGA-67-6	1.1561644	0	1.8168045	2.9160753	0.9569488	1.6656202	0	0.1561377
TCGA- MN	3.2191781	0	1.4682184	3.5530533	0.548338	2.4422004	0	0.148869
TCGA-L4-A	1.1917808	0	1.7035437	1.1983056	1.1095612	0.5253673	0.1582077	0.187261
TCGA-05-4	3.7506849	0	1.9063134	1.9088897	1.2372578	0.4480552	0.4213714	0.5085301

TCGA-55-8	1.8465753	0	1.6076262	1.7163764	1.1585309	0.6321751	0.2017593	0
TCGA-55-6	3.5424658	1	2.1508846	0.1858665	0.9176991	1.7337448	0.4918018	0
TCGA-55-8	1.9232877	1	1.8915358	0	0.623586	2.0469799	0	0.0652966
TCGA-86-7	2.7315068	0	1.6176746	1.8547537	1.0971388	2.3094964	0.2609891	0.3745109
TCGA-38-4	4.0876712	1	1.4845506	0.5582676	1.0199865	1.0295589	0.092884	0
TCGA-55-A	1.7260274	0	1.3325074	3.6687772	0.5637439	1.0496308	0	0.0301242
TCGA-55-8	2.4328767	0	2.3600552	0.2200823	1.4615286	1.0891591	0.3142905	0.0683269
TCGA-44-7	1.5260274	1	1.7539468	1.3288365	0.9540474	1.2956055	0.1365852	0.0282862
TCGA-55-8	0.1315068	0	2.1030612	1.8128678	0.6180507	0.9668734	0	0
TCGA-97-A	1.4794521	0	1.878843	2.091666	0.5579737	0.7324863	0.16221	0
TCGA-MP-	7.1780822	1	1.9715895	2.3728685	0.9091971	0.7941871	0.164915	0.0680516
TCGA-86-7	2.9369863	0	2.0425393	1.1763228	1.8981309	1.8738132	0.5512945	0
TCGA-86-8	0.3808219	0	2.0939995	2.0315364	1.234256	1.486045	0	0
TCGA-05-4	1.6712329	0	1.9181954	1.3225627	1.3172462	1.9787814	0	0
TCGA-86-6	1.030137	1	1.5921102	0.5684213	0.7420062	1.7838757	0	0
TCGA-67-6	0.3863014	0	1.5804832	2.1687704	0.372395	1.0107798	0	0
TCGA-73-4	2.5260274	1	2.0090963	0.4404207	1.0279325	0.494262	0.1897929	0.0398403
TCGA-64-1	4.7342466	0	2.6168987	3.0935602	0.8443857	0.7005285	0.6946574	0
TCGA-55-6	3.7780822	1	1.2462256	1.4207249	0.6461627	0.7910222	0.2708279	0
TCGA-L9-A	1.5479452	0	2.2095469	0.251204	1.2220008	1.2889491	0.1292171	0.0267293
TCGA-55-1	4.0520548	0	1.9398284	1.7389407	1.499425	0.4111008	0.6203987	0.2053925
TCGA-55-6	8.9342466	0	1.8042188	1.5808208	0.8519988	0.8815863	0.258338	0
TCGA-86-8	1.9205479	0	1.9466934	1.8627093	1.3131297	0.7278769	0.5497185	0.0524159
TCGA-91-6	0.8849315	0	1.7633691	0.3234277	1.0590783	0.5851548	0	0
TCGA-55-A	1.8876712	0	2.3723392	0.3353689	1.5730351	1.3761237	0.826518	0
TCGA-05-5	1.2520548	0	1.3905576	1.2634551	0.8622728	0.5506052	0.1570437	0.4506966
TCGA-05-4	1.1671233	0	2.2492936	4.5497246	1.2385423	0.7633266	0.4520682	0.1505597
TCGA- MN	0.230137	0	1.7369944	2.0037821	1.1486738	2.1800205	0	0.0301242
TCGA-95-A	1.7835616	0	2.7350455	0.1781091	1.7614129	1.5012608	0.3320491	0.0367495
TCGA-49-4	1.1726027	1	1.6898806	1.0394894	0.7494914	1.20989	0.1497796	0
TCGA-05-4	2.0027397	1	2.5726957	1.8821735	1.8080967	0.0832474	0.8663937	0.0167816
TCGA-86-8	1.2164384	1	1.5029925	0.5381408	0.3609822	1.8124572	0.1996252	0.198997
TCGA-44-3	3.0958904	0	1.3197913	1.2092038	0.6254575	0.7542889	0	0.0254541
TCGA-62-8	3.5315068	0	1.2441564	0.9007219	1.057069	0.6720179	0	0
TCGA-78-7	10.794521	0	2.5740773	3.0372944	1.2371354	1.4958998	0.1432628	0.0873271
TCGA-78-8	0.8794521	1	2.2282646	1.2024491	0.6280737	1.2863489	0.5131875	0.0404015
TCGA-38-4	0.969863	1	1.3237736	0.9281239	0.7737432	1.3840498	0.199374	0.4688439
TCGA-73-7	4.1945205	1	2.0141767	1.8699109	1.8002476	0.9266826	0.4815056	0.0668125
TCGA-50-6	0.9205479	1	1.6578686	0.6009358	1.2013829	0.9581374	0.5934972	0
TCGA-67-3	1.0547945	0	2.0831793	1.2368905	1.3925372	0.7117592	0	0
TCGA-71-8	0.5753425	1	2.602148	1.2739932	1.0836559	1.440527	0	0.1104969
TCGA-50-5	1.7890411	1	2.2624632	1.6271399	1.6804144	0.933346	0.228357	0
TCGA-78-7	3.2082192	1	2.0268709	0.2611095	2.385072	1.1911836	0	0

TCGA-44-5	2.7753425	0	2.3331375	2.8740887	0.6053995	1.0047531	0.4291071	0.6663021
TCGA-44-2	3.7452055	0	1.4382396	1.8471551	0.8085086	1.0943712	0.247563	0.0354832
TCGA-86-8	2.7205479	0	1.7166837	1.7546738	1.0592168	1.2110745	0.2027625	0.0214797
TCGA-S2-A	1.4054795	0	1.3938006	0.9920433	0.7108781	0.703898	0.1482183	0
TCGA-MP-	0.8410959	1	1.3981314	2.4735791	0.865998	0.614003	0	0.5043662
TCGA-50-7	0.8438356	1	1.7775354	1.5944052	1.0515113	1.0248161	0.1596292	0.6413611
TCGA-05-5	1.8356164	0	2.3353403	2.2410775	0.6928735	1.206018	0.2226809	0.0472592
TCGA-44-8	1.0547945	0	1.6428864	1.7995018	0.7755142	1.8274532	0	0.0224743
TCGA-05-4	1.8328767	0	2.2715751	0.229957	0.7838338	0.9918257	0.5949307	0
TCGA-55-1	5.6575342	0	1.8434613	3.597984	1.6282603	0.9216648	0.2152437	0.0455828
TCGA-99-8	0.1205479	0	1.5760383	0.5304703	0.5840004	0.8119642	0.1964811	0.1206174
TCGA-67-6	0.4767123	0	1.9225785	2.3595493	0.9141052	1.2576745	0	0
TCGA-97-7	2.1232877	0	1.6104166	0.8509594	0.8251343	1.4724358	0	0.0281448
TCGA-05-4	2.1671233	0	2.0016581	1.1972993	0.7974281	1.1200201	0.6082415	0.0425043
TCGA-50-6	3.4739726	1	1.8877202	0.2168584	0.4326388	1.3411888	0	0
TCGA-50-6	1.0136986	1	2.0246034	1.2797684	1.0123541	0.8577422	0	1.287945
TCGA-53-7	2.5452055	1	2.4218831	3.0441843	0.9831669	1.3946239	0.5465611	0
TCGA-62-A	6.0246575	0	2.5370227	3.8245438	1.4722278	0.9164002	0.1828194	0
TCGA-50-5	5.0136986	1	1.8536364	1.1604685	0.7363018	0.5251668	0	0

AC032011.1	AL162632.3	LINC02518	LINC00592	AL031600.2	AP000346.1	AC012409.4	riskScore	risk			
0.064331	0.0614306	0.363115	1.0467006		0	0.1764504	0.3605328	1.837029 high			
0.0563061		0	0.4142438	0.6289135	0	0.2947825		0	1.4779859 high		
0.2699906	0.6071054	1.4440839	1.1230702	0.4287856	0.57415		0	2.4338245 high			
0.2415962	0.0812033		0	0.4156505	0.3858175	0.119821		0	1.004778 low		
0.3925372	0.1367164	1.5661819	0.2229282		0	0.0697022		0	3.7342447 high		
0.3811726	0.1680001	0.1247252	0.5311693	0.725872	0.2445218			0	0.7724869 low		
0.193204	0.2189671	0.0839282	0.2352369	0.5197432	0.1667154			0	0.7521256 low		
	0	0.1889073		0	0.1385524		0		0	2.1287923 high	
	0	0.4188916		0	0.590626	0.1998765		0		0	1.2250488 high
0.117695	0.5319678	0.1228715	0.2330291	1.3230241	0.3131297			0	0.4811575 low		
0.0799754	0.0762854	0.1624679	1.3598023	0.6556266		0	1.045443	4.0909035 high			
0.09815		0	0.1981169	0.1646576	0.2358496	0.0706641	0.2856981	0.4322334 low			
0.1088924	0.0529723	1.3999367	0.0382957	0.4808855	0.1886541	0.3148705		1.1046307 high			
0.1107641	0.4248146	0.115566		0	0.4882065		0		0	0.5089751 low	
0.2716647		0	0.1483484	1.2981304	0.8352773	0.2886539			0	0.4476429 low	
0.2851063	0.108491	0.1560082	0.248049		0		0	0.2263852	0.6715002 low		
	0	0.2503553	0.1871343	0.2413521		0		0		0	1.5964998 high
0.3920976	0.2266318		0	0.2056427		0		0	0.3533233	1.594431 high	
0.2683146	0.6592852	0.2276179	0.1437199	1.7971583	0.4324784	2.8524783		0.6197737 low			
0.232538	0.2895984	0.0852887	0.3337672		0	0.169412			0	0.9626496 low	
	0	0	0.2125071	0.1431322		0		0	0.5580717	1.9866165 high	
0.1071522		0	0.0756011	0.2981891		0	0.0518593	0.2141248	1.1805251 high		
0.329181	0.2574934		0	0.6767187	0.6014113	0.8507194			0	0.3508829 low	
0.0309716	0.2211965		0	0.2376861	0.6305919	0.0440443	0.1829465	1.8796794 high			
0.1273691		0	0.4197546	0.2319239	0.158854	0.1782366			0	0.5230586 low	
0.3609822		0		0	0.4421208			0		0	0.1601595 low
0.3460775	0.0522768	0.2164859	0.2784606	0.9834588	0.5276709	0.3115031		0.1870868 low			
0.2178511	0.2467121		0	0.5340616	0.5787938	0.1281614			0	0.4459347 low	
0.0835197	0.0797024		0	0.2686739		0	0.2262619			0	0.4720201 low
0.1050075	0.2821432	0.559345	0.6029319		0	0.2811938			0	1.0729096 high	
0.1880211	0.2131295		0	0.1566555		0	0.2609891	0.230449	1.2679868 high		
0.1170301	0.0569996	0.5274708	0.0813396		0	0.679694	0.3365119	0.3723819 low			
0.0316775	0.5718224		0	1.0892269	0.2914857		0	0.3516283	1.1817552 high		
0.9602646	0.879758	0.2926247	0.174449	0.993155	0.4895429	3.0595456		30.012552 high			
0.075738	0.4435006	0.2933705	0.4152178	0.626953	0.1068843			0	0.6801715 low		
0.0918014		0	0	0.1832006	0.4129441			0		0	0.7046534 low
	0	0.2040155	1.1149666	1.9226926		0	0.1551016	0.5808208	9.6009645 high		
0.067501	0.2697513	0.0705267	0.1573025	0.8847933	0.3092926	0.2005044		0.2506054 low			
0.1190241	0.1673579		0	0.2711866	0.9015717	0.166844	0.3420996	0.5663867 low			
0.0713508	0.3122005		0	0.1432628	1.1881477			0		0	0.2234173 low
0.2028878	0.1003049	0.2115107	0.0729976		0	0.147437			0	1.5846758 high	
0.1867541	0.2862898		0	0.0885487	0.7630716	0.3362834			0	0.0992063 low	

0.24123	0	0	0.1700532	0	0	1.0925457	2.9702002 high
0	0.1700532	0	0.0842004	0.2877087	0	0.8610819	0.563515 low
0.169412	0.187261	0.0612923	0.5425057	0.1437852	0.082975	0.175301	2.1612698 high
0.0905825	0.1680001	0.1833277	0.6988408	0	0.2445218	0.4886178	1.7101053 high
0.0931545	0.531369	0.5044679	1.98167	0.153546	0.0450235	0.3525326	9.6653901 high
0.0676387	0.1264443	0.3798427	0.6115501	0	0	0	3.6134237 high
0.0360461	0	0	0.353662	0.3276874	0.1479579	0	0.8746885 low
0.064331	0	0.1314578	0	0.2990101	0.0461418	0	0.5318282 low
0.439038	0	0	0.1296127	0.7572793	0	0.4226636	1.8417255 high
0.0377336	0.1389455	0.0777899	0.0766959	0.3419857	0.2496274	0	0.6020551 low
0	0.0299829	0	0.1051417	0	0	0.1851054	1.6960858 high
0.0444641	0.4724878	0	0.1471764	0.2113861	0.0629506	0.8304176	1.3374245 high
0.2156165	0.2062681	0.116764	0.7108781	0	0.0805212	0	1.1951917 high
0	0.5566011	0	0.2260152	0.9976176	0.1900459	0.3864799	0.4780244 low
0.1340895	0.0439044	0.0947766	0.5541468	0	0.0652966	0.2653169	2.1910747 high
0	0.0890913	0	0.1265765	0	0	0	0.3760539 low
0	0.1642714	0	0.1202192	0	0.3453964	0	0.5643762 low
0.0889557	0.3132459	0	0.2322924	0	0.1251221	0	1.5694991 high
0.3441467	0.1184926	0.0875986	0.8329305	0.2028878	0.4687397	0	1.8695615 high
0.0574156	0.054779	0	0.223299	0.2689134	0.429964	0	0.7570053 low
0.0617071	0.115566	0.1263121	0.7580473	0	0.2473199	0	1.2734821 high
0.0803848	0.0766959	0	0.2108875	1.1086248	0.5752155	0	0.1402272 low
0.0429244	0.0809305	0.2505978	0.0588014	0	0.4273916	0	0.4984167 low
0.1335635	0.4212637	0	0.0930192	0.4075159	0.4033768	0	0.9133367 low
0	0.1547128	0	0.5345597	0	0	0	1.807309 high
0.0543623	0.3292958	0	0.178874	0.255682	0	0	0.9190599 low
0.2061431	0.6650289	0.2147465	0.0742314	0	0	0.9675375	8.3443878 high
0.422233	0.2950177	0.0990932	0.458067	1.2463473	0.2555612	0	0.1605141 low
0.3353689	0.1152996	0.4485838	0.5650122	0	0.3203114	0	3.7265309 high
0.0654344	0.4135941	0	0.4674882	0	0.0469799	0.1945914	1.409458 high
0	0.1423481	0	0.1531568	0	0	0.4202937	3.2798424 high
0.2497487	0.2389091	0	1.3101074	0.817787	0.2814312	0.9562797	0.1906702 low
0.2662768	0.1330374	0	0.0654344	0.2270017	0.3111543	0	0.5943335 low
0.0522768	0.0499095	0	0.0714881	0.640806	0.144438	0.2984237	0.5090026 low
0	0.2231754	0.0647449	0.293959	0	0.3924273	0	1.0134427 low
0.1890338	0.1807844	0	0.7426961	0	0	0	1.0898535 high
0.0515809	0.0492125	1.1981169	0.2930173	0.2434254	0	0	0.3207518 low
0.1112985	0.582267	1.1750455	2.4058018	0.6846392	0.3627785	0.9987732	4.3437158 high
0.3905576	0.1031283	1.2015711	0.279293	0.4773665	0.5291713	0.5691022	2.5292552 high
0.2607483	0.3056787	0	0.2266318	0.9996393	0.3587338	0.6925165	0.6203181 low
0.2856981	0.2734562	0.2973675	0.2934882	0.3518544	0	0	2.3383884 high
0.0600474	0.1392075	0.0626744	0.1781091	0.514804	0.2033891	0	0.4244175 low
0.2256451	0.0458623	0.5806279	0.4317834	0.2276179	0.82546	0.8817429	0.381726 low

0.0413832	0.3529845	0.3140584	0.4218023	0.7937711	0.2716647	0.2398868	0.1754234 low
0	0.4785059	0	0.1206174	0.2142492	0	0	2.0082881 high
0.0519985	0.0976108	0	0.5397296	0.2450088	0	0	1.1509943 high
0.2126316	0.7054914	0	0.5223571	0.3415304	0.1545832	0.2198345	1.6677184 high
0	0.1678717	0.3455099	2.5908894	0	0.0436245	0	7.5140829 high
0.0832474	0.2606279	0	0.1664584	0.2012575	0	0	0.5225816 low
0	0.3149865	0.093425	0.7307486	1.0910566	0.3489122	0	0.0840711 low
0	0.0787464	0.3173041	0.1651723	0.3750672	0	0	0.5442044 low
0.1428708	0.5811102	0	0.3628906	0	0.1994996	1.1984313	1.1611362 high
0.1163648	0.3105728	0	0.4287856	0	0.1631122	0.834873	3.2326901 high
0.2174789	0.3317052	0	0.2464689	0.3490255	0.2073931	1.0087737	2.5528186 high
0.4662356	0.0923428	0.1009776	0.162339	0	0.0695647	0.2817873	0.9519871 low
0.0839282	0.2970153	0.4603756	0.4107753	0	0.1182268	0	1.4996628 high
0.1244605	0.5721136	0.5871729	0.3427823	0.2035144	0.1184926	0	1.4830676 high
0.1895399	0.1232689	0	0.2149951	0.3057955	0	0	0.7382568 low
0	0.635708	0.4579619	0.821547	0.5369481	0.0890913	0	1.7730975 high
0.2126316	0.2033891	0	0.1493894	0.8488783	0.1545832	0	0.4682511 low
0.2674759	0.7200157	0.1914363	0.7249994	0	0	0.5083272	16.580807 high
0	0.2244108	0.1673579	0	0	0	0	1.5774327 high
0.1011122	0.403595	1.3316479	1.6214776	0	0	0.202261	28.000335 high
0.0528332	0.4006472	0	0.4148932	0	0.0747794	0	2.0411438 high
0.1056781	0.1952216	0.1102296	0.0371713	0	0.2172307	0	0.6582044 low
0.6164048	0.8292813	0.7020369	0.0435545	2.6274434	0.7987556	0.6371929	0.0371357 low
0.0735461	0.4838803	0.6286336	4.3966938	0	0.1036655	0.7225535	105.54487 high
0.2311866	0.1512094	0.164915	0.354565	0	0.2204538	0	0.9447688 low
0.2765562	0.1384213	0.527771	2.2218771	0.6164518	0	0	4.6211114 high
0	0.0676387	0	0.0963968	0	0.0999011	0	0.7529104 low
0.3447149	0.3718376	0.9157886	0.6623878	0	0.4694693	0.3102238	3.8974465 high
0.2078929	0	0.0758749	1.2749474	0	0.1982427	0	2.5178169 high
0.03492	0.5528689	0.0720372	0.3817264	0	0.0494914	0.3838287	1.9356001 high
0.0847446	0.230326	0.3255015	0.0588014	0.3846025	0.2295879	0.8100729	1.3892502 high
0.0618453	0.369941	0	0	0	0.4590119	0	1.3258626 high
0	0.2085173	0	2.7945614	0	0.2078929	0.4202937	3.1179505 high
0.1156992	0.4833644	0.2321695	0.9819987	0	0.5616927	0.6035967	4.0725541 high
0.7029235	0.0625362	0.1963552	0.4514353	0	0.2211965	0	1.5726712 high
0.1430015	0.2311866	0.1491293	0.3223897	0	0.1522483	0.2166101	0.2872711 low
0.1367164	0.0668125	0.7882681	0.1845977	0.8090851	0.1910572	0.3885753	0.2031448 low
0.0825664	0.153546	0	0.0571383	0	0.3238888	0	0.4247189 low
0	0.2668765	0	0.1343524	0.4420146	0.3826119	0	0.1240232 low
0.2773897	0.3819478	1.075738	0.2263852	0.2367069	0.2028878	0	2.2633043 high
0	0	0	0.2702299	0	0.1183597	0	1.7368935 high
0.169027	0.0558898	0	0.0405418	0	0.2353594	0	0.1781068 low
0.3795101	0.281075	0.2106382	0.1418251	0.4641455	0.2151194	0.5537537	0.7370721 low

0.2878269	0.3249258	0	0.0850167	0.2901885	0	0	0.811815 low
0.3678183	0.1150332	0	0.8463128	0	0.1147668	0.444985	6.1754803 high
0.0287106	0.0806577	0	0.3361691	0.4912887	0.5442875	0	0.4480249 low
0.533962	0.4499575	0.093425	0.2336427	0	0.2963106	0.4834676	3.909751 high
0.1764504	0.3196179	0	0.0630887	0.4094724	0.1278974	0.8553519	1.9421919 high
0.175812	0.1680001	0.6245221	0.2363395	0	0.1273691	0.488515	2.0850652 high
0.4056657	0.072586	0.1545832	0	0.6286336	0.1071522	0.4185678	0.7279752 low
0.0716254	0.0683269	0.1457427	0	0	0.1952216	0.3965979	1.9495729 high
0.6964984	0.6529683	0	0.0753729	1.3203884	1.106594	1.194066	0.1983789 low
0.1645289	0.1571731	0.3246954	0.8044666	0	0.6004601	0	0.7678022 low
0.1514692	0.2509616	0.3002995	0.1460035	0.4170559	0.3606452	0	0.3001405 low
0.2081426	0.1356662	0	0.4003193	0	0.1025909	0	2.1464827 high
0.067088	0.2947825	0	0.6672109	0	0	0.6705684	3.4664463 high
0.0871912	0.3750672	0.1765781	0.7886857	0.3942946	0	0	0.4646364 low
0.0654344	0.3891262	0.1335635	0.2525368	0.5541468	0	0	1.1588502 high
0.4370146	0.0596322	0.1276333	0.3827226	0.2906603	0.4627853	0.3506104	0.7932162 low
0.1118326	0.3582837	0	0.2218152	0	0	0	1.232784 high
0.0943712	0	0	0.2458607	0	0.9008765	0	0.4993893 low
0.3645724	0	0	0.4802651	0.446468	0.4953882	0	0.3440886 low
0.0745054	0.1054099	0.9571717	0.3687125	0.7384652	0.249506	0.4109923	0.708763 low
0.0603933	0.8061591	0	0.1411712	2.179734	1.1069848	1.3977208	0.010428 low
0	0.1228715	0	0.0896337	0	0.1801479	0	0.9251738 low
0.0787464	0	0	0.2539893	0	0.1108977	0.4315695	2.1763285 high
0	0.2529	0	0	0	0	0	0.9995176 low
0.116897	0.1644001	0	0.5208497	0.2784606	0.0842004	0	0.8772279 low
0.0451634	0.0432045	0.0930192	0.285343	0	0.1840898	0.2607483	1.0305324 high
0.0790196	0.0754642	0.3052118	0.3458505	0.3609822	0	0	2.2585649 high
0.1528973	0	0	0.5212518	0.6453328	0.0746424	0	0.3441084 low
0.1067503	0.3289513	0.1112985	0.2760797	0	0.348799	0	1.3421758 high
0.2667566	0.0681893	0.1454818	0.5735684	0.3289513	0.1007086	0.3959402	0.872786 low
0.0339339	0.0640551	0	0.1133004	0.1634987	0.0946415	0	0.5874559 low
0	0.2510828	0	0.1852323	1.8140171	0.2788174	0	0.0937125 low
0.4760187	0.2210728	0.4469972	0.1236662	1.2109499	0	0.3450557	0.147471 low
0.1828194	0.0457225	0.5096452	0.0970714	0.2270017	0.4699901	0	0.1860516 low
0	0.0588014	0.1257833	0.5647196	0.2867629	0.0869196	0	0.1489708 low
0.2235461	0.3646845	0.3564819	0.0328061	0	0	0	3.0280899 high
0.0588014	0.2123826	0.605684	0.4557546	0	0	0.3320491	1.4858034 high
0.587941	0.1852323	0	0.1357975	0	0.4953882	0	1.5624743 high
0.2606279	0.249506	0.1147668	0.0387171	0.8474357	0.0791562	0.7983409	0.6876395 low
0.1269729	0.3369689	0	0.2504766	0.3011194	0.5447821	0.3630028	0.6825065 low
0.3006509	0.3248106	0	0.0641931	0	0.305562	0	1.9864506 high
0.2565276	0.086512	0	0.1231364	0	0	0	2.1907652 high
0	0.0626744	0	0.045443	0	0	0	0.5271719 low

0.2018847	0	0	0.1416944	0.6838312	0.0604625	0	1.0092571 low
0.1868808	0.2336427	0	0.3618806	0.7637515	0.0918014	0.3637878	1.3420215 high
0	0.1487389	0	0.20989	0.3645724	0.7222037	0	0.3530572 low
0.2558028	0.5298709	0	0.4003193	0.2579761	0.3520805	0	1.8093232 high
0.1015156	0.6160753	0	0.0705267	0.4518573	0.1426095	0	0.938814 low
0.1521185	0.5098478	0	0.8186052	0.3567073	0	0	5.4466391 high
0	0	0.9253918	3.3532245	0	0	0	6.4471136 high
0.2655569	0.1326426	0.1448295	1.2349304	0	0.1940871	0	2.3068418 high
0.2856981	0.2734562	0	0.1245928	0.1512094	0.1696685	0	3.4076586 high
0.2820246	0.1263121	0.0706641	0.6570912	0	0.2692726	0.2007554	1.6574667 high
0	0.4307135	0	1.2144357	0.167101	0.0492125	0.2033891	0.5299237 low
0.0735461	0.0702519	0	0.23842	0.3379965	0.6058736	0	0.1987646 low
1.0324535	0.4392508	0	0.3294107	0	0.1458731	0	12.455056 high
0.2865263	0.4319973	0	0.0708015	0.9441088	0.3929768	0	0.287927 low
0.1388145	0.0275787	0.2251515	1.1955996	0	0.1939609	0.3234277	0.4561428 low
0.0965317	0.260387	0.6135317	1.8080555	0.1587248	0.0918014	0	4.2528562 high
0.2234225	0.1319845	0.4330663	0	0	0.1315895	0.5025343	0.6812703 low
0.3386811	0.1164978	0.6351507	0.3817264	0	0	0	1.6899033 high
0.1651723	0	0.5084287	0.3218127	0	0.0807941	0	1.4828032 high
0.191689	0.063779	0	0.2174789	0.7800581	0.1826923	0.3726178	0.1308307 low
0.103934	0.0992279	0.2092662	0.3871419	0	0.1458731	0	0.8590923 low
0.2990101	0.2862898	0.11112985	0.2118844	0.4722798	0	0	1.4598453 high
0.1307991	0.1248575	0.0923428	0.2041407	0	0	0.2590615	1.6446825 high
0.0805212	0.1497796	0	0.055751	0	0.4081684	0	0.3724493 low
0.6178627	0.0451634	0.1886541	0.3248106	0.4192153	0.1925729	0.2723816	2.2561514 high
0	0.1558787	0	0.0581086	0	0	0	2.0394978 high
0.0856966	0.6088092	0.3286067	0.548338	0	0	0.4653999	2.3641274 high
0.2240403	0.0750534	0	0.0545012	1.2709475	0.3997727	0	0.0526554 low
0.2157407	0.2063932	4.1856525	3.6802456	0	0.1064823	0.4162993	87.563515 high
0	0.6332917	0	0.450591	0.5348585	0	1.2529	4.7094164 high
0	1.0878701	0.5788904	0.7215039	0	0	0	2.1571233 high
0.3306731	0.2187192	1.230449	1.124262	0.5193406	0.1665869	0.3415304	2.3381597 high
0.1858665	0.9718102	1.340961	4.0897693	0	0.1770886	0.891808	289.3748 high
0.1513393	0.0980152	0	0.235482	0	0.0374525	0	0.824336 low
0.9632275	0.1865006	0.6797841	0.3479302	3.3505727	0.3124328	2.5281545	0.3420444 low
0.5856997	0.7476882	0	0.642563	2.6886504	1.0571152	2.135305	0.3501442 low
0.293135	0	0	0.072586	0	0	0	1.5849235 high
0.4103411	0.1882743	0	0.2023864	0	0.554638	0	0.0810846 low
0	0.1440464	0.0648829	0.0430645	0.4151096	0.1298765	0	0.7559167 low
0.1870076	0.1099623	0	0.251204	0	0	0	1.017707 high
0.2149951	0.5454742	0.0676387	1.9164766	0.1580785	0.1773438	0.6518668	7.4057109 high
0.1423481	0.0695647	0	0.4393572	0	0	0	1.6537072 high
0.1624679	0.0797024	0	0.0578315	0	0.117695	0	0.9430651 low

	0	0.0923428	0.1953476	0	0	0	0	1.047872 high
	0	0.1565261	0	0.0773797	0	0.1560082	0	0.3970622 low
0.1978654	0	0.1406478	0.047678	0.3186928	0	0.9367416	0	1.0420465 high
0.1811662	0.2905424	0	0.8139351	0.4195389	0.1928254	0.2726205	0	0.9306333 low
	0	0.0972062	0.1062143	0	0.2441564	0.2730981	0	0.4954564 low
0.2072682	0.0521376	0.1119661	0.1104969	0	0	0	0	1.1970234 high
0.0918014	0.3930867	0	0.8935175	0.4131608	0	0.4944668	0	5.4680533 high
0.1026581	0.2886539	0.0736832	0.800786	0.325444	0.6191782	0	0	0.6183704 low
	0.077243	0.2107629	0	0.154972	0.3534362	0	0	0.174767 low
	0	0.3515153	0.4468914	0.1103632	0	0	0	1.2438267 high
0.1584663	0.1514692	0	0.3401634	0	0.0775165	0	0	0.8710251 low
0.1263121	0.1207501	0.2526579	0.6177686	0.2997136	0.0911243	0	0	0.8927401 low
0.0350608	0.3314759	0.8670266	0.0945063	0	0.0978804	0	0	3.4123649 high
0.1800205	0.1197767	0	0.6419775	2.5752881	0.2383385	1.0616149	0	0.1424998 low
0.3362834	0	0	0.2826177	0	0.1040682	0	0	1.6572621 high
0.8105665	0.1937087	0	0	0	0.1471764	0.5548344	0	4.6568578 high
0.1727435	0.5104556	0	0.3850444	0	0	0.4809889	0	3.6808608 high
	0	0.1401242	0	0.4513298	0	0.2046416	0	0.3538543 low
0.1126334	0.054779	0	0.3231971	0	0.3666999	0	0	0.4685912 low
0.0691522	0.2188432	0.0721744	0.0942361	0	0.0496308	0	0	1.1002038 high
0.1997509	0.2778658	0	0.0717627	0.4591168	0.2770326	0	0	0.2843823 low
0.1166309	0.2148708	0	0.429964	0	0	0	0	0.9491242 low
	0.141302	0.1350094	0	0	0.1973622	0	0	0.5499977 low
0.0469799	0.1304038	1.5536063	0.4229865	0.2225573	0.2491419	0.269871	0	1.4480873 high
0.3061456	0.1044709	0.1142337	0.1830736	0.8441446	0.153546	0	0	0.331601 low
0.0713508	0.1953476	0.1453514	0.0972062	0	0.3663642	0.9609565	0	1.5250116 high
	0	0.1824381	0	0.5231605	0.2368293	0	0.2867629	1.0372248 high
0.3179986	0	0.4262111	0.0792928	0	0	0.5957901	0	0.7146649 low
0.1448295	0.4533332	0	0.4465739	0	0.138028	0	0	1.6455522 high
0.9498711	0.0485152	0.1042024	0.5535571	0.2397646	0.6895675	0	0	2.4105153 high
0.1174291	0.1389455	0.288772	0.3359405	0	0.0846086	0.3377682	0	6.7175232 high
0.2244108	0.0751903	1.4643546	3.4672273	0.3596336	0.782073	0.7639214	0	7.3785761 high
0.3135941	0.6047356	0	0.2732175	0.2055176	0	0.249506	0	2.7623105 high
	0	0.3012364	0	0.4755	0	0.1579492	0	1.63257 high
0.2062681	0.3708338	0	1.4829515	0	0	0	0	1.4937499 high
	0	0.1628545	0	0.4814023	0	0.0506062	0	2.1532441 high
0.6769895	0.8603669	0	1.4662356	0	0	0	0	6.2066835 high
0.1547128	0.5174269	0	1.08882	0.3623296	0.6456094	0	0	0.4058906 low
	0	0.1634987	0	0.0610158	0	0	0	1.3424542 high
0.2875905	0.0739573	0.1574318	0.4634132	0.3541136	0.8365705	0	0	0.3963637 low
	0.417488	0.0565835	0.4338141	0.3653566	0.7092906	0.5042644	1.2046416	2.4958187 high
0.0629506	0	0.1285574	0	0.7442472	0.3981314	0.3533233	0	0.3220103 low
0.0462815	0.2085173	0.5630605	0.1532865	0	0	0	0	0.7766535 low

0.2998308	0.0391384	0.0846086	0.2125071	0.1962293	0.0582472	0	0.6170099 low
0.1430015	0.3761792	0	0.1470461	0.3366262	0.455439	0	0.7646485 low
0.1235338	0.5407217	0.0871912	0.4089292	0	0.0600474	0	6.2436883 high
0.2048919	0.1012467	0.1106305	0.1091599	0.4697818	0.4651909	0	0.6632291 low
0.0364682	0.2273714	0	0.3173041	0	0	0	1.5875679 high
0.1130337	0.3903375	0	0.5337627	0	0	0	1.28754 high
0.0873271	0.16221	0.6060631	0.0899049	0.5583656	0.6172044	0	0.2782878 low
0.2051422	0.1961034	0.1108977	0.177854	0.4704067	0.1491293	0	0.7662477 low
0	0.1933302	0	0.4989657	0	0.1926992	0	0.8220398 low
0.064607	0.1774714	0.1318528	0.0880059	0.2998308	0	0.8911081	1.3853512 high
0.0657102	0.6574571	0.1340895	0.3651326	0.5559143	0.1798932	0	0.6564584 low
0.2006299	0.4539653	0.3213509	0.8475961	0.3797318	0.117828	0	0.7568185 low
0.3702759	0.231801	0	0.0706641	0.4532278	0	0.295723	0.9394457 low
0.3610946	0	0	0.1972364	0	0.4908781	0.5284713	1.4049306 high
0	0.3017047	0.711583	0.1160986	0	0.2313095	0	2.4845708 high
0	0.1443075	0	0.0536674	0.354565	0.2107629	0	0.5654839 low
0.1830736	0.107554	0	0.6219932	0.6286336	0.0546401	0.2244108	0.8164719 low
0	0.1840898	0	0.1350094	0	0	0	0.1552151 low
0	0	0.2216915	0.2848694	0	0	0	0.9273461 low
0.2390314	0.2703495	0	0.5316684	0.2409859	2.0648484	0	0.2352977 low
0.1987456	0.1170301	0	0.2169825	0.2003788	0	0.4514353	0.4965322 low
0	0.1939609	0	0	0	0	0	0.6292348 low
0.0994972	0.1840898	0	0.3720606	0.2389091	0.3841604	0	0.2346419 low
0	0.3474389	0.261952	0.0913952	0.971516	0.1835818	0.6712028	0.991138 low
0.0698396	0.3603081	0.1423481	0.3476657	0.5854433	0	0.945308	0.864843 low
0	0	0	0.1654296	0.2906603	0.0881416	0.3507235	1.1158405 high
0	0.3448285	0.1991226	0.790355	0	0	0.5265697	0.4118455 low
0.5342609	0.3469852	0	0.1889073	0	0.606537	0	1.7066152 high
0.2620723	0.4643546	0.1428708	1.5094425	0	0.1914363	0.3893465	2.9333888 high
0.0594937	0.0843364	0.178874	0.0411028	0	0	0	1.0977883 high
0.1545832	0.2027625	1.3828886	0.7649406	0.296663	0.5049763	0	0.1445762 low
0	0.6769895	0.1627257	0.3049782	0	0.1129003	0	6.0977929 high
0.1634987	0.0233263	0	0.1298765	0	0.0347792	0.1462642	1.6521428 high
0	0	0	0.0791562	0.5001902	0	0	0.5388708 low
0.0568609	0.2056427	0	1.4964623	0	0	0	0.6816882 low
0.5783108	0	0	0.154324	0	0	0	1.3111753 high
0	0.0764223	0.1624679	1.0681205	0.6557182	0.2172307	0	1.4651252 high
0.2790552	0.7385517	0.3991164	0.2991274	0.5474498	0	0	0.6840195 low
0	0.2468337	0	0.4421208	0	0.1282934	0	0.7183699 low
0	0.1491293	0	0.2584586	0.3654686	0.1129003	0.7746711	0.7822347 low
0.2990101	0.1971105	0	0.0740944	0	0.4094724	0	0.8027096 low
0.3378823	0.3944044	0.3514022	0.3469852	0	0	0	0.9615961 low
0.104605	0	0.2105136	0	0	0.2801249	0	0.3560182 low

0.0930192	0.2516888	0.1881477	1.0051844	0	0.1309309	0	1.251842 high
0.0342157	0	0	0.0468403	0.1650436	0.2278643	0.200881	0.5397735 low
0.041243	0.1150332	0.0850167	0.13711	0	0.1146335	0	0.6080782 low
0	0.0561673	0.1204847	0.0406821	0.2751263	0.1618232	0.3323929	0.899584 low
0	0.0496308	0	0.1713349	0	0.0735461	0.5429019	1.5447145 high
0.1243281	0.1187584	0	0.3542265	0	0	0	1.6360139 high
0.0897693	0.2431816	0	0.2343787	0.7206287	0	0	0.340563 low
0.0850167	0.1948435	0.0886844	0.4594316	0.3852654	0.2822619	0.6470842	1.7734398 high
0.3355976	0.3917677	0	0.8555114	0	0	0	2.4164574 high
0	0.4988636	1.0252415	3.2422366	0	0.6128714	0	1.4308464 high
0.1894134	0.1811662	0.0822938	0.0275787	0.359746	0.2141248	0	1.1986703 high
0	0.0820213	0	0.144699	0.3893465	0.2326608	0	0.3012426 low
0.1058122	0.1955996	0	0.0735461	0.2535053	0	0	1.4121117 high
0.2993619	0.380951	0	0.5991273	0.1958515	0.269512	0.2379308	1.3525246 high
0	0.0821576	0	0.276318	0	0	0	1.9156918 high
0.1074201	1.0226874	0	0.6637545	0	0.1508196	0	14.133876 high
0.5869809	0.1170301	0.3184615	0.1393385	0	0	0.2433035	0.7738428 low
0.272501	0.5319678	0	0.6188025	0.3360548	0.2894804	0	0.2623888 low
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