

**Supplementary Figure S1 Bubble plot of the correlation between pan cancer carcinogenic pathway and DDIT3 expression.**

**Supplementary Figure S2 Functional analysis for the darkred module genes**

Functional analysis of genes residing within the darkred module based on gene sets using (A) KEGG, (B) GO-BP, (C) GO-CC, and (D) GO-MF.

**Supplementary Figure S3 Functional analysis for the yellow module genes**

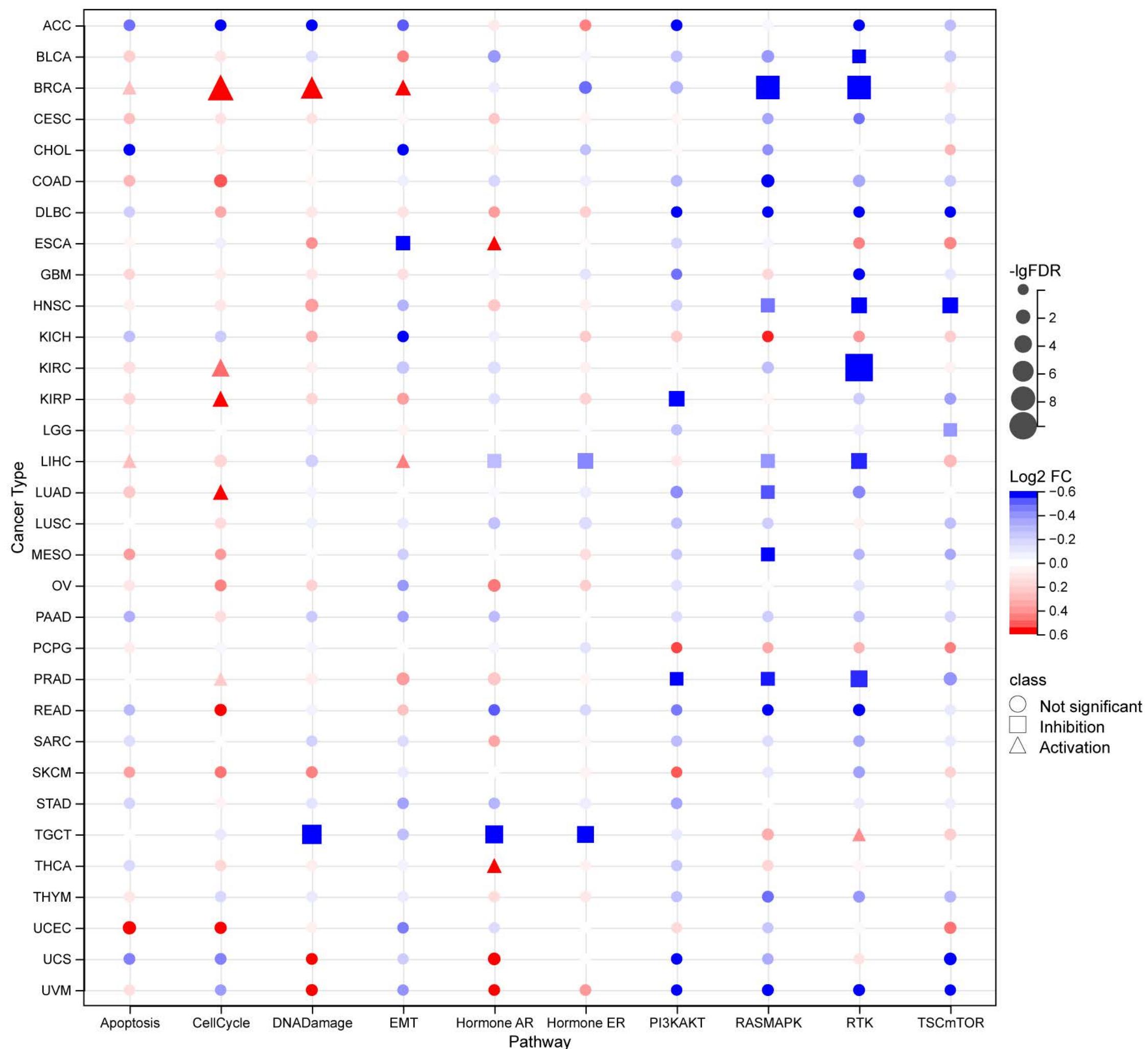
Functional analysis of genes residing within the yellow module based on gene sets using (A) KEGG, (B) GO-BP, (C) GO-CC, and (D) GO-MF.

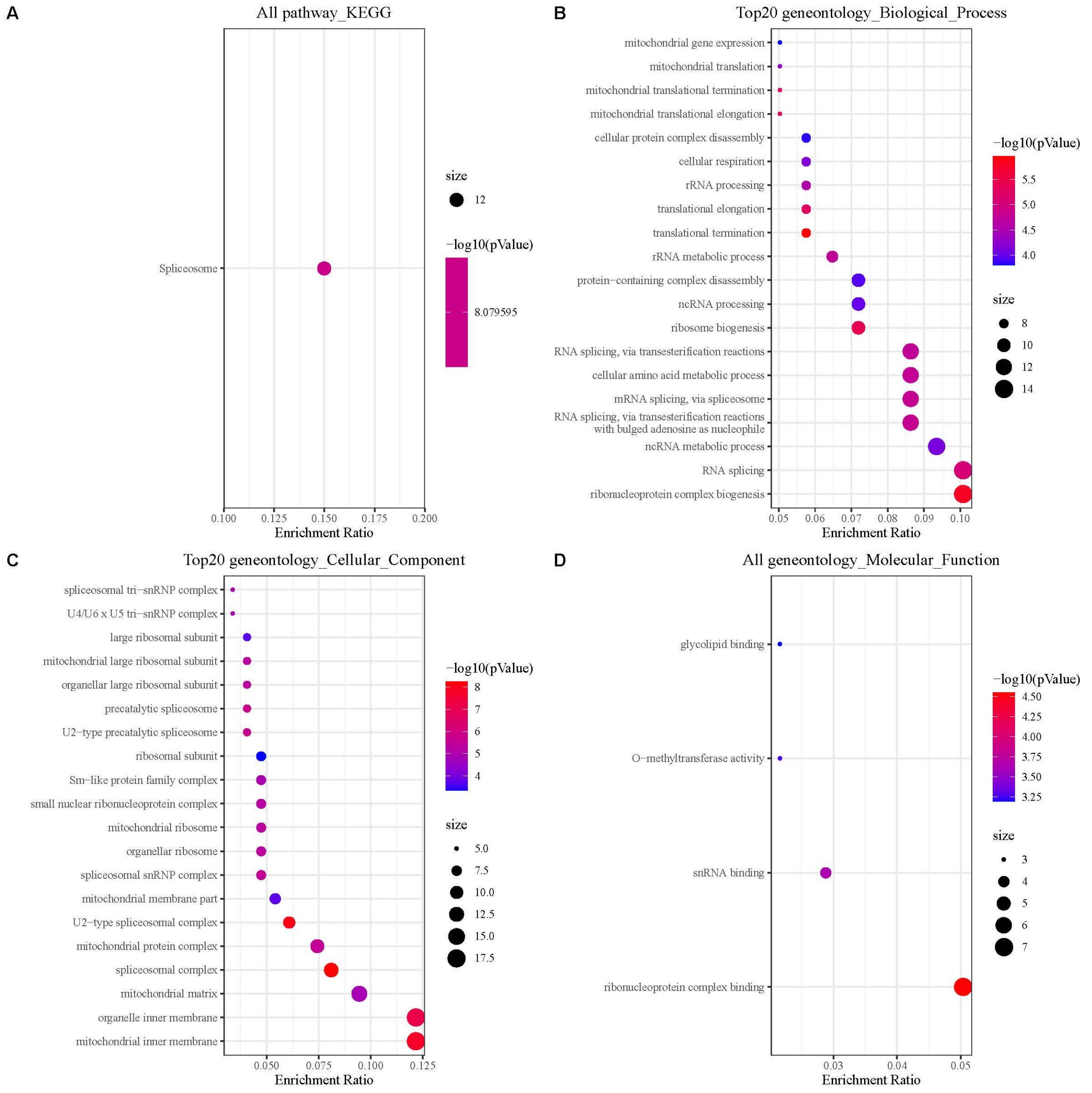
**Supplementary Figure S4 Screening of DDIT3-related genes in breast cancer.**

(A) Coefficient profiles as obtained from the LASSO regression model. (B) Cross-validation process employed for the selection of tuning parameters within the LASSO regression model.

**Supplementary Figure S5 DDIT3-related risk score and mutation landscape.**

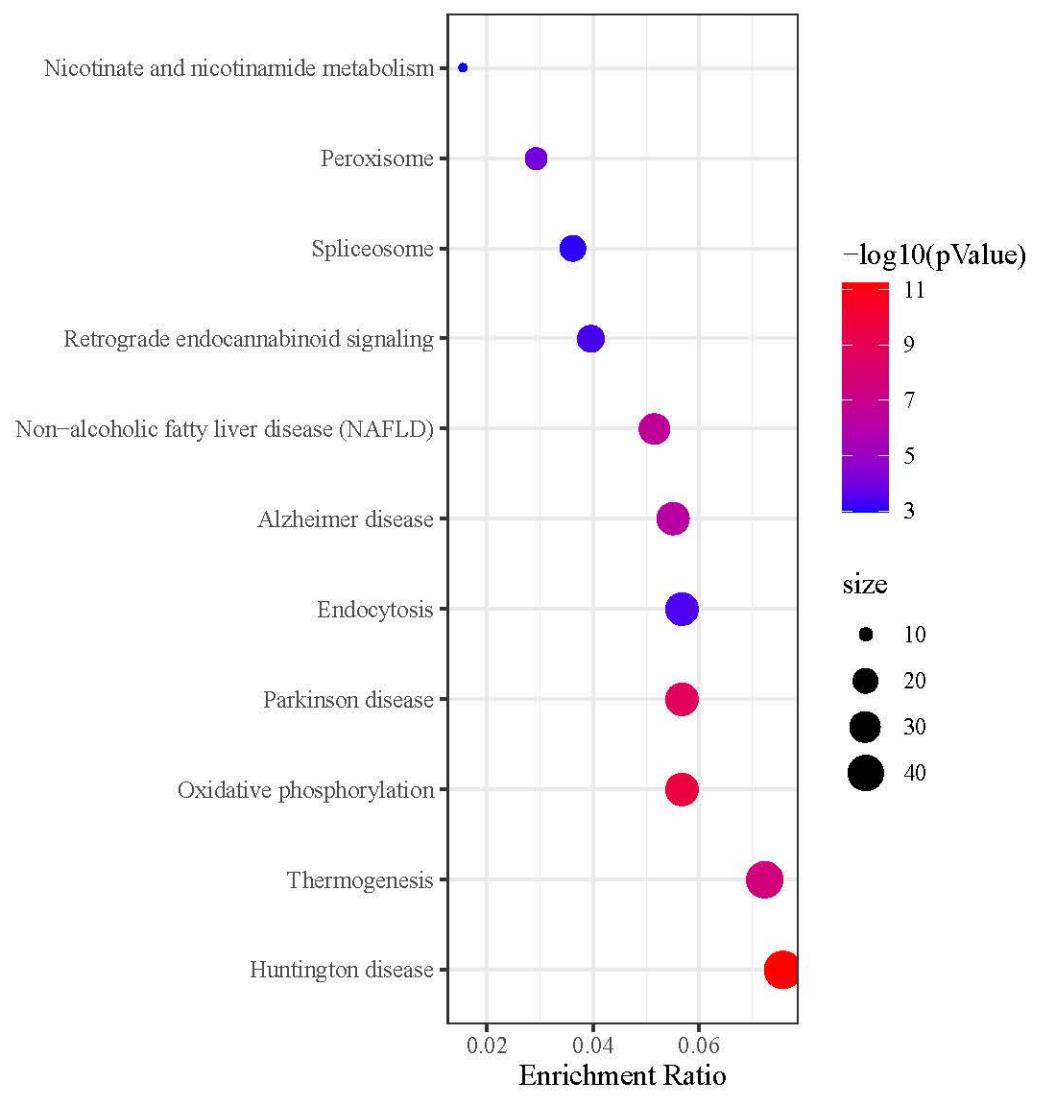
Comparative analysis of the mutation landscape between cohorts classified as having high and low risk scores.





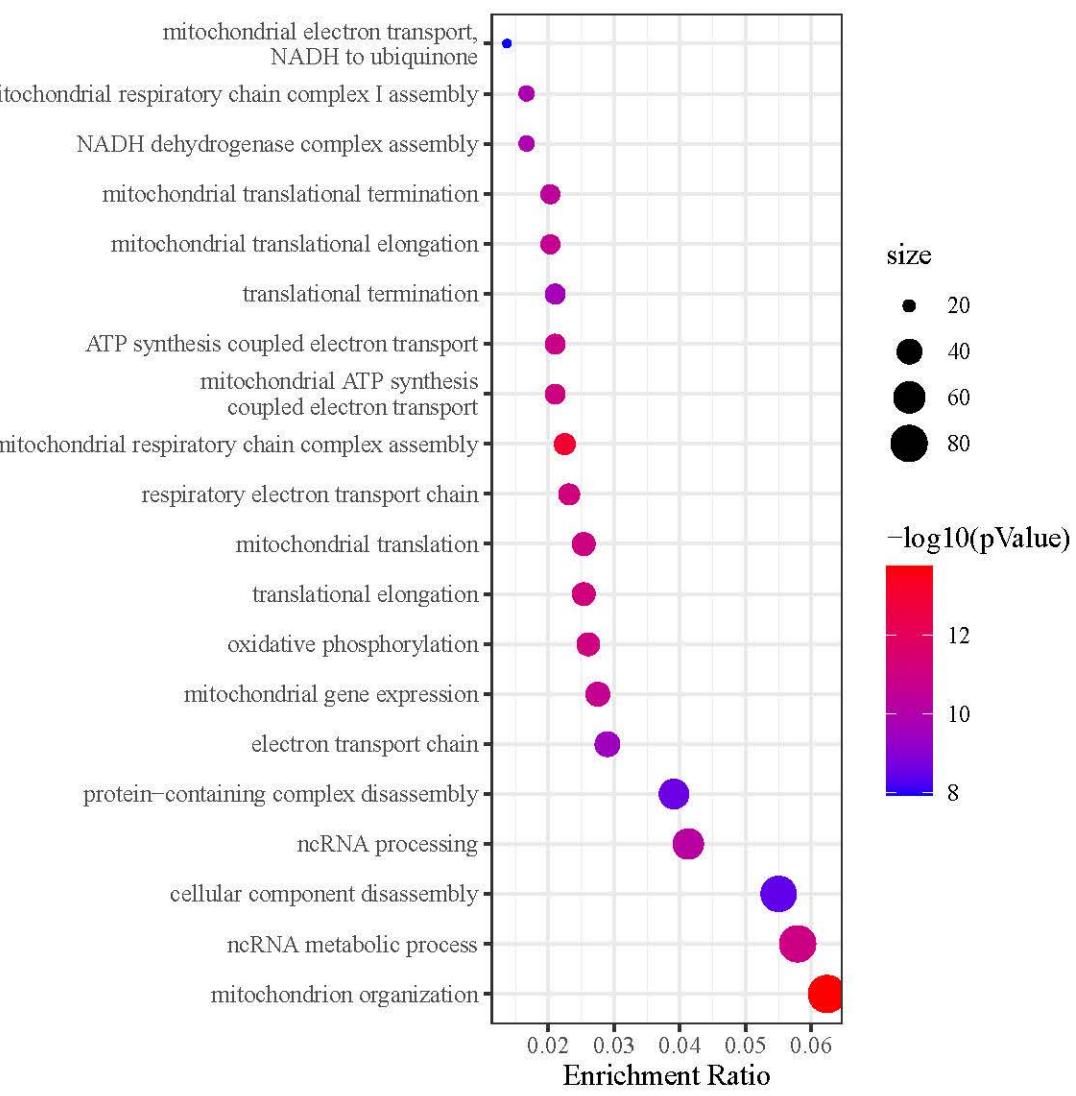
A

All pathway\_KEGG



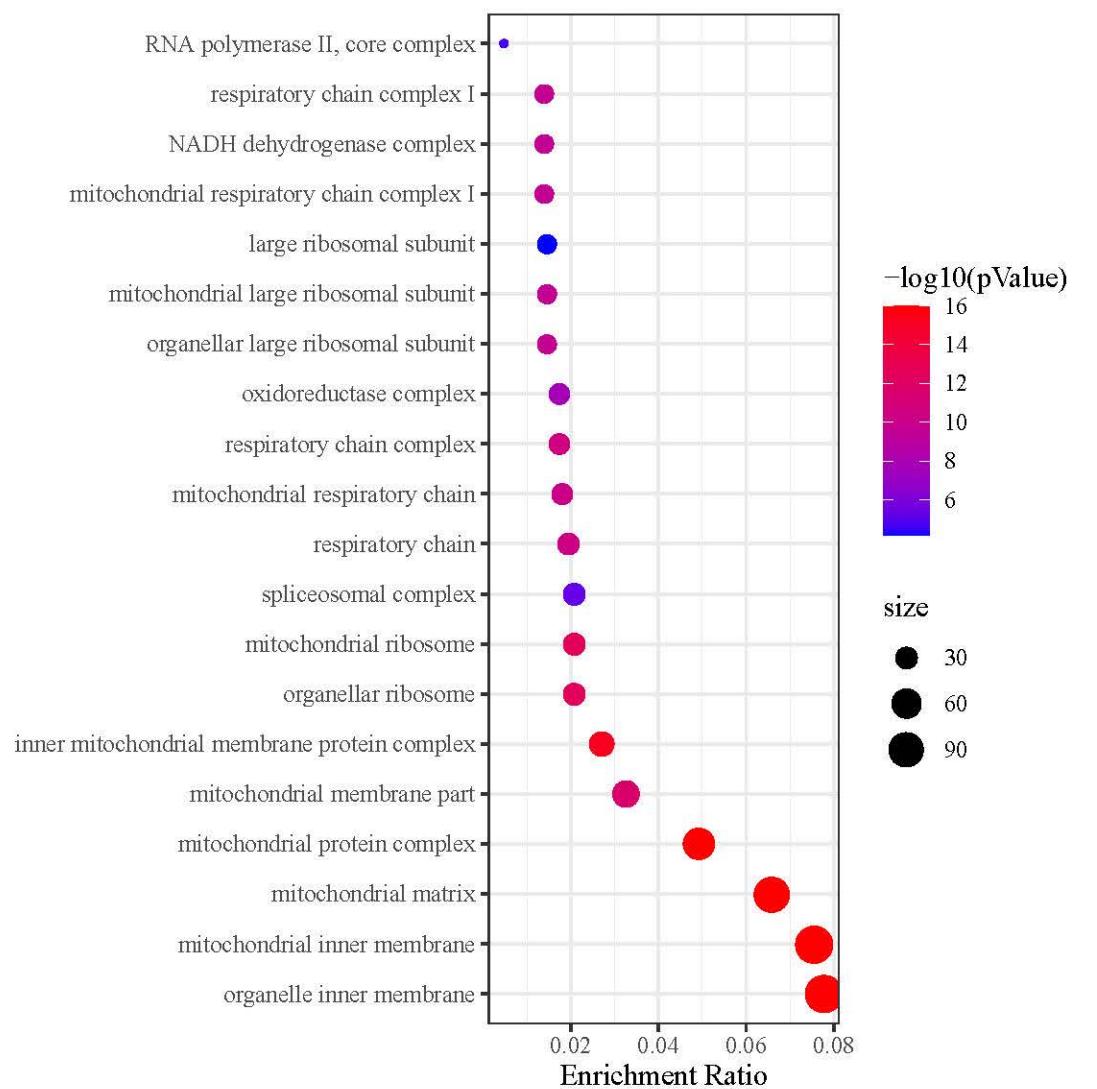
B

Top20 geneontology\_Biological\_Process



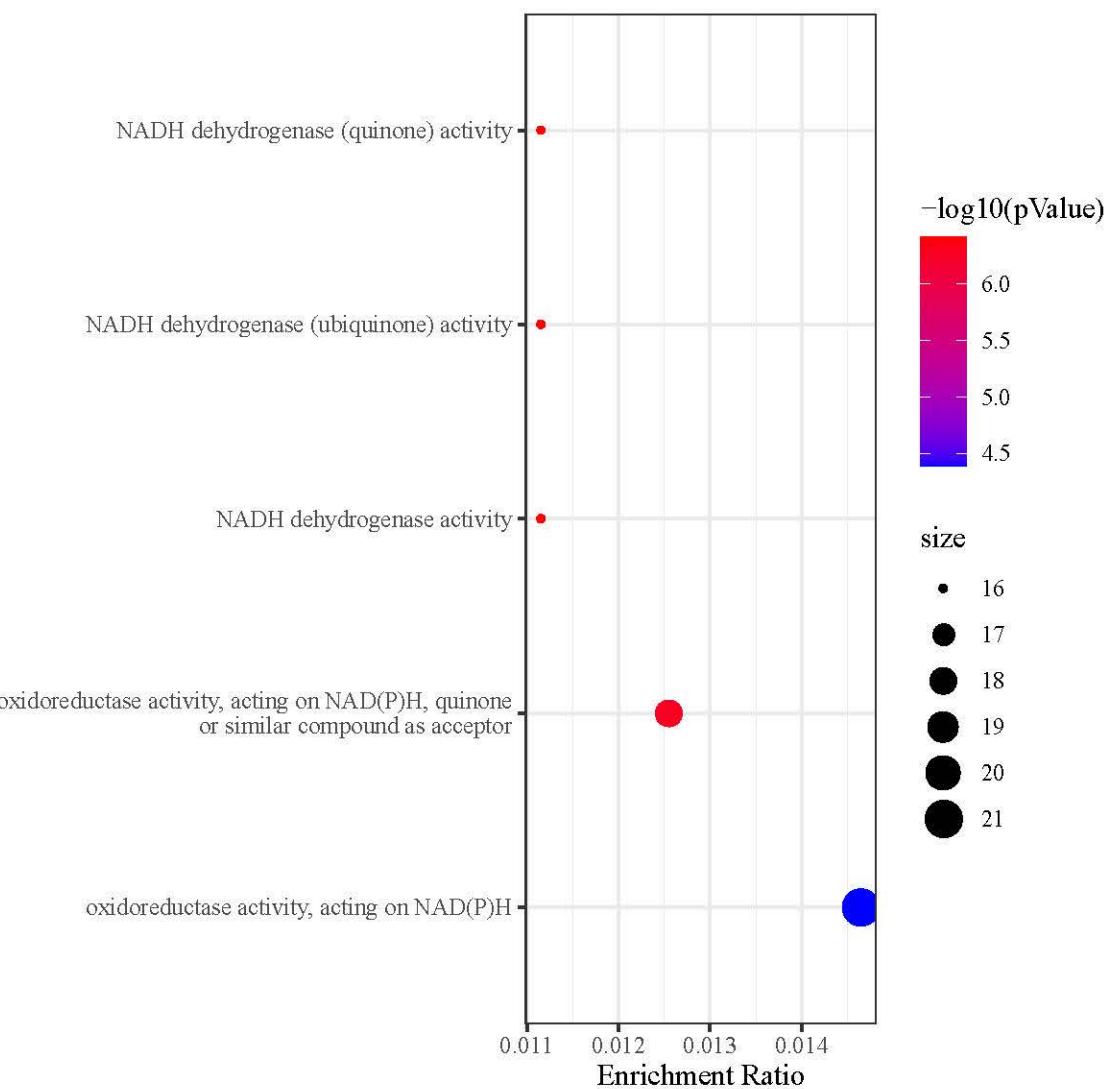
C

Top20 geneontology\_Cellular\_Component

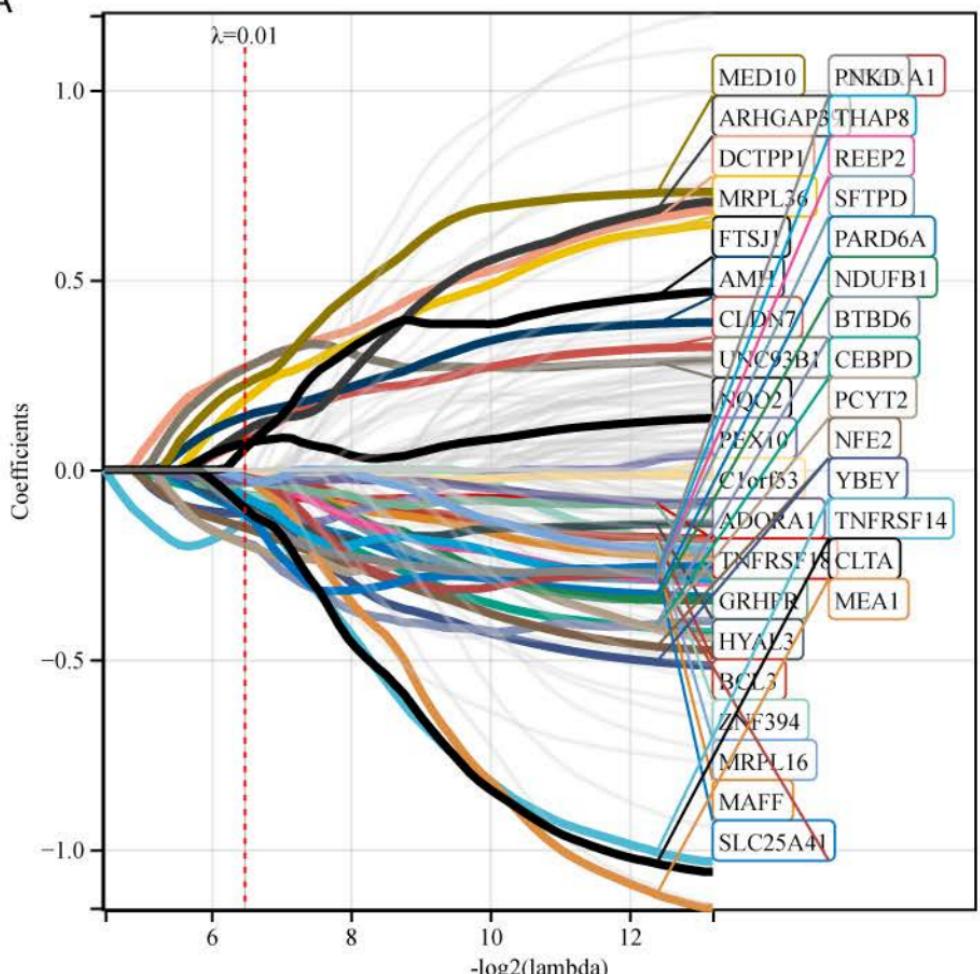


D

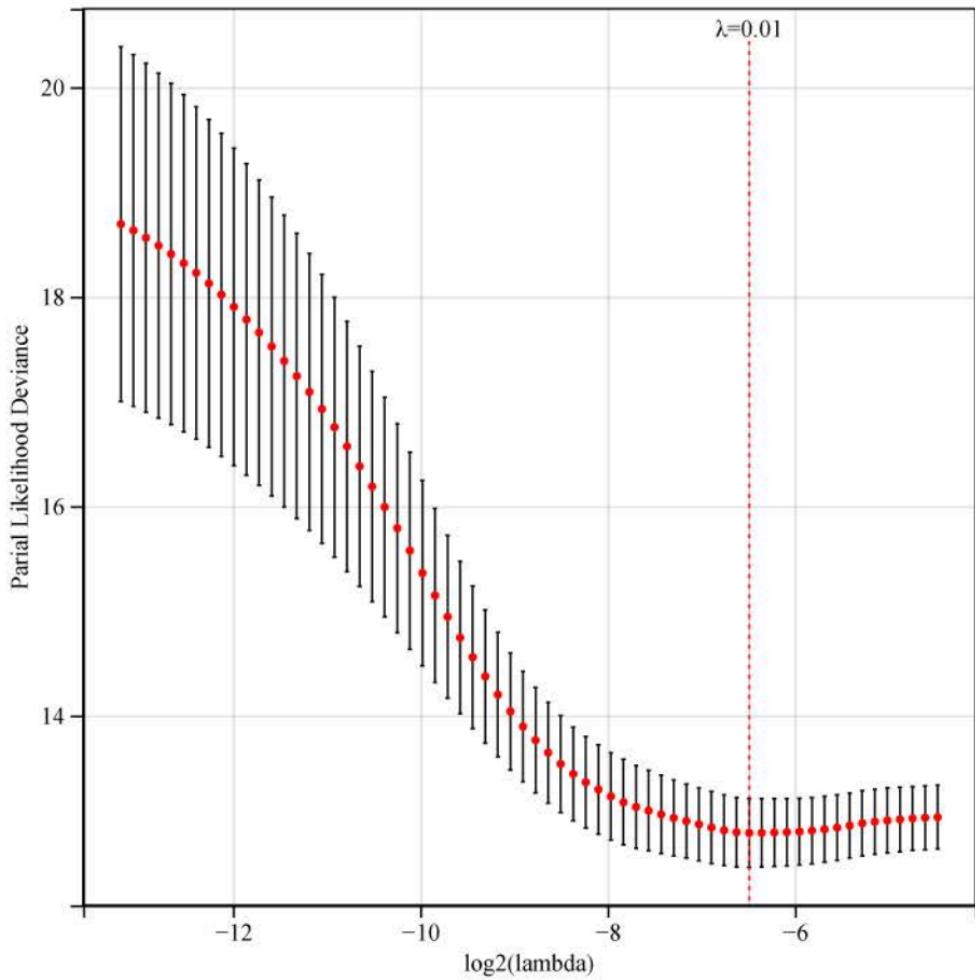
All geneontology\_Molecular\_Function

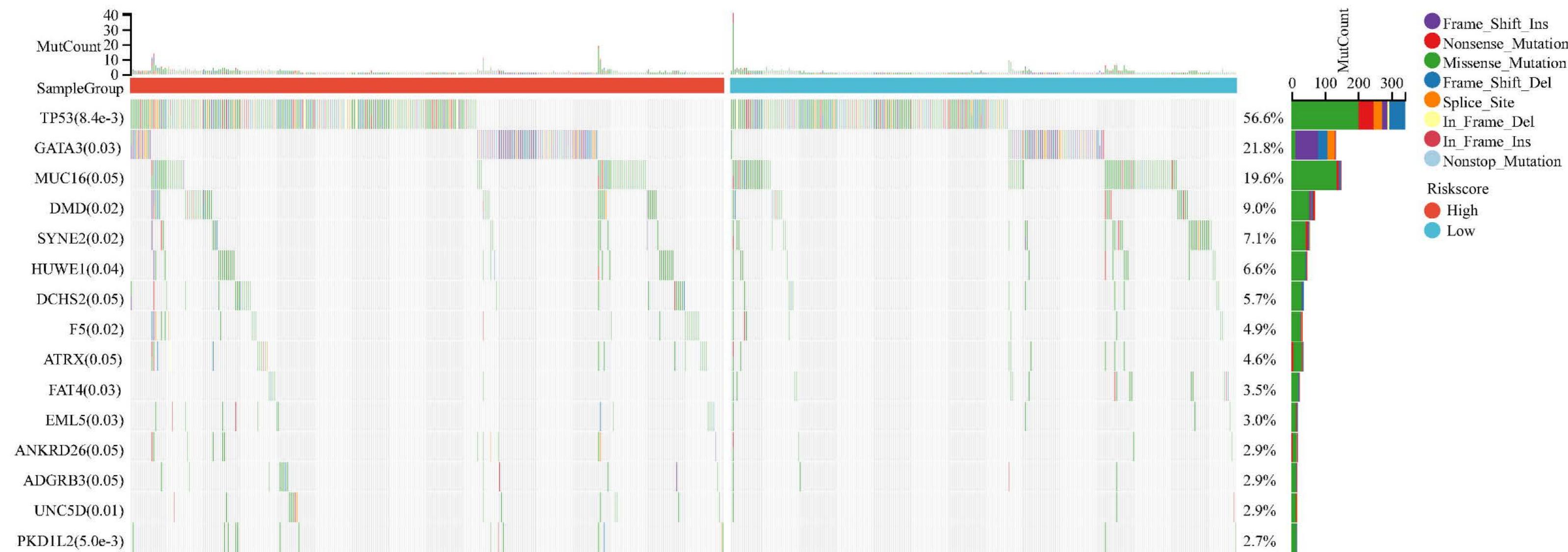


A



B





**Supplementary Table 1. Gene set of pathway associated with immunotherapy.**

Pathway	Genes
IFN- $\gamma$ _signature	TIGIT; CD27; CD8A; PDCD1LG2; LAG3; CD274; CXCR6; CMKLR1; NKG7; CCL5; PSMB10; IDO1; CXCL9; HLA-DQA1; CD276; STAT1; HLA-DRB1; HLA-E
APM_signal	B2M; HLA-A; HLA-B; HLA-C; TAP1; TAP2
Base_excision_repair	FEN1; LIG1; NEIL3; PARP1; PARP2; PCNA; POLE; POLE2; UNG
Cell_cycle	BUB1; BUB1B; CCNA2; CCNB2; CCNE1; CCNE2; CDC20; CDC25A; CDC25C; CDC6; CDK1; CDK2; CDKN2A; DBF4; E2F1; E2F2; ESPL1; MAD2L1; MAD2L2; MCM2; MCM4; MCM6; MCM7; ORC1; ORC6; PCNA; PLK1; SKP2; SMC3; TFDP1; TTK; YWHAB
DNA_replication	DNA2; FEN1; LIG1; MCM2; MCM4; MCM6; MCM7; PCNA; POLA2; POLE; POLE2; PRIM1; PRIM2; RFC2; RFC3; RFC4; RFC5; RNASEH2A; RPA1; RPA3
Fanconi_anemia_pathway	BLM; BRCA1; BRCA2; BRIP1; EME1; ERCC4; FANCA; FANCB; FANCD2; FANCI; PALB2; RAD51; RAD51C; RMI1; RMI2; RPA1; RPA3; TOP3A; UBE2T
Homologous_recombination	BLM; BRCA2; EME1; RAD51; RAD51C; RAD54L; RPA1; RPA3; TOP3A; XRCC2; XRCC3
MicroRNAs_in_cancer	BRCA1; CCNE1; CCNE2; CDC25A; CDC25C; CDCA5; CDKN2A; DNMT1; E2F1; E2F2; EZH2; KIF23; STMN1; TRIM71
Mismatch_repair	EXO1; LIG1; PCNA; RFC2; RFC3; RFC4; RFC5; RPA1; RPA3
Nucleotide_excision_repair	CETN2; ERCC4; LIG1; PCNA; POLE; POLE2; RFC2; RFC3; RFC4; RFC5; RPA1; RPA3
Oocyte_meiosis	AURKA; BUB1; CCNB2; CCNE1; CCNE2; CDC20; CDC25C; CDK1; CDK2; ESPL1; FBXO5; MAD2L1; MAD2L2; PLK1; SGOL1; SMC3; YWHAB
p53_signaling_pathway	CCNB2; CCNE1; CCNE2; CDK1; CDK2; CDKN2A; GTSE1; PPM1D; RFWD2; RRM2
Progesterone-mediated_oocyte_maturation	BUB1; CCNA2; CCNB2; CDC25A; CDC25C; CDK1; CDK2; MAD2L1; MAD2L2; PLK1
Proteasome	IFNG; PSMA4; PSMB2; PSMB4; PSMC4; PSMD4; PSMD7
Pyrimidine_metabolism	CTPS1; DTYMK; POLA2; POLE; POLE2; PRIM1; PRIM2; RRM2; TYMS
Spliceosome	HNRNPM; LSM3; LSM4; LSM5; MAGOHB; PRPF19; SF3B2; SF3B3; SF3B4; SNRNP40; SNRPA1; SNRPC; USP39; WBP11
Systemic_lupus_erythematosus	H2AFV; H2AFZ; HIST1H2AB; HIST1H2AG; HIST1H2AH; HIST1H2AI; HIST1H2AM; HIST1H2BC; HIST1H2BD; HIST1H2BF; HIST1H2BJ; HIST1H2BK; HIST1H2BL; HIST1H2BN; HIST1H2BO; HIST1H3B; HIST1H3D; HIST1H3H; HIST1H4A; HIST1H4B; HIST2H2AB; HIST2H2AC; HIST2H2BE; HIST2H2BF; HIST
Viral_carcinogenesis	CCNA2; CCNE1; CCNE2; CDC20; CDK1; CDK2; CDKN2A; CREB3L4; GTF2E1; HIST1H2BC; HIST1H2BD; HIST1H2BF; HIST1H2BJ; HIST1H2BK; HIST1H2BL; HIST1H2BN; HIST1H2BO; HIST1H4A; HIST1H4B; HIST2H2BE; HIST2H2BF; SKP2; YWHAB

**Supplementary Table 2. Survival related genes filtered by univariate Cox regression analysis.**

Tag	HR	Lower	Upper	Likelihood	logrank	Wald
TNFRSF14	0.672222	0.559277	0.807975	4.10E-05	2.69E-05	2.32E-05
CEBPD	0.778446	0.676163	0.896201	0.000456	0.000482	0.000493
UBXN11	0.724392	0.596718	0.879382	0.001046	0.001097	0.001117
YBEY	0.673315	0.530175	0.855101	0.00123	0.001212	0.001118
DCTPP1	1.506211	1.175919	1.929277	0.001003	0.001249	0.001183
NUDT14	0.714118	0.580386	0.878663	0.001675	0.001533	0.001459
BTBD6	0.62818	0.471538	0.836858	0.002244	0.001617	0.001488
ZNF394	0.509512	0.33535	0.774126	0.002006	0.001679	0.00158
HYI	0.689225	0.546161	0.869765	0.001876	0.001704	0.001716
TNFRSF18	0.835118	0.743931	0.937481	0.002064	0.002193	0.002256
UBXN1	0.710746	0.571336	0.884173	0.003096	0.002271	0.002176
MBLAC1	0.678998	0.528805	0.871848	0.00198	0.002391	0.002404
ARHGEF1	0.68954	0.541998	0.877246	0.002764	0.002628	0.002477
CCDC28B	0.719862	0.580358	0.892899	0.002361	0.002874	0.002784
NFE2	0.73804	0.60347	0.902619	0.001577	0.00295	0.003101
ZDHHC8	0.724549	0.58334	0.89994	0.004084	0.00357	0.003578
ZNF219	0.774279	0.651271	0.92052	0.003743	0.003682	0.003753
MZT2A	0.777944	0.655251	0.92361	0.004439	0.004163	0.004139
TNIP1	0.67787	0.517959	0.887152	0.005991	0.004685	0.004623
FAU	0.714015	0.565699	0.901216	0.006319	0.005102	0.004576
TMEM91	0.732493	0.586883	0.91423	0.005197	0.005815	0.005906
PCYT2	0.705437	0.548869	0.906667	0.007036	0.006811	0.006427
BCL3	0.767409	0.634078	0.928778	0.009183	0.007212	0.006553
PARD6A	0.773599	0.640936	0.933721	0.007669	0.007629	0.007486
MAFF	0.762213	0.623075	0.932423	0.007196	0.008253	0.008282
XRCC1	0.6922	0.526026	0.91087	0.010872	0.009047	0.008628
NDUFB1	0.676832	0.505035	0.90707	0.008461	0.009192	0.008979
NAB2	0.751076	0.605625	0.93146	0.009882	0.009201	0.009148
GSTT2B	0.833589	0.725997	0.957126	0.00787	0.009598	0.009839
PEX10	0.683523	0.512485	0.911643	0.010637	0.009763	0.009611
SFTPД	0.776945	0.641697	0.940699	0.00653	0.009957	0.009697
WDR83	0.706665	0.541839	0.921629	0.011258	0.010472	0.0104
C1orf53	0.793042	0.663718	0.947565	0.01022	0.010624	0.010682
BATF	0.854657	0.757476	0.964307	0.012488	0.01074	0.010769
EFHD2	0.754065	0.607448	0.936069	0.011384	0.010833	0.010502
SNAPC2	0.756093	0.609923	0.937292	0.01202	0.010994	0.010748
USF2	0.694308	0.524502	0.91909	0.012515	0.011241	0.010786
ERF	0.743205	0.59006	0.936098	0.013229	0.01185	0.011707
REEP2	0.813455	0.69175	0.956571	0.010057	0.01237	0.012528
TRMT2A	0.724447	0.563111	0.932007	0.013257	0.012453	0.01215
HINT2	0.733579	0.57448	0.936738	0.014621	0.013439	0.012995
HYAL3	0.74632	0.591714	0.941323	0.012511	0.013476	0.013491
C6orf226	0.775982	0.633982	0.949787	0.014287	0.014185	0.013911
ZBTB17	0.712018	0.54321	0.933286	0.016961	0.014274	0.013892
ZNF358	0.775703	0.633715	0.949504	0.015449	0.01436	0.013805
DDAH2	0.787093	0.649955	0.953167	0.0166	0.014378	0.014245
MEA1	0.67922	0.499541	0.923527	0.014865	0.014402	0.013608

CCDC107	0.761406	0.612135	0.947078	0.015199	0.014521	0.014352
THAP8	0.687883	0.50983	0.92812	0.014998	0.014687	0.014363
NDUFAB1	1.502098	1.083132	2.083125	0.014086	0.01513	0.014744
MIIP	0.744533	0.587107	0.944172	0.014669	0.015449	0.014935
DUS1L	0.745245	0.588143	0.944311	0.016509	0.015642	0.014919
GSTK1	0.717515	0.548028	0.939419	0.018246	0.016137	0.015756
MZT2B	0.830455	0.713758	0.966233	0.017415	0.016322	0.016192
RPS6KA1	0.74243	0.582048	0.947004	0.018566	0.016752	0.016463
MAP1LC3A	0.81095	0.682914	0.96299	0.018048	0.016926	0.016844
ERCC1	0.679499	0.495158	0.932469	0.018602	0.016945	0.016712
DAXX	0.699399	0.521985	0.937114	0.019215	0.017117	0.016618
RFNG	0.741863	0.580531	0.94803	0.019044	0.017434	0.017008
GRHPR	0.684106	0.499725	0.936515	0.019351	0.018236	0.017822
ADORA1	0.874395	0.781763	0.978003	0.016745	0.018505	0.01881
INO80B	0.717562	0.544228	0.946102	0.017998	0.018697	0.018637
SART1	0.777153	0.630198	0.958375	0.023209	0.018806	0.018397
CPSF4	0.679878	0.493287	0.937049	0.020304	0.018982	0.018414
CCDC12	0.731796	0.563867	0.949737	0.019463	0.019103	0.01889
GRINA	1.301298	1.043844	1.622251	0.019164	0.019624	0.019209
EIF3G	0.727012	0.556395	0.949948	0.021092	0.019925	0.019479
RING1	0.710835	0.533481	0.947148	0.022066	0.020043	0.019766
CADM4	0.828271	0.706283	0.971328	0.023699	0.020274	0.020458
MTA1	0.740076	0.57428	0.953738	0.022213	0.02035	0.020019
UNC93B1	1.282872	1.039222	1.583646	0.021298	0.021045	0.020451
CORT	0.562334	0.343189	0.921415	0.01872	0.022261	0.022324
AIMP2	1.441809	1.055215	1.970039	0.020586	0.022432	0.021595
PIGU	1.415374	1.051908	1.904427	0.023812	0.022547	0.021781
ACAA1	0.738121	0.568322	0.958652	0.024626	0.023031	0.022813
CLASRP	0.760096	0.600159	0.962654	0.023598	0.023041	0.022862
SLC25A41	0.575668	0.357214	0.927719	0.017273	0.02318	0.023322
ODF3B	0.829113	0.705168	0.974845	0.022402	0.023444	0.023308
EML3	0.745776	0.577497	0.963091	0.027134	0.025027	0.024563
SIRT7	0.726583	0.548639	0.962241	0.026515	0.026202	0.025846
NDUFB7	0.784412	0.634315	0.970027	0.029437	0.027025	0.025039
SNRNP70	0.759707	0.595405	0.969348	0.026631	0.027283	0.027081
FADS3	0.770675	0.611517	0.971256	0.026801	0.027371	0.027309
AGTRAP	0.767442	0.606604	0.970925	0.029354	0.027728	0.027394
KCTD17	0.780673	0.626225	0.973213	0.028843	0.027848	0.02771
IRF7	0.84495	0.72718	0.981793	0.029826	0.027999	0.027815
MRPL36	1.349434	1.033761	1.761503	0.030655	0.0284	0.027512
ZRSR2	0.701773	0.511705	0.962439	0.031874	0.028715	0.027983
PNKD	0.771933	0.61231	0.973168	0.029047	0.028766	0.028518
ZBTB48	0.72248	0.539623	0.9673	0.030387	0.029302	0.029016
CHMP6	0.766688	0.603452	0.97408	0.034079	0.03015	0.029634
SERTAD1	0.780253	0.623903	0.975785	0.031325	0.030189	0.029644
AKAP8L	0.72908	0.548049	0.969908	0.030992	0.030353	0.030023
RFXANK	0.752898	0.582863	0.972535	0.031447	0.030433	0.029765
PICK1	0.759454	0.591992	0.974286	0.031782	0.030538	0.030393
TECPRI	0.725941	0.543174	0.970205	0.031514	0.030837	0.030436

CLDN7	1.24838	1.021356	1.525865	0.024698	0.031628	0.030287
SNRNP35	0.718713	0.531827	0.971272	0.033219	0.031691	0.031581
RBFA	0.728651	0.545803	0.972755	0.032339	0.031836	0.031766
SMARCD3	0.826578	0.694361	0.983972	0.031483	0.032101	0.032221
USE1	0.735741	0.556017	0.973558	0.033038	0.032253	0.031754
CCDC106	0.825888	0.693287	0.98385	0.034201	0.032258	0.032169
DMAP1	0.716528	0.527609	0.973092	0.033825	0.032875	0.032791
POLR2I	0.731616	0.549265	0.974506	0.032633	0.033425	0.032637
MTFP1	0.776317	0.614614	0.980562	0.034115	0.033671	0.033613
GPX1	0.774997	0.612563	0.980503	0.038799	0.034786	0.033667
MRPL16	0.700146	0.503553	0.973491	0.036884	0.035226	0.034029
AMH	1.149816	1.009178	1.310054	0.041046	0.035642	0.035975
SDHA	1.364184	1.022137	1.820694	0.035787	0.03646	0.034976
MED10	1.399937	1.020894	1.919714	0.036651	0.03747	0.036769
TMEM160	0.81197	0.667351	0.98793	0.036849	0.037906	0.037406
RBM42	0.769873	0.601228	0.985823	0.040879	0.038934	0.038159
PNKP	0.76304	0.590483	0.986023	0.040658	0.039019	0.03868
ARHGAP39	1.245208	1.010813	1.533957	0.039945	0.039411	0.0393
ANAPC11	0.785568	0.624857	0.987613	0.041754	0.039711	0.038763
NAT14	0.825918	0.6882	0.991195	0.040412	0.039963	0.03988
TRIP6	0.855653	0.737157	0.993198	0.045419	0.0402	0.040393
COPS6	0.718184	0.524676	0.983061	0.045551	0.04055	0.03877
SARS2	0.719046	0.523689	0.98728	0.040891	0.041783	0.041439
ZGPAT	0.803744	0.651088	0.992192	0.041302	0.042322	0.042062
FKBP2	0.844914	0.718083	0.994145	0.047356	0.042496	0.042283
ASPSCR1	0.763798	0.588637	0.991082	0.042726	0.042748	0.042625
PPFIA3	1.195257	1.004555	1.422161	0.047307	0.044226	0.044304
PAOX	0.773854	0.602842	0.993377	0.046761	0.044525	0.044208
CROCC	0.813886	0.665633	0.995158	0.045652	0.044681	0.044721
MPST	0.796733	0.638188	0.994666	0.046178	0.045167	0.044728
TRMT1	0.769954	0.596087	0.994536	0.045428	0.045538	0.045293
EMID1	0.868412	0.755649	0.998002	0.043456	0.046607	0.046795
FTSJ1	1.479993	1.007408	2.174272	0.044849	0.046982	0.045763
CNPY3	0.743532	0.555507	0.995198	0.049838	0.04763	0.046335
IFI27L1	0.784307	0.616818	0.997275	0.047468	0.047726	0.047454
NQO2	1.322097	1.002061	1.744344	0.049821	0.048581	0.048322
PKN3	0.811458	0.659237	0.998829	0.047237	0.048693	0.048722
NDUFAF3	0.772568	0.597854	0.998339	0.051353	0.049151	0.048533
PRKCZ	0.769242	0.592092	0.999396	0.052593	0.049661	0.049473
CLTA	0.716551	0.514934	0.997108	0.050968	0.049766	0.048026
ABHD8	0.794914	0.631948	0.999906	0.052056	0.049943	0.049906
CCDC9	0.803602	0.645804	0.999956	0.055765	0.049949	0.049954