

Tumour	Significant Genes (sample count, pvalue)	OncoGene Significance
LGG	MRPS17 , (16, 8.6e-07); EGFR , (11, 9.8e-07); SEC61G , (10, 1.0e-06); * VOPP1 , * LANCL2 , (9, 1.09e-06);	
OV	* ZNF709 , ZNF788, (38, 4.6e-08); * ZNF763 , (34, 5.9e-08); * CYP4F3 , OR10H5, PGLYRP2, OR1I1, WIZ, OR10H2, OR10H3, CYP4F2, * CYP4F11 , * AKAP8L , ILVBL, RASAL3, CYP4F22, OR10H1, SYDE1, EPHX3, * NOTCH3 , CYP4F12, (20, 1.06e-07); * ZNF564 , RTBDN, IER2, ZNF333, RLN3, * RFX1 , OR7A17, ZNF799, PALM3, GADD45GIP1, * MAST1 , ZNF442, ZNF443, ZNF563, * AKAP8 , OR7A10, * BRD4 , * CCDC130 , NFIX, KLF1, * ZNF44 , * CACNA1A , CTD-2192J16.24, * CCDC105 , CALR, RAD23A, OR7C1, CTD-3105H18.16, DCAF15, OR7A5, NACC1, TRMT1, ZNF625, WDR83, SYCE2, ZNF878, DAND5, DNASE2, NDUFB7, TNPO2, PRKACA, GIPC1, ZNF791, PTGER1, LYL1, C19orf67, ASF1B, FARSA, C19orf53, NANOS3, STX10, ASNA1, LPHN1, HOOK2, PODNL1, C19orf43, MRI1, OR7C2, CD97, DDX39A, * SLC1A6 , ZNF20, SAMD1, CC2D1A, AC090427.1, * ZNF433 , ZSWIM4, * ZNF490 , GCDH, CASP14, *RNASEH2A, WDR83OS, hsa-mir-1199, IL27RA, ZNF136, PKN1, EMR2, *TECR, ZNF844, *CLEC17A, C19orf57, FBXW9, DNAJB1, DHPS, EMR3, MAN2B1, * BEST2 , PRDX2, JUNB, AC008686.1, (19, 1.06e-07); SLC44A2, ZNF439, ZNF440, ECSIT, DKFPZ761J1410, * ZNF627 , ZNF69, * ELOF1 , AC011475.1, PRKCSH, CARM1, LDLR, SPC24, YIPF2, CCDC159, C19orf52, SWSAP1, KANK2, TMEM205, ZNF653, * RAB3D , * RGL3 , ZNF491, ZNF441, C19orf38, ZNF823, CCDC151, AC05, * SMARCA4 , * TSPAN16 , DOCK6, C19orf80, TMED1, CNN1, QTRT1, ELAVL3, IFL3, CCNE1, * C19orf12 , PLEKH1, (17, 1.12e-07); S1PR5, OCEL1, P2RY11, * EPOR , ATG4D, ICAM5, TMEM221, SMIM7, SLC27A1, RAB8A, * FAM129C , DDA1, KRI1, CTC-429P9.4, USE1, SIN3B, RAVER1, MED26, DNMT1, ANO8, NWD1, EPS15L1, S1PR2, CHERP, MRPL4, * GTPBP3 , F2RL3, MVB12A, TMEM38A, CIB3, PPAN-P2RY11, BST2, CTD-2521M24.10, AP1M2, ABHD8, NXNL1, C19orf44, AP1M1, FAM32A, PLVAP, USHBP1, SLC35E1, ZGLP1, EIF3G, OR10H4, CDKN2D, NR2F6, AC010646.3, ICAM3, ICAM1, PPAN, CALR3, PGLS, FDX1L, HSH2D, CDC37, KLF2, TPM4, MRPL34, TYK2, * ANGPTL6 , KEAP1, ICAM4, * PDE4A , BABAM1, ANKLE1, * OC90 , * URI1 , * DNM2 , POP4, VSTM2B, (16, 1.15e-07); * CPAMD8 , * MYO9B , * C19orf66 , HAUS8, RDH8, (15, 3.9e-05); UQCRFS1, * COL5A3 , * ZNF536 , * WDR88 , CEBPA, LRP3, AC020952.1, * SLC7A10 , HOXC4, (14, 4.5e-4)	0.047
ESCA	FGF4, * FGF3 , * PPFIA1 , FADD, * CTTN , AP001888.1, * ANO1 , (16, 2.2e-07); FGF19, ORAOV1, * CCND1 , (15, 2.2e-07);	0.002
HNSC	* PPFIA1 , FADD, * CTTN , * ANO1 , (29, 0.003); FGF4, * FGF3 , FGF19, ORAOV1, AP001888.1, * CCND1 , (28, 0.004);	1.90E-05
UCEC	EFNA3, (14, 7.1e-08);	2.10E-05
SKCM	* NARS2 , (33, 6.0e-08); UCP2, P2RY2, INTS4, AQP11, * P4HA3 , COA4, TPBGL, * NEU3 , MRPL48, RPS3, POLD3, MOGAT2, * GAB2 , * C2CD3 , * UVRAG , SPCS2, * B3GNT6 , * FAM168A , THRSP, OR2AT4, DNAJB13, SERPINH1, OMP, * LRRC32 , GDPD5, * GDPD4 , NDUFC2, * ARHGEF17 , KCNE3, AAMDC, * DGAT2 , * ACER3 , XRA1, KLHL35, * PAK1 , PPME1, P2RY6, * SLC02B1 , DKFPZ434E1119, * C11orf30 , * RELT , CLNS1A, * CAPN5 , RSF1, WNT11, * RP11-111M22.2 , CHRD12, * PGM2L1 , USP35, RAB6A, TSKU, UCP3, LIPT2, KCTD14, * MAP6 , KCTD21, ALG8, * PAAF1 , ARRB1, PLEKHB1, NDUFC2-KCTD14, PRKRIR, (32, 6.4e-08); * RNF169 , (31, 6.7e-08); * RAB38 , CCDC81, PRSS23, * MYO7A , * TENM4 , * ME3 , * GRM5 , * TMEM135 , FZD4, * TYR , (30, 7.0e-08); * SYTL2 , TMEM126B, CCDC89, CREBFZ, TMEM126A, (29, 7.3e-08); PICALM, * DLG2 , AP000974.1, ANKRD42, * CCDC90B , EED, C11orf82, * PCF11 , FAM181B, * C11orf73 , * PRCP , RAB30, CCDC83, (28, 7.6e-08); * CTSC , (25, 8.5e-08); TNFAIP8L3, HDC, AP4E1, MYEF2, MYO5C, CTXN2, CEP152, TMOD3, * SEMA6D , COPS2, FKSG62, DUT, LYSMD2, TRPM7, GALK2, CYP19A1, BCL2L10, FAM227B, SLC27A2, SPPL2A, * SLC12A1 , TMOD2, MAPK6, FGF7, GABPB1, GLDN, ATP8B4, USP8, SECISBP2L, USP50, DTWD1, SLC24A5, EID1, GNB5, DMXL2, * LEO1 , SHC4, SCG3, FBN1, (24, 9.0e-05);	
GBM	ZNF479, (50, 1.1ee-08); ZNF716, (48, 1.9ee-08); POM121L12, (47, 2.2ee-08); * COBL , * GRB10 , DDC, * IKZF1 , (43, 3.8e-08); LRIG3, * RP11-362K2.2 , * FIGNL1 , C7orf72, VWC2, (42, 4.2e-08); * ZPBP , (41, 4.5e-08); PKD1L1, C7orf69, C7orf57, C7orf65, UPP1, * SUN3 , HUS1, AC004899.1, * SRGAP1 , RP11-272B17.2, * SLC16A7 , C12orf66, XPOT, * CAND1 , AC025262.1, * DPY19L2 , AVPR1A, TMEM5, * TBC1D30 , GNS, * TBK1 , C12orf56, * RASSF3 , (38, 5.7e-08); * TSPAN31 , CDK4, * TNS3 , * ABCA13 , AC090673.2, LLPH, TMBIM4, * MON2 , * HELB , * WIF1 , * GRIP1 , * USP15 , IRAK3, * LEMD3 , HMGA2, * MSRB3 , C12orf61, RP11-366L20.2, * FAM19A2 , (37, 6.1e-08); CTDSP2, RP11-571M6.15, TSFM, METTL21B, METTL1, * CYP27B1 , IFNG, (36, 6.4e-08); XRCC6BP1, AC006455.1, AC073188.1, IL22, * PPM1H , * IL26 , * DYRK2 , MDM1, (35, 6.8e-08); * AVIL , * AGAP2 , OS9, AGAP2-AS1, IGFBP3, RAMP3, ADCY1, AC096582.1, * IGFBP1 , ZNF727, ZNF736, ZNF273, ZNF138, ZNF107, * ZNF680 , ZNF679, ZNF117, ERV3-1, MDM2, (34, 7.2e-08); AC011294.3, ZNF92, * AC124890.1 , (33, 7.6e-08); * RAP1B , (32, 8.e-08); * MARCH9 , (31, 8.4e-08); * AQP1 , * SLC35E3 , (30, 8.7e-08); * NUP107 , (29, 8.6e-07); * NUCD3 , NACAD, CCM2, MYO1G, TBRG4, PPIA, YKT6, POLD2, ZMIZ2, * GCK , TMED4, MYL7, DDX56, AEBP1, NPC1L1, POLM, OGDH, * CAMK2B , (28, 4.1e-05);	
LUSC	HTR3C, DVL3, MAP6D1, HTR3E, ABCC5, * KLHL24 , ECE2, ABCF3, PARL, ALG3, YEATS2, VWA5B2, AP2M1, HTR3D, CAMK2N2, (30, 6.3e-08); CHRD, POLR2H, * ATP11B , RP11-433C9.2, KLHL6, LAMP3, FAM131A, * MCCC1 , EIF4G1, THPO, CLCN2, * EIF2B5 , MAGEF1, EPHB3, * PSMD2 , * DCUN1D1 , (29, 6.5e-08); * B3GNT5 , EHHADH, MAP3K13, * C3orf70 , * MCF2L2 , * CCDC39 , SOX2, * VPS8 , TTC14, * TMEM41A , * GPR160 , * TNIK , RPL22L1, PLD1, TNFSF10, TMEM212, PP13439, * EIF5A2 , GHSR, SLC2A2, AC092964.1, * NCEH1 , AC007919.2, AC092964.2, * FNDC3B , (28, 6.8e-08); DNAJC19, * FXR1 , MRPL47, ACTL6A, * PIK3CA , * SAMD7 , * ZMAT3 , USP13, * MECOM , SI, SKIL, * ECT2 , * NDUFB5 , BCHE, SEC62, * KCNMB2 , * CLDN11 , * SLITRK3 , LRRC31, * MYNN , PRKCI, * KCNMB3 , * SPATA16 , GNB4, * TBL1XR1 , LRRIQ4, ACTRT3, LRRC34, ZNF639, * SLC7A14 , * PHC3 , MFN1, (27, 7.1e-08); * SERPINI1 , * SERPINI2 , * PEX5L , * WDR49 , * NLGN1 , * ZBBX , * GOLIM4 , * NAALADL2 , * PDCD10 , (26, 7.4e-08); * LIPH , TRA2B, C3orf65, * IGF2BP2 , SENP2, (25, 7.6e-08); ETV5, RTP1, RFC4, AHSG, EIF4A2, DGKG, HRG, KNG1, ADIPOQ, TBCCD1, RPL39L, CRYGS, DNAJB11, FETUB, ST6GAL1, ARL14, (21, 5.2e-05);	
BRCA	ZFP41, (8, 3.4e-07);	