

Supplementary Table 3. Biological functions of the genes with altered expression in MCF-7/ADR cells as determined by Ingenuity Pathway analysis*

Category	Function	Function annotation†	P‡	Gene symbol	No. of molecules
Cancer	Neoplasia	Neoplasia	3.95×10^{-26}	<i>ABAT, ABCB1, ADA, AGR2, AKAP12, AKR1C2, AKT3, ANK3, ANXA1, ANXA9, AREG, AXL, B2M, BEX1, BMP7, BRIP1, BST2, C6ORF211, CA2, CALD1, CCNE2, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), CLDN3, CLGN, COL1A1, COL4A1, COL4A2, COL6A1, COL6A2, CREBBP, CRIP1, CTGF, CXCL12, CYP1B1, CYR61, DNAJB4, DNAJC15, ECM1, EGFR, EML1 (includes EG:2009), EMP1, EPCAM, ESR1, EXT1, FABP5, FBN2 (includes EG:2201), FHL2, FN1, FSCN1, FYN, GATA3, GDF15, GFRA1, GJA1, GLUL, GPX1, GPX3, GREB1, GSTP1, HEY2, HIST1H2BD, HIST1H4H (includes EG:8365), HIST2H2AA3, HIST2H2BE, HMGA1, HMGA2, HNMT, ICAM1, ID2, IGF2BP3, IGFBP3, IGFBP5, IL7R, INSIG1, IRS1, ISOC1, KLF4, KRT7, KRT19, LOX, MAFB, MCAM, MGP, MLPH, MMP1 (includes EG:4312), MSLN, MSMB, MSX2, MT1E, MUC1, MYB (includes EG:4602), MYL9 (includes EG:10398), NNMT, NOV, NPY1R, NUAK1, OLFM1, PDGFC, PDZK1, PIK3R1, PIK3R3, PPP1R3D, PRAME (includes EG:23532), PSAT1, RB1, RET, RRAS, S100A14, S100P, SCUBE2, SDC2, SEMA3C, SERPINA3, SERPINE1, SERPINE2, SERTAD2, SGK3, SLC12A8, SLC16A3, SLC19A2, SLC24A3, SLC39A6, SOCS2, SPARC, SRPX, STXBP6, TACSTD2, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, TM4SF1, TPD52L1, TUBB2A, VAV3, VIM, WNT5A, WWOX, ZYX</i>	144
Cancer	Tumorigenesis	Tumorigenesis	8.79×10^{-26}	<i>ABAT, ABCB1, ADA, AGR2, AKAP12, AKR1C2, AKT3, ANK3, ANXA1, ANXA9, AREG, AXL, B2M, BEX1, BMP7, BRIP1, BST2, C6ORF211, CA2, CALD1, CCNE2, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), CLDN3, CLGN, COL1A1, COL4A1, COL4A2, COL6A1, COL6A2, CREBBP, CRIP1, CTGF, CXADR, CXCL12, CYP1B1, CYR61, DNAJB4, DNAJC15, ECM1, EGFR, EML1 (includes EG:2009), EMP1, EPCAM, ESR1, EXT1, FABP5, FBN2 (includes EG:2201), FHL2, FN1, FSCN1, FYN, GATA3, GDF15, GFRA1, GJA1, GLUL, GPX1, GPX3, GREB1, GSTP1, HEY2, HIST1H2BD, HIST1H4H (includes EG:8365), HIST2H2AA3, HIST2H2BE, HMGA1, HMGA2, HNMT, ICAM1, ID2, IGF2BP3, IGFBP3, IGFBP5, IL7R, INSIG1, IRS1, ISOC1, KLF4, KRT7, KRT19, LOX, MAFB, MCAM, MGP, MLPH, MMP1 (includes EG:4312), MSLN, MSMB, MSX2, MT1E, MUC1, MYB (includes EG:4602), MYL9 (includes EG:10398), NNMT, NOV, NPY1R, NUAK1, OLFM1, PDGFC, PDZK1, PIK3R1, PIK3R3, PPP1R3D, PRAME (includes EG:23532), PSAT1, RB1, RET, RRAS, S100A14, S100P, SCUBE2, SDC2, SEMA3C, SERPINA3, SERPINE1, SERPINE2,</i>	145

				<i>SERTAD2, SGK3, SLC12A8, SLC16A3, SLC19A2, SLC24A3, SLC39A6, SOCS2, SPARC, SRPX, STXBP6, TACSTD2, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, TM4SF1, TPD52L1, TUBB2A, VAV3, VIM, WNT5A, WWOX, ZYX</i>	
Cancer	Tumorigenesis	Tumorigenesis of organ	3.07×10^{-3}	<i>AREG, CDKN2A</i>	2
Cancer	Cancer	Cancer	2.23×10^{-25}	<i>ABAT, ABCB1, ADA, AGR2, AKAP12, AKR1C2, AKT3, ANK3, ANXA1, ANXA9, AREG, AXL, B2M, BEX1, BMP7, BRIP1, BST2, C6ORF211, CA2, CALD1, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), CLDN3, CLGN, COL1A1, COL4A1, COL4A2, COL6A1, COL6A2, CREBBP, CRIP1, CTGF, CXCL12, CYP1B1, CYR61, DNAJB4, DNAJC15, ECM1, EGFR, EMP1, EPCAM, ESR1, EXT1, FABP5, FBN2 (includes EG:2201), FHL2, FN1, FSCN1, FYN, GATA3, GDF15, GFRA1, GJA1, GLUL, GPX1, GPX3, GREB1, GSTP1, HEY2, HIST1H2BD, HIST1H4H (includes EG:8365), HIST2H2AA3, HIST2H2BE, HMGA1, HMGA2, HNMT, ICAM1, ID2, IGF2BP3, IGFBP3, IGFBP5, IL7R, INSIG1, IRS1, KLF4, KRT7, KRT19, LOX, MAFB, MCAM, MGP, MLPH, MMP1 (includes EG:4312), MSLN, MSMB, MSX2, MT1E, MUC1, MYB (includes EG:4602), MYL9 (includes EG:10398), NNMT, NPY1R, NUAK1, OLFM1, PDGFC, PDZK1, PIK3R1, PIK3R3, PPP1R3D, PRAME (includes EG:23532), PSAT1, RB1, RET, RRAS, S100A14, S100P, SCUBE2, SDC2, SEMA3C, SERPINA3, SERPINE1, SERPINE2, SERTAD2, SGK3, SLC12A8, SLC16A3, SLC19A2, SLC24A3, SLC39A6, SOCS2, SPARC, SRPX, STXBP6, TACSTD2, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, TM4SF1, TPD52L1, TUBB2A, VAV3, VIM, WNT5A, WWOX, ZYX</i>	140

Cancer	Primary tumor	Primary tumor	7.41×10^{-21}	<i>ABAT, ABCB1, ADA, AGR2, AKAP12, AKT3, ANXA1, ANXA9, AREG, AXL, B2M, BEX1, BMP7, C6ORF211, CA2, CD24, CDH1, CDH2, CDKN2A, CEACAM6</i> (includes EG:4680), <i>CLDN3, COL1A1, COL4A1, COL4A2, CREBBP, CTGF, CYP1B1, CYR61, ECM1, EGFR, EPCAM, ESR1, EXT1, FABP5, FBN2</i> (includes EG:2201), <i>FHL2, FN1, FSCN1, FYN, GATA3, GDF15, GFRA1, GJA1, GPX1, GPX3, GSTP1, HIST2H2AA3, HMGA1, HMGA2, HNMT, ID2, IGFBP3, IGFBP5, INSIG1, ISOC1, KLF4, KRT7, KRT19, LOX, MAFB, MCAM, MGP, MSLN, MSMB, MSX2, MT1E, MUC1, MYL9</i> (includes EG:10398), <i>NNMT, NOV, NPY1R, OLFM1, PDGFC, PDZK1, PIK3R1, PIK3R3, PPP1R3D, PRAME</i> (includes EG:23532), <i>PSAT1, RB1, RET, RRAS, SCUBE2, SDC2, SEMA3C, SERPINE2, SGK3, SLC19A2, SLC24A3, SLC39A6, SOCS2, SPARC, STXBP6, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, TM4SF1, TUBB2A, VAV3, VIM, WNT5A, WWOX, ZYX</i>	105
Cancer	Tumor	Tumor	9.06×10^{-21}	<i>ABAT, ABCB1, ADA, AGR2, AKAP12, AKT3, ANXA1, ANXA9, AREG, AXL, B2M, BEX1, BMP7, C6ORF211, CA2, CD24, CDH1, CDH2, CDKN2A, CEACAM6</i> (includes EG:4680), <i>CLDN3, COL1A1, COL4A1, COL4A2, CREBBP, CTGF, CYP1B1, CYR61, ECM1, EGFR, EPCAM, ESR1, EXT1, FABP5, FBN2</i> (includes EG:2201), <i>FHL2, FN1, FSCN1, FYN, GATA3, GDF15, GFRA1, GJA1, GPX1, GPX3, GSTP1, HIST2H2AA3, HMGA1, HMGA2, HNMT, ID2, IGFBP3, IGFBP5, IL7R, INSIG1, ISOC1, KLF4, KRT7, KRT19, LOX, MAFB, MCAM, MGP, MSLN, MSMB, MSX2, MT1E, MUC1, MYL9</i> (includes EG:10398), <i>NNMT, NOV, NPY1R, OLFM1, PDGFC, PDZK1, PIK3R1, PIK3R3, PPP1R3D, PRAME</i> (includes EG:23532), <i>PSAT1, RB1, RET, RRAS, SCUBE2, SDC2, SEMA3C, SERPINE2, SGK3, SLC19A2, SLC24A3, SLC39A6, SOCS2, SPARC, STXBP6, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, TM4SF1, TUBB2A, VAV3, VIM, WNT5A, WWOX, ZYX</i>	106
Cancer	Malignant tumor	Malignant tumor	3.21×10^{-20}	<i>ABAT, ABCB1, ADA, AGR2, AKAP12, AKT3, ANXA1, ANXA9, AREG, AXL, B2M, BEX1, BMP7, C6ORF211, CA2, CD24, CDH1, CDH2, CDKN2A, CEACAM6</i> (includes EG:4680), <i>CLDN3, COL1A1, COL4A1, COL4A2, CREBBP, CYP1B1, CYR61, ECM1, EGFR, EPCAM, ESR1, EXT1, FABP5, FBN2</i> (includes EG:2201), <i>FHL2, FN1, FSCN1, FYN, GATA3, GDF15, GFRA1, GJA1, GPX1, GPX3, GSTP1, HIST2H2AA3, HMGA1, HMGA2, HNMT, ID2, IGFBP3, IGFBP5, INSIG1, KLF4, KRT7, KRT19, LOX, MAFB, MCAM, MGP, MSLN, MSMB, MSX2, MT1E, MUC1, MYL9</i> (includes EG:10398), <i>NNMT, NOV, NPY1R, OLFM1, PDGFC, PDZK1, PIK3R1, PIK3R3, PPP1R3D, PRAME</i> (includes EG:23532), <i>PSAT1, RB1, RET, RRAS, SCUBE2, SDC2, SEMA3C, SERPINE2, SGK3, SLC19A2, SLC24A3, SLC39A6, SOCS2, SPARC, STXBP6, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, TM4SF1, TUBB2A, VAV3, VIM, WNT5A, WWOX, ZYX</i>	102

Cancer	Carcinoma	Carcinoma	4.48×10^{-18}	<i>ABAT, ABCB1, AGR2, AKT3, ANXA1, ANXA9, AREG, AXL, B2M, BEX1, BMP7, C6ORF211, CA2, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), CLDN3, COL1A1, CREBBP, CYP1B1, CYR61, ECM1, EGFR, EPCAM, ESR1, FABP5, FBN2 (includes EG:2201), FHL2, FN1, FSCN1, FYN, GATA3, GDF15, GFRA1, GJA1, GPX3, GSTP1, HIST2H2AA3, HMGA2, IGFBP3, IGFBP5, INSIG1, KRT7, KRT19, LOX, MAFB, MCAM, MGP, MSLN, MSMB, MSX2, MT1E, MUC1, MYL9 (includes EG:10398), NNMT, NPY1R, PDZK1, PIK3R1, PRAME (includes EG:23532), PSAT1, RB1, RET, RRAS, SCUBE2, SEMA3C, SERPINE2, SLC19A2, SLC39A6, SOCS2, TFF1, TFF3, TFPI2, TGFB1, TM4SF1, TUBB2A, VAV3, VIM, WNT5A, WWOX</i>	81
Cancer	Ovarian cancer	Ovarian cancer	1.16×10^{-16}	<i>ABCB1, ADA, AGR2, AKT3, BMP7, CD24, CDH1, CDKN2A, CEACAM6 (includes EG:4680), CLDN3, COL4A1, EGFR, EPCAM, ESR1, FN1, FYN, GPX3, HIST2H2AA3, HMGA1, HMGA2, IGF2BP3, MCAM, MGP, MMP1 (includes EG:4312), MSLN, MUC1, PRAME (includes EG:23532), PSAT1, RET, RRAS, S100P, SEMA3C, SLC16A3, TACSTD2, TFF3, TFPI2, TUBB2A, VAV3</i>	38
Cancer	Colorectal cancer	Colorectal cancer	5.18×10^{-15}	<i>AKR1C2, AKT3, ANK3, BEX1, BMP7, BST2, CA2, CALD1, CDH2, CDKN2A, CLDN3, CLGN, COL4A1, COL4A2, COL6A1, COL6A2, CRIP1, CTGF, CXCL12, CYP1B1, EGFR, EMP1, EPCAM, FBN2 (includes EG:2201), FN1, FSCN1, FYN, GDF15, GJA1, GLUL, GREB1, HIST1H2BD, HIST1H4H (includes EG:8365), ICAM1, ID2, IGFBP5, INSIG1, IRS1, MAFB, MCAM, MLPH, NUAK1, RET, SERPINA3, SERPINE1, SERTAD2, SLC12A8, SLC16A3, SRPX, STXBP6, TFF3, TM4SF1, TPD52L1, TUBB2A</i>	54
Cancer	Breast carcinoma	Breast carcinoma	4.51×10^{-11}	<i>ABAT, ABCB1, AGR2, ANXA1, ANXA9, AREG, BEX1, BMP7, C6ORF211, CDH2, CEACAM6 (includes EG:4680), ECM1, EGFR, ESR1, FBN2 (includes EG:2201), FSCN1, GATA3, GFRA1, IGFBP3, MT1E, MUC1, NPY1R, PRAME (includes EG:23532), RB1, SCUBE2, SLC19A2, SLC39A6, TFF1, TUBB2A</i>	29
Cancer	Mammary tumor	Mammary tumor	1.15×10^{-10}	<i>ABAT, ABCB1, AGR2, AKT3, ANXA1, ANXA9, AREG, BEX1, BMP7, C6ORF211, CDH2, CEACAM6 (includes EG:4680), ECM1, EGFR, ESR1, FBN2 (includes EG:2201), FSCN1, GATA3, GFRA1, IGFBP3, MT1E, MUC1, NPY1R, PRAME (includes EG:23532), RB1, RET, SCUBE2, SLC19A2, SLC39A6, TFF1, TUBB2A</i>	31
Cancer	Breast cancer	Breast cancer	1.58×10^{-10}	<i>ABAT, ABCB1, ADA, AGR2, AKR1C2, AKT3, ANXA1, ANXA9, AREG, BEX1, BMP7, BRIP1, C6ORF211, CDH2, CDKN2A, CEACAM6 (includes EG:4680), CLDN3, CLGN, COL1A1, CYR61, ECM1, EGFR, ESR1, FBN2 (includes EG:2201), FSCN1, FYN, GATA3, GFRA1, HEY2, HIST2H2BE, HMGA1, IGFBP3, KLF4, MT1E, MUC1, NPY1R, PRAME (includes EG:23532), RB1, RET, S100A14, SCUBE2, SGK3, SLC19A2, SLC39A6, SPARC, TACSTD2, TFF1, TGFB1, TUBB2A, WWOX</i>	50

Cancer	Pancreatic tumor	Pancreatic tumor	3.40×10^{-9}	<i>AXL, CDKN2A, CREBBP, CYR61, EGFR, FN1, IGFBP3, IGFBP5, KLF4, KRT7, KRT19, MSLN, MSX2, PIK3R1, RET, SERPINE2, TFF1, TUBB2A</i>	18
Cancer	Digestive organ tumor	Digestive organ tumor	3.56×10^{-9}	<i>ABCB1, AKT3, ANXA1, AXL, CA2, CDH1, CDKN2A, COL1A1, CREBBP, CYR61, EGFR, FN1, FSCN1, FYN, GDF15, GJA1, GSTP1, IGFBP3, IGFBP5, INSIG1, KLF4, KRT7, KRT19, MSLN, MSX2, NNMT, PIK3R1, RB1, RET, SERPINE2, TFF1, TM4SF1, TUBB2A, VIM, WWOX</i>	35
Cancer	Genital tumor	Genital tumor	1.36×10^{-8}	<i>ABAT, ABCB1, ADA, AKT3, CD24, CLDN3, EGFR, EPCAM, ESR1, FHL2, FYN, GATA3, GDF15, GPX3, HIST2H2AA3, HMGA2, IGFBP3, KRT7, MCAM, MGP, MSLN, MSMB, MUC1, MYL9 (includes EG:10398), PSAT1, RB1, RET, RRAS, SEMA3C, TFF3, TFPI2, TUBB2A, VAV3</i>	33
Cancer	Pancreatic cancer	Pancreatic cancer	6.91×10^{-8}	<i>AXL, CDH2, CDKN2A, CREBBP, CYR61, EGFR, FBN2 (includes EG:2201), FN1, FYN, IGFBP3, IGFBP5, KLF4, KRT7, KRT19, MSLN, MSX2, PIK3R1, RET, SERPINE2, TFF1, TUBB2A</i>	21
Cancer	Gonadal tumor	Gonadal tumor	7.95×10^{-7}	<i>ABCB1, ADA, CD24, CLDN3, EGFR, EPCAM, GPX3, HIST2H2AA3, HMGA2, MCAM, MGP, MSLN, MUC1, RET, RRAS, SEMA3C, TFPI2, TUBB2A, VAV3</i>	19
Cancer	Ovarian tumor	Ovarian tumor	1.73×10^{-6}	<i>ABCB1, CD24, CLDN3, EGFR, EPCAM, GPX3, HIST2H2AA3, HMGA2, MCAM, MGP, MSLN, MUC1, RET, RRAS, SEMA3C, TFPI2, TUBB2A, VAV3</i>	18
Cancer	Non-small cell lung cancer	Non-small cell lung cancer	2.45×10^{-6}	<i>AKT3, B2M, CDH1, CDKN2A, DNAJB4, EGFR, EPCAM, FN1, FYN, MAFB, MSLN, RB1, RET, S100P, TUBB2A</i>	15
Cancer	Prostate cancer	Prostate cancer	6.19×10^{-6}	<i>ABAT, AKT3, CDH1, COL1A1, CYP1B1, DNAJC15, EGFR, EPCAM, ESR1, FHL2, FYN, GATA3, GDF15, GPX3, IGFBP3, IL7R, KRT7, MMP1 (includes EG:4312), MSMB, MUC1, MYL9 (includes EG:10398), PSAT1, RB1, RET, TFF3, TUBB2A, WWOX</i>	27
Cancer	Clear-cell ovarian carcinoma	Clear-cell ovarian carcinoma	7.74×10^{-6}	<i>CLDN3, EPCAM, GPX3, HMGA2, MGP, MUC1</i>	6
Cancer	Colon cancer	Colon cancer	8.99×10^{-6}	<i>AKT3, BMP7, BST2, CA2, CALD1, CLDN3, COL6A1, COL6A2, CRIP1, EGFR, FN1, FSCN1, GDF15, GJA1, HIST1H2BD, HIST1H4H (includes EG:8365), INSIG1, MCAM, SERPINA3, SERPINE1, STXBP6, TFF3, TPD52L1, TUBB2A</i>	24
Cancer	Carcinoma in situ	Carcinoma in situ	1.60×10^{-5}	<i>FHL2, GATA3, GPX3, HMGA2, IGFBP3, KRT7, MUC1, MYL9 (includes EG:10398), PSAT1, TFF3</i>	10
Cancer	Prostatic intraepithelial neoplasia	Prostatic intraepithelial neoplasia	2.72×10^{-5}	<i>ESR1, FHL2, GATA3, GPX3, IGFBP3, KRT7, MUC1, MYL9 (includes EG:10398), PSAT1, TFF3</i>	10
Cancer	Uterine tumor	Uterine tumor	3.42×10^{-5}	<i>CD24, CDH1, CDH2, CYP1B1, CYR61, EGFR, ESR1, HMGA1, NNMT, IGFBP5, OLFM1, PIK3R3, PPP1R3D, RB1, RET, SLC24A3, STXBP6, TFAP2C, TGFB1, TUBB2A, ZYX</i>	21

Cancer	Pancreatic adenocarcinoma	Pancreatic adenocarcinoma	3.67×10^{-5}	<i>AXL, CYR61, IGFBP3, KRT7, KRT19, MSLN, SERPINE2</i>	7
Cancer	Head and neck tumor	Head and neck tumor	3.69×10^{-5}	<i>ABAT, AKT3, ANXA1, CDKN2A, CREBBP, EGFR, EPCAM, ESR1, FABP5, FN1, FYN, ID2, IGFBP5, INSIG1, PDGFC, PIK3R1, RB1, RET, TGFB1, TUBB2A</i>	20
Cancer	Non-small-cell lung carcinoma	Non-small-cell lung carcinoma	4.92×10^{-5}	<i>AKT3, B2M, CDKN2A, EGFR, EPCAM, FN1, FYN, MAFB, MSLN, RB1, RET, TUBB2A</i>	12
Cancer	Prostatic carcinoma	Prostatic carcinoma	5.27×10^{-5}	<i>ABAT, AKT3, EGFR, EPCAM, ESR1, FHL2, FYN, GATA3, GDF15, GPX3, IGFBP3, KRT7, MSMB, MUC1, MYL9 (includes EG:10398), PSAT1, RB1, RET, TFF3, TUBB2A</i>	20
Cancer	Oral cancer	Oral cancer	6.50×10^{-5}	<i>CDKN2A, CREBBP, EGFR, FN1, IGFBP5, INSIG1, PIK3R1, RET</i>	8
Cancer	Prostatic intraepithelial tumor	Prostatic intraepithelial tumor	8.92×10^{-5}	<i>FHL2, GATA3, GPX3, IGFBP3, KRT7, MUC1, MYL9 (includes EG:10398), PSAT1, TFF3</i>	9
Cancer	Serous ovarian carcinoma	Serous ovarian carcinoma	9.31×10^{-5}	<i>CD24, CLDN3, EPCAM, GPX3, HIST2H2AA3, MCAM, MGP, RRAS, SEMA3C, TFPI2, VAV3</i>	11
Cancer	Lung carcinoma	Lung carcinoma	9.52×10^{-5}	<i>ABCB1, AKT3, B2M, CDKN2A, CLDN3, EGFR, EPCAM, FN1, FYN, LOX, MAFB, MSLN, RB1, RET, TUBB2A</i>	15
Cancer	Leiomyoma	Leiomyoma	9.97×10^{-5}	<i>CD24, CTGF, CYP1B1, CYR61, EGFR, ESR1, HNMT, IGFBP5, ISOC1, NOV, OLFM1, PIK3R3, PPP1R3D, SLC24A3, STXBP6, TFAP2C, TGFB1, ZYX</i>	18
Cancer	Endometrioid carcinoma	Endometrioid carcinoma	1.19×10^{-4}	<i>CLDN3, EPCAM, GPX3, HMGA2, MGP</i>	5
Cancer	Neuroendocrine tumor	Neuroendocrine tumor	1.25×10^{-4}	<i>AXL, CYR61, EGFR, IGFBP3, KRT7, KRT19, MSLN, MSX2, RET, SERPINE2, TUBB2A</i>	11
Cancer	Benign tumor	Benign tumor	2.26×10^{-4}	<i>CD24, CTGF, CYP1B1, CYR61, EGFR, ESR1, FYN, GSTP1, HNMT, IGFBP5, ISOC1, NOV, OLFM1, PIK3R3, PPP1R3D, RET, SLC24A3, STXBP6, TFAP2C, TGFB1, ZYX</i>	21
Cancer	Peripheral T-cell lymphoma	Peripheral T-cell lymphoma	2.76×10^{-4}	<i>COL4A1, COL4A2, CYR61, FN1, NNMT</i>	5
Cancer	Endocrine gland tumor	Endocrine gland tumor	3.03×10^{-4}	<i>ABCB1, AXL, CYR61, EGFR, IGFBP3, KRT7, KRT19, MSLN, MSX2, RET, SERPINE2, TGFB1, TUBB2A</i>	13
Cancer	Neuroendocrine carcinoma	Neuroendocrine carcinoma	3.62×10^{-4}	<i>AXL, CYR61, EGFR, IGFBP3, KRT7, KRT19, MSLN, MSX2, SERPINE2</i>	9
Cancer	Pancreatic carcinoma	Pancreatic carcinoma	4.17×10^{-4}	<i>AXL, CYR61, IGFBP3, KRT7, KRT19, MSLN, MSX2, SERPINE2</i>	8
Cancer	Lung cancer	Lung cancer	8.27×10^{-4}	<i>ABCB1, AKT3, B2M, CDH1, CDKN2A, CLDN3, DNAJB4, EGFR, EPCAM, FN1, FYN, LOX, MAFB, MSLN, RB1, RET, S100P, TUBB2A</i>	18

Cancer	Adenocarcinoma	Adenocarcinoma	1.12×10^{-3}	<i>ANXA1, CDH1, CLDN3, EGFR, ESR1, GJA1, GSTP1, LOX, RET, SERPINE2, TM4SF1, TUBB2A, VIM</i>	13
Cancer	Sarcoma	Sarcoma	1.15×10^{-3}	<i>ABCB1, CDH1, CDKN2A, EGFR, EXT1, FYN, HMGA1, RB1, RET, SDC2, SPARC, TUBB2A</i>	12
Cancer	Acinar-cell carcinoma	Acinar-cell carcinoma	1.21×10^{-3}	<i>CREBBP, FN1, IGFBP5, PIK3R1</i>	4
Cancer	Esophageal carcinoma	Esophageal carcinoma	1.21×10^{-3}	<i>EGFR, GSTP1, RB1, WWOX</i>	4
Cancer	Serous ovarian carcinoma process	Serous ovarian carcinoma process	1.28×10^{-3}	<i>CLDN3, MSLN, MUC1</i>	3
Cancer	Lymphoma	Lymphoma	1.57×10^{-3}	<i>ABCB1, ADA, AKT3, CDKN2A, COL4A1, COL4A2, CYR61, FN1, FYN, ID2, NNMT, RET, TUBB2A</i>	13
Cancer	Osteosarcoma	Osteosarcoma	1.68×10^{-3}	<i>CDKN2A, FYN, RB1, SDC2, SPARC</i>	5
Cancer	Neuroepithelial tumor	Neuroepithelial tumor	1.91×10^{-3}	<i>AKAP12, AKT3, CDKN2A, EGFR, ESR1, ID2, PDGFC, RB1, RET</i>	9
Cancer	Hyperproliferation	Hyperproliferation	1.92×10^{-3}	<i>AREG, CDH1, EGFR, ESR1, MMP1 (includes EG:4312), RET</i>	6
Cancer	Salivary gland tumor	Salivary gland tumor	1.92×10^{-3}	<i>CREBBP, FN1, IGFBP5, INSIG1, PIK3R1, RET</i>	6
Cancer	Papillary thyroid carcinoma	Papillary thyroid carcinoma	2.10×10^{-3}	<i>EGFR, FN1, RET, TFF3, VIM, WNT5A</i>	6
Cancer	Bone tumor	Bone tumor	2.42×10^{-3}	<i>ADA, CDKN2A, EXT1, FYN, RB1, RET, SDC2, SPARC</i>	8
Cancer	Ductal carcinoma	Ductal carcinoma	2.59×10^{-3}	<i>ABAT, ANXA1, ANXA9, CDH2, ECM1, FBN2 (includes EG:2201), MT1E</i>	7
Cancer	Glioma	Glioma	2.74×10^{-3}	<i>AKT3, CDKN2A, EGFR, ESR1, ID2, PDGFC, RB1, RET</i>	8
Cancer	Mucoepidermoid carcinoma	Mucoepidermoid carcinoma	2.94×10^{-3}	<i>CREBBP, FN1, IGFBP5, INSIG1, PIK3R1</i>	5
Cancer	T-cell non-Hodgkin lymphoma	T-cell non-Hodgkin lymphoma	2.95×10^{-3}	<i>AKT3, COL4A1, COL4A2, CYR61, FN1, NNMT</i>	6
Cancer	Ductal carcinoma in situ	Ductal carcinoma in situ	3.07×10^{-3}	<i>ANXA1, ESR1</i>	2
Cancer	Malignancy	Malignancy of tumor	3.07×10^{-3}	<i>EGFR, ID2</i>	2
Cancer	Brain tumor	Brain tumor	3.08×10^{-3}	<i>ABAT, AKT3, CDKN2A, EGFR, ESR1, FYN, ID2, PDGFC, RB1, RET, TGFB1, TUBB2A</i>	12
Cancer	Astrocytoma	Astrocytoma	3.20×10^{-3}	<i>AKT3, EGFR, ESR1, ID2, PDGFC, RET</i>	6

Cancer	Ovarian carcinoma	Ovarian carcinoma	3.20×10^{-3}	<i>ABCB1, EGFR, EPCAM, HMGA2, MSLN, TUBB2A</i>	6
Cancer	Lymphatic node tumor	Lymphatic node tumor	4.48×10^{-3}	<i>AKT3, CDKN2A, COL4A1, COL4A2, CYR61, FN1, FYN, ID2, NNMT, TUBB2A</i>	10
Cancer	Glioblastoma	Glioblastoma	4.76×10^{-3}	<i>AKT3, EGFR, ESR1, PDGFC, RET</i>	5
Cancer	Prolactinoma	Prolactinoma	5.04×10^{-3}	<i>EGFR, ESR1</i>	2
Cancer	Esophageal cancer	Esophageal cancer	5.28×10^{-3}	<i>ADA, EGFR, GSTP1, RB1, RET, TUBB2A, WWOX</i>	7
Cancer	Transformation	Transformation	5.77×10^{-3}	<i>AXL, CDH1, EGFR, ESR1, IRS1, NRCAM, PIK3R1, RASIP1, RET, TFF3</i>	10
Cancer	Transformation	Transformation of cells	1.47×10^{-2}	<i>AXL, EGFR, ESR1, IRS1, NRCAM, PIK3R1, RASIP1, RET, TFF3</i>	9
Cancer	Transformation	Transformation of eukaryotic cells	1.57×10^{-2}	<i>AXL, EGFR, ESR1, IRS1, NRCAM, PIK3R1, RASIP1, TFF3</i>	8
Cancer	Mucinous ovarian carcinoma	Mucinous ovarian carcinoma	6.46×10^{-3}	<i>EPCAM, GPX3, MGP</i>	3
Cancer	Cholangiocarcinoma	Cholangiocarcinoma	6.65×10^{-3}	<i>ANXA1, EGFR, TM4SF1, VIM</i>	4
Cancer	Cervical carcinoma	Cervical carcinoma	7.27×10^{-3}	<i>CDH1, CDH2, RB1, TGFB1, TUBB2A</i>	5
Cancer	Squamous cell tumor	Squamous cell tumor	7.32×10^{-3}	<i>CDKN2A, EGFR, FYN, GPX1, GSTP1, HMGA2, PDZK1, SOCS2, TUBB2A, WWOX</i>	10
Cancer	Leiomyosarcoma	Leiomyosarcoma	7.40×10^{-3}	<i>CDKN2A, FYN, HMGA1, RET</i>	4
Cancer	Uterine leiomyoma	Uterine leiomyoma	8.33×10^{-3}	<i>CD24, CYR61, EGFR, HNMT, IGFBP5, OLFM1, PIK3R3, PPP1R3D, SLC24A3, STXBP6, TFAP2C, ZYX</i>	12
Cancer	Colorectal tumor	Colorectal tumor	9.49×10^{-3}	<i>AKT3, EGFR, FSCN1, GDF15, GJA1, INSIG1, RET, TUBB2A</i>	8
Cancer	Renal cancer	Renal cancer	9.69×10^{-3}	<i>ABCB1, CDH1, EGFR, EPCAM, FN1, IL7R, RB1, RET, TUBB2A</i>	9
Cancer	Chemotherapy resistance	Chemotherapy resistance of tumor cell lines	1.03×10^{-2}	<i>AXL, EGFR</i>	2
Cancer	Myosarcoma	Myosarcoma	1.06×10^{-2}	<i>CDKN2A, FYN, HMGA1, RET, TUBB2A</i>	5
Cancer	Non-hodgkin lymphoma	Non-hodgkin lymphoma	1.13×10^{-2}	<i>AKT3, CDKN2A, COL4A1, COL4A2, CYR61, FN1, FYN, NNMT, TUBB2A</i>	9
Cancer	Cervical cancer	Cervical cancer	1.18×10^{-2}	<i>CDH1, CDH2, EGFR, RB1, RET, TGFB1, TUBB2A</i>	7
Cancer	Metastasis	Metastasis	1.21×10^{-2}	<i>ABCB1, CYP1B1, EGFR, FYN, TUBB2A, VIM</i>	6
Cancer	Chondrosarcoma	Chondrosarcoma	1.35×10^{-2}	<i>EXT1, FYN</i>	2

Cancer	Renal-cell carcinoma	Renal-cell carcinoma	1.35×10^{-2}	<i>ABCB1, CDH1, EGFR, FN1, RB1, RET, TUBB2A</i>	7
Cancer	Squamous-cell carcinoma	Squamous-cell carcinoma	1.63×10^{-2}	<i>CDKN2A, EGFR, FYN, GSTP1, HMGA2, PDZK1, SOCS2, TUBB2A, WWOX</i>	9
Reproductive system disease	Ovarian cancer	Ovarian cancer	1.16×10^{-16}	<i>ABCB1, ADA, AGR2, AKT3, BMP7, CD24, CDH1, CDKN2A, CEACAM6 (includes EG:4680), CLDN3, COL4A1, EGFR, EPCAM, ESR1, FN1, FYN, GPX3, HIST2H2AA3, HMGA1, HMGA2, IGF2BP3, MCAM, MGP, MMP1 (includes EG:4312), MSLN, MUC1, PRAME (includes EG:23532), PSAT1, RET, RRAS, S100P, SEMA3C, SLC16A3, TACSTD2, TFF3, TFPI2, TUBB2A, VAV3</i>	38
Reproductive system disease	Breast carcinoma	Breast carcinoma	4.51×10^{-11}	<i>ABAT, ABCB1, AGR2, ANXA1, ANXA9, AREG, BEX1, BMP7, C6ORF211, CDH2, CEACAM6 (includes EG:4680), ECM1, EGFR, ESR1, FBN2 (includes EG:2201), FSCN1, GATA3, GFRA1, IGFBP3, MT1E, MUC1, NPY1R, PRAME (includes EG:23532), RB1, SCUBE2, SLC19A2, SLC39A6, TFF1, TUBB2A</i>	29
Reproductive system disease	Mammary tumor	Mammary tumor	1.15×10^{-10}	<i>ABAT, ABCB1, AGR2, AKT3, ANXA1, ANXA9, AREG, BEX1, BMP7, C6ORF211, CDH2, CEACAM6 (includes EG:4680), ECM1, EGFR, ESR1, FBN2 (includes EG:2201), FSCN1, GATA3, GFRA1, IGFBP3, MT1E, MUC1, NPY1R, PRAME (includes EG:23532), RB1, RET, SCUBE2, SLC19A2, SLC39A6, TFF1, TUBB2A</i>	31
Reproductive system disease	Breast cancer	Breast cancer	1.58×10^{-10}	<i>ABAT, ABCB1, ADA, AGR2, AKR1C2, AKT3, ANXA1, ANXA9, AREG, BEX1, BMP7, BRIP1, C6ORF211, CDH2, CDKN2A, CEACAM6 (includes EG:4680), CLDN3, CLGN, COL1A1, CYR61, ECM1, EGFR, ESR1, FBN2 (includes EG:2201), FSCN1, FYN, GATA3, GFRA1, HEY2, HIST2H2BE, HMGA1, IGFBP3, KLF4, MT1E, MUC1, NPY1R, PRAME (includes EG:23532), RB1, RET, S100A14, SCUBE2, SGK3, SLC19A2, SLC39A6, SPARC, TACSTD2, TFF1, TGFB1, TUBB2A, WWOX</i>	50
Reproductive system disease	Genital tumor	Genital tumor	1.36×10^{-8}	<i>ABAT, ABCB1, ADA, AKT3, CD24, CLDN3, EGFR, EPCAM, ESR1, FHL2, FYN, GATA3, GDF15, GPX3, HIST2H2AA3, HMGA2, IGFBP3, KRT7, MCAM, MGP, MSLN, MSMB, MUC1, MYL9 (includes EG:10398), PSAT1, RB1, RET, RRAS, SEMA3C, TFF3, TFPI2, TUBB2A, VAV3</i>	33
Reproductive system disease	Endometriosis	Endometriosis	4.03×10^{-8}	<i>ANK3, BCAT1, CCR10, COL6A1, COL6A2, CTGF, CXADR, CYR61, EGFR, ESR1, FN1, GATA3, GPX3, GSTP1, HMGA2, ICAM1, IL18, MMP1 (includes EG:4312), MPPED2 (includes EG:744), MT1E, MUC1, OLFM1, PIK3R1, SDC2, SLC16A1, STC1, TGFB1, TGFB1, UCHL1, VIM</i>	30
Reproductive system disease	Reproductive system disorder	Reproductive system disorder	6.95×10^{-8}	<i>ANK3, BCAT1, CCR10, COL6A1, COL6A2, CRIP1, CTGF, CXADR, CYR61, EGFR, EML1 (includes EG:2009), ESR1, FN1, GATA3, GPX1, GPX3, GSTP1, HMGA2, ICAM1, IL18, MMP1 (includes EG:4312), MPPED2 (includes EG:744), MT1E, MUC1, OLFM1, OXTR, PIK3R1, SDC2, SLC16A1, STC1, STXBP6, TGFB1, TGFB1, UCHL1, VIM, XIST</i>	36

Reproductive system disease	Gonadal tumor	Gonadal tumor	7.95×10^{-7}	<i>ABCB1, ADA, CD24, CLDN3, EGFR, EPCAM, GPX3, HIST2H2AA3, HMGA2, MCAM, MGP, MSLN, MUC1, RET, RRAS, SEMA3C, TFPI2, TUBB2A, VAV3</i>	19
Reproductive system disease	Ovarian tumor	Ovarian tumor	1.73×10^{-6}	<i>ABCB1, CD24, CLDN3, EGFR, EPCAM, GPX3, HIST2H2AA3, HMGA2, MCAM, MGP, MSLN, MUC1, RET, RRAS, SEMA3C, TFPI2, TUBB2A, VAV3</i>	18
Reproductive system disease	Prostate cancer	Prostate cancer	6.19×10^{-6}	<i>ABAT, AKT3, CDH1, COL1A1, CYP1B1, DNAJC15, EGFR, EPCAM, ESR1, FHL2, FYN, GATA3, GDF15, GPX3, IGFBP3, IL7R, KRT7, MMP1 (includes EG:4312), MSMB, MUC1, MYL9 (includes EG:10398), PSAT1, RB1, RET, TFF3, TUBB2A, WWOX</i>	27
Reproductive system disease	Clear-cell ovarian carcinoma	Clear-cell ovarian carcinoma	7.74×10^{-6}	<i>CLDN3, EPCAM, GPX3, HMGA2, MGP, MUC1</i>	6
Reproductive system disease	Prostatic intraepithelial neoplasia	Prostatic intraepithelial neoplasia	2.72×10^{-5}	<i>ESR1, FHL2, GATA3, GPX3, IGFBP3, KRT7, MUC1, MYL9 (includes EG:10398), PSAT1, TFF3</i>	10
Reproductive system disease	Uterine tumor	Uterine tumor	3.42×10^{-5}	<i>CD24, CDH1, CDH2, CYP1B1, CYR61, EGFR, ESR1, HMGA1, HNMT, IGFBP5, OLFM1, PIK3R3, PPP1R3D, RB1, RET, SLC24A3, STXBP6, TFAP2C, TGFB1, TUBB2A, ZYX</i>	21
Reproductive system disease	Prostatic carcinoma	Prostatic carcinoma	5.27×10^{-5}	<i>ABAT, AKT3, EGFR, EPCAM, ESR1, FHL2, FYN, GATA3, GDF15, GPX3, IGFBP3, KRT7, MSMB, MUC1, MYL9 (includes EG:10398), PSAT1, RB1, RET, TFF3, TUBB2A</i>	20
Reproductive system disease	Prostatic intraepithelial tumor	Prostatic intraepithelial tumor	8.92×10^{-5}	<i>FHL2, GATA3, GPX3, IGFBP3, KRT7, MUC1, MYL9 (includes EG:10398), PSAT1, TFF3</i>	9
Reproductive system disease	Serous ovarian carcinoma	Serous ovarian carcinoma	9.31×10^{-5}	<i>CD24, CLDN3, EPCAM, GPX3, HIST2H2AA3, MCAM, MGP, RRAS, SEMA3C, TFPI2, VAV3</i>	11
Reproductive system disease	Endometrioid carcinoma	Endometrioid carcinoma	1.19×10^{-4}	<i>CLDN3, EPCAM, GPX3, HMGA2, MGP</i>	5
Reproductive system disease	Serous ovarian carcinoma process	Serous ovarian carcinoma process	1.28×10^{-3}	<i>CLDN3, MSLN, MUC1</i>	3
Reproductive system disease	Ductal carcinoma	Ductal carcinoma	2.59×10^{-3}	<i>ABAT, ANXA1, ANXA9, CDH2, ECM1, FBN2 (includes EG:2201), MT1E</i>	7
Reproductive system disease	Ductal carcinoma in situ	Ductal carcinoma in situ	3.07×10^{-3}	<i>ANXA1, ESR1</i>	2

Reproductive system disease	Ovarian carcinoma	Ovarian carcinoma	3.20×10^{-3}	<i>ABCB1, EGFR, EPCAM, HMGA2, MSLN, TUBB2A</i>	6
Reproductive system disease	Mucinous ovarian carcinoma	Mucinous ovarian carcinoma	6.46×10^{-3}	<i>EPCAM, GPX3, MGP</i>	3
Reproductive system disease	Cervical carcinoma	Cervical carcinoma	7.27×10^{-3}	<i>CDH1, CDH2, RB1, TGFB1, TUBB2A</i>	5
Reproductive system disease	Uterine leiomyoma	Uterine leiomyoma	8.33×10^{-3}	<i>CD24, CYR61, EGFR, HNMT, IGFBP5, OLFM1, PIK3R3, PPP1R3D, SLC24A3, STXBP6, TFAP2C, ZYX</i>	12
Reproductive system disease	Adenomyosis	Adenomyosis	9.16×10^{-3}	<i>CRIP1, ESR1, GPX1, STXBP6, XIST</i>	5
Reproductive system disease	Cervical cancer	Cervical cancer	1.18×10^{-2}	<i>CDH1, CDH2, EGFR, RB1, RET, TGFB1, TUBB2A</i>	7
Cellular growth and proliferation	Proliferation	Proliferation of cells	1.38×10^{-15}	<i>AKT3, ANXA1, AREG, AXL, BCAT1, BDNF, BIN1, BMP7, BST2, BTG3, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), COL1A1, COL4A1, COL4A2, CRIP1, CTGF, CXCL12, CYR61, DNAJB4, ECM1, EGFR, EIF5A, EMP1, EMP3, EPCAM, ESR1, FABP5, FKBP1B, FN1, FOSL1, FSCN1, HMGA1, ICAM1, ID2, IFI16, IGF2BP3, IGFBP3, IGFBP5, IGFBP6, IL18, IL7R, INSIG1, IRS1, KLF4, LAMB1, LOX, MGP, MSX2, MYB (includes EG:4602), NCOA3, NDN, NOV, OXTR, PDGFC, PDZK1, PTPRK, RB1, RET, RLN2, S100P, SDC2, SERPINE1, SOCS2, SPARC, STC1, TACSTD2, TBX2, TFPI2, TGFB1, TGFB1I1, TGFB1I, TNFRSF6B, UGT2B15, WNT5A</i>	78
Cellular growth and proliferation	Proliferation	Proliferation of eukaryotic cells	1.92×10^{-13}	<i>AKT3, ANXA1, AREG, AXL, BDNF, BMP7, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), COL1A1, COL4A1, COL4A2, CTGF, CXCL12, CYR61, DNAJB4, ECM1, EGFR, EPCAM, ESR1, FABP5, FKBP1B, FN1, FSCN1, HMGA1, ICAM1, ID2, IFI16, IGF2BP3, IGFBP3, IGFBP5, IGFBP6, IL18, IL7R, IRS1, LAMB1, LOX, MGP, MSX2, MYB (includes EG:4602), NCOA3, NOV, OXTR, PDGFC, PTPRK, RB1, RET, RLN2, S100P, SDC2, SERPINE1, SOCS2, SPARC, STC1, TFPI2, TGFB1, TNFRSF6B, UGT2B15, WNT5A</i>	61

Cellular growth and proliferation	Proliferation	Proliferation of cell lines	6.99×10^{-13}	<i>AKT3, ANXA1, AREG, AXL, BDNF, BMP7, CD24, CDH1, CDKN2A, CEACAM6 (includes EG:4680), COL1A1, CTGF, CXCL12, CYR61, DNAJB4, EGFR, EPCAM, ESR1, FABP5, FKBP1B, FN1, FSCN1, HMGA1, ID2, IFI16, IGF2BP3, IGFBP3, IGFBP5, IGFBP6, IRS1, LOX, MSX2, MYB (includes EG:4602), NCOA3, NOV, OXTR, RB1, RET, S100P, SDC2, SERPINE1, SOCS2, SPARC, TFPI2, TGFB1, TNFRSF6B, UGT2B15, WNT5A</i>	48
Cellular growth and proliferation	Proliferation	Proliferation of tumor cell lines	1.67×10^{-9}	<i>AKT3, ANXA1, AREG, AXL, BDNF, BMP7, CD24, CDH1, CDKN2A, CEACAM6 (includes EG:4680), COL1A1, CTGF, CXCL12, CYR61, DNAJB4, EGFR, EPCAM, ESR1, FABP5, HMGA1, ID2, IGF2BP3, IGFBP3, IGFBP5, MSX2, MYB (includes EG:4602), NCOA3, NOV, RB1, RET, SERPINE1, SPARC, TFPI2, TGFB1, UGT2B15, WNT5A</i>	36
Cellular growth and proliferation	Proliferation	Proliferation of normal cells	4.10×10^{-6}	<i>AXL, BMP7, CD24, CDH2, CDKN2A, COL4A1, COL4A2, CTGF, CXCL12, ECM1, EGFR, FN1, HMGA1, ICAM1, IGFBP3, IGFBP5, IL18, IL7R, LAMB1, MGP, PDGFC, PTPRK, RLN2, SERPINE1, SPARC, STC1, TGFB1</i>	27
Cellular growth and proliferation	Proliferation	Proliferation of carcinoma cell lines	4.22×10^{-6}	<i>AKT3, ANXA1, CXCL12, CYR61, DNAJB4, EGFR, NCOA3, RB1, TGFB1, WNT5A</i>	10
Cellular growth and proliferation	Proliferation	Proliferation of connective tissue cells	6.22×10^{-6}	<i>CDKN2A, CTGF, FN1, IGFBP3, IGFBP5, IL18, PDGFC, RLN2, STC1, TGFB1</i>	10
Cellular growth and proliferation	Proliferation	Proliferation of chondrocytes	1.79×10^{-5}	<i>CTGF, IGFBP3, IL18, STC1</i>	4
Cellular growth and proliferation	Proliferation	Proliferation of breast cancer cell lines	9.31×10^{-5}	<i>BMP7, CDKN2A, CYR61, EGFR, ESR1, ID2, IGFBP3, MYB (includes EG:4602), NCOA3, SERPINE1, TGFB1</i>	11
Cellular growth and proliferation	Proliferation	Proliferation of lung cancer cell lines	1.97×10^{-4}	<i>AKT3, ANXA1, AXL, CYR61, DNAJB4, EGFR, NCOA3, RB1</i>	8
Cellular growth and proliferation	Proliferation	Proliferation of breast cell lines	4.02×10^{-4}	<i>CDH1, IGFBP3, LOX, TGFB1</i>	4
Cellular growth and proliferation	Proliferation	Proliferation of squamous cell carcinoma cell lines	4.02×10^{-4}	<i>ANXA1, CDH1, EGFR, EPCAM</i>	4
Cellular growth and proliferation	Proliferation	Proliferation of pancreatic cancer cell lines	1.45×10^{-3}	<i>CEACAM6 (includes EG:4680), HMGA1, MSX2, NCOA3</i>	4

Cellular growth and proliferation	Proliferation	Proliferation of thyroid tumor cell lines	1.74×10^{-3}	CXCL12, TGFB1, WNT5A	3
Cellular growth and proliferation	Proliferation	Proliferation of lung cell lines	2.28×10^{-3}	CDKN2A, FN1, IFI16	3
Cellular growth and proliferation	Proliferation	Proliferation of endothelial cells	3.19×10^{-3}	AXL, CDH2, COL4A1, COL4A2, ECM1, FN1, MGP	7
Cellular growth and proliferation	Proliferation	Proliferation of colon cancer cell lines	5.20×10^{-3}	AREG, CD24, ID2, RB1, TGFB1	5
Cellular growth and proliferation	Proliferation	Proliferation of epithelial cells	6.18×10^{-3}	EGFR, LAMB1, PTPRK, SPARC, TGFB1	5
Cellular growth and proliferation	Proliferation	Proliferation of epidermal cells	6.46×10^{-3}	EGFR, PTPRK, TGFB1	3
Cellular growth and proliferation	Proliferation	Proliferation of T lymphocytes	1.28×10^{-2}	CD24, CDKN2A, CXCL12, FN1, ICAM1, IL18, IL7R, TGFB1	8
Cellular growth and proliferation	Proliferation	Proliferation of endocrine cells	1.35×10^{-2}	FN1, TGFB1	2
Cellular growth and proliferation	Proliferation	Proliferation of fibroblast cell lines	1.68×10^{-2}	CDKN2A, CXCL12, FN1, S100P, TGFB1	5
Cellular growth and proliferation	Growth	Growth of cells	1.60×10^{-12}	ABCB1, AKT3, ANXA1, AREG, BMP7, CALCR, CDA, CDH1, CDKN2A, COL6A1, COL6A2, CTGF, CXADR, CXCL12, CYR61, DBN1, DKK3, DNAJB4, ECM1, EGFR, EIF5A, ESR1, FBN2 (includes EG:2201), FHL2, FN1, FOSL1, GDF15, GFRA1, GJA1, GPX1, GPX3, GSTP1, HMGA1, HMGA2, ID2, IFI16, IGFBP3, IGFBP5, IRS1, KLF4, LAMB1, MCAM, MSX2, MUC1, MYL9 (includes EG:10398), NCOA3, NDN, NOV, NRCAM, OXTR, PDCD10, PEG10, PIK3R1, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, SERPINE1, SERPINE2, SERTAD2, SPARC, TFAP2C, TFF3, TGFB1, TNFRSF6B, UBE2L6, XBP1, ZYX	69

Cellular growth and proliferation	Growth	Growth of eukaryotic cells	6.99×10^{-12}	<i>ABCB1, AKT3, ANXA1, AREG, BMP7, CALCR, CDH1, CDKN2A, CTGF, CXADR, CYR61, DKK3, DNAJB4, EGFR, EIF5A, ESR1, FHL2, FN1, FOSL1, GDF15, GFRA1, GJA1, HMGA1, HMGA2, ID2, IFI16, IGFBP3, IGFBP5, IRS1, KLF4, MCAM, MSX2, MYL9 (includes EG:10398), NCOA3, NOV, NRCAM, OXTR, PDCD10, PEG10, PIK3R1, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, TFAP2C, TFF3, TGFB1, XBP1</i>	49
Cellular growth and proliferation	Growth	Growth of cell lines	5.80×10^{-11}	<i>ABCB1, AKT3, ANXA1, AREG, BMP7, CALCR, CDH1, CDKN2A, CTGF, CXADR, CYR61, DKK3, DNAJB4, EGFR, EIF5A, ESR1, FN1, FOSL1, GDF15, GJA1, HMGA1, HMGA2, IFI16, IGFBP3, IGFBP5, IRS1, KLF4, MSX2, MYL9 (includes EG:10398), NCOA3, NOV, NRCAM, OXTR, PDCD10, PIK3R1, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, TFAP2C, TGFB1, XBP1</i>	43
Cellular growth and proliferation	Growth	Growth of tumor cell lines	1.55×10^{-10}	<i>ABCB1, AKT3, ANXA1, CDH1, CDKN2A, CTGF, CXADR, CYR61, DKK3, DNAJB4, EGFR, EIF5A, ESR1, FOSL1, GDF15, GJA1, HMGA1, HMGA2, IFI16, IGFBP3, IGFBP5, IRS1, KLF4, MSX2, MYL9 (includes EG:10398), NCOA3, NOV, PDCD10, PIK3R1, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, TFAP2C, TGFB1, XBP1</i>	37
Cellular growth and proliferationn	Growth	Growth of breast cancer cell lines	3.81×10^{-6}	<i>CDH1, CYR61, EGFR, ESR1, GJA1, HMGA1, IGFBP3, IGFBP5, IRS1, NCOA3, TFAP2C, TGFB1, XBP1</i>	13
Cellular growth and proliferation	Growth	Growth of neuroblastoma cell lines	2.29×10^{-4}	<i>EGFR, HMGA2, IGFBP5, RET</i>	4
Cellular growth and proliferation	Growth	Arrest in growth of eukaryotic cells	8.52×10^{-4}	<i>AKT3, BMP7, CDKN2A, KLF4, PRAME (includes EG:23532), RB1, TGFB1</i>	7
Cellular growth and proliferation	Growth	Growth of epithelial cells	8.74×10^{-4}	<i>AREG, EGFR, FHL2, ID2, RB1</i>	5
Cellular growth and proliferation	Growth	Growth of cancer cells	1.20×10^{-3}	<i>CDKN2A, GFRA1, MCAM, PEG10, RB1, TFF3, TGFB1</i>	7
Cellular growth and proliferation	Growth	Growth of tumor cells	1.34×10^{-3}	<i>CDKN2A, GFRA1, HMGA2, MCAM, PEG10, RB1, TFF3, TGFB1</i>	8

Cellular growth and proliferation	Growth	Growth of fibroblast cell lines	1.66×10^{-3}	<i>EGFR, FN1, IGFBP3, IRS1, NRCAM, RET, TGFB1</i>	7
Cellular growth and proliferation	Growth	Growth of epidermal cells	5.41×10^{-3}	<i>EGFR, FHL2, ID2</i>	3
Cellular growth and proliferation	Growth	Arrest in growth of cell lines	6.18×10^{-3}	<i>AKT3, BMP7, KLF4, PRAME (includes EG:23532), TGFB1</i>	5
Cellular growth and proliferation	Growth	Growth of carcinoma cell lines	6.19×10^{-3}	<i>AKT3, ANXA1, DNAJB4, IGFBP3, NOV, PIK3R1</i>	6
Cellular growth and proliferation	Growth	Growth of endometrial cancer cell lines	7.45×10^{-3}	<i>CYR61, ESR1</i>	2
Cellular growth and proliferation	Growth	Growth of squamous cell carcinoma cell lines	1.03×10^{-2}	<i>CDKN2A, EGFR</i>	2
Cellular growth and proliferation	Growth	Growth of prostate cancer cell lines	1.14×10^{-2}	<i>CXADR, IFI16, IGFBP5, PDCD10, TGFB1</i>	5
Cellular growth and proliferation	Growth	Growth of lung cancer cell lines	1.48×10^{-2}	<i>AKT3, ANXA1, DNAJB4, IGFBP3, PIK3R1</i>	5
Cellular growth and proliferation	Growth	Growth of normal cells	1.51×10^{-2}	<i>AREG, CDKN2A, EGFR, FHL2, ID2, IGFBP5, RB1, TGFB1</i>	8
Cellular growth and proliferation	Growth	Growth of connective tissue cells	1.52×10^{-2}	<i>AREG, CDKN2A, TGFB1</i>	3
Cellular growth and proliferation	Growth	Growth of pancreatic cancer cell lines	1.52×10^{-2}	<i>IGFBP3, MSX2, NCOA3</i>	3
Cellular growth and proliferation	Colony formation	Colony formation by cells	2.83×10^{-5}	<i>ANKRD1, CDA, CDKN2A, CEACAM5 (includes EG:1048), CXCL12, CYR61, EGFR, IFI16, IGFBP3, IRS1, NCOA3, PIK3R1, PLAGL1, RB1, SPARC, TGFB1, TGFB1I1</i>	17
Cellular growth and proliferation	Colony formation	Colony formation by eukaryotic cells	1.45×10^{-4}	<i>ANKRD1, CDA, CDKN2A, CEACAM5 (includes EG:1048), CYR61, EGFR, IFI16, IGFBP3, IRS1, NCOA3, PIK3R1, RB1, SPARC, TGFB1, TGFB1I1</i>	15

Cellular growth and proliferation	Colony formation	Colony formation by cell lines	2.31×10^{-4}	<i>ANKRD1, CDKN2A, CEACAM5</i> (includes EG:1048), <i>CYR61, EGFR, IFI16, IGFBP3, IRS1, NCOA3, PIK3R1, RB1, SPARC, TGFB1</i>	13
Cellular growth and proliferation	Colony formation	Colony formation by tumor cell lines	3.45×10^{-4}	<i>ANKRD1, CDKN2A, CEACAM5</i> (includes EG:1048), <i>CYR61, EGFR, IFI16, IGFBP3, NCOA3, RB1, SPARC, TGFB1</i>	11
Cellular growth and proliferation	Quantity	Affects quantity of tumor cell lines	1.92×10^{-3}	<i>CDKN2A, ESR1, IGFBP3, IGFBP5, IGFBP6, KLF4</i>	6
Cellular growth and proliferation	Quantity	Affects quantity of prostate cancer cell lines	5.04×10^{-3}	<i>IGFBP3, IGFBP5</i>	2
Cellular growth and proliferation	Induction	Induction of T lymphocytes	3.07×10^{-3}	<i>IL18, MUC1</i>	2
Cellular growth and proliferation	Co-stimulation	Co-stimulation of T lymphocytes	1.35×10^{-2}	<i>CD24, FN1</i>	2
Cellular growth and proliferation	Formation	Formation of endothelial cells	1.35×10^{-2}	<i>FN1, SERPINE1</i>	2
Cellular movement	Invasion	Invasion of cells	2.41×10^{-15}	<i>AKT3, ANXA1, AREG, AXL, BMP7, CD24, CDH1, CDH2, CLDN3, CXCL12, DKK3, DNAJB4, EGFR, ESR1, FABP5, FN1, FSCN1, FXYD5, GDF15, HMGA1, ID2, IL18, LOX, MCAM, MMP1</i> (includes EG:4312), <i>NCOA3, NOV, NUAK1, SPARC, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, VIM, WNT5A</i>	36
Cellular movement	Invasion	Invasion of eukaryotic cells	2.19×10^{-14}	<i>AKT3, ANXA1, AREG, AXL, BMP7, CD24, CDH1, CLDN3, CXCL12, DKK3, DNAJB4, EGFR, ESR1, FABP5, FN1, FSCN1, FXYD5, GDF15, HMGA1, IL18, LOX, MCAM, MMP1</i> (includes EG:4312), <i>NCOA3, NOV, NUAK1, SPARC, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, VIM, WNT5A</i>	34
Cellular movement	Invasion	Invasion of cell lines	2.41×10^{-14}	<i>AKT3, ANXA1, AREG, AXL, BMP7, CD24, CDH1, CLDN3, CXCL12, DKK3, DNAJB4, EGFR, ESR1, FABP5, FN1, FSCN1, FXYD5, GDF15, HMGA1, LOX, MMP1</i> (includes EG:4312), <i>NCOA3, NOV, NUAK1, SPARC, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, VIM, WNT5A</i>	32
Cellular movement	Invasion	Invasion of tumor cell lines	5.51×10^{-14}	<i>AKT3, ANXA1, AXL, BMP7, CD24, CDH1, CLDN3, CXCL12, DKK3, DNAJB4, EGFR, ESR1, FABP5, FN1, FXYD5, GDF15, HMGA1, LOX, MMP1</i> (includes EG:4312), <i>NCOA3, NOV, NUAK1, SPARC, TFAP2C, TFF1, TFF3, TFPI2, TGFB1, VIM, WNT5A</i>	30

Cellular movement	Invasion	Invasion of prostate cancer cell lines	5.23×10^{-6}	<i>BMP7, CDH1, FABP5, FN1, MMP1</i> (includes EG:4312), <i>NCOA3, VIM</i>	7
Cellular movement	Invasion	Invasion of colon cancer cell lines	1.50×10^{-4}	<i>ANXA1, EGFR, NUAK1, TFF1, TFF3</i>	5
Cellular movement	Invasion	Invasion of epithelial cell lines	1.67×10^{-4}	<i>AREG, CDH1, EGFR, FSCN1</i>	4
Cellular movement	Invasion	Invasion of carcinoma cell lines	2.76×10^{-4}	<i>AKT3, DNAJB4, TGFB1, VIM, WNT5A</i>	5
Cellular movement	Invasion	Invasion of breast cell lines	3.94×10^{-4}	<i>EGFR, FXYD5, TGFB1</i>	3
Cellular movement	Invasion	Invasion of lung cancer cell lines	1.72×10^{-3}	<i>AKT3, AXL, DNAJB4, TGFB1</i>	4
Cellular movement	Invasion	Invasion of hepatoma cell lines	2.28×10^{-3}	<i>MMP1</i> (includes EG:4312), <i>TFPI2, TGFB1</i>	3
Cellular movement	Invasion	Invasion of breast cancer cell lines	3.88×10^{-3}	<i>CDH1, ESR1, FN1, FXYD5, LOX, MMP1</i> (includes EG:4312), <i>SPARC</i>	7
Cellular movement	Invasion	Invasion of melanoma cells	7.45×10^{-3}	<i>CDH1, MCAM</i>	2
Cellular movement	Invasion	Invasion of bone cancer cell lines	1.03×10^{-2}	<i>DKK3, NOV</i>	2
Cellular movement	Invasion	Invasion of thyroid tumor cell lines	1.03×10^{-2}	<i>VIM, WNT5A</i>	2
Cellular movement	Movement	Movement of cells	1.56×10^{-9}	<i>AKT3, ANXA1, AREG, AXL, BDNF, CDH1, CDH2, CDKN2A, COL4A1, COL4A2, CTGF, CXCL12, CYR61, EGFR, FN1, FYN, GJA1, ICAM1, ICAM2, ID2, IGFBP3, IGFBP5, IGFBP6, IL18, LAMB1, LOX, MGP, MSX2, NCOA3, NOV, PTPRK, RASGRP1, RET, RLN2, S100P, SERPINA3, SERPINE1, STC1, TFF3, TGFB1, TGM2, TNFRSF6B, WNT5A</i>	43
Cellular movement	Movement	Movement of eukaryotic cells	1.13×10^{-8}	<i>AKT3, ANXA1, AXL, BDNF, CDH1, CDH2, CDKN2A, COL4A1, COL4A2, CTGF, CXCL12, CYR61, EGFR, FN1, FYN, GJA1, ICAM1, ICAM2, IGFBP3, IGFBP5, IGFBP6, IL18, LOX, MGP, MSX2, NCOA3, NOV, RASGRP1, RET, RLN2, S100P, SERPINA3, SERPINE1, STC1, TFF3, TGFB1, TGM2, TNFRSF6B, WNT5A</i>	39

Cellular movement	Movement	Movement of cell lines	1.79×10^{-5}	<i>AKT3, CDH1, CDH2, CDKN2A, CXCL12, EGFR, FN1, FYN, GJA1, ICAM1, IGFBP6, LOX, MSX2, NCOA3, NOV, RET, RLN2, SERPINE1, TFF3, TGFB1, TNFRSF6B, WNT5A</i>	22
Cellular movement	Movement	Movement of tumor cell lines	2.19×10^{-5}	<i>AKT3, CDH1, CDH2, CDKN2A, CXCL12, EGFR, FN1, FYN, GJA1, IGFBP6, MSX2, NCOA3, NOV, RET, RLN2, SERPINE1, TGFB1, WNT5A</i>	18
Cellular movement	Movement	Movement of neuroblastoma cell lines	2.29×10^{-4}	<i>CXCL12, EGFR, RET</i>	3
Cellular movement	Movement	Movement of rhabdomyosarcoma cell lines	5.28×10^{-4}	<i>CXCL12, IGFBP6</i>	2
Cellular movement	Movement	Movement of kidney cell lines	9.99×10^{-3}	<i>FN1, ICAM1, RET, TFF3</i>	4
Cellular movement	Migration	Migration of cells	3.34×10^{-9}	<i>AKT3, ANXA1, AREG, AXL, BDNF, CDH1, CDH2, CDKN2A, COL4A1, COL4A2, CTGF, CXCL12, CYR61, EGFR, FN1, FYN, GJA1, ICAM1, ICAM2, ID2, IGFBP3, IGFBP5, IGFBP6, IL18, LAMB1, LOX, MGP, MSX2, NCOA3, NOV, PTPRK, RASGRP1, RET, RLN2, S100P, SERPINA3, SERPINE1, STC1, TGFB1, TGM2, TNFRSF6B, WNT5A</i>	42
Cellular movement	Migration	Migration of eukaryotic cells	2.40×10^{-8}	<i>AKT3, ANXA1, AXL, BDNF, CDH1, CDH2, CDKN2A, COL4A1, COL4A2, CTGF, CXCL12, CYR61, EGFR, FN1, FYN, GJA1, ICAM1, ICAM2, IGFBP3, IGFBP5, IGFBP6, IL18, LOX, MGP, MSX2, NCOA3, NOV, RASGRP1, RET, RLN2, S100P, SERPINA3, SERPINE1, STC1, TGFB1, TGM2, TNFRSF6B, WNT5A</i>	38
Cellular movement	Migration	Migration of normal cells	1.30×10^{-7}	<i>ANXA1, AXL, BDNF, CDH1, CDH2, COL4A1, COL4A2, CTGF, CXCL12, CYR61, EGFR, FN1, ICAM1, ICAM2, IGFBP3, IGFBP5, IL18, MGP, RASGRP1, RLN2, S100P, SERPINA3, SERPINE1, STC1, TGFB1, TGM2, TNFRSF6B</i>	27
Cellular movement	Migration	Migration of endothelial cells	2.54×10^{-7}	<i>AXL, COL4A1, COL4A2, CXCL12, CYR61, FN1, ICAM1, IGFBP3, MGP, S100P, SERPINE1, STC1, TGFB1</i>	13
Cellular movement	Migration	Migration of cell lines	4.25×10^{-5}	<i>AKT3, CDH1, CDH2, CDKN2A, CXCL12, EGFR, FN1, FYN, GJA1, ICAM1, IGFBP6, LOX, MSX2, NCOA3, NOV, RET, RLN2, SERPINE1, TGFB1, TNFRSF6B, WNT5A</i>	21
Cellular movement	Migration	Migration of tumor cell lines	6.22×10^{-5}	<i>AKT3, CDH1, CDH2, CDKN2A, CXCL12, EGFR, FN1, FYN, GJA1, IGFBP6, MSX2, NCOA3, NOV, RLN2, SERPINE1, TGFB1, WNT5A</i>	17
Cellular movement	Migration	Migration of myeloid cells	6.84×10^{-5}	<i>ANXA1, CXCL12, FN1, ICAM1, SERPINE1, TGFB1, TGM2</i>	7
Cellular movement	Migration	Migration of neutrophils	3.97×10^{-4}	<i>ANXA1, FN1, ICAM1, SERPINE1, TGFB1</i>	5
Cellular movement	Migration	Migration of granulocytes	4.78×10^{-4}	<i>ANXA1, CXCL12, FN1, ICAM1, SERPINE1, TGFB1</i>	6

Cellular movement	Migration	Lymphocyte migration	6.53×10^{-4}	CXCL12, EGFR, FN1, ICAM1, ICAM2, IL18, RASGRP1, TGM2, TNFRSF6B	9
Cellular movement	Migration	Migration of mononuclear leukocytes	8.93×10^{-4}	CXCL12, EGFR, FN1, ICAM1, ICAM2, IL18, RASGRP1, TGFB1, TGM2, TNFRSF6B	10
Cellular movement	Migration	Migration of breast cancer cell lines	1.54×10^{-3}	CDH2, CXCL12, EGFR, FN1, GJA1, RLN2, SERPINE1	7
Cellular movement	Migration	Migration of leukocytes	1.64×10^{-3}	ANXA1, CXCL12, EGFR, FN1, ICAM1, ICAM2, IL18, RASGRP1, RLN2, SERPINA3, SERPINE1, TGFB1, TGM2, TNFRSF6B	14
Cellular movement	Migration	Migration of bone cancer cell lines	1.74×10^{-3}	FN1, FYN, NOV	3
Cellular movement	Migration	Migration of lung cancer cell lines	2.28×10^{-3}	AKT3, CDKN2A, TGFB1	3
Cellular movement	Migration	Migration of carcinoma cell lines	2.91×10^{-3}	AKT3, TGFB1, WNT5A	3
Cellular movement	Migration	Migration of breast cell lines	3.64×10^{-3}	EGFR, LOX, TGFB1	3
Cellular movement	Migration	Migration of prostate cancer cell lines	7.40×10^{-3}	CDH1, CXCL12, FN1, NCOA3	4
Cellular movement	Cell movement	Cell movement of cell lines	1.68×10^{-8}	BMP7, CCR10, CD24, CDH2, CLDN3, CXCL12, DKK3, DNAJB4, EGFR, ESR1, FN1, FOSL1, FSCN1, KRT19, NRCAM, RET, RLN2, SERPINE1, SPARC, TFF3, TGFB1, WNT5A	22
Cellular movement	Cell movement	Cell movement of tumor cell lines	1.95×10^{-7}	BMP7, CD24, CDH2, CLDN3, CXCL12, DKK3, DNAJB4, EGFR, ESR1, FN1, FOSL1, KRT19, RLN2, SERPINE1, SPARC, TGFB1, WNT5A	17
Cellular movement	Cell movement	Cell movement of eukaryotic cells	5.03×10^{-7}	ANXA1, BDNF, BMP7, CCR10, CD24, CDH2, CLDN3, CXCL12, DKK3, DNAJB4, EGFR, ESR1, FN1, FOSL1, FSCN1, ICAM1, KRT19, MSN, NRCAM, RET, RLN2, SERPINA3, SERPINE1, SPARC, TFF3, TGFB1, TNFRSF6B, WNT5A	27
Cellular movement	Cell movement	Cell movement	5.31×10^{-7}	ANXA1, BDNF, BMP7, CALD1, CCR10, CD24, CDH2, CLDN3, CXCL12, CYR61, DKK3, DNAJB4, EGFR, ESR1, FN1, FOSL1, FSCN1, ICAM1, KRT19, MSN, NRCAM, RET, RLN2, SERPINA3, SERPINE1, SPARC, TFF3, TGFB1, TNFRSF6B, VIM, WNT5A	31
Cellular movement	Cell movement	Cell movement of lung cancer cell lines	3.94×10^{-4}	DNAJB4, FN1, FOSL1	3

Cellular movement	Cell movement	Cell movement of breast cancer cell lines	1.49×10^{-3}	<i>ESR1, KRT19, SERPINE1, SPARC, TGFB1</i>	5
Cellular movement	Cell movement	Cell movement of leukemia cells	3.07×10^{-3}	<i>CXCL12, TGFB1</i>	2
Cellular movement	Cell movement	Cell movement of lung cell lines	5.04×10^{-3}	<i>FOSL1, TFF3</i>	2
Cellular movement	Cell movement	Cell movement of pancreatic cancer cell lines	5.04×10^{-3}	<i>CDH2, TGFB1</i>	2
Cellular movement	Cell movement	Cell movement of carcinoma cell lines	1.35×10^{-2}	<i>DNAJB4, FOSL1</i>	2
Cellular movement	Scattering	Scattering of cell lines	1.06×10^{-5}	<i>EGFR, FN1, MSX2, RET, TFF3</i>	5
Cellular movement	Scattering	Scattering of tumor cell lines	3.94×10^{-4}	<i>EGFR, MSX2, RET</i>	3
Cellular movement	Scattering	Scattering of neuroblastoma cell lines	5.28×10^{-4}	<i>EGFR, RET</i>	2
Cellular movement	Scattering	Scattering of kidney cell lines	7.45×10^{-3}	<i>FN1, TFF3</i>	2
Cellular movement	Chemotaxis	Chemotaxis of granule cells	5.28×10^{-4}	<i>BDNF, CXCL12</i>	2
Cellular movement	Chemotaxis	Chemotaxis of rhabdomyosarcoma cell lines	1.56×10^{-3}	<i>CXCL12, FN1</i>	2
Cellular movement	Chemotaxis	Chemotaxis of microglia	5.04×10^{-3}	<i>TGFB1, TNFRSF6B</i>	2
Cellular movement	Chemotaxis	Chemotaxis	1.49×10^{-2}	<i>ANXA1, BDNF, CCR10, CXCL12, CYR61, EGFR, FN1, FOSL1, RLN2, SERPINA3, TFF3, TGFB1, TNFRSF6B</i>	13
Cellular movement	Locomotion	Locomotion of eukaryotic cells	5.28×10^{-4}	<i>CXCL12, TGFB1</i>	2
Cellular movement	Adhesion	Adhesion of t lymphocytes	2.12×10^{-3}	<i>CXCL12, FN1, ICAM1, ICAM2, RASGRP1</i>	5

Gastrointestinal disease	Colorectal cancer	Colorectal cancer	5.18×10^{-15}	<i>AKR1C2, AKT3, ANK3, BEX1, BMP7, BST2, CA2, CALD1, CDH2, CDKN2A, CLDN3, CLGN, COL4A1, COL4A2, COL6A1, COL6A2, CRIP1, CTGF, CXCL12, CYP1B1, EGFR, EMP1, EPCAM, FBN2 (includes EG:2201), FN1, FSCN1, FYN, GDF15, GJA1, GLUL, GREB1, HIST1H2BD, HIST1H4H (includes EG:8365), ICAM1, ID2, IGFBP5, INSIG1, IRS1, MAFB, MCAM, MLPH, NUAK1, RET, SERPINA3, SERPINE1, SERTAD2, SLC12A8, SLC16A3, SRPX, STXBP6, TFF3, TM4SF1, TPD52L1, TUBB2A</i>	54
Gastrointestinal disease	Pancreatic tumor	Pancreatic tumor	3.40×10^{-9}	<i>AXL, CDKN2A, CREBBP, CYR61, EGFR, FN1, IGFBP3, IGFBP5, KLF4, KRT7, KRT19, MSLN, MSX2, PIK3R1, RET, SERPINE2, TFF1, TUBB2A</i>	18
Gastrointestinal disease	Digestive organ tumor	Digestive organ tumor	3.56×10^{-9}	<i>ABCB1, AKT3, ANXA1, AXL, CA2, CDH1, CDKN2A, COL1A1, CREBBP, CYR61, EGFR, FN1, FSCN1, FYN, GDF15, GJA1, GSTP1, IGFBP3, IGFBP5, INSIG1, KLF4, KRT7, KRT19, MSLN, MSX2, NNMT, PIK3R1, RB1, RET, SERPINE2, TFF1, TM4SF1, TUBB2A, VIM, WWOX</i>	35
Gastrointestinal disease	Colon cancer	Colon cancer	8.99×10^{-6}	<i>AKT3, BMP7, BST2, CA2, CALD1, CLDN3, COL6A1, COL6A2, CRIP1, EGFR, FN1, FSCN1, GDF15, GJA1, HIST1H2BD, HIST1H4H (includes EG:8365), INSIG1, MCAM, SERPINA3, SERPINE1, STXBP6, TFF3, TPD52L1, TUBB2A</i>	24
Gastrointestinal disease	Pancreatic adenocarcinoma	Pancreatic adenocarcinoma	3.67×10^{-5}	<i>AXL, CYR61, IGFBP3, KRT7, KRT19, MSLN, SERPINE2</i>	7
Gastrointestinal disease	Pancreatic carcinoma	Pancreatic carcinoma	4.17×10^{-4}	<i>AXL, CYR61, IGFBP3, KRT7, KRT19, MSLN, MSX2, SERPINE2</i>	8
Gastrointestinal disease	Acinar-cell carcinoma	Acinar-cell carcinoma	1.21×10^{-3}	<i>CREBBP, FN1, IGFBP5, PIK3R1</i>	4
Gastrointestinal disease	Esophageal carcinoma	Esophageal carcinoma	1.21×10^{-3}	<i>EGFR, GSTP1, RB1, WWOX</i>	4
Gastrointestinal disease	Salivary gland tumor	Salivary gland tumor	1.92×10^{-3}	<i>CREBBP, FN1, IGFBP5, INSIG1, PIK3R1, RET</i>	6
Gastrointestinal disease	Mucoepidermoid carcinoma	Mucoepidermoid carcinoma	2.94×10^{-3}	<i>CREBBP, FN1, IGFBP5, INSIG1, PIK3R1</i>	5
Gastrointestinal disease	Esophageal cancer	Esophageal cancer	5.28×10^{-3}	<i>ADA, EGFR, GSTP1, RB1, RET, TUBB2A, WWOX</i>	7
Gastrointestinal disease	Cholangiocarcinoma	Cholangiocarcinoma	6.65×10^{-3}	<i>ANXA1, EGFR, TM4SF1, VIM</i>	4
Gastrointestinal disease	Colorectal tumor	Colorectal tumor	9.49×10^{-3}	<i>AKT3, EGFR, FSCN1, GDF15, GJA1, INSIG1, RET, TUBB2A</i>	8

Cellular development	Developmental process	Developmental process of tumor cell lines	3.60×10^{-11}	<i>ABCB1, AKT3, ANXA1, BMP7, CDH1, CDKN2A, CREBBP, CTGF, CXADR, CYR61, DKK3, DNAJB4, EGFR, EIF5A, ESR1, FOSL1, GDF15, GJA1, HMGA1, HMGA2, ID2, IFI16, IGFBP3, IGFBP5, IRS1, KLF4, MSX2, MUC1, MYB (includes EG:4602), MYL9 (includes EG:10398), NCOA3, NOV, PDCD10, PIK3R1, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, SNCA, TFAP2C, TGFB1, XBP1</i>	43
Cellular development	Developmental process	Developmental process of epithelial cells	5.97×10^{-7}	<i>ANXA1, AREG, BMP7, CDH1, COL4A1, CREBBP, EGFR, FHL2, FN1, ID2, RB1, TGFB1</i>	12
Cellular development	Developmental process	Developmental process of epidermal cells	1.76×10^{-4}	<i>ANXA1, CDH1, EGFR, FHL2, FN1, ID2, TGFB1</i>	7
Cellular development	Developmental process	Developmental process of fibroblast cell lines	3.35×10^{-4}	<i>BMP7, EGFR, FN1, IGFBP3, IRS1, MGP, NRCAM, RET, TGFB1</i>	9
Cellular development	Developmental process	Developmental process of keratinocytes	4.78×10^{-4}	<i>ANXA1, CDH1, EGFR, FN1, ID2, TGFB1</i>	6
Cellular development	Developmental process	Developmental process of cancer cells	4.95×10^{-4}	<i>CDKN2A, GFRA1, MCAM, MUC1, PEG10, RB1, TFF3, TGFB1</i>	8
Cellular development	Developmental process	Developmental process of tumor cells	5.67×10^{-4}	<i>CDKN2A, GFRA1, HMGA2, MCAM, MUC1, PEG10, RB1, TFF3, TGFB1</i>	9
Cellular development	Developmental process	Developmental process of kidney cell lines	1.31×10^{-3}	<i>BMP7, CALCR, ESR1, OXTR, TGFB1</i>	5
Cellular development	Developmental process	Developmental process of endothelial cells	1.44×10^{-3}	<i>AXL, FN1, IGFBP3, MGP, STC1, TGFB1</i>	6
Cellular development	Developmental process	Developmental process of carcinoma cell lines	1.54×10^{-3}	<i>AKT3, ANXA1, DNAJB4, IGFBP3, NOV, PIK3R1, TGFB1</i>	7
Cellular development	Developmental process	Developmental process of mononuclear leukocytes	2.78×10^{-3}	<i>CDKN2A, CXCL12, GATA3, GFRA1, IFI16, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B</i>	11

Cellular development	Developmental process	Developmental process of lung cancer cell lines	3.74×10^{-3}	<i>AKT3, ANXA1, DNAJB4, IGFBP3, PIK3R1, TGFB1</i>	6
Cellular development	Developmental process	Developmental process of prostate cancer cell lines	3.74×10^{-3}	<i>BMP7, CXADR, IFI16, IGFBP5, PDCD10, TGFB1</i>	6
Cellular development	Developmental process	Developmental process of myeloid leukemia cells	5.04×10^{-3}	<i>GFRA1, MUC1</i>	2
Cellular development	Developmental process	Developmental process of thymocytes	5.04×10^{-3}	<i>CDKN2A, IL7R</i>	2
Cellular development	Developmental process	Developmental process of epithelial cell lines	6.65×10^{-3}	<i>CALCR, ESR1, ID2, TGFB1</i>	4
Cellular development	Developmental process	Developmental process of blood cells	7.17×10^{-3}	<i>CDKN2A, CXCL12, EGFR, GATA3, GFRA1, IFI16, IL18, IL7R, MAFB, MUC1, MYB (includes EG:4602), RAC2, TGFB1, TNFRSF6B</i>	14
Cellular development	Developmental process	Developmental process of bone cancer cell lines	8.90×10^{-3}	<i>NOV, PLAGL1, SNCA</i>	3
Cellular development	Developmental process	Developmental process of fibroblasts	1.03×10^{-2}	<i>BMP7, CDKN2A, TGFB1</i>	3
Cellular development	Developmental process	Developmental process of muscle cells	1.68×10^{-2}	<i>IGFBP3, IGFBP5, RB1, TGFB1</i>	4
Cellular development	Growth	Growth of tumor cell lines	1.55×10^{-10}	<i>ABCB1, AKT3, ANXA1, CDH1, CDKN2A, CTGF, CXADR, CYR61, DKK3, DNAJB4, EGFR, EIF5A, ESR1, FOSL1, GDF15, GJA1, HMGA1, HMGA2, IFI16, IGFBP3, IGFBP5, IRS1, KLF4, MSX2, MYL9 (includes EG:10398), NCOA3, NOV, PDCD10, PIK3R1, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, TFAP2C, TGFB1, XBP1</i>	37
Cellular development	Growth	Growth of breast cancer cell lines	3.81×10^{-6}	<i>CDH1, CYR61, EGFR, ESR1, GJA1, HMGA1, IGFBP3, IGFBP5, IRS1, NCOA3, TFAP2C, TGFB1, XBP1</i>	13
Cellular development	Growth	Growth of neuroblastoma cell lines	2.29×10^{-4}	<i>EGFR, HMGA2, IGFBP5, RET</i>	4

Cellular development	Growth	Growth of epithelial cells	8.74×10^{-4}	AREG, EGFR, FHL2, ID2, RB1	5
Cellular development	Growth	Growth of cancer cells	1.20×10^{-3}	CDKN2A, GFRA1, MCAM, PEG10, RB1, TFF3, TGFB1	7
Cellular development	Growth	Growth of tumor cells	1.34×10^{-3}	CDKN2A, GFRA1, HMGA2, MCAM, PEG10, RB1, TFF3, TGFB1	8
Cellular development	Growth	Growth of fibroblast cell lines	1.66×10^{-3}	EGFR, FN1, IGFBP3, IRS1, NRCAM, RET, TGFB1	7
Cellular development	Growth	Growth of epidermal cells	5.41×10^{-3}	EGFR, FHL2, ID2	3
Cellular development	Growth	Growth of carcinoma cell lines	6.19×10^{-3}	AKT3, ANXA1, DNAJB4, IGFBP3, NOV, PIK3R1	6
Cellular development	Growth	Growth of endometrial cancer cell lines	7.45×10^{-3}	CYR61, ESR1	2
Cellular development	Growth	Growth of squamous cell carcinoma cell lines	1.03×10^{-2}	CDKN2A, EGFR	2
Cellular development	Growth	Growth of prostate cancer cell lines	1.14×10^{-2}	CXADR, IFI16, IGFBP5, PDCD10, TGFB1	5
Cellular development	Growth	Growth of lung cancer cell lines	1.48×10^{-2}	AKT3, ANXA1, DNAJB4, IGFBP3, PIK3R1	5
Cellular development	Growth	Growth of pancreatic cancer cell lines	1.52×10^{-2}	IGFBP3, MSX2, NCOA3	3
Cellular development	Differentiation	Differentiation of cells	6.22×10^{-6}	ANXA1, AREG, BMP7, CDH1, CDKN2A, COL4A1, CREBBP, CTGF, CXCL12, FN1, GATA3, ID2, IFI16, IGFBP3, IGFBP5, IGFBP6, IL18, KLF4, MAFB, MGP, MUC1, MYB (includes EG:4602), PIK3R1, PRAME (includes EG:23532), RB1, SNCA, STC1, TGFB1	28
Cellular development	Differentiation	Differentiation of cell lines	1.61×10^{-4}	BMP7, CREBBP, CTGF, FN1, ID2, IGFBP5, IGFBP6, MGP, MUC1, MYB (includes EG:4602), PIK3R1, PRAME (includes EG:23532), RB1, SNCA, TGFB1	15
Cellular development	Differentiation	Differentiation of T lymphocytes	2.81×10^{-4}	CDKN2A, CXCL12, GATA3, IL18, MUC1, TGFB1	6
Cellular development	Differentiation	Differentiation of epithelial cells	1.18×10^{-3}	ANXA1, CDH1, COL4A1, CREBBP, FN1, TGFB1	6

Cellular development	Differentiation	Differentiation of tumor cell lines	2.99×10^{-3}	<i>CREBBP, CTGF, IGFBP5, MUC1, MYB</i> (includes EG:4602), <i>PIK3R1, PRAME</i> (includes EG:23532), <i>RB1, SNCA, TGFB1</i>	10
Cellular development	Differentiation	Differentiation of mononuclear leukocytes	4.13×10^{-3}	<i>CDKN2A, CXCL12, GATA3, IFI16, IL18, MUC1, TGFB1</i>	7
Cellular development	Differentiation	Differentiation of th2 cells	5.04×10^{-3}	<i>GATA3, TGFB1</i>	2
Cellular development	Differentiation	Differentiation of keratinocytes	6.65×10^{-3}	<i>ANXA1, CDH1, FN1, TGFB1</i>	4
Cellular development	Differentiation	Differentiation of chondrocytes	1.03×10^{-2}	<i>STC1, TGFB1</i>	2
Cellular development	Differentiation	Differentiation of leukemia cell lines	1.28×10^{-2}	<i>CREBBP, MUC1, MYB</i> (includes EG:4602), <i>PIK3R1, RB1, TGFB1</i>	6
Cellular development	Differentiation	Differentiation of muscle cells	1.34×10^{-2}	<i>IGFBP3, RB1, TGFB1</i>	3
Cellular development	Differentiation	Differentiation of fibroblast cell lines	1.52×10^{-2}	<i>BMP7, MGP, TGFB1</i>	3
Cellular development	Differentiation	Differentiation of blood cells	1.69×10^{-2}	<i>CDKN2A, CXCL12, GATA3, IFI16, IL18, MAFB, MUC1, MYB</i> (includes EG:4602), <i>TGFB1</i>	9
Cellular development	Development	Development of cells	4.69×10^{-4}	<i>BDNF, BMP7, CDH1, CDKN2A, CREBBP, CXCL12, CYR61, EGFR, FN1, GATA3, ID2, IL18, IL7R, MAP7, MUC1, RAC2, RRAS, STC1, TGFB1, TNFRSF6B</i>	20
Cellular development	Development	T cell development	1.70×10^{-3}	<i>CDKN2A, CXCL12, GATA3, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B</i>	9
Cellular development	Development	Development of blood cells	2.13×10^{-3}	<i>CDKN2A, CXCL12, EGFR, GATA3, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B</i>	10
Cellular development	Epithelial-mesenchymal transition	Epithelial-mesenchymal transition of fibroblasts	5.28×10^{-4}	<i>BMP7, TGFB1</i>	2
Cellular development	Epithelial-mesenchymal transition	Epithelial-mesenchymal transition of hepatocytes	5.28×10^{-4}	<i>BMP7, TGFB1</i>	2
Cellular development	Epithelial-mesenchymal transition	Epithelial-mesenchymal transition of kidney cell lines	3.07×10^{-3}	<i>BMP7, TGFB1</i>	2

Cellular development	Transdifferentiation	Transdifferentiation of epithelial cells	1.56×10^{-3}	<i>COL4A1, TGFB1</i>	2
Cellular development	Transdifferentiation	Transdifferentiation of mesenchymal cells	1.56×10^{-3}	<i>COL4A1, TGFB1</i>	2
Cellular development	Apoptosis	Apoptosis of T lymphocytes	2.71×10^{-3}	<i>CDKN2A, CXCL12, IL7R, RAC2, TGFB1, TNFRSF6B</i>	6
Cellular development	Induction	Induction of T lymphocytes	3.07×10^{-3}	<i>IL18, MUC1</i>	2
Cellular development	Tubulation	Tubulation of endothelial cells	5.41×10^{-3}	<i>AXL, IGFBP3, MGP</i>	3
Cellular development	Shape change	Shape change of fibroblasts	7.45×10^{-3}	<i>CDKN2A, FN1</i>	2
Cellular development	Activation-induced cell death	Activation-induced cell death of T lymphocytes	1.35×10^{-2}	<i>RAC2, TGFB1</i>	2
Genetic disorder	Pancreatic tumor	Pancreatic tumor	3.40×10^{-9}	<i>AXL, CDKN2A, CREBBP, CYR61, EGFR, FN1, IGFBP3, IGFBP5, KLF4, KRT7, KRT19, MSLN, MSX2, PIK3R1, RET, SERPINE2, TFF1, TUBB2A</i>	18
Genetic disorder	Pancreatic cancer	Pancreatic cancer	6.91×10^{-8}	<i>AXL, CDH2, CDKN2A, CREBBP, CYR61, EGFR, FBN2 (includes EG:2201), FN1, FYN, IGFBP3, IGFBP5, KLF4, KRT7, KRT19, MSLN, MSX2, PIK3R1, RET, SERPINE2, TFF1, TUBB2A</i>	21
Genetic disorder	Non-small cell lung cancer	Non-small cell lung cancer	2.45×10^{-6}	<i>AKT3, B2M, CDH1, CDKN2A, DNAJB4, EGFR, EPCAM, FN1, FYN, MAFB, MSLN, RB1, RET, S100P, TUBB2A</i>	15
Genetic disorder	Prostate cancer	Prostate cancer	6.19×10^{-6}	<i>ABAT, AKT3, CDH1, COL1A1, CYP1B1, DNAJC15, EGFR, EPCAM, ESR1, FHL2, FYN, GATA3, GDF15, GPX3, IGFBP3, IL7R, KRT7, MMP1 (includes EG:4312), MSMB, MUC1, MYL9 (includes EG:10398), PSAT1, RB1, RET, TFF3, TUBB2A, WWOX</i>	27
Genetic disorder	Alzheimer disease	Alzheimer disease	7.95×10^{-6}	<i>ABAT, AKAP2, ANK3, BDNF, BIN1, CDC42EP3, CDKN2A, COL4A1, COL4A2, CTGF, CXCL12, DNAJC15, EML1 (includes EG:2009), ESR1, EXT1, GATA3, GREB1, GRHL2, HMGA1, ICAM1, IGFBP3, IGFBP6, LEPREL1, LPHN2, MAP1B, MMP1 (includes EG:4312), NEFH, PLTP, RAB3B, RNASET2, SCD, SELENBP1, SERPINA3, SERPINE2, SLC12A8, SNCA, SPARC, STXBP6, TGFB1, TGM2, TUBB2A, UBB, UCHL1, WWOX</i>	44
Genetic disorder	Colon cancer	Colon cancer	8.99×10^{-6}	<i>AKT3, BMP7, BST2, CA2, CALD1, CLDN3, COL6A1, COL6A2, CRIP1, EGFR, FN1, FSCN1, GDF15, GJA1, HIST1H2BD, HIST1H4H (includes EG:8365), INSIG1, MCAM, SERPINA3, SERPINE1, STXBP6, TFF3, TPD52L1, TUBB2A</i>	24

Genetic disorder	Prostatic intraepithelial neoplasia	Prostatic intraepithelial neoplasia	2.72×10^{-5}	<i>ESR1, FHL2, GATA3, GPX3, IGFBP3, KRT7, MUC1, MYL9</i> (includes EG:10398), <i>PSAT1, TFF3</i>	10
Genetic disorder	Genetic disorder	Genetic disorder	2.73×10^{-5}	<i>ABAT, ABCB1, ADA, AGA, AKAP2, AKAP12, AKR1B1, AKR1C2, AKT3, AMMECR1, ANK3, APPBP2, ATXN1, AXL, B2M, BCAS3, BCAT1, BDNF, BIN1, BMP7, BRIP1, BST2, C6ORF97, CA2, CALCR, CALD1, CD24, CDA, CDC42EP3, CDH1, CDH2, CDKN2A, CDS1, CELSR2, CHST6, CLDN3, COCH, COL1A1, COL21A1, COL4A1, COL4A2, COL6A1, COL6A2, CREBBP, CRIP1, CRYAB, CTGF, CXCL12, CYP1B1, CYR61, DFNA5, DKK3, DNAJB4, DNAJC15, ECM1, EFEMP2, EGFR, EGR3, EML1</i> (includes EG:2009), <i>ENOSF1, EPCAM, ESR1, EXT1, FBN2</i> (includes EG:2201), <i>FHL2, FN1, FNDC3B, FSCN1, FYN, GATA3, GDF15, GFRA1, GJA1, GNA14, GNAL, GPX3, GREB1, GRHL2, HEG1, HIST1H2BD, HIST1H4H</i> (includes EG:8365), <i>HIST2H2AA3, HMGA1, HMGA2, HNMT, ICAM1, ID4, IGF2BP2, IGFBP3, IGFBP5, IGFBP6, IL18, IL7R, INSIG1, IRS1, ISOC1, KIAA1324, KLF4, KRT7, KRT19, KYNU, LAMB1, LDHB, LEPREL1, LOX, LPCAT1, LPHN2, LXN, MAFB, MAP1B, MCAM, MLLT11, MLPH, MMP1</i> (includes EG:4312), <i>MPPED2</i> (includes EG:744), <i>MSLN, MSMB, MSX2, MT1E, MTUS1, MUC1, MYL9</i> (includes EG:10398), <i>NCOA3, NDN, NEBL, NEFH, NRCAM, NSMAF, OLFM1, ORC5L, PBX1, PDGFC, PDLIM4, PIK3R1, PIK3R3, PLAGL1, PLTP, PSAT1, PSMB9, PTRF, RAB3B, RAC2, RAI14, RAMP3, RASGRP1, RB1, RBM47, RBPMS, RCN1, RET, RNASET2, S100P, SCD, SCHIP1, SCUBE2, SDC2, SELENBP1, SEMA3C, SERPINA3, SERPINB7, SERPINE1, SERPINE2, SGCE, SH3YL1, SIDT1, SLC12A2, SLC12A8, SLC19A2, SLC24A3, SLC27A6, SLC6A14, SNCA, SNRPN, SOX3, SPARC, SRPX, STXBP6, TACSTD2, TFF1, TFF3, TFP12, TGFB1, TGM2, TMF1, TNFRSF6B, TPD52L1, TPM2, TUBB2A, TUBB2B, UBB, UCHL1, UGT2B15, VAV3, VIM, WWOX, XBP1, ZNF395</i>	198
Genetic disorder	Pancreatic adenocarcinoma	Pancreatic adenocarcinoma	3.67×10^{-5}	<i>AXL, CYR61, IGFBP3, KRT7, KRT19, MSLN, SERPINE2</i>	7
Genetic disorder	Non-small-cell lung carcinoma	Non-small-cell lung carcinoma	4.92×10^{-5}	<i>AKT3, B2M, CDKN2A, EGFR, EPCAM, FN1, FYN, MAFB, MSLN, RB1, RET, TUBB2A</i>	12
Genetic disorder	Prostatic carcinoma	Prostatic carcinoma	5.27×10^{-5}	<i>ABAT, AKT3, EGFR, EPCAM, ESR1, FHL2, FYN, GATA3, GDF15, GPX3, IGFBP3, KRT7, MSMB, MUC1, MYL9</i> (includes EG:10398), <i>PSAT1, RB1, RET, TFF3, TUBB2A</i>	20
Genetic disorder	Prostatic intraepithelial tumor	Prostatic intraepithelial tumor	8.92×10^{-5}	<i>FHL2, GATA3, GPX3, IGFBP3, KRT7, MUC1, MYL9</i> (includes EG:10398), <i>PSAT1, TFF3</i>	9

Genetic disorder	Lung carcinoma	Lung carcinoma	9.52×10^{-5}	<i>ABCB1, AKT3, B2M, CDKN2A, CLDN3, EGFR, EPCAM, FN1, FYN, LOX, MAFB, MSLN, RB1, RET, TUBB2A</i>	15
Genetic disorder	Pancreatic carcinoma	Pancreatic carcinoma	4.17×10^{-4}	<i>AXL, CYR61, IGFBP3, KRT7, KRT19, MSLN, MSX2, SERPINE2</i>	8
Genetic disorder	Osteoporosis	Osteoporosis	6.49×10^{-4}	<i>CA2, CALCR, COL1A1, ESR1, PDLIM4</i>	5
Genetic disorder	Lung cancer	Lung cancer	8.27×10^{-4}	<i>ABCB1, AKT3, B2M, CDH1, CDKN2A, CLDN3, DNAJB4, EGFR, EPCAM, FN1, FYN, LOX, MAFB, MSLN, RB1, RET, S100P, TUBB2A</i>	18
Genetic disorder	Light chain associated amyloidosis	Light chain-associated amyloidosis	9.96×10^{-4}	<i>CXCL12, IGFBP5, RB1, TGFB1</i>	4
Genetic disorder	Dupuytren contracture	Dupuytren contracture	1.20×10^{-3}	<i>COL1A1, COL21A1, COL4A1, COL4A2, COL6A1, COL6A2, MMP1 (includes EG:4312)</i>	7
Genetic disorder	Acinar-cell carcinoma	Acinar-cell carcinoma	1.21×10^{-3}	<i>CREBBP, FN1, IGFBP5, PIK3R1</i>	4
Genetic disorder	Serous ovarian carcinoma process	Serous ovarian carcinoma process	1.28×10^{-3}	<i>CLDN3, MSLN, MUC1</i>	3
Genetic disorder	Bethlem myopathy	Bethlem myopathy	1.56×10^{-3}	<i>COL6A1, COL6A2</i>	2
Genetic disorder	Cutis laxa, recessive type 1	Cutis laxa, recessive type 1	1.56×10^{-3}	<i>EFEMP2, LOX</i>	2
Genetic disorder	Ullrich congenital muscular dystrophy	Ullrich congenital muscular dystrophy	1.56×10^{-3}	<i>COL6A1, COL6A2</i>	2
Genetic disorder	Nonsyndromic hearing impairment	Nonsyndromic hearing impairment	2.28×10^{-3}	<i>COCH, DFNA5, GRHL2</i>	3
Genetic disorder	Gelatinous drop-like corneal dystrophy	Gelatinous drop-like corneal dystrophy	3.07×10^{-3}	<i>TACSTD2, TGFB1</i>	2
Genetic disorder	Osteoarthritis	Osteoarthritis	6.62×10^{-3}	<i>CRYAB, EFEMP2, IL18, MMP1 (includes EG:4312), TFPI2, TGFB1</i>	6
Genetic disorder	Congenital central hypoventilation syndrome	Congenital central hypoventilation syndrome	7.45×10^{-3}	<i>BDNF, RET</i>	2

Genetic disorder	Progressive supranuclear palsy	Progressive supranuclear palsy	9.07×10^{-3}	<i>ABAT, SERPINA3, TGM2, UBB</i>	4
Genetic disorder	Postmenopausal osteoporosis	Postmenopausal osteoporosis	1.35×10^{-2}	<i>CALCR, ESR1</i>	2
Cell death	Cell death	Cell death of eukaryotic cells	1.33×10^{-8}	<i>ABCB1, AKT3, ANKRD1, ANXA1, ATMIN, B2M, BDNF, BMP7, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), COL4A2, CRYAB, CTGF, CXCL12, CYR61, DKK3, DNAJC15, DUSP4, EGFR, EMP1, EMP3, ESR1, FHL2, FN1, GDF15, GPX1, HMGA2, ID2, IFI16, IGFBP3, IGFBP5, IGFBP6, IL18, IL7R, LGALS3BP, MSN, MUC1, MYB (includes EG:4602), NCOA3, NUAK1, PDZK1, PLAGL1, PRAME (includes EG:23532), RAC2, RB1, RET, S100P, SDC2, SNCA, SPARC, SRPX, TFF3, TGFB1, TGM2, TNFRSF6B, WWOX, XBP1</i>	60
Cell death	Cell death	Cell death of cell lines	2.32×10^{-8}	<i>ABCB1, AKT3, ANKRD1, ATMIN, B2M, BDNF, BMP7, CD24, CDH1, CDKN2A, CEACAM6 (includes EG:4680), CRYAB, CTGF, CXCL12, CYR61, DKK3, DNAJC15, DUSP4, EGFR, EMP1, EMP3, ESR1, FHL2, FN1, GDF15, ID2, IFI16, IGFBP3, IGFBP5, IGFBP6, IL18, LGALS3BP, MSN, MUC1, MYB (includes EG:4602), NCOA3, NUAK1, PDZK1, PLAGL1, PRAME (includes EG:23532), RB1, RET, S100P, SDC2, SNCA, SPARC, SRPX, TFF3, TGFB1, TGM2, TNFRSF6B, WWOX, XBP1</i>	53
Cell death	Cell death	Cell death of tumor cell lines	3.22×10^{-8}	<i>ABCB1, AKT3, ANKRD1, ATMIN, B2M, BDNF, BMP7, CD24, CDH1, CDKN2A, CEACAM6 (includes EG:4680), CRYAB, CTGF, CXCL12, CYR61, DKK3, DNAJC15, DUSP4, EGFR, ESR1, FN1, GDF15, IFI16, IGFBP3, IGFBP5, IGFBP6, LGALS3BP, MSN, MUC1, MYB (includes EG:4602), NCOA3, NUAK1, PDZK1, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, SNCA, SPARC, SRPX, TFF3, TGFB1, TGM2, TNFRSF6B, WWOX, XBP1</i>	47
Cell death	Cell death	Cell death	1.20×10^{-7}	<i>ABCB1, AKT3, ANKRD1, ANXA1, ATMIN, B2M, BDNF, BMP7, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), COL4A2, CREBBP, CRYAB, CTGF, CXCL12, CYR61, DKK3, DNAJC15, DUSP4, EGFR, EIF5A, EMP1, EMP3, ESR1, FHL2, FN1, GDF15, GPX1, GSTP1, HMGA2, ICAM2, ID2, IFI16, IGFBP3, IGFBP5, IGFBP6, IL18, IL7R, LGALS3BP, MSN, MUC1, MYB (includes EG:4602), NCOA3, NUAK1, PDZK1, PLAGL1, PRAME (includes EG:23532), RAC2, RB1, RET, S100P, SDC2, SNCA, SPARC, SRPX, TFF1, TFF3, TGFB1, TGM2, TNFRSF6B, TPD52L1, WWOX, XBP1, ZNF443</i>	67
Cell death	Cell death	Cell death of breast cancer cell lines	1.18×10^{-4}	<i>B2M, CDH1, CTGF, CYR61, DNAJC15, DUSP4, EGFR, ESR1, FN1, IGFBP3, IGFBP5, MUC1, NCOA3, TGFB1, XBP1</i>	15
Cell death	Cell death	Cell death of bone cancer cell lines	9.31×10^{-4}	<i>ATMIN, CDH1, CDKN2A, DKK3, IGFBP5, PLAGL1, SDC2</i>	7

Cell death	Cell death	Cell death of normal cells	1.23×10^{-3}	<i>ANXA1, BMP7, CDH1, CDH2, CDKN2A, COL4A2, CRYAB, CTGF, CXCL12, EGFR, FN1, GPX1, IL18, IL7R, RAC2, RB1, SNCA, TGFB1, TNFRSF6B</i>	19
Cell death	Cell death	Cell death of neuroblastoma cell lines	1.28×10^{-2}	<i>BDNF, IGFBP3, IGFBP5, RET, SNCA, TGFB1</i>	6
Cell death	Apoptosis	Apoptosis of tumor cell lines	1.31×10^{-7}	<i>AKT3, ANKRD1, ATMIN, B2M, BMP7, CD24, CDH1, CDKN2A, CEACAM6 (includes EG:4680), CRYAB, CTGF, CXCL12, CYR61, DKK3, DNAJC15, DUSP4, EGFR, ESR1, FN1, GDF15, IFI16, IGFBP3, IGFBP5, IGFBP6, LGALS3BP, MUC1, MYB (includes EG:4602), NCOA3, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, SPARC, SRPX, TFF3, TGFB1, TGM2, TNFRSF6B, WWOX, XBP1</i>	41
Cell death	Apoptosis	Apoptosis of eukaryotic cells	1.34×10^{-7}	<i>AKT3, ANKRD1, ANXA1, ATMIN, B2M, BMP7, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), COL4A2, CRYAB, CTGF, CXCL12, CYR61, DKK3, DNAJC15, DUSP4, EGFR, ESR1, FHL2, FN1, GDF15, HMGA2, ID2, IFI16, IGFBP3, IGFBP5, IGFBP6, IL18, IL7R, LGALS3BP, MUC1, MYB (includes EG:4602), NCOA3, PLAGL1, PRAME (includes EG:23532), RAC2, RB1, RET, SDC2, SNCA, SPARC, SRPX, TFF3, TGFB1, TGM2, TNFRSF6B, WWOX, XBP1</i>	51
Cell death	Apoptosis	Apoptosis	3.98×10^{-7}	<i>AKT3, ANKRD1, ANXA1, ATMIN, B2M, BMP7, CD24, CDH1, CDH2, CDKN2A, CEACAM6 (includes EG:4680), COL4A2, CREBBP, CRYAB, CTGF, CXCL12, CYR61, DKK3, DNAJC15, DUSP4, EGFR, EIF5A, ESR1, FHL2, FN1, GDF15, GPX1, GSTP1, HMGA2, ICAM2, ID2, IFI16, IGFBP3, IGFBP5, IGFBP6, IL18, IL7R, LGALS3BP, MUC1, MYB (includes EG:4602), NCOA3, PLAGL1, PRAME (includes EG:23532), RAC2, RB1, RET, SDC2, SNCA, SPARC, SRPX, TFF1, TFF3, TGFB1, TGM2, TNFRSF6B, TPD52L1, WWOX, XBP1, ZNF443</i>	59
Cell death	Apoptosis	Apoptosis of cell lines	5.22×10^{-7}	<i>AKT3, ANKRD1, ATMIN, B2M, BMP7, CD24, CDH1, CDKN2A, CEACAM6 (includes EG:4680), CRYAB, CTGF, CXCL12, CYR61, DKK3, DNAJC15, DUSP4, EGFR, ESR1, FHL2, FN1, GDF15, ID2, IFI16, IGFBP3, IGFBP5, IGFBP6, IL18, LGALS3BP, MUC1, MYB (includes EG:4602), NCOA3, PLAGL1, PRAME (includes EG:23532), RB1, RET, SDC2, SPARC, SRPX, TFF3, TGFB1, TGM2, TNFRSF6B, WWOX, XBP1</i>	44
Cell death	Apoptosis	Apoptosis of breast cancer cell lines	1.09×10^{-4}	<i>B2M, CDH1, CTGF, CYR61, DUSP4, EGFR, ESR1, FN1, IGFBP3, IGFBP5, MUC1, NCOA3, TGFB1, XBP1</i>	14
Cell death	Apoptosis	Apoptosis of normal cells	9.54×10^{-4}	<i>ANXA1, BMP7, CDH1, CDH2, CDKN2A, COL4A2, CRYAB, CTGF, CXCL12, EGFR, FN1, IL7R, RAC2, RB1, SNCA, TGFB1, TNFRSF6B</i>	17
Cell death	Apoptosis	Apoptosis of prostate cancer cell lines	1.44×10^{-3}	<i>BMP7, CDKN2A, DKK3, EGFR, GDF15, IGFBP3, IGFBP5, NCOA3</i>	8

Cell death	Apoptosis	Apoptosis of epithelial cells	1.44×10^{-3}	<i>CDH1, CRYAB, EGFR, FN1, RB1, TGFB1</i>	6
Cell death	Apoptosis	Apoptosis of bone cancer cell lines	1.75×10^{-3}	<i>ATMIN, CDH1, CDKN2A, IGFBP5, PLAGL1, SDC2</i>	6
Cell death	Apoptosis	Apoptosis of t lymphocytes	2.71×10^{-3}	<i>CDKN2A, CXCL12, IL7R, RAC2, TGFB1, TNFRSF6B</i>	6
Cell death	Apoptosis	Apoptosis of carcinoma cell lines	4.04×10^{-3}	<i>AKT3, CDKN2A, DNAJC15, IGFBP3, IGFBP6, TGFB1</i>	6
Cell death	Apoptosis	Apoptosis of glomerular cells	7.45×10^{-3}	<i>CTGF, TGFB1</i>	2
Cell death	Apoptosis	Apoptosis of lung cancer cell lines	9.08×10^{-3}	<i>AKT3, CDKN2A, IGFBP3, IGFBP6, RB1, WWOX</i>	6
Cell death	Apoptosis	Apoptosis of embryonic cells	1.03×10^{-2}	<i>BMP7, TNFRSF6B</i>	2
Cell death	Apoptosis	Apoptosis of bladder cancer cell lines	1.71×10^{-2}	<i>EGFR, IFI16</i>	2
Cell death	Survival	Survival of cell lines	2.69×10^{-4}	<i>ABCB1, AXL, BDNF, CA2, CDH1, CDKN2A, CLDN3, CXCL12, EGFR, HMGA1, ID2, IGFBP3, KRT19, MCAM, NUAK1, PPP1R3D, PTPRK, S100P, SNCA, TGFB1, TGM2</i>	21
Cell death	Survival	Survival of eukaryotic cells	6.12×10^{-4}	<i>ABCB1, AXL, BDNF, CA2, CDH1, CDKN2A, CLDN3, CXCL12, EGFR, FN1, HMGA1, ICAM1, ID2, IGFBP3, KRT19, MCAM, NUAK1, PPP1R3D, PTPRK, S100P, SNCA, TGFB1, TGM2</i>	23
Cell death	Survival	Survival of tumor cell lines	6.95×10^{-4}	<i>AXL, BDNF, CA2, CDH1, CDKN2A, CLDN3, CXCL12, EGFR, HMGA1, ID2, IGFBP3, KRT19, MCAM, NUAK1, PPP1R3D, PTPRK, SNCA, TGM2</i>	18
Cell death	Survival	Survival of neuroblastoma cell lines	1.74×10^{-3}	<i>BDNF, SNCA, TGM2</i>	3
Cell death	Survival	Survival of breast cancer cell lines	3.59×10^{-3}	<i>CA2, HMGA1, ID2, IGFBP3, KRT19</i>	5
Cell death	Anoikis	Anoikis of dermal cells	5.28×10^{-4}	<i>EGFR, FN1</i>	2
Cell death	Cell viability	Cell viability of eukaryotic cells	9.47×10^{-4}	<i>CDH2, CXCL12, FKBP1B, FN1, GDF15, GSTP1, IGFBP3, PBX1, SNCA, STC1</i>	10
Cell death	Cell viability	Cell viability of cell lines	1.23×10^{-2}	<i>CDH2, FKBP1B, GDF15, GSTP1, PBX1, SNCA, STC1</i>	7

Cell death	Cell viability	Cell viability of tumor cell lines	1.51×10^{-2}	<i>CDH2, GDF15, GSTP1, PBX1, SNCA, STC1</i>	6
Cell death	Activation-induced cell death	Activation-induced cell death of T lymphocytes	1.35×10^{-2}	<i>RAC2, TGFB1</i>	2
Cell-cell signaling and interaction	Adhesion	Adhesion of cells	2.75×10^{-8}	<i>ANXA1, ANXA9, BMP7, CDH1, CDH2, CTGF, CXCL12, CYR61, DKK3, EGFR, EPCAM, FN1, FXYD5, ICAM1, ICAM2, IL18, IL32, LAMB1, LGALS3BP, LOXL2, MCAM, MPZL2, PLTP, PTPRK, RASGRP1, RLN2, RRAS, SDC2, SERPINE1, TGFB1, TGFB1I1, TGFB1I, TGM2, ZYX</i>	34
Cell-cell signaling and interaction	Adhesion	Adhesion of eukaryotic cells	6.91×10^{-7}	<i>ANXA1, BMP7, CDH1, CDH2, CTGF, CXCL12, CYR61, DKK3, EGFR, EPCAM, FN1, FXYD5, ICAM1, ICAM2, LGALS3BP, MCAM, PLTP, RASGRP1, RLN2, RRAS, SDC2, TGFB1, TGM2</i>	23
Cell-cell signaling and interaction	Adhesion	Adhesion of cell lines	6.27×10^{-6}	<i>ANXA1, BMP7, CDH1, CDH2, CXCL12, DKK3, EGFR, FN1, FXYD5, ICAM1, LGALS3BP, RLN2, RRAS, SDC2, TGFB1, TGM2</i>	16
Cell-cell signaling and interaction	Adhesion	Adhesion of cancer cells	1.34×10^{-5}	<i>CDH2, CXCL12, EGFR, FN1, ICAM1, MCAM</i>	6
Cell-cell signaling and interaction	Adhesion	Adhesion of tumor cell lines	8.77×10^{-5}	<i>ANXA1, BMP7, CDH1, CDH2, CXCL12, DKK3, EGFR, FN1, FXYD5, ICAM1, RLN2, SDC2</i>	12
Cell-cell signaling and interaction	Adhesion	Adhesion of normal cells	1.37×10^{-4}	<i>ANXA1, CDH1, CDH2, CTGF, CXCL12, CYR61, EPCAM, FN1, ICAM1, ICAM2, PLTP, RASGRP1, RLN2, TGFB1</i>	14
Cell-cell signaling and interaction	Adhesion	Adhesion of lymphocytes	1.07×10^{-3}	<i>CXCL12, FN1, ICAM1, ICAM2, RASGRP1, TGFB1</i>	6
Cell-cell signaling and interaction	Adhesion	Adhesion of leukocytes	1.19×10^{-3}	<i>ANXA1, CXCL12, FN1, ICAM1, ICAM2, PLTP, RASGRP1, RLN2, TGFB1</i>	9
Cell-cell signaling and interaction	Adhesion	Adhesion of T lymphocytes	2.12×10^{-3}	<i>CXCL12, FN1, ICAM1, ICAM2, RASGRP1</i>	5
Cell-cell signaling and interaction	Adhesion	Adhesion of epithelial cells	3.18×10^{-3}	<i>CDH1, EPCAM, FN1, ICAM1</i>	4
Cell-cell signaling and interaction	Adhesion	Adhesion of endothelial cells	4.35×10^{-3}	<i>CDH2, FN1, ICAM1, ICAM2, TGFB1</i>	5

Cell–cell signaling and interaction	Adhesion	Adhesion of connective tissue cells	5.41×10^{-3}	<i>CDH2, CTGF, ICAM1</i>	3
Cell–cell signaling and interaction	Adhesion	Adhesion of embryonic cell lines	5.41×10^{-3}	<i>FN1, ICAM1, RRAS</i>	3
Cell–cell signaling and interaction	Adhesion	Adhesion of endothelial cell lines	5.68×10^{-3}	<i>ANXA1, CDH2, CXCL12, ICAM1, TGFB1</i>	5
Cell–cell signaling and interaction	Adhesion	Adhesion of lung cancer cell lines	7.45×10^{-3}	<i>CXCL12, ICAM1</i>	2
Cell–cell signaling and interaction	Adhesion	Adhesion of leukemia cell lines	1.10×10^{-2}	<i>CDH1, FN1, ICAM1, RLN2</i>	4
Cell–cell signaling and interaction	Adhesion	Adhesion of leukocyte cell lines	1.34×10^{-2}	<i>ANXA1, TGFB1, TGM2</i>	3
Cell–cell signaling and interaction	Adhesion	Adhesion of melanoma cells	1.35×10^{-2}	<i>CDH2, MCAM</i>	2
Cell–cell signaling and interaction	Adhesion	Adhesion of keratinocytes	1.71×10^{-2}	<i>CDH1, FN1</i>	2
Cell–cell signaling and interaction	Binding	Binding of cells	8.98×10^{-6}	<i>AXL, CALCR, CD24, CEACAM5 (includes EG:1048), CXADR, CXCL12, FN1, GFRA1, ICAM1, IGFBP3, IGFBP5, LGALS3BP, MSN, OXTR, PIK3R1, SDC2, SERPINE1, SPARC, TGFB1, TGM2</i>	20
Cell–cell signaling and interaction	Binding	Binding of cell lines	1.07×10^{-5}	<i>AXL, CALCR, CD24, CEACAM5 (includes EG:1048), CXADR, CXCL12, FN1, ICAM1, IGFBP3, IGFBP5, LGALS3BP, OXTR, PIK3R1, SDC2, SERPINE1, TGFB1</i>	16
Cell–cell signaling and interaction	Binding	Binding of eukaryotic cells	1.52×10^{-5}	<i>AXL, CALCR, CD24, CEACAM5 (includes EG:1048), CXADR, CXCL12, FN1, GFRA1, ICAM1, IGFBP3, IGFBP5, LGALS3BP, MSN, OXTR, PIK3R1, SDC2, SERPINE1, SPARC, TGFB1</i>	19
Cell–cell signaling and interaction	Binding	Binding of fibroblasts	3.16×10^{-5}	<i>IGFBP3, LGALS3BP, SDC2, SPARC</i>	4

Cell-cell signaling and interaction	Binding	Binding of tumor cell lines	1.11×10^{-3}	<i>CEACAM5</i> (includes EG:1048), <i>CXADR</i> , <i>CXCL12</i> , <i>IGFBP5</i> , <i>LGALS3BP</i> , <i>OXTR</i> , <i>PIK3R1</i> , <i>SERPINE1</i> , <i>TGFB1</i>	9
Cell-cell signaling and interaction	Binding	Binding of normal cells	2.35×10^{-3}	<i>CD24</i> , <i>CXCL12</i> , <i>FN1</i> , <i>ICAM1</i> , <i>IGFBP3</i> , <i>LGALS3BP</i> , <i>MSN</i> , <i>SDC2</i> , <i>SERPINE1</i> , <i>SPARC</i>	10
Cell-cell signaling and interaction	Binding	Binding of breast cancer cell lines	5.41×10^{-3}	<i>IGFBP5</i> , <i>OXTR</i> , <i>SERPINE1</i>	3
Cell-cell signaling and interaction	Binding	Binding of blood platelets	1.18×10^{-2}	<i>CD24</i> , <i>FN1</i> , <i>ICAM1</i>	3
Cell-cell signaling and interaction	Binding	Binding of endothelial cell lines	1.43×10^{-2}	<i>AXL</i> , <i>FN1</i> , <i>ICAM1</i> , <i>PIK3R1</i>	4
Cell-cell signaling and interaction	Binding	Binding of cancer cells	1.71×10^{-2}	<i>CD24</i> , <i>GFRA1</i>	2
Cell-cell signaling and interaction	Activation	Activation of cells	3.16×10^{-5}	<i>ADA</i> , <i>AREG</i> , <i>B2M</i> , <i>CD24</i> , <i>CDH1</i> , <i>CEACAM5</i> (includes EG:1048), <i>CXCL12</i> , <i>EGFR</i> , <i>FN1</i> , <i>ICAM1</i> , <i>ICAM2</i> , <i>MMP1</i> (includes EG:4312), <i>NUAK1</i> , <i>RLN2</i> , <i>SDC2</i> , <i>TGFB1</i> , <i>TGM2</i> , <i>TNFRSF6B</i>	18
Cell-cell signaling and interaction	Activation	Activation of cell lines	4.34×10^{-4}	<i>B2M</i> , <i>CDH1</i> , <i>MMP1</i> (includes EG:4312), <i>NUAK1</i> , <i>TGFB1</i> , <i>TGM2</i> , <i>TNFRSF6B</i>	7
Cell-cell signaling and interaction	Activation	Activation of eukaryotic cells	4.93×10^{-4}	<i>ADA</i> , <i>AREG</i> , <i>B2M</i> , <i>CDH1</i> , <i>CEACAM5</i> (includes EG:1048), <i>CXCL12</i> , <i>EGFR</i> , <i>FN1</i> , <i>MMP1</i> (includes EG:4312), <i>NUAK1</i> , <i>RLN2</i> , <i>SDC2</i> , <i>TGFB1</i> , <i>TGM2</i> , <i>TNFRSF6B</i>	15
Cell-cell signaling and interaction	Activation	Activation of muscle cells	1.56×10^{-3}	<i>TGFB1</i> , <i>TGM2</i>	2
Cell-cell signaling and interaction	Activation	Activation of kidney cell lines	1.03×10^{-2}	<i>TGFB1</i> , <i>TGM2</i>	2
Cell-cell signaling and interaction	Activation	Activation of leukocyte cell lines	1.35×10^{-2}	<i>B2M</i> , <i>TNFRSF6B</i>	2
Cell-cell signaling and interaction	Activation	Activation of epithelial cells	1.71×10^{-2}	<i>AREG</i> , <i>EGFR</i>	2

Cell-cell signaling and interaction	Attachment	Attachment of eukaryotic cells	1.80×10^{-4}	<i>EGFR, FN1, SERPINE1, TFP12, TGFB1, TGFB1</i>	6
Cell-cell signaling and interaction	Attachment	Attachment of fibroblasts	1.56×10^{-3}	<i>FN1, TGFB1</i>	2
Cell-cell signaling and interaction	Attachment	Attachment of cell lines	2.76×10^{-3}	<i>EGFR, FN1, SERPINE1, TGFB1</i>	4
Cell-cell signaling and interaction	Attachment	Attachment of normal cells	6.46×10^{-3}	<i>FN1, TFP12, TGFB1</i>	3
Cell-cell signaling and interaction	Attachment	Attachment of breast cancer cell lines	7.45×10^{-3}	<i>EGFR, SERPINE1</i>	2
Cell-cell signaling and interaction	Attachment	Attachment of tumor cell lines	8.90×10^{-3}	<i>EGFR, SERPINE1, TGFB1</i>	3
Cell-cell signaling and interaction	Contact growth inhibition	Contact growth inhibition	1.99×10^{-4}	<i>BMP7, CD24, CDH1, CDKN2A, EGFR, IFI16, TGFB1</i>	7
Cell-cell signaling and interaction	Contact growth inhibition	Contact growth inhibition of eukaryotic cells	6.87×10^{-4}	<i>BMP7, CD24, CDH1, EGFR, IFI16, TGFB1</i>	6
Cell-cell signaling and interaction	Contact growth inhibition	Contact growth inhibition of cell lines	2.12×10^{-3}	<i>BMP7, CD24, CDH1, EGFR, IFI16</i>	5
Cell-cell signaling and interaction	Contact growth inhibition	Contact growth inhibition of tumor cell lines	5.30×10^{-3}	<i>BMP7, CD24, CDH1, IFI16</i>	4
Cell-cell signaling and interaction	Contact growth inhibition	Contact growth inhibition of colon cancer cell lines	1.03×10^{-2}	<i>CD24, CDH1</i>	2
Cell-cell signaling and interaction	Formation	Formation of cell-matrix contacts	1.56×10^{-3}	<i>CDH1, FN1</i>	2
Cell-cell signaling and interaction	Quantity	Quantity of intercellular junctions	1.56×10^{-3}	<i>CDH1, GDF15</i>	2

Cell-cell signaling and interaction	Induction	Induction of T lymphocytes	3.07×10^{-3}	<i>IL18, MUC1</i>	2
Cell-cell signaling and interaction	Association	Association of cells	7.45×10^{-3}	<i>IGFBP3, TGFB1</i>	2
Cell-cell signaling and interaction	Co-stimulation	Co-stimulation of T lymphocytes	1.35×10^{-2}	<i>CD24, FN1</i>	2
Cell-cell signaling and interaction	Detachment	Detachment of eukaryotic cells	1.52×10^{-2}	<i>ANXA1, FN1, IGFBP3</i>	3
Cell-cell signaling and interaction	Communication	Communication of cells	1.52×10^{-2}	<i>AREG, BST2, CEACAM6 (includes EG:4680), ESR1, GDF15, GJA1, IL18, STC1, TFAP2C, WNT5A, ZYX</i>	11
Tissue development	Adhesion	Adhesion of cells	2.75×10^{-8}	<i>ANXA1, ANXA9, BMP7, CDH1, CDH2, CTGF, CXCL12, CYR61, DKK3, EGFR, EPCAM, FN1, FXYD5, ICAM1, ICAM2, IL18, IL32, LAMB1, LGALS3BP, LOXL2, MCAM, MPZL2, PLTP, PTPRK, RASGRP1, RLN2, RRAS, SDC2, SERPINE1, TGFB1, TGFB1I1, TGFBI, TGM2, ZYX</i>	34
Tissue development	Adhesion	Adhesion of eukaryotic cells	6.91×10^{-7}	<i>ANXA1, BMP7, CDH1, CDH2, CTGF, CXCL12, CYR61, DKK3, EGFR, EPCAM, FN1, FXYD5, ICAM1, ICAM2, LGALS3BP, MCAM, PLTP, RASGRP1, RLN2, RRAS, SDC2, TGFB1, TGM2</i>	23
Tissue development	Adhesion	Adhesion of cell lines	6.27×10^{-6}	<i>ANXA1, BMP7, CDH1, CDH2, CXCL12, DKK3, EGFR, FN1, FXYD5, ICAM1, LGALS3BP, RLN2, RRAS, SDC2, TGFB1, TGM2</i>	16
Tissue development	Adhesion	Adhesion of cancer cells	1.34×10^{-5}	<i>CDH2, CXCL12, EGFR, FN1, ICAM1, MCAM</i>	6
Tissue development	Adhesion	Adhesion of tumor cell lines	8.77×10^{-5}	<i>ANXA1, BMP7, CDH1, CDH2, CXCL12, DKK3, EGFR, FN1, FXYD5, ICAM1, RLN2, SDC2</i>	12
Tissue development	Adhesion	Adhesion of normal cells	1.37×10^{-4}	<i>ANXA1, CDH1, CDH2, CTGF, CXCL12, CYR61, EPCAM, FN1, ICAM1, ICAM2, PLTP, RASGRP1, RLN2, TGFB1</i>	14
Tissue development	Adhesion	Adhesion of lymphocytes	1.07×10^{-3}	<i>CXCL12, FN1, ICAM1, ICAM2, RASGRP1, TGFB1</i>	6
Tissue development	Adhesion	Adhesion of leukocytes	1.19×10^{-3}	<i>ANXA1, CXCL12, FN1, ICAM1, ICAM2, PLTP, RASGRP1, RLN2, TGFB1</i>	9
Tissue development	Adhesion	Adhesion of T lymphocytes	2.12×10^{-3}	<i>CXCL12, FN1, ICAM1, ICAM2, RASGRP1</i>	5
Tissue development	Adhesion	Adhesion of epithelial cells	3.18×10^{-3}	<i>CDH1, EPCAM, FN1, ICAM1</i>	4

Tissue development	Adhesion	Adhesion of endothelial cells	4.35×10^{-3}	<i>CDH2, FN1, ICAM1, ICAM2, TGFB1</i>	5
Tissue development	Adhesion	Adhesion of connective tissue cells	5.41×10^{-3}	<i>CDH2, CTGF, ICAM1</i>	3
Tissue development	Adhesion	Adhesion of embryonic cell lines	5.41×10^{-3}	<i>FN1, ICAM1, RRAS</i>	3
Tissue development	Adhesion	Adhesion of endothelial cell lines	5.68×10^{-3}	<i>ANXA1, CDH2, CXCL12, ICAM1, TGFB1</i>	5
Tissue development	Adhesion	Adhesion of lung cancer cell lines	7.45×10^{-3}	<i>CXCL12, ICAM1</i>	2
Tissue development	Adhesion	Adhesion of leukemia cell lines	1.10×10^{-2}	<i>CDH1, FN1, ICAM1, RLN2</i>	4
Tissue development	Adhesion	Adhesion of leukocyte cell lines	1.34×10^{-2}	<i>ANXA1, TGFB1, TGM2</i>	3
Tissue development	Adhesion	Adhesion of melanoma cells	1.35×10^{-2}	<i>CDH2, MCAM</i>	2
Tissue development	Adhesion	Adhesion of keratinocytes	1.71×10^{-2}	<i>CDH1, FN1</i>	2
Tissue development	Aggregation	Aggregation of tumor cell lines	4.71×10^{-4}	<i>BMP7, CDH1, CEACAM5 (includes EG:1048), ICAM1, TGFB1</i>	5
Tissue development	Aggregation	Aggregation of tumor cells	1.56×10^{-3}	<i>CDH2, CEACAM5 (includes EG:1048)</i>	2
Tissue development	Aggregation	Aggregation of cells	2.42×10^{-3}	<i>BMP7, CDH1, CDH2, CEACAM5 (includes EG:1048), EGFR, EPCAM, ICAM1, TGFB1</i>	8
Tissue development	Aggregation	Aggregation of lymphoma cell lines	5.04×10^{-3}	<i>ICAM1, TGFB1</i>	2
Tissue development	Aggregation	Aggregation of colon cancer cell lines	7.45×10^{-3}	<i>CDH1, CEACAM5 (includes EG:1048)</i>	2
Tissue development	Aggregation	Aggregation of fibroblast cell lines	7.45×10^{-3}	<i>CDH2, EPCAM</i>	2
Tissue development	Formation	Formation of endothelial tube	2.37×10^{-3}	<i>COL4A1, FN1, SERPINE1, TGFB1</i>	4

Cell morphology	Morphology	Morphology of eukaryotic cells	4.26×10^{-7}	<i>BMP7, CDH1, CDKN2A, DKK3, EGFR, FN1, FYN, GDF15, IGFBP3, IL18, PEG10, PLAGL1, RB1, SDC2, SERPINE1, TGFB1, VIM</i>	17
Cell morphology	Morphology	Morphology of cells	6.84×10^{-7}	<i>BMP7, CDH1, CDKN2A, DKK3, EGFR, FN1, FYN, GDF15, IGFBP3, IL18, MUC1, PEG10, PLAGL1, RB1, SDC2, SERPINE1, TGFB1, VIM</i>	18
Cell morphology	Morphology	Morphology of cell lines	3.86×10^{-6}	<i>BMP7, CDH1, CDKN2A, DKK3, EGFR, FYN, GDF15, IGFBP3, PEG10, RB1, SDC2, SERPINE1, TGFB1, VIM</i>	14
Cell morphology	Morphology	Morphology of tumor cell lines	1.57×10^{-4}	<i>BMP7, CDH1, CDKN2A, DKK3, EGFR, GDF15, RB1, SDC2, VIM</i>	9
Cell morphology	Morphology	Morphology of bone cancer cell lines	5.04×10^{-3}	<i>DKK3, RB1</i>	2
Cell morphology	Morphology	Morphology of cancer cells	1.03×10^{-2}	<i>EGFR, PLAGL1</i>	2
Cell morphology	Morphology	Morphology of embryonic cell lines	1.35×10^{-2}	<i>IGFBP3, SERPINE1</i>	2
Cell morphology	Morphology	Morphology of gonadal cell lines	1.35×10^{-2}	<i>PEG10, SERPINE1</i>	2
Cell morphology	Mineralization	Mineralization of eukaryotic cells	1.74×10^{-3}	<i>AXL, SNCA, TGFB1</i>	3
Cell morphology	Mineralization	Mineralization of cell lines	7.45×10^{-3}	<i>SNCA, TGFB1</i>	2
Cell morphology	Polarity	Polarity of eukaryotic cells	5.04×10^{-3}	<i>CXCL12, EGFR</i>	2
Cell morphology	Tubulation	Tubulation of endothelial cells	5.41×10^{-3}	<i>AXL, IGFBP3, MGP</i>	3
Cell morphology	Tubulation	Tubulation of eukaryotic cells	1.68×10^{-2}	<i>AXL, CXCL12, IGFBP3, MGP</i>	4
Cell morphology	Length	Length of neurites	7.45×10^{-3}	<i>BDNF, ESR1</i>	2
Cell morphology	Shape change	Shape change of fibroblasts	7.45×10^{-3}	<i>CDKN2A, FN1</i>	2
Cell morphology	Polarization	Polarization of leukocytes	7.62×10^{-3}	<i>CXCL12, FN1, TGFB1</i>	3
Respiratory disease	Non-small cell lung cancer	Non-small cell lung cancer	2.45×10^{-6}	<i>AKT3, B2M, CDH1, CDKN2A, DNAJB4, EGFR, EPCAM, FN1, FYN, MAFB, MSLN, RB1, RET, S100P, TUBB2A</i>	15

Respiratory disease	Respiratory disorder	Respiratory disorder	1.36×10^{-5}	<i>BDNF, CD24, CTSZ</i> (includes EG:1522), <i>EGFR, ESR1, FN1, GATA3, GLUL, HIST1H2AC, HIST1H2BD, HIST1H2BK, HIST2H2AA3, HIST2H2BE, HMGA2, HNMT, IGFBP3, IL18, MGP, PDGFC, RET, S100P, SPARC, TGFB1, XBP1, ZYX</i>	25
Respiratory disease	Severe acute respiratory syndrome	Severe acute respiratory syndrome	2.05×10^{-5}	<i>CD24, CTSZ</i> (includes EG:1522), <i>GLUL, HIST1H2AC, HIST1H2BD, HIST1H2BK, HIST2H2AA3, HIST2H2BE, PDGFC, S100P, SPARC, ZYX</i>	12
Respiratory disease	Non-small-cell lung carcinoma	Non-small-cell lung carcinoma	4.92×10^{-5}	<i>AKT3, B2M, CDKN2A, EGFR, EPCAM, FN1, FYN, MAFB, MSLN, RB1, RET, TUBB2A</i>	12
Respiratory disease	Lung carcinoma	Lung carcinoma	9.52×10^{-5}	<i>ABCB1, AKT3, B2M, CDKN2A, CLDN3, EGFR, EPCAM, FN1, FYN, LOX, MAFB, MSLN, RB1, RET, TUBB2A</i>	15
Respiratory disease	Lung cancer	Lung cancer	8.27×10^{-4}	<i>ABCB1, AKT3, B2M, CDH1, CDKN2A, CLDN3, DNAJB4, EGFR, EPCAM, FN1, FYN, LOX, MAFB, MSLN, RB1, RET, S100P, TUBB2A</i>	18
Respiratory disease	Congenital central hypoventilation syndrome	Congenital central hypoventilation syndrome	7.45×10^{-3}	<i>BDNF, RET</i>	2
Neurological disease	Alzheimer disease	Alzheimer disease	7.95×10^{-6}	<i>ABAT, AKAP2, ANK3, BDNF, BIN1, CDC42EP3, CDKN2A, COL4A1, COL4A2, CTGF, CXCL12, DNAJC15, EML1</i> (includes EG:2009), <i>ESR1, EXT1, GATA3, GREB1, GRHL2, HMGA1, ICAM1, IGFBP3, IGFBP6, LEPREL1, LPHN2, MAP1B, MMP1</i> (includes EG:4312), <i>NEFH, PLTP, RAB3B, RNASET2, SCD, SELENBP1, SERPINA3, SERPINE2, SLC12A8, SNCA, SPARC, STXBP6, TGFB1, TGM2, TUBB2A, UBB, UCHL1, WWOX</i>	44
Neurological disease	Neurological disorder	Neurological disorder of rodents	6.49×10^{-4}	<i>ATXN1, BDNF, IL18, SNCA, TGFB1</i>	5
Neurological disease	Neurological disorder	Neurological disorder of mice	1.72×10^{-3}	<i>ATXN1, BDNF, SNCA, TGFB1</i>	4
Neurological disease	Neurological disorder	Neurological disorder	1.93×10^{-3}	<i>ABAT, ABCB1, AGA, AKAP2, AKAP12, AKR1B1, AKR1C2, AKT3, ANK3, ATXN1, BCAS3, BDNF, BIN1, C6ORF97, CA2, CD24, CDC42EP3, CDH2, CDKN2A, CDS1, COCH, COL4A1, COL4A2, CREBBP, CRYAB, CTGF, CXCL12, DFNA5, DKK3, DNAJC15, DUSP4, ECM1, EGFR, EGR3, EML1</i> (includes EG:2009), <i>ENOSF1, ESR1, EXT1, FHL2, FNDC3B, FYN, GATA3, GFRA1, GJA1, GNA14, GNAL, GREB1, GRHL2, HIST2H2AA3, HMGA1, ICAM1, ID4, IGFBP3, IGFBP6, IL18, IL7R, ISOC1, LAMB1, LDHB, LEPREL1, LPCAT1, LPHN2, MAP1B, MGP, MLLT11, MMP1</i> (includes EG:4312), <i>MREG, MT1E, MUC1, NCOA3, NDN, NEFH, NSMAF, OLFM1, ORC5L, PBX1, PIK3R1, PLTP, PSMB9, RAB3B, RAI14, RAMP3, RCN1, RET, RNASET2, S100P, SCD, SCHIP1, SCUBE2, SDC2, SELENBP1, SERPINA3,</i>	115

				<i>SERPINE1, SERPINE2, SGCE, SLC12A2, SLC12A8, SLC16A1, SLC24A3, SLC27A6, SNCA, SOX3, SPARC, SRPX, STXBP6, TGFB1, TGM2, TUBB2A, TUBB2B, UBB, UCHL1, VAV3, VIM, WWOX, XBP1</i>	
Neurological disease	Neurological deficiency	Neurological deficiency	1.56×10^{-3}	<i>MGP, TGFB1</i>	2
Neurological disease	Nonsyndromic hearing impairment	Nonsyndromic hearing impairment	2.28×10^{-3}	<i>COCH, DFNA5, GRHL2</i>	3
Neurological disease	Glioma	Glioma	2.74×10^{-3}	<i>AKT3, CDKN2A, EGFR, ESR1, ID2, PDGFC, RB1, RET</i>	8
Neurological disease	Brain tumor	Brain tumor	3.08×10^{-3}	<i>ABAT, AKT3, CDKN2A, EGFR, ESR1, FYN, ID2, PDGFC, RB1, RET, TGFB1, TUBB2A</i>	12
Neurological disease	Astrocytoma	Astrocytoma	3.20×10^{-3}	<i>AKT3, EGFR, ESR1, ID2, PDGFC, RET</i>	6
Neurological disease	Glioblastoma	Glioblastoma	4.76×10^{-3}	<i>AKT3, EGFR, ESR1, PDGFC, RET</i>	5
Neurological disease	Congenital central hypoventilation syndrome	Congenital central hypoventilation syndrome	7.45×10^{-3}	<i>BDNF, RET</i>	2
Neurological disease	Progressive supranuclear palsy	Progressive supranuclear palsy	9.07×10^{-3}	<i>ABAT, SERPINA3, TGM2, UBB</i>	4
Cellular compromise	Adhesion	Adhesion of cancer cells	1.34×10^{-5}	<i>CDH2, CXCL12, EGFR, FN1, ICAM1, MCAM</i>	6
Cellular compromise	Adhesion	Adhesion of melanoma cells	1.35×10^{-2}	<i>CDH2, MCAM</i>	2
Cellular compromise	Aggregation	Aggregation of tumor cells	1.56×10^{-3}	<i>CDH2, CEACAM5 (includes EG:1048)</i>	2
Tumor morphology	Adhesion	Adhesion of cancer cells	1.34×10^{-5}	<i>CDH2, CXCL12, EGFR, FN1, ICAM1, MCAM</i>	6
Tumor morphology	Adhesion	Adhesion of melanoma cells	1.35×10^{-2}	<i>CDH2, MCAM</i>	2
Tumor morphology	Aggregation	Aggregation of tumor cells	1.56×10^{-3}	<i>CDH2, CEACAM5 (includes EG:1048)</i>	2

Tumor morphology	Invasion	Invasion of melanoma cells	7.45×10^{-3}	CDH1, MCAM	2
Tumor morphology	Morphology	Morphology of cancer cells	1.03×10^{-2}	EGFR, PLAGL1	2
Cell cycle	Cell division process	Cell division process of eukaryotic cells	1.43×10^{-5}	AXL, BMP7, CALCR, CCNE2, CDH1, CDKN2A, COL1A1, CYR61, DNAJB4, EGFR, ESR1, FN1, GDF15, GNAI1, HMGA1, ID2, IFI16, IGFBP3, IGFBP5, IRS1, KRT19, MYB (includes EG:4602), NCOA3, PLAGL1, RB1, SPARC, TGFB1, XBP1	28
Cell cycle	Cell division process	Cell division process	3.11×10^{-5}	AXL, BCAT1, BMP7, BOP1, BTG3, CALCR, CCNE2, CDH1, CDKN2A, COL1A1, CREBBP, CXCL12, CYR61, DHRS2 (includes EG:10202), DNAJB4, EGFR, ESR1, FN1, FOSL1, GDF15, GNAI1, HMGA1, ID2, IFI16, IGFBP3, IGFBP5, IRS1, KRT19, MYB (includes EG:4602), NCOA3, PLAGL1, PTPRK, RB1, SPARC, TGFB1, TPD52L1, UBB, XBP1	38
Cell cycle	Cell division process	Cell division process of cell lines	1.58×10^{-4}	AXL, CALCR, CDH1, CDKN2A, CYR61, DNAJB4, EGFR, ESR1, GDF15, GNAI1, HMGA1, IFI16, IGFBP3, IGFBP5, IRS1, KRT19, MYB (includes EG:4602), NCOA3, PLAGL1, RB1, SPARC, TGFB1, XBP1	23
Cell cycle	Cell division process	Cell division process of tumor cell lines	2.00×10^{-4}	AXL, CDH1, CDKN2A, CYR61, DNAJB4, EGFR, ESR1, GDF15, GNAI1, HMGA1, IFI16, IGFBP3, IGFBP5, KRT19, MYB (includes EG:4602), NCOA3, PLAGL1, RB1, TGFB1, XBP1	20
Cell cycle	Cell division process	Cell division process of thyrocytes	5.28×10^{-4}	CDKN2A, TGFB1	2
Cell cycle	Cell division process	Arrest in cell division process of carcinoma cell lines	6.52×10^{-4}	CDKN2A, CYR61, IFI16, RB1	4
Cell cycle	Cell division process	Cell division process of breast cancer cell lines	6.88×10^{-4}	CDH1, CDKN2A, EGFR, ESR1, GDF15, IGFBP5, KRT19, XBP1	8
Cell cycle	Cell division process	Cell division process of carcinoma cell lines	1.01×10^{-3}	CDKN2A, CYR61, DNAJB4, IFI16, RB1	5
Cell cycle	Cell division process	Arrest in cell division process of eukaryotic cells	1.20×10^{-3}	CALCR, CDKN2A, COL1A1, CYR61, EGFR, FN1, GDF15, IFI16, IGFBP5, KRT19, PLAGL1, RB1, TGFB1, XBP1	14

Cell cycle	Cell division process	Arrest in cell division process of normal cells	1.31×10^{-3}	<i>CDKN2A, COL1A1, FN1, RB1, TGFB1</i>	5
Cell cycle	Cell division process	Arrest in cell division process of breast cancer cell lines	2.94×10^{-3}	<i>EGFR, GDF15, IGFBP5, KRT19, XBP1</i>	5
Cell cycle	Cell division process	Arrest in cell division process of tumor cell lines	3.29×10^{-3}	<i>CDKN2A, CYR61, EGFR, GDF15, IFI16, IGFBP5, KRT19, PLAGL1, RB1, TGFB1, XBP1</i>	11
Cell cycle	Cell division process	Cell division process of lung cell lines	3.64×10^{-3}	<i>CDKN2A, NCOA3, SPARC</i>	3
Cell cycle	Cell division process	Entry into cell division process	4.97×10^{-3}	<i>CDH1, CDKN2A, CREBBP, FN1, NCOA3, RB1, TGFB1</i>	7
Cell cycle	Cell division process	Arrest in cell division process of cell lines	5.19×10^{-3}	<i>CALCR, CDKN2A, CYR61, EGFR, GDF15, IFI16, IGFBP5, KRT19, PLAGL1, RB1, TGFB1, XBP1</i>	12
Cell cycle	Cell division process	Cell division process of normal cells	5.61×10^{-3}	<i>BMP7, CCNE2, CDKN2A, COL1A1, FN1, ID2, RB1, TGFB1</i>	8
Cell cycle	Cell division process	Cell division process of epithelial cell lines	9.99×10^{-3}	<i>CALCR, CDKN2A, IRS1, TGFB1</i>	4
Cell cycle	Cell division process	Cell division process of lung cancer cell lines	1.10×10^{-2}	<i>CDKN2A, CYR61, DNAJB4, RB1</i>	4
Cell cycle	Cell division process	Arrest in cell division process of lung cancer cell lines	1.18×10^{-2}	<i>CDKN2A, CYR61, RB1</i>	3
Cell cycle	Cell cycle progression	Arrest in cell cycle progression of eukaryotic cells	8.92×10^{-5}	<i>CDKN2A, COL1A1, EGFR, IFI16, KRT19, PLAGL1, RB1, TGFB1, XBP1</i>	9
Cell cycle	Cell cycle progression	Cell cycle progression of cell lines	1.56×10^{-4}	<i>CDKN2A, DNAJB4, EGFR, ESR1, IFI16, IRS1, KRT19, PLAGL1, RB1, TGFB1, XBP1</i>	11

Cell cycle	Cell cycle progression	Cell cycle progression	2.22×10^{-4}	<i>AXL, BMP7, BOP1, BTG3, CDKN2A, COL1A1, CXCL12, CYR61, DHRS2 (includes EG:10202), DNAJB4, EGFR, ESR1, FOSL1, IFI16, IGFBP3, IRS1, KRT19, PLAGL1, PTPRK, RB1, SPARC, TGFB1, UBB, XBP1</i>	24
Cell cycle	Cell cycle progression	Arrest in cell cycle progression of blood cells	2.29×10^{-4}	<i>CDKN2A, COL1A1, TGFB1</i>	3
Cell cycle	Cell cycle progression	Cell cycle progression of eukaryotic cells	2.32×10^{-4}	<i>CDKN2A, COL1A1, DNAJB4, EGFR, ESR1, IFI16, IRS1, KRT19, PLAGL1, RB1, TGFB1, XBP1</i>	12
Cell cycle	Cell cycle progression	Arrest in cell cycle progression of tumor cell lines	2.82×10^{-4}	<i>CDKN2A, EGFR, IFI16, KRT19, PLAGL1, RB1, XBP1</i>	7
Cell cycle	Cell cycle progression	Cell cycle progression of tumor cell lines	6.09×10^{-4}	<i>CDKN2A, DNAJB4, EGFR, ESR1, IFI16, KRT19, PLAGL1, RB1, XBP1</i>	9
Cell cycle	Cell cycle progression	Cell cycle progression of carcinoma cell lines	1.28×10^{-3}	<i>DNAJB4, IFI16, RB1</i>	3
Cell cycle	Cell cycle progression	Arrest in cell cycle progression of carcinoma cell lines	1.56×10^{-3}	<i>IFI16, RB1</i>	2
Cell cycle	Cell cycle progression	Cell cycle progression of bone cancer cell lines	2.28×10^{-3}	<i>CDKN2A, PLAGL1, RB1</i>	3
Cell cycle	Cell cycle progression	Cell cycle progression of fibroblast cell lines	2.91×10^{-3}	<i>CDKN2A, IRS1, RB1</i>	3
Cell cycle	Cell cycle progression	Arrest in cell cycle progression of bone cancer cell lines	3.07×10^{-3}	<i>CDKN2A, PLAGL1</i>	2
Cell cycle	Cell cycle progression	Cell cycle progression of t lymphocytes	5.04×10^{-3}	<i>CDKN2A, TGFB1</i>	2

Cell cycle	Cell cycle progression	Cell cycle progression of normal cells	1.20×10^{-2}	<i>CDKN2A, COL1A1, RB1, TGFB1</i>	4
Cell cycle	Cell cycle progression	Cell cycle progression of lung cancer cell lines	1.35×10^{-2}	<i>DNAJB4, RB1</i>	2
Cell cycle	G1 phase	G1 phase of normal cells	2.29×10^{-4}	<i>CCNE2, FN1, ID2, RB1</i>	4
Cell cycle	G1 phase	G1 phase of eukaryotic cells	3.04×10^{-4}	<i>CCNE2, CDKN2A, CYR61, EGFR, FN1, GDF15, HMGA1, ID2, NCOA3, RB1, TGFB1</i>	11
Cell cycle	G1 phase	G1 phase of epithelial cells	3.07×10^{-3}	<i>ID2, RB1</i>	2
Cell cycle	G1 phase	Arrest in G1 phase of eukaryotic cells	5.60×10^{-3}	<i>CDKN2A, CYR61, EGFR, FN1, GDF15, RB1, TGFB1</i>	7
Cell cycle	G1 phase	G1 phase of tumor cell lines	9.17×10^{-3}	<i>CDKN2A, CYR61, EGFR, GDF15, HMGA1, NCOA3, TGFB1</i>	7
Cell cycle	G1 phase	Arrest in G1 phase of normal cells	1.35×10^{-2}	<i>FN1, RB1</i>	2
Cell cycle	G1/S phase transition	G1/S phase transition of normal cells	3.94×10^{-4}	<i>CCNE2, FN1, ID2</i>	3
Cell cycle	G1/S phase transition	G1/S phase transition of eukaryotic cells	1.59×10^{-3}	<i>CCNE2, CDKN2A, FN1, HMGA1, ID2, NCOA3</i>	6
Cell cycle	S phase	Entry into S phase of cell lines	8.74×10^{-4}	<i>CDH1, CDKN2A, NCOA3, RB1, TGFB1</i>	5
Cell cycle	S phase	Entry into S phase of colon cancer cell lines	1.56×10^{-3}	<i>CDH1, CDKN2A</i>	2
Cell cycle	S phase	Entry into S phase of lung cell lines	1.56×10^{-3}	<i>CDKN2A, NCOA3</i>	2
Cell cycle	S phase	S phase of cell lines	7.07×10^{-3}	<i>CDH1, CDKN2A, EGFR, NCOA3, RB1, TGFB1</i>	6
Cell cycle	S phase	S phase of tumor cell lines	9.85×10^{-3}	<i>CDH1, CDKN2A, EGFR, RB1, TGFB1</i>	5

Cell cycle	S phase	Entry into S phase of fibroblast cell lines	1.03×10^{-2}	<i>CDKN2A, NCOA3</i>	2
Cell cycle	Interphase	Entry into interphase of eukaryotic cells	1.18×10^{-3}	<i>CDH1, CDKN2A, FN1, NCOA3, RB1, TGFB1</i>	6
Cell cycle	Interphase	Interphase of eukaryotic cells	1.34×10^{-3}	<i>CALCR, CCNE2, CDH1, CDKN2A, CYR61, EGFR, FN1, GDF15, HMGA1, ID2, IGFBP5, MYB (includes EG:4602), NCOA3, RB1, TGFB1</i>	15
Cell cycle	Interphase	Interphase of normal cells	2.94×10^{-3}	<i>CCNE2, CDKN2A, FN1, ID2, RB1</i>	5
Cell cycle	Interphase	Interphase of tumor cell lines	5.74×10^{-3}	<i>CDH1, CDKN2A, CYR61, EGFR, GDF15, HMGA1, IGFBP5, MYB (includes EG:4602), NCOA3, RB1, TGFB1</i>	11
Cell cycle	Interphase	Interphase of cell lines	9.76×10^{-3}	<i>CALCR, CDH1, CDKN2A, CYR61, EGFR, GDF15, HMGA1, IGFBP5, MYB (includes EG:4602), NCOA3, RB1, TGFB1</i>	12
Cell cycle	Interphase	Interphase of breast cancer cell lines	9.85×10^{-3}	<i>CDH1, CDKN2A, EGFR, GDF15, IGFBP5</i>	5
Cell cycle	Interphase	Interphase of carcinoma cell lines	1.03×10^{-2}	<i>CDKN2A, CYR61, RB1</i>	3
Cell cycle	Interphase	Interphase of epithelial cell lines	1.03×10^{-2}	<i>CALCR, CDKN2A, TGFB1</i>	3
Cell cycle	Cell stage	Cell stage	2.26×10^{-3}	<i>AXL, BCAT1, BMP7, BTG3, CALCR, CCNE2, CDH1, CDKN2A, CREBBP, CXCL12, CYR61, EGFR, FN1, GDF15, GNAI1, HMGA1, ID2, IGFBP3, IGFBP5, MYB (includes EG:4602), NCOA3, RB1, SPARC, TGFB1, TPD52L1</i>	25
Cell cycle	Cell stage	Entry into cell stage	2.59×10^{-3}	<i>CDH1, CDKN2A, CREBBP, FN1, NCOA3, RB1, TGFB1</i>	7
Cell cycle	Cell stage	Cell stage of eukaryotic cells	3.92×10^{-3}	<i>BMP7, CALCR, CCNE2, CDH1, CDKN2A, CYR61, EGFR, FN1, GDF15, GNAI1, HMGA1, ID2, IGFBP5, MYB (includes EG:4602), NCOA3, RB1, TGFB1</i>	17
Cell cycle	Cell stage	Cell stage of normal cells	5.79×10^{-3}	<i>BMP7, CCNE2, CDKN2A, FN1, ID2, RB1</i>	6
Cell cycle	Mitogenesis	Mitogenesis	8.55×10^{-3}	<i>AXL, CYR61, EGFR, IGFBP3, SPARC, TGFB1</i>	6
Connective tissue development and function	Proliferation	Proliferation of chondrocytes	1.79×10^{-5}	<i>CTGF, IGFBP3, IL18, STC1</i>	4

Connective tissue development and function	Proliferation	Proliferation of fibroblast cell lines	1.68×10^{-2}	<i>CDKN2A, CXCL12, FN1, S100P, TGFB1</i>	5
Connective tissue development and function	Binding	Binding of fibroblasts	3.16×10^{-5}	<i>IGFBP3, LGALS3BP, SDC2, SPARC</i>	4
Connective tissue development and function	Attachment	Attachment of fibroblasts	1.56×10^{-3}	<i>FN1, TGFB1</i>	2
Connective tissue development and function	Growth	Growth of fibroblast cell lines	1.66×10^{-3}	<i>EGFR, FN1, IGFBP3, IRS1, NRCAM, RET, TGFB1</i>	7
Connective tissue development and function	Cell cycle progression	Cell cycle progression of fibroblast cell lines	2.91×10^{-3}	<i>CDKN2A, IRS1, RB1</i>	3
Connective tissue development and function	Aggregation	Aggregation of fibroblast cell lines	7.45×10^{-3}	<i>CDH2, EPCAM</i>	2
Connective tissue development and function	Shape change	Shape change of fibroblasts	7.45×10^{-3}	<i>CDKN2A, FN1</i>	2
Connective tissue development and function	S phase	Entry into S phase of fibroblast cell lines	1.03×10^{-2}	<i>CDKN2A, NCOA3</i>	2
Connective tissue development and function	Differentiation	Differentiation of chondrocytes	1.03×10^{-2}	<i>STC1, TGFB1</i>	2
Connective tissue development and function	Differentiation	Differentiation of fibroblast cell lines	1.52×10^{-2}	<i>BMP7, MGP, TGFB1</i>	3

Infectious disease	Severe acute respiratory syndrome	Severe acute respiratory syndrome	2.05×10^{-5}	<i>CD24, CTSZ (includes EG:1522), GLUL, HIST1H2AC, HIST1H2BD, HIST1H2BK, HIST2H2AA3, HIST2H2BE, PDGFC, S100P, SPARC, ZYX</i>	12
Hematopoiesis	Migration	Migration of myeloid cells	6.84×10^{-5}	<i>ANXA1, CXCL12, FN1, ICAM1, SERPINE1, TGFB1, TGM2</i>	7
Hematopoiesis	Migration	Migration of neutrophils	3.97×10^{-4}	<i>ANXA1, FN1, ICAM1, SERPINE1, TGFB1</i>	5
Hematopoiesis	Migration	Migration of granulocytes	4.78×10^{-4}	<i>ANXA1, CXCL12, FN1, ICAM1, SERPINE1, TGFB1</i>	6
Hematopoiesis	Differentiation	Differentiation of T lymphocytes	2.81×10^{-4}	<i>CDKN2A, CXCL12, GATA3, IL18, MUC1, TGFB1</i>	6
Hematopoiesis	Differentiation	Differentiation of mononuclear leukocytes	4.13×10^{-3}	<i>CDKN2A, CXCL12, GATA3, IFI16, IL18, MUC1, TGFB1</i>	7
Hematopoiesis	Differentiation	Differentiation of Th2 cells	5.04×10^{-3}	<i>GATA3, TGFB1</i>	2
Hematopoiesis	Development	T cell development	1.70×10^{-3}	<i>CDKN2A, CXCL12, GATA3, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B</i>	9
Hematopoiesis	Development	Development of blood cells	2.13×10^{-3}	<i>CDKN2A, CXCL12, EGFR, GATA3, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B</i>	10
Hematopoiesis	Apoptosis	Apoptosis of T lymphocytes	2.71×10^{-3}	<i>CDKN2A, CXCL12, IL7R, RAC2, TGFB1, TNFRSF6B</i>	6
Hematopoiesis	Induction	Induction of T lymphocytes	3.07×10^{-3}	<i>IL18, MUC1</i>	2
Hematopoiesis	Activation-induced cell death	Activation-induced cell death of T lymphocytes	1.35×10^{-2}	<i>RAC2, TGFB1</i>	2
Immune cell trafficking	Migration	Migration of myeloid cells	6.84×10^{-5}	<i>ANXA1, CXCL12, FN1, ICAM1, SERPINE1, TGFB1, TGM2</i>	7
Immune cell trafficking	Migration	Migration of neutrophils	3.97×10^{-4}	<i>ANXA1, FN1, ICAM1, SERPINE1, TGFB1</i>	5
Immune cell trafficking	Migration	Migration of granulocytes	4.78×10^{-4}	<i>ANXA1, CXCL12, FN1, ICAM1, SERPINE1, TGFB1</i>	6
Immune cell trafficking	Migration	Lymphocyte migration	6.53×10^{-4}	<i>CXCL12, EGFR, FN1, ICAM1, ICAM2, IL18, RASGRP1, TGM2, TNFRSF6B</i>	9
Immune cell trafficking	Migration	Migration of mononuclear leukocytes	8.93×10^{-4}	<i>CXCL12, EGFR, FN1, ICAM1, ICAM2, IL18, RASGRP1, TGFB1, TGM2, TNFRSF6B</i>	10
Immune cell trafficking	Migration	Migration of leukocytes	1.64×10^{-3}	<i>ANXA1, CXCL12, EGFR, FN1, ICAM1, ICAM2, IL18, RASGRP1, RLN2, SERPINA3, SERPINE1, TGFB1, TGM2, TNFRSF6B</i>	14

Immune cell trafficking	Adhesion	Adhesion of lymphocytes	1.07×10^{-3}	<i>CXCL12, FN1, ICAM1, ICAM2, RASGRP1, TGFB1</i>	6
Immune cell trafficking	Adhesion	Adhesion of leukocytes	1.19×10^{-3}	<i>ANXA1, CXCL12, FN1, ICAM1, ICAM2, PLTP, RASGRP1, RLN2, TGFB1</i>	9
Immune cell trafficking	Adhesion	Adhesion of T lymphocytes	2.12×10^{-3}	<i>CXCL12, FN1, ICAM1, ICAM2, RASGRP1</i>	5
Cardiovascular system development and function	Neovascularization	Neovascularization	7.99×10^{-5}	<i>CXCL12, ICAM1, TGFB1, TNFRSF6B</i>	4
Cardiovascular system development and function	Neovascularization	Neovascularization of animal	1.56×10^{-3}	<i>CXCL12, TNFRSF6B</i>	2
Cardiovascular system development and function	Cardiovascular process	Cardiovascular process of coronary artery	8.61×10^{-4}	<i>EMP1, EMP3, ICAM1, IFI16, RB1, TUBB6</i>	6
Cardiovascular system development and function	Cardiovascular process	Cardiovascular process of blood vessel	2.28×10^{-3}	<i>COL4A2, EMP1, EMP3, ICAM1, IFI16, RB1, TGFB1, TUBB6</i>	8
Cardiovascular system development and function	Cardiovascular process	Cardiovascular process of animal	6.46×10^{-3}	<i>CXCL12, CYR61, TNFRSF6B</i>	3
Cardiovascular system development and function	Formation	Formation of endothelial tube	2.37×10^{-3}	<i>COL4A1, FN1, SERPINE1, TGFB1</i>	4
Cardiovascular system development and function	Formation	Formation of endothelial cells	1.35×10^{-2}	<i>FN1, SERPINE1</i>	2
Cardiovascular system development and function	Proliferation	Proliferation of endothelial cells	3.19×10^{-3}	<i>AXL, CDH2, COL4A1, COL4A2, ECM1, FN1, MGP</i>	7

Cardiovascular system development and function	Adhesion	Adhesion of endothelial cells	4.35×10^{-3}	<i>CDH2, FN1, ICAM1, ICAM2, TGFB1</i>	5
Cardiovascular system development and function	Adhesion	Adhesion of endothelial cell lines	5.68×10^{-3}	<i>ANXA1, CDH2, CXCL12, ICAM1, TGFB1</i>	5
Cardiovascular system development and function	Tubulation	Tubulation of endothelial cells	5.41×10^{-3}	<i>AXL, IGFBP3, MGP</i>	3
Cardiovascular system development and function	Development	Development of blood vessel	6.52×10^{-3}	<i>COL1A1, COL4A2, CYR61, ECM1, FN1, IL18, SERPINE1, SPARC</i>	8
Cardiovascular system development and function	Angiogenesis	Angiogenesis	1.35×10^{-2}	<i>COL4A2, CYR61, ECM1, FN1, IL18, SERPINE1, SPARC</i>	7
Cardiovascular system development and function	Binding	Binding of endothelial cell lines	1.43×10^{-2}	<i>AXL, FN1, ICAM1, PIK3R1</i>	4
Organismal development	Neovascularization	Neovascularization	7.99×10^{-5}	<i>CXCL12, ICAM1, TGFB1, TNFRSF6B</i>	4
Organismal development	Neovascularization	Neovascularization of animal	1.56×10^{-3}	<i>CXCL12, TNFRSF6B</i>	2
Organismal development	Development	Development of blood vessel	6.52×10^{-3}	<i>COL1A1, COL4A2, CYR61, ECM1, FN1, IL18, SERPINE1, SPARC</i>	8
Organismal development	Angiogenesis	Angiogenesis	1.35×10^{-2}	<i>COL4A2, CYR61, ECM1, FN1, IL18, SERPINE1, SPARC</i>	7
Tissue morphology	Quantity	Quantity of cells	8.16×10^{-5}	<i>BDNF, BMP7, CDKN2A, CXCL12, ESR1, FN1, IGFBP3, IGFBP5, IGFBP6, KLF4, RB1, SNCA, STC1, TGFB1</i>	14
Tissue morphology	Quantity	Quantity of tumor cell lines	1.92×10^{-3}	<i>CDKN2A, ESR1, IGFBP3, IGFBP5, IGFBP6, KLF4</i>	6
Tissue morphology	Quantity	Quantity of prostate cancer cell lines	5.04×10^{-3}	<i>IGFBP3, IGFBP5</i>	2

Dermatologic al diseases and conditions	Burn	Burn	1.37×10^{-4}	<i>COL1A1, COL21A1, COL4A1, COL4A2, COL6A1, COL6A2, ESR1</i>	7
Dermatologic al diseases and conditions	Cutis laxa, recessive type 1	Cutis laxa, recessive type 1	1.56×10^{-3}	<i>EFEMP2, LOX</i>	2
Dermatologic al diseases and conditions	Dermatological disorder	Dermatological disorder	1.40×10^{-2}	<i>ADA, AREG, CDKN2A, COL1A1, COL21A1, COL4A1, COL4A2, COL6A1, COL6A2, ECM1, EFEMP2, EGFR, ESR1, FN1, FYN, GDF15, GJA1, LOX, MMP1 (includes EG:4312), TGFB1, TUBB2A, ZNF395</i>	22
DNA replication, recombination, and repair	Synthesis	Synthesis of DNA	1.44×10^{-4}	<i>AREG, BMP7, CDKN2A, CXCL12, CYR61, EGFR, ESR1, FN1, IGFBP3, IGFBP5, PIK3R1, RB1, TFPI2, TGFB1</i>	14
DNA replication, recombination, and repair	Bending	Bending of DNA	5.28×10^{-4}	<i>ESR1, TFF1</i>	2
DNA replication, recombination, and repair	Incorporation	Incorporation of thymidine	2.12×10^{-3}	<i>CDKN2A, EGFR, FN1, IGFBP5, TGFB1</i>	5
DNA replication, recombination, and repair	Fragmentation	Fragmentation of DNA	3.40×10^{-3}	<i>ABCB1, CDKN2A, CXCL12, GPX1, TFF3, TGFB1, TPD52L1</i>	7
DNA replication, recombination, and repair	Metabolism	Metabolism of DNA	9.46×10^{-3}	<i>ABCB1, AREG, CDKN2A, CXCL12, GPX1, KRT7, NT5E, ORC5L, PDGFC, TFF3, TGFB1, TPD52L1</i>	12
Cellular function and maintenance	Contact growth inhibition	Contact growth inhibition	1.99×10^{-4}	<i>BMP7, CD24, CDH1, CDKN2A, EGFR, IFI16, TGFB1</i>	7
Cellular function and maintenance	Contact growth inhibition	Contact growth inhibition of eukaryotic cells	6.87×10^{-4}	<i>BMP7, CD24, CDH1, EGFR, IFI16, TGFB1</i>	6
Cellular function and maintenance	Contact growth inhibition	Contact growth inhibition of cell lines	2.12×10^{-3}	<i>BMP7, CD24, CDH1, EGFR, IFI16</i>	5

Cellular function and maintenance	Contact growth inhibition	Contact growth inhibition of tumor cell lines	5.30×10^{-3}	<i>BMP7, CD24, CDH1, IFI16</i>	4
Cellular function and maintenance	Contact growth inhibition	Contact growth inhibition of colon cancer cell lines	1.03×10^{-2}	<i>CD24, CDH1</i>	2
Cellular function and maintenance	Differentiation	Differentiation of T lymphocytes	2.81×10^{-4}	<i>CDKN2A, CXCL12, GATA3, IL18, MUC1, TGFB1</i>	6
Cellular function and maintenance	Differentiation	Differentiation of Th2 cells	5.04×10^{-3}	<i>GATA3, TGFB1</i>	2
Cellular function and maintenance	Organization	Organization of cytoskeleton	3.23×10^{-4}	<i>CXCL12, FN1, FSCN1, ICAM1, PLS3, SDC2</i>	6
Cellular function and maintenance	Organization	Organization of actin cytoskeleton	3.64×10^{-3}	<i>CXCL12, FN1, FSCN1, ICAM1</i>	4
Cellular function and maintenance	Metabolic process	Metabolic process of tumor cell lines	5.28×10^{-4}	<i>EPCAM, SNCA</i>	2
Cellular function and maintenance	Development	T cell development	1.70×10^{-3}	<i>CDKN2A, CXCL12, GATA3, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B</i>	9
Cellular function and maintenance	Apoptosis	Apoptosis of T lymphocytes	2.71×10^{-3}	<i>CDKN2A, CXCL12, IL7R, RAC2, TGFB1, TNFRSF6B</i>	6
Cellular function and maintenance	Cytostasis	Cytostasis of colon cancer cell lines	2.91×10^{-3}	<i>CD24, CDH1, TGFB1</i>	3
Cellular function and maintenance	Cytostasis	Cytostasis of eukaryotic cells	3.66×10^{-3}	<i>BMP7, CD24, CDH1, CDKN2A, EGFR, ID2, IFI16, TGFB1</i>	8
Cellular function and maintenance	Cytostasis	Cytostasis of epithelial cell lines	5.04×10^{-3}	<i>EGFR, TGFB1</i>	2
Cellular function and maintenance	Cytostasis	Cytostasis of tumor cell lines	7.07×10^{-3}	<i>BMP7, CD24, CDH1, EGFR, IFI16, TGFB1</i>	6

Cellular function and maintenance	Cytostasis	Cytostasis of normal cells	1.03×10^{-2}	<i>CDKN2A, ID2, TGFB1</i>	3
Cellular function and maintenance	Induction	Induction of T lymphocytes	3.07×10^{-3}	<i>IL18, MUC1</i>	2
Cellular function and maintenance	Activation-induced cell death	Activation-induced cell death of T lymphocytes	1.35×10^{-2}	<i>RAC2, TGFB1</i>	2
Cell-mediated immune response	Differentiation	Differentiation of T lymphocytes	2.81×10^{-4}	<i>CDKN2A, CXCL12, GATA3, IL18, MUC1, TGFB1</i>	6
Cell-mediated immune response	Differentiation	Differentiation of Th2 cells	5.04×10^{-3}	<i>GATA3, TGFB1</i>	2
Cell-mediated immune response	Development	T cell development	1.70×10^{-3}	<i>CDKN2A, CXCL12, GATA3, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B</i>	9
Cell-mediated immune response	Adhesion	Adhesion of T lymphocytes	2.12×10^{-3}	<i>CXCL12, FN1, ICAM1, ICAM2, RASGRP1</i>	5
Cell-mediated immune response	Apoptosis	Apoptosis of T lymphocytes	2.71×10^{-3}	<i>CDKN2A, CXCL12, IL7R, RAC2, TGFB1, TNFRSF6B</i>	6
Cell-mediated immune response	Induction	Induction of T lymphocytes	3.07×10^{-3}	<i>IL18, MUC1</i>	2
Cell-mediated immune response	Activation-induced cell death	Activation-induced cell death of T lymphocytes	1.35×10^{-2}	<i>RAC2, TGFB1</i>	2
Hematological system development and function	Differentiation	Differentiation of T lymphocytes	2.81×10^{-4}	<i>CDKN2A, CXCL12, GATA3, IL18, MUC1, TGFB1</i>	6
Hematological system development and function	Differentiation	Differentiation of mononuclear leukocytes	4.13×10^{-3}	<i>CDKN2A, CXCL12, GATA3, IFI16, IL18, MUC1, TGFB1</i>	7

Hematological system development and function	Differentiation	Differentiation of Th2 cells	5.04×10^{-3}	GATA3, TGFB1	2
Hematological system development and function	Differentiation	Differentiation of blood cells	1.69×10^{-2}	CDKN2A, CXCL12, GATA3, IFI16, IL18, MAFB, MUC1, MYB (includes EG:4602), TGFB1	9
Hematological system development and function	Adhesion	Adhesion of lymphocytes	1.07×10^{-3}	CXCL12, FN1, ICAM1, ICAM2, RASGRP1, TGFB1	6
Hematological system development and function	Adhesion	Adhesion of leukocytes	1.19×10^{-3}	ANXA1, CXCL12, FN1, ICAM1, ICAM2, PLTP, RASGRP1, RLN2, TGFB1	9
Hematological system development and function	Adhesion	Adhesion of T lymphocytes	2.12×10^{-3}	CXCL12, FN1, ICAM1, ICAM2, RASGRP1	5
Hematological system development and function	Adhesion	Adhesion of leukocyte cell lines	1.34×10^{-2}	ANXA1, TGFB1, TGM2	3
Hematological system development and function	Development	T cell development	1.70×10^{-3}	CDKN2A, CXCL12, GATA3, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B	9
Hematological system development and function	Development	Development of blood cells	2.13×10^{-3}	CDKN2A, CXCL12, EGFR, GATA3, IL18, IL7R, MUC1, RAC2, TGFB1, TNFRSF6B	10
Hematological system development and function	Apoptosis	Apoptosis of T lymphocytes	2.71×10^{-3}	CDKN2A, CXCL12, IL7R, RAC2, TGFB1, TNFRSF6B	6
Hematological system development and function	Induction	Induction of T lymphocytes	3.07×10^{-3}	IL18, MUC1	2

Hematological system development and function	Binding	Binding of blood platelets	1.18×10^{-2}	<i>CD24, FN1, ICAM1</i>	3
Hematological system development and function	Proliferation	Proliferation of T lymphocytes	1.28×10^{-2}	<i>CD24, CDKN2A, CXCL12, FN1, ICAM1, IL18, IL7R, TGFB1</i>	8
Hematological system development and function	Activation	Activation of leukocyte cell lines	1.35×10^{-2}	<i>B2M, TNFRSF6B</i>	2
Hematological system development and function	Activation-induced cell death	Activation-induced cell death of T lymphocytes	1.35×10^{-2}	<i>RAC2, TGFB1</i>	2
Hematological system development and function	Co-stimulation	Co-stimulation of T lymphocytes	1.35×10^{-2}	<i>CD24, FN1</i>	2
Cellular assembly and organization	Organization	Organization of cytoskeleton	3.23×10^{-4}	<i>CXCL12, FN1, FSCN1, ICAM1, PLS3, SDC2</i>	6
Cellular assembly and organization	Organization	Organization of actin cytoskeleton	3.64×10^{-3}	<i>CXCL12, FN1, FSCN1, ICAM1</i>	4
Cellular assembly and organization	Rearrangement	Rearrangement of cytoskeleton	9.96×10^{-4}	<i>CTGF, CXCL12, RET, TGFB1</i>	4
Cellular assembly and organization	Association	Association of cytoskeleton	1.56×10^{-3}	<i>CXCL12, FXYD5</i>	2
Cellular assembly and organization	Formation	Formation of cell-matrix contacts	1.56×10^{-3}	<i>CDH1, FN1</i>	2
Cellular assembly and organization	Formation	Formation of cellular inclusion bodies	1.71×10^{-2}	<i>SNCA, WWOX</i>	2
Cellular assembly and organization	Quantity	Quantity of intercellular junctions	1.56×10^{-3}	<i>CDH1, GDF15</i>	2

Cellular assembly and organization	Quantity	Quantity of neurites	5.04×10^{-3}	ESR1, SNCA	2
Cellular assembly and organization	Quantity	Quantity of plasma membrane projections	1.03×10^{-2}	BST2, ESR1, SNCA	3
Cellular assembly and organization	Development	Development of filaments	7.45×10^{-3}	CRYAB, TGFB1	2
Connective tissue disorders	Connective tissue disorder	Connective tissue disorder	3.55×10^{-4}	ABCB1, AKAP2, AREG, B2M, BCAT1, BDNF, BTG3, C6ORF97, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, COL1A1, COL21A1, COL4A1, COL4A2, COL6A1, COL6A2, COMMD10, CRYAB, CXCL12, EFEMP2, ESR1, FN1, FYN, GALNT3, GLUL, GPD1L, HIST1H2AC, HNMT, ICAM1, IL18, IL7R, IRS1, LDHB, LOX, MMP1 (includes EG:4312), MPPED2 (includes EG:744), MTUS1, NCOA3, NEBL, PBX1, PIK3R3, PTPRK, RB1, RCN1, SERPINA3, SERTAD2, SLC24A3, SNCA, STXBP6, TFPI2, TGFB1, TPD52L1, TPM2, TUBB2A, UBB, UBE2L6, VAV3, VIM, WWOX	65
Connective tissue disorders	Dupuytren contracture	Dupuytren contracture	1.20×10^{-3}	COL1A1, COL21A1, COL4A1, COL4A2, COL6A1, COL6A2, MMP1 (includes EG:4312)	7
Connective tissue disorders	Synovitis	Synovitis	1.74×10^{-3}	B2M, EFEMP2, TGFB1	3
Connective tissue disorders	Rheumatic disease	Rheumatic disease	2.20×10^{-3}	ABCB1, AKAP2, AREG, B2M, BCAT1, BDNF, BTG3, C6ORF97, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, COL4A1, COL6A1, COMMD10, CRYAB, CXCL12, EFEMP2, ESR1, FN1, FYN, GLUL, GPD1L, HIST1H2AC, HNMT, ICAM1, IL18, IL7R, IRS1, LDHB, LOX, MMP1 (includes EG:4312), MPPED2 (includes EG:744), MTUS1, NCOA3, NEBL, PBX1, PIK3R3, PTPRK, RB1, RCN1, SERPINA3, SERTAD2, SLC24A3, SNCA, STXBP6, TFPI2, TGFB1, TPD52L1, TPM2, TUBB2A, UBB, UBE2L6, VAV3, VIM, WWOX	59
Connective tissue disorders	Arthritis	Arthritis	2.23×10^{-3}	ABCB1, AKAP2, AREG, B2M, BCAT1, BDNF, BTG3, C6ORF97, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, COL4A1, COL6A1, COMMD10, CRYAB, CXCL12, EFEMP2, FYN, GLUL, GPD1L, HIST1H2AC, HNMT, ICAM1, IL18, IL7R, IRS1, LDHB, LOX, MMP1 (includes EG:4312), MPPED2 (includes EG:744), MTUS1, NCOA3, NEBL, PBX1, PIK3R3, PTPRK, RB1, RCN1, SERPINA3, SERTAD2, SLC24A3, SNCA, STXBP6, TFPI2, TGFB1, TPD52L1, TPM2, TUBB2A, UBB, UBE2L6, VAV3, VIM, WWOX	57

Connective tissue disorders	Osteoarthritis	Osteoarthritis	6.62×10^{-3}	<i>CRYAB, EFEMP2, IL18, MMP1</i> (includes EG:4312), <i>TFPI2, TGFB1</i>	6
Skeletal and muscular disorders	Skeletal and muscular disorder	Skeletal and muscular disorder	3.75×10^{-4}	<i>ABAT, ABCB1, AKAP2, AKT3, AREG, ATXN1, B2M, BCAT1, BDNF, BIN1, BTG3, C6ORF97, CA2, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, CDS1, COCH, COL1A1, COL4A1, COL6A1, COL6A2, COMMD10, CRYAB, CXCL12, DKK3, EFEMP2, EML1</i> (includes EG:2009), <i>ESR1, FN1, FYN, GJA1, GLUL, GNAL, GPD1L, HIST1H2AC, HIST2H2AA3, HNMT, ICAM1, ID4, IL18, IL7R, IRS1, ISOC1, LAMB1, LDHB, LOX, MAP1B, MGP, MMP1</i> (includes EG:4312), <i>MPPED2</i> (includes EG:744), <i>MSX2, MT1E, MTUS1, NCOA3, NEBL, PBX1, PDCD10, PDLIM4, PIK3R3, PSMB9, PTPRK, RAI14, RAMP3, RB1, RCN1, S100P, SCD, SCHIP1, SCUBE2, SERPINA3, SERPINE1, SERTAD2, SGCE, SLC16A1, SLC24A3, SNCA, SRPX, STXBP6, TFPI2, TGFB1, TGM2, TPD52L1, TPM2, TUBB2A, TUBB2B, UBB, UBE2L6, UCHL1, VAV3, VIM, WWOX</i>	96
Skeletal and muscular disorders	Skeletal and muscular disorder	Skeletal and muscular disorder of humans	9.99×10^{-3}	<i>ATXN1, BDNF, CRYAB, SNCA</i>	4
Skeletal and muscular disorders	Osteoporosis	Osteoporosis	6.49×10^{-4}	<i>CA2, CALCR, COL1A1, ESR1, PDLIM4</i>	5
Skeletal and muscular disorders	Bethlem myopathy	Bethlem myopathy	1.56×10^{-3}	<i>COL6A1, COL6A2</i>	2
Skeletal and muscular disorders	Ullrich congenital muscular dystrophy	Ullrich congenital muscular dystrophy	1.56×10^{-3}	<i>COL6A1, COL6A2</i>	2
Skeletal and muscular disorders	Osteosarcoma	Osteosarcoma	1.68×10^{-3}	<i>CDKN2A, FYN, RB1, SDC2, SPARC</i>	5
Skeletal and muscular disorders	Rheumatic disease	Rheumatic disease	2.20×10^{-3}	<i>ABCB1, AKAP2, AREG, B2M, BCAT1, BDNF, BTG3, C6ORF97, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, COL4A1, COL6A1, COMMD10, CRYAB, CXCL12, EFEMP2, ESR1, FN1, FYN, GLUL, GPD1L, HIST1H2AC, HNMT, ICAM1, IL18, IL7R, IRS1, LDHB, LOX, MMP1</i> (includes EG:4312), <i>MPPED2</i> (includes EG:744), <i>MTUS1, NCOA3, NEBL, PBX1, PIK3R3, PTPRK, RB1, RCN1, SERPINA3, SERTAD2, SLC24A3, SNCA, STXBP6, TFPI2, TGFB1, TPD52L1, TPM2, TUBB2A, UBB, UBE2L6, VAV3, VIM, WWOX</i>	59

Skeletal and muscular disorders	Arthritis	Arthritis	2.23×10^{-3}	<i>ABCB1, AKAP2, AREG, B2M, BCAT1, BDNF, BTG3, C6ORF97, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, COL4A1, COL6A1, COMMD10, CRYAB, CXCL12, EFEMP2, FYN, GLUL, GPD1L, HIST1H2AC, HNMT, ICAM1, IL18, IL7R, IRS1, LDHB, LOX, MMP1 (includes EG:4312), MPPE2 (includes EG:744), MTUS1, NCOA3, NEBL, PBX1, PIK3R3, PTPRK, RB1, RCN1, SERPINA3, SERTAD2, SLC24A3, SNCA, STXBP6, TFPI2, TGFB1, TPD52L1, TPM2, TUBB2A, UBB, UBE2L6, VAV3, VIM, WWOX</i>	57
Skeletal and muscular disorders	Bone tumor	Bone tumor	2.42×10^{-3}	<i>ADA, CDKN2A, EXT1, FYN, RB1, RET, SDC2, SPARC</i>	8
Skeletal and muscular disorders	Osteoarthritis	Osteoarthritis	6.62×10^{-3}	<i>CRYAB, EFEMP2, IL18, MMP1 (includes EG:4312), TFPI2, TGFB1</i>	6
Skeletal and muscular disorders	Leiomyosarcoma	Leiomyosarcoma	7.40×10^{-3}	<i>CDKN2A, FYN, HMGA1, RET</i>	4
Skeletal and muscular disorders	Progressive supranuclear palsy	Progressive supranuclear palsy	9.07×10^{-3}	<i>ABAT, SERPINA3, TGM2, UBB</i>	4
Skeletal and muscular disorders	Myosarcoma	Myosarcoma	1.06×10^{-2}	<i>CDKN2A, FYN, HMGA1, RET, TUBB2A</i>	5
Skeletal and muscular disorders	Chondrosarcoma	Chondrosarcoma	1.35×10^{-2}	<i>EXT1, FYN</i>	2
Skeletal and muscular disorders	Postmenopausal osteoporosis	Postmenopausal osteoporosis	1.35×10^{-2}	<i>CALCR, ESR1</i>	2
Inflammatory response	Migration	Migration of neutrophils	3.97×10^{-4}	<i>ANXA1, FN1, ICAM1, SERPINE1, TGFB1</i>	5
Inflammatory response	Binding	Binding of blood platelets	1.18×10^{-2}	<i>CD24, FN1, ICAM1</i>	3
Reproductive system development and function	Proliferation	Proliferation of breast cell lines	4.02×10^{-4}	<i>CDH1, IGFBP3, LOX, TGFB1</i>	4
Reproductive system development and function	Morphology	Morphology of gonadal cell lines	1.35×10^{-2}	<i>PEG10, SERPINE1</i>	2

Immunologic al disease	Immunodeficienc y	Immunodeficien cy	4.78×10^{-4}	<i>ADA, B2M, IL7R, MLPH, RAC2, RASGRP1</i>	6
Immunologic al disease	Immunological disorder	Immunological disorder	8.94×10^{-3}	<i>ABAT, ABCB1, ADA, ADCY1, AKAP2, ANK3, APPBP2, AREG, ATXN1, B2M, BCAT1, BTG3, C6ORF97, CALCR, CALD1, CD24, CDA, CDH1, CDH2, COL4A1, COMMD10, CXCL12, ESR1, FNDC3B, FYN, GFRA1, GLUL, GNA14, GNAL, GPD1L, GRHL2, HIST1H2AC, HIST1H4H (includes EG:8365), HNMT, ICAM1, IGFBP3, IL18, IL7R, IRS1, KRT7, LDHB, LOX, MLPH, MPPE2 (includes EG:744), MTUS1, NCOA3, NEBL, NRCAM, PBX1, PIK3R3, PSMB9, PTPRK, RAC2, RAI14, RASGRP1, RB1, RCN1, RET, SERPINA3, SERTAD2, SLC24A3, SNCA, SNRPN, STXBP6, TGFB1, TGM2, TNFRSF6B, TPD52L1, TPM2, TUBB2A, UBE2L6, VAV3, VIM, WNT5A, WWOX, ZNF395</i>	76
Behavior	Locomotion	Locomotion of eukaryotic cells	5.28×10^{-4}	<i>CXCL12, TGFB1</i>	2
Gene expression	Bending	Bending of dna	5.28×10^{-4}	<i>ESR1, TFF1</i>	2
Gene expression	Activation	Activation of ap1 response element	9.15×10^{-4}	<i>ESR1, IL18, MMP1 (includes EG:4312)</i>	3
Gene expression	Activation	Activation of CAGA box	7.45×10^{-3}	<i>CREBBP, TGFB1</i>	2
Gene expression	Activation	Activation of SMAD binding sequence	1.03×10^{-2}	<i>TGFB1, TGFB1I1</i>	2
Gene expression	Activation	Activation of TGF-β response element	1.03×10^{-2}	<i>BMP7, TGFB1</i>	2
Gene expression	Expression	Expression of SP1 binding site	4.47×10^{-3}	<i>ESR1, RB1, TGFB1</i>	3
Gene expression	Transcription	Transcription of chromatin templates	5.04×10^{-3}	<i>CREBBP, ESR1</i>	2
Gene expression	Transcription	Transcription of AP1 consensus site	1.03×10^{-2}	<i>ESR1, TGFB1</i>	2
Gene expression	Transcription	Transcription of TATA box	1.35×10^{-2}	<i>ESR1, RB1</i>	2
Gene expression	Binding	Binding of AP1 consensus site	1.18×10^{-2}	<i>CXCL12, GPX1, TGFB1</i>	3
Gene expression	Repression	Repression of gene	1.55×10^{-2}	<i>ESR1, FHL2, FOXG1, RB1</i>	4

Nervous system development and function	Chemotaxis	Chemotaxis of granule cells	5.28×10^{-4}	<i>BDNF, CXCL12</i>	2
Nervous system development and function	Chemotaxis	Chemotaxis of microglia	5.04×10^{-3}	<i>TGFB1, TNFRSF6B</i>	2
Nervous system development and function	Quantity	Quantity of neuritis	5.04×10^{-3}	<i>ESR1, SNCA</i>	2
Nervous system development and function	Length	Length of neuritis	7.45×10^{-3}	<i>BDNF, ESR1</i>	2
Nervous system development and function	Development	Development of nervous system	8.70×10^{-3}	<i>FOXG1, GSTP1, MAFB, PDGFC, SNCA, SOX3, ZIC1</i>	7
Metabolic disease	Light chain associated amyloidosis	Light chain-associated amyloidosis	9.96×10^{-4}	<i>CXCL12, IGFBP5, RB1, TGFB1</i>	4
Metabolic disease	Gestational diabetes mellitus	Gestational diabetes mellitus	7.45×10^{-3}	<i>IRS1, PIK3R1</i>	2
Metabolic disease	Diabetes	Diabetes of mice	1.03×10^{-2}	<i>B2M, IGFBP3</i>	2
Carbohydrate metabolism	Synthesis	Synthesis of polysaccharide	1.18×10^{-3}	<i>CTGF, EXT1, IGFBP3, IRS1, TGFB1, UGDH</i>	6
Carbohydrate metabolism	Synthesis	Synthesis of proteoglycan	4.47×10^{-3}	<i>CTGF, EXT1, TGFB1</i>	3
Carbohydrate metabolism	Synthesis	Synthesis of carbohydrate	6.28×10^{-3}	<i>CTGF, EXT1, IGFBP3, IRS1, TGFB1, UGDH, XBP1</i>	7
Carbohydrate metabolism	Accumulation	Accumulation of doxorubicin	3.07×10^{-3}	<i>ABCB1, DNAJC15</i>	2
Carbohydrate metabolism	Release	Release of polysaccharide	3.07×10^{-3}	<i>FN1, IL18</i>	2
Carbohydrate metabolism	Release	Release of carbohydrate	7.62×10^{-3}	<i>ABCB1, FN1, IL18</i>	3

Carbohydrate metabolism	Metabolism	Metabolism of N-acetylglucosamine	1.35×10^{-2}	<i>CHST6, EXTL2</i>	2
Embryonic development	Transdifferentiation	Transdifferentiation of mesenchymal cells	1.56×10^{-3}	<i>COL4A1, TGFB1</i>	2
Embryonic development	Adhesion	Adhesion of embryonic cell lines	5.41×10^{-3}	<i>FN1, ICAM1, RRAS</i>	3
Embryonic development	Apoptosis	Apoptosis of embryonic cells	1.03×10^{-2}	<i>BMP7, TNFRSF6B</i>	2
Embryonic development	Morphology	Morphology of embryonic cell lines	1.35×10^{-2}	<i>IGFBP3, SERPINE1</i>	2
Inflammatory disease	Adult onset still's disease	Adult-onset Still's disease	1.56×10^{-3}	<i>ICAM1, IL18</i>	2
Inflammatory disease	Synovitis	Synovitis	1.74×10^{-3}	<i>B2M, EFEMP2, TGFB1</i>	3
Inflammatory disease	Rheumatic disease	Rheumatic disease	2.20×10^{-3}	<i>ABCB1, AKAP2, AREG, B2M, BCAT1, BDNF, BTG3, C6ORF97, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, COL4A1, COL6A1, COMMD10, CRYAB, CXCL12, EFEMP2, ESR1, FN1, FYN, GLUL, GPD1L, HIST1H2AC, HNMT, ICAM1, IL18, IL7R, IRS1, LDHB, LOX, MMP1 (includes EG:4312), MPPED2 (includes EG:744), MTUS1, NCOA3, NEBL, PBX1, PIK3R3, PTPRK, RB1, RCN1, SERPINA3, SERTAD2, SLC24A3, SNCA, STXBP6, TFPI2, TGFB1, TPD52L1, TPM2, TUBB2A, UBB, UBE2L6, VAV3, VIM, WWOX</i>	59
Inflammatory disease	Arthritis	Arthritis	2.23×10^{-3}	<i>ABCB1, AKAP2, AREG, B2M, BCAT1, BDNF, BTG3, C6ORF97, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, COL4A1, COL6A1, COMMD10, CRYAB, CXCL12, EFEMP2, FYN, GLUL, GPD1L, HIST1H2AC, HNMT, ICAM1, IL18, IL7R, IRS1, LDHB, LOX, MMP1 (includes EG:4312), MPPED2 (includes EG:744), MTUS1, NCOA3, NEBL, PBX1, PIK3R3, PTPRK, RB1, RCN1, SERPINA3, SERTAD2, SLC24A3, SNCA, STXBP6, TFPI2, TGFB1, TPD52L1, TPM2, TUBB2A, UBB, UBE2L6, VAV3, VIM, WWOX</i>	57
Inflammatory disease	Osteoarthritis	Osteoarthritis	6.62×10^{-3}	<i>CRYAB, EFEMP2, IL18, MMP1 (includes EG:4312), TFPI2, TGFB1</i>	6

Inflammatory disease	Inflammatory disorder	Inflammatory disorder	8.03×10^{-3}	<i>ABCB1, ADA, AKAP2, AREG, ATXN1, B2M, BCAS3, BCAT1, BDNF, BTG3, C6ORF97, CA2, CALCR, CALD1, CD24, CDA, CDH1, CDH2, CDKN2A, COCH, COL4A1, COL4A2, COL6A1, COMMD10, CREBBP, CRYAB, CXCL12, DKK3, EFEMP2, EGFR, EML1 (includes EG:2009), ESR1, FN1, FYN, GATA3, GFRA1, GLUL, GPD1L, HIST1H2AC, HNMT, ICAM1, IGFBP3, IL18, IL7R, IRS1, KYNU, LDHB, LEPREL1, LOX, LPHN2, MMP1 (includes EG:4312), MPPED2 (includes EG:744), MTUS1, MUC1, NCOA3, NEBL, ORC5L, PBX1, PIK3R3, PTPRK, PTRF, RB1, RBM47, RCN1, SELENBP1, SERPINA3, SERPINB7, SERPINE1, SERPINE2, SERTAD2, SIDT1, SLC24A3, SNCA, SNRPN, STXBP6, TFP12, TGFB1, TNFRSF6B, TPD52L1, TPM2, TUBB2A, UBB, UBE2L6, VAV3, VIM, WWOX, ZNF395</i>	87
Respiratory system development and function	S phase	Entry into S phase of lung cell lines	1.56×10^{-3}	<i>CDKN2A, NCOA3</i>	2
Respiratory system development and function	Proliferation	Proliferation of lung cell lines	2.28×10^{-3}	<i>CDKN2A, FN1, IFI16</i>	3
Skeletal and muscular system development and function	Activation	Activation of muscle cells	1.56×10^{-3}	<i>TGFB1, TGM2</i>	2
Skeletal and muscular system development and function	Ossification	Ossification	4.15×10^{-3}	<i>BMP7, EXT1, MGP, SPARC</i>	4
Skeletal and muscular system development and function	Differentiation	Differentiation of muscle cells	1.34×10^{-2}	<i>IGFBP3, RB1, TGFB1</i>	3
Nucleic acid metabolism	Incorporation	Incorporation of thymidine	2.12×10^{-3}	<i>CDKN2A, EGFR, FN1, IGFBP5, TGFB1</i>	5
Small molecule biochemistry	Incorporation	Incorporation of thymidine	2.12×10^{-3}	<i>CDKN2A, EGFR, FN1, IGFBP5, TGFB1</i>	5

Small molecule biochemistry	Metabolism	Metabolism of steroid	2.91×10^{-3}	<i>AKR1C2, UGT2B15, WWOX</i>	3
Small molecule biochemistry	Metabolism	Metabolism of N-acetylglucosamine	1.35×10^{-2}	<i>CHST6, EXTL2</i>	2
Small molecule biochemistry	Accumulation	Accumulation of doxorubicin	3.07×10^{-3}	<i>ABCB1, DNAJC15</i>	2
Small molecule biochemistry	Synthesis	Synthesis of proteoglycan	4.47×10^{-3}	<i>CTGF, EXT1, TGFB1</i>	3
Small molecule biochemistry	Quantity	Quantity of eicosanoid	1.52×10^{-2}	<i>EGFR, IL18, TGFB1</i>	3
Auditory disease	Nonsyndromic hearing impairment	Nonsyndromic hearing impairment	2.28×10^{-3}	<i>COCH, DFNA5, GRHL2</i>	3
Lipid metabolism	Metabolism	Metabolism of steroid	2.91×10^{-3}	<i>AKR1C2, UGT2B15, WWOX</i>	3
Lipid metabolism	Quantity	Quantity of eicosanoid	1.52×10^{-2}	<i>EGFR, IL18, TGFB1</i>	3
Drug metabolism	Accumulation	Accumulation of doxorubicin	3.07×10^{-3}	<i>ABCB1, DNAJC15</i>	2
Molecular transport	Accumulation	Accumulation of doxorubicin	3.07×10^{-3}	<i>ABCB1, DNAJC15</i>	2
Molecular transport	Uptake	Uptake of phosphate	3.07×10^{-3}	<i>STC1, TGFB1</i>	2
Molecular transport	Quantity	Quantity of eicosanoid	1.52×10^{-2}	<i>EGFR, IL18, TGFB1</i>	3
Ophthalmic disease	Gelatinous drop-like corneal dystrophy	Gelatinous drop-like corneal dystrophy	3.07×10^{-3}	<i>TACSTD2, TGFB1</i>	2
Organismal survival	Survival	Survival of humans	4.70×10^{-3}	<i>EGFR, HMGA2, MYB (includes EG:4602), SERPINE1</i>	4
Cell signaling	Insulin-like growth factor receptor signaling pathway	Insulin-like growth factor receptor signaling pathway	5.04×10^{-3}	<i>IRS1, PIK3R1</i>	2

Hair and skin development and function	Cytostasis	Cytostasis of epithelial cell lines	5.04×10^{-3}	<i>EGFR, TGFB1</i>	2
Hair and skin development and function	Proliferation	Proliferation of epidermal cells	6.46×10^{-3}	<i>EGFR, PTPRK, TGFB1</i>	3
Hair and skin development and function	Differentiation	Differentiation of keratinocytes	6.65×10^{-3}	<i>ANXA1, CDH1, FN1, TGFB1</i>	4
Hair and skin development and function	Adhesion	Adhesion of keratinocytes	1.71×10^{-2}	<i>CDH1, FN1</i>	2
Endocrine system disorders	Gestational diabetes mellitus	Gestational diabetes mellitus	7.45×10^{-3}	<i>IRS1, PIK3R1</i>	2
Endocrine system disorders	Diabetes	Diabetes of mice	1.03×10^{-2}	<i>B2M, IGFBP3</i>	2
Hematological disease	Hematological disorder	Hematological disorder	9.69×10^{-3}	<i>ADA, AKT3, B2M, BCAT1, BRIP1, CALCR, CALD1, COL4A1, CREBBP, CTGF, CXCL12, DUSP4, EGFR, ESR1, FN1, FYN, ICAM1, IGFBP3, IL7R, ORC5L, RET, SERPINE1, TGFB1</i>	23
Renal and urological disease	Renal cancer	Renal cancer	9.69×10^{-3}	<i>ABCB1, CDH1, EGFR, EPCAM, FN1, IL7R, RB1, RET, TUBB2A</i>	9
Renal and urological disease	Renal-cell carcinoma	Renal-cell carcinoma	1.35×10^{-2}	<i>ABCB1, CDH1, EGFR, FN1, RB1, RET, TUBB2A</i>	7
Renal and urological system development and function	Activation	Activation of kidney cell lines	1.03×10^{-2}	<i>TGFB1, TGM2</i>	2
Protein synthesis	Expression	Expression of protein	1.06×10^{-2}	<i>EGFR, EIF5A, FN1, PAIP1, TGFB1</i>	5
Free radical scavenging	Generation	Generation of reactive oxygen species	1.10×10^{-2}	<i>CRYAB, GPX1, MUC1, TGFB1</i>	4
Post-translational modification	Acetylation	Acetylation of protein	1.71×10^{-2}	<i>CREBBP, SNCA</i>	2

* EG = Entrez Gene number.

†Function is defined as the cellular, physiological, or disease process associated with the gene(s) displayed. Functional annotation was provided by Ingenuity Pathway Analysis software (Redwood City, CA). No. of molecules indicates the number of genes from our dataset that have the function listed.

‡ Ingenuity Pathways Analysis software (Redwood City, CA) was used to determine which biological relationships exist between the genes present in Supplementary Tables 1 and 2. The reference set for this analysis was the Ingenuity Knowledge Base (Genes only), and the network analysis was set to direct relationships. All data sources were used to obtain data from patients and all human cell lines and tissues. The stringent filter was also used. Seventy molecules per network and 25 networks were used for this analysis. The p-value associated with Functional Analysis for a dataset was calculated using the Fisher Exact Test. All statistical tests were two-sided.