

Supplementary Materials

Table S1. The relative expression level of 229 distinct proteins found in the scramble of CCA cell lines

| Protein Name | Accession number | ^a Scramble KKU-213A | ^b Scramble KKU-100 |
|--|------------------|-----------------------------------|----------------------------------|
| 2-oxoglutarate dehydrogenase, mitochondrial | A0A0D9SFS3 | 17.87760 | 18.38777 |
| 5-azacytidine induced 1 | B2RN10 | 17.63603 | 16.14597 |
| Activating signal cointegrator 1 complex subunit 3 | Q8N3C0 | 14.59243 | 15.96103 |
| AD 2 | Q9Y6Z1 | 15.15167 | 14.29617 |
| ADAM metallopeptidase with thrombospondin type 1 motif, 13, isoform CRA_a | A0A024R8G4 | 19.39700 | 19.87283 |
| Adenomatous polyposis coli protein isoform t | A0A224AV41 | 15.61146 | 16.76630 |
| Adhesion G protein-coupled receptor B2 | A2A3C6 | 17.09320 | 16.84173 |
| ALS2 | A0A2X0SFA0 | 15.71283 | 15.95290 |
| Alternative protein ZNF606 | L8ECK5 | 18.01287 | 17.70510 |
| Amiloride-sensitive sodium channel subunit alpha 1 | C5HTY8 | 15.53853 | 15.90037 |
| Arf-GAP with Rho-GAP domain, ANK repeat and PH domain-containing protein 3 | G5E9Y3 | 16.08333 | 16.77900 |
| ARHGEF12 | A0A2X0SFF5 | 16.04597 | 15.10513 |
| ATPase MORC2 | Q9Y6X9 | 14.79150 | 17.62440 |
| ATP-binding cassette sub-family A member 10 | Q8WWZ4 | 17.59753 | 16.79400 |
| ATP-binding cassette sub-family A member 2 | A0A087W XK5 | 13.94003 | 14.19603 |
| ATP-binding cassette, sub-family C, member 3 isoform MRP3 variant | Q59H05 | 16.05830 | 14.61843 |
| Aurora kinase A-interacting protein | Q9NWT8 | 15.30900 | 16.01690 |
| BAG family molecular chaperone regulator 3 | C9JFK9 | 15.89267 | 14.33363 |
| BAH and coiled-coil domain-containing protein 1 | F8WBW8 | 15.41506 | 17.83900 |
| Beta-klotho | Q86Z14 | 16.48230 | 15.33950 |
| Brain my037 protein | Q9H3I9 | 13.24126 | 12.97372 |
| Brain specific protein, isoform CRA_a | A0A024R702 | 18.11290 | 18.01560 |
| Breast carcinoma amplified sequence 2 variant | Q53HE3 | 14.73707 | 15.91413 |
| C10orf54 protein | Q2TA85 | 14.29267 | 15.93750 |
| C6orf134 protein | Q8TAV6 | 16.22963 | 14.86110 |
| Calcium and integrin-binding protein 1 | Q99828 | 17.53020 | 15.07003 |
| Calpain-12 | Q6ZSI9 | 18.27563 | 19.84010 |
| CCR4-NOT transcription complex, subunit 4, isoform CRA_b | A0A024R781 | 17.79387 | 17.79763 |
| CDC42 | P60953 | 18.18233 | 16.82080 |
| Centaurin-alpha 2 protein variant | Q53HQ4 | 15.02563 | 15.27843 |
| Centromere protein F | P49454 | 18.46990 | 16.87707 |

Table S1. The relative expression level of 229 distinct proteins found in the scramble of CCA cell lines (cont.)

| Protein Name | Accession number | ^a Scramble KKU-213A | ^b Scramble KKU-100 |
|---|------------------|-----------------------------------|----------------------------------|
| Centrosomal protein 350kDa, isoform CRA_a | A0A024R914 | 15.37743 | 16.94347 |
| Centrosomal protein of 120 kDa | Q8N960 | 18.17633 | 19.09570 |
| Centrosomal protein of 135 kDa | Q66GS9 | 16.49937 | 15.45380 |
| Chromobox homolog 4 (Pc class homolog, Drosophila), isoform CRA_a | A0A0S2Z5B2 | 13.94140 | 17.20417 |
| Chronic lymphocytic leukemia up-regulated protein 1 | Q5K131 | 18.04747 | 16.93763 |
| Cilia- and flagella-associated protein 47 | Q6ZTR5 | 17.62390 | 16.24830 |
| ciliary rootlet coiled-coil protein 2 | H7BZ55 | 16.08763 | 16.99510 |
| Cis-aconitate decarboxylase | H7C1Q4 | 15.31716 | 15.37867 |
| Coiled-coil domain-containing protein 188 | H7C350 | 14.41050 | 13.98323 |
| Collagen, type IV, alpha 4, isoform CRA_a | J3KNM7 | 17.25637 | 16.62053 |
| Collagen, type VI, alpha 3 | D9ZGF2 | 16.37223 | 15.43660 |
| COMM domain-containing protein 7 | V9GY66 | 15.81030 | 14.28113 |
| CREB3 regulatory factor | Q8IUR6 | 17.62947 | 17.87150 |
| cyclin-Y-like protein 3 | P0C7X3 | 13.08580 | 13.41513 |
| Dedicator of cytokinesis 1 | B2RUU3 | 17.06073 | 17.23113 |
| Dedicator of cytokinesis 8, isoform CRA_a | D3DRH1 | 18.32153 | 15.47723 |
| Dedicator of cytokinesis protein 3 | Q8IZD9 | 16.42507 | 14.05907 |
| Delta globin | A0N071 | 16.63513 | 15.81870 |
| Disintegrin and metalloproteinase domain-containing protein 11 | K7EKA8 | 13.82340 | 15.20453 |
| Dmx-like 2, isoform CRA_c | A0A024R5V2 | 19.17487 | 15.25170 |
| DNA excision repair protein ERCC-6 | Q03468 | 14.87870 | 15.02820 |
| DNA topoisomerase 1 | P11387 | 13.65473 | 15.18317 |
| DNA-repair protein complementing XP-G cells variant | Q59FZ7 | 16.87553 | 16.00773 |
| E1A-binding protein p400 | Q96L91 | 13.67566 | 16.80097 |
| E3 ubiquitin-protein ligase RBBP6 | A0A2R8Y880 | 17.89767 | 13.64367 |
| E3 ubiquitin-protein ligase RNF213 | A0A0A0MTR7 | 18.60923 | 19.21367 |
| E3 ubiquitin-protein ligase TRAF7 | Q6Q0C0 | 14.88397 | 18.38587 |
| Ectonucleotide pyrophosphatase/phosphodiesterase family member 2 | E7EUF1 | 15.43807 | 14.42127 |
| EF-hand calcium-binding domain-containing protein 13 | Q8IY85 | 16.58067 | 14.71837 |
| EMILIN-2 | Q9BXX0 | 15.78950 | 16.04143 |
| Endothelin-converting enzyme-like 1 | O95672 | 15.84047 | 18.22840 |
| Envoplakin | Q92817 | 17.29983 | 18.32663 |

Table S1. The relative expression level of 229 distinct proteins found in the scramble of CCA cell lines (cont.)

| Protein Name | Accession number | ^a Scramble KKU-213A | ^b Scramble KKU-100 |
|--|------------------|-----------------------------------|----------------------------------|
| Eukaryotic translation initiation factor 4 gamma, 3 variant | Q59GJ0 | 24.22767 | 22.38883 |
| Family with sequence similarity 184, member A | Q6P9G8 | 15.25993 | 15.02410 |
| FBLN7 protein | Q8IY13 | 17.98420 | 17.12727 |
| F-box/WD repeat-containing protein 11 | Q9UKB1 | 15.32986 | 13.77667 |
| Fc receptor-like protein 2 | Q96LA5 | 22.58750 | 22.53563 |
| FEX1 | Q0PNF2 | 15.73710 | 16.98583 |
| FLJ00077 protein | Q9H7J7 | 16.64933 | 13.34683 |
| G protein-coupled receptor 6 | E7EP76 | 15.31173 | 14.27837 |
| GCN1 general control of amino-acid synthesis 1-like 1 (Yeast), isoform CRA_b | A0A024RBS1 | 14.81610 | 12.99043 |
| General transcription factor II-I repeat domain-containing protein 2B | A0A494C1N8 | 15.99300 | 15.64023 |
| Glycoprotein receptor gp330/megalin | Q7Z5C1 | 16.99430 | 16.54913 |
| Golgin subfamily A member 8S | H3BPF8 | 16.43907 | 13.92670 |
| GPI mannosyltransferase 2 | Q9NUD9 | 14.42080 | 15.28547 |
| GTP-binding protein 1 | H0YH10 | 18.26700 | 15.61947 |
| Guanine nucleotide-binding protein-like 3 | Q9BVP2 | 15.71447 | 19.19287 |
| HCFC1 | A0A223PQH6 | 18.57467 | 16.97147 |
| HCG1990170, isoform CRA_a | D6W4X8 | 24.53713 | 20.85010 |
| HCG2009921, isoform CRA_c | Q96MT5 | 14.43366 | 16.01797 |
| HCG2036600, isoform CRA_a | D3DNY4 | 17.74387 | 18.13080 |
| HCG2039413, isoform CRA_a | A0A024RE15 | 16.93923 | 17.15690 |
| HCG401289, isoform CRA_e | A8MT14 | 13.20563 | 15.64897 |
| Heat shock 105kDa/110kDa protein 1, isoform CRA_a | A0A024RDQ0 | 16.22890 | 15.96347 |
| HERC2-like protein 3 | Q9BVR0 | 14.71810 | 12.78303 |
| Histone acetyltransferase p300 | Q09472 | 15.70146 | 17.28517 |
| Histone deacetylase 5 | Q9UQL6 | 15.72247 | 16.61137 |
| Histone H1.2 | P16403 | 14.06360 | 14.74320 |
| Histone PARylation factor 1 | Q9NWX4 | 17.84790 | 16.72517 |
| Histone-lysine N-methyltransferase KMT5C | A0A0D9SF94 | 14.14720 | 14.02173 |
| Histone-lysine N-methyltransferase SETD1B | A0A0A0MQV9 | 16.24203 | 13.90673 |
| IG c109_heavy_IGHV1-69_IGHD5-24_IGHJ4 | A0A5C2GDC2 | 14.32250 | 14.08840 |
| IG c1159_heavy_IGHV4-31_IGHD3-22_IGHJ4 | A0A5C2GTJ0 | 13.59333 | 13.92013 |
| IG c1235_heavy_IGHV3-74_IGHD2-8_IGHJ6 | A0A5C2GPC0 | 13.69417 | 16.14020 |

Table S1. The relative expression level of 229 distinct proteins found in the scramble of CCA cell lines (cont.)

| Protein Name | Accession number | ^a Scramble KKU-213A | ^b Scramble KKU-100 |
|--|------------------|-----------------------------------|----------------------------------|
| IG c1531_heavy_IGHV3-74_IGHD3-3_IGHJ4 | A0A5C2GVM4 | 16.59933 | 18.09440 |
| IG c780_heavy_IGHV3-11_IGHD4-23_IGHJ4 | A0A5C2GQG4 | 16.48320 | 16.31570 |
| IGH + IGL c482_heavy_IGHV1-69_IGHD4-23_IGHJ3 | A0A5C2GBQ8 | 16.84727 | 17.51350 |
| IGH c505_heavy__IGHV2-5_IGHD6-19_IGHJ4 | A0A5C2GB92 | 13.49936 | 15.63530 |
| Inhibitor of Bruton tyrosine kinase | E9PDR5 | 14.20103 | 15.63863 |
| Inositol 1,4,5-trisphosphate receptor type 3 | Q14573 | 17.08020 | 17.53343 |
| Insulin-like growth factor 2 mRNA-binding protein 3 | O00425 | 18.44683 | 17.82880 |
| Interferon-induced protein with tetratricopeptide repeats 1 | P09914 | 12.31237 | 12.55710 |
| Interleukin-34 | J3QQT3 | 14.13210 | 14.31483 |
| Intraflagellar transport 81 homolog (Chlamydomonas), isoform CRA_c | A0A024RBJ1 | 17.00607 | 18.40990 |
| Kallikrein-8 | A0A0A0MQY9 | 12.50293 | 12.89183 |
| Kelch repeat and BTB (POZ) domain containing 4 variant | Q53HN0 | 15.44260 | 17.56830 |
| Keratocan | O60938 | 15.84103 | 14.07130 |
| KIAA0423, isoform CRA_a | G3XAE9 | 16.53570 | 17.97470 |
| KIAA0649, isoform CRA_a | A0A024R8E9 | 14.87946 | 16.80693 |
| KIF26A variant protein | B0IIR4 | 17.37247 | 17.87057 |
| Kinesin-like protein KIF19 | Q2TAC6 | 16.63303 | 18.37177 |
| La-related protein 6 | Q9BRS8 | 14.78993 | 17.18317 |
| Large tegument protein | A0A126LAX2 | 17.44143 | 16.35543 |
| Leucine rich repeat containing 24 | G3V1D8 | 17.60197 | 15.57600 |
| Leucine-, glutamate- and lysine-rich protein 1 | J3KP02 | 13.99303 | 15.94800 |
| Leucine-rich PPR-motif containing | E5KNY5 | 18.22730 | 15.60623 |
| Lymphocyte antigen 75 | O60449 | 17.46047 | 16.89870 |
| Lysine-specific demethylase 6B | O15054 | 15.78803 | 14.22957 |
| Matrix metalloproteinase 15 (Membrane-inserted), isoform CRA_a | A0A024R6U8 | 15.80653 | 14.85597 |
| Matrix-remodelling associated 8, isoform CRA_a | A0A024R0A0 | 13.47153 | 16.93940 |
| Metastasis-suppressor KiSS-1 | A0A0D9SES6 | 17.80073 | 16.97213 |
| Microtubule-associated tumor suppressor candidate 2 | Q5JR59 | 18.45887 | 16.33460 |
| Molybdenum cofactor synthesis 1, isoform CRA_c | A0A024RD17 | 14.27760 | 14.20430 |
| MORC family CW-type zinc finger protein 4 | Q8TE76 | 15.35043 | 16.66670 |
| Mucin 5AC, oligomeric mucus/gel-forming | A7Y9J9 | 15.19480 | 17.36117 |
| Mucin-16 | Q8WXI7 | 16.23690 | 16.68887 |

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|---|------------------|-----------------------------------|----------------------------------|
| Myosin-10 | P35580 | 16.25173 | 16.91530 |
| Myotubularin-related protein 6 | Q9Y217 | 15.29543 | 14.81380 |
| N-acetylaspartate synthetase | Q8N9F0 | 13.55867 | 13.87167 |
| NAD-dependent protein deacetylase sirtuin-3, mitochondrial | E9PM52 | 14.40500 | 14.95453 |
| NADH dehydrogenase (Ubiquinone) 1 beta subcomplex, 10, 22kDa, isoform CRA_b | A8K761 | 13.96456 | 12.50590 |
| Neurobeachin | Q5T321 | 15.61820 | 14.75657 |
| Neurofilament heavy polypeptide | P12036 | 19.27670 | 17.23753 |
| Neurogenic locus notch homolog protein 4 | A0A1U9X943 | 15.20473 | 15.55113 |
| Neurologin 3 isoform A | X5DNV3 | 16.57070 | 18.54673 |
| NMDA receptor regulated 1-like | A4FU51 | 14.40600 | 16.02517 |
| NOELIN3_V2 | Q6IMJ0 | 16.00263 | 15.12617 |
| NRAP protein | A0AVL2 | 17.68630 | 17.09530 |
| Nuclear factor 1 | Q7Z3K7 | 15.87430 | 15.30123 |
| Nuclear protein MDM1 | Q8TC05 | 15.67590 | 15.73000 |
| Nuclear receptor-interacting protein 3 | E9PIW3 | 18.00467 | 17.19533 |
| Nucleolar protein 8, isoform CRA_a | A0A024R250 | 14.18330 | 17.07353 |
| Olfactory receptor 1A1 | Q9P1Q5 | 18.71137 | 20.03993 |
| PDZ domain-containing RING finger protein 4 | Q6ZMN7 | 16.82587 | 14.73747 |
| Pecanex-like protein 2 | A6NKB5 | 16.40663 | 16.47937 |
| Phosphatidylinositol 4-kinase alpha | P42356 | 14.30720 | 15.06517 |
| PLA2R1 protein | B7ZML4 | 16.47977 | 14.66623 |
| Pleckstrin homology domain-containing family G member 1 | A0A2R8Y2B1 | 14.63725 | 14.47950 |
| PLEKHG3 | A0A2X0SFH4 | 18.76337 | 18.87670 |
| Postreplication repair protein hRAD18p variant | Q53H10 | 18.58957 | 19.58557 |
| Pre-mRNA-processing factor 40 homolog A | O75400 | 17.30100 | 16.85277 |
| PREX1 | A0A2X0SFH1 | 16.83563 | 16.29473 |
| Probable ATP-dependent RNA helicase DDX6 | P26196 | 13.92507 | 15.68617 |
| Programmed cell death 11, isoform CRA_a | A0A3B3IUD7 | 16.57407 | 18.52967 |
| Prohibitin | D6RBK0 | 13.96413 | 14.79397 |
| Protein Daple | Q9P219 | 15.43177 | 17.67827 |
| Protein dopey-1 | Q5TA12 | 14.01883 | 16.26517 |
| Protein FAM131C | Q96AQ9 | 12.22923 | 13.09577 |

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| Protein Name | Accession number | ^a Scramble KKU-213A | ^b Scramble KKU-100 |
|--|------------------|-----------------------------------|----------------------------------|
| Protein mono-ADP-ribosyltransferase PARP4 | Q9UKK3 | 16.54940 | 15.60237 |
| PURA protein | Q2NLC9 | 17.27140 | 17.25970 |
| RAD50-interacting protein 1 | Q6NUQ1 | 14.95506 | 12.60600 |
| Ras association (RalGDS/AF-6) and pleckstrin homology domains 1, isoform CRA_b | C9K0J5 | 15.56210 | 14.15890 |
| Ras protein-specific guanine nucleotide-releasing factor 1, isoform CRA_b | Q8IUU5 | 15.53436 | 17.56273 |
| Retinal-specific phospholipid-transporting ATPase ABCA4 | P78363 | 15.82810 | 16.77223 |
| Retinitis pigmentosa 1-like 1 protein | Q8IWN7 | 16.67670 | 14.04800 |
| Rho guanine nucleotide exchange factor 12 | E9PMR6 | 16.54123 | 15.66653 |
| Rho guanine nucleotide exchange factor 17 | H0YFZ8 | 20.57300 | 17.17850 |
| Rhodopsin | V9H017 | 17.62013 | 18.56350 |
| Ribosomal protein L15 | E7ENU7 | 16.84987 | 14.67147 |
| RNA-binding protein 25 | P49756 | 16.56587 | 17.44307 |
| Rootletin | Q5TZA2 | 18.92520 | 16.37253 |
| RRP12-like protein | Q5JTH9 | 16.64320 | 14.70807 |
| SAPS domain family, member 3, isoform CRA_c | A0A024R5F3 | 15.86840 | 16.32363 |
| SARDH protein | Q05BT1 | 15.34596 | 14.85590 |
| Serine/threonine-protein kinase MRCK gamma | Q6DT37 | 16.92260 | 16.37440 |
| Serine/threonine-protein kinase ULK4 | Q96C45 | 16.22613 | 14.61367 |
| SLC12A2 protein | B7ZM24 | 15.55646 | 15.77110 |
| SOGA | F2FB34 | 16.84667 | 17.00327 |
| Solute carrier 16 (Monocarboxylic acid transporters), member 8, isoform CRA_a | A0A024R1N8 | 17.47080 | 17.54893 |
| SOS1 | Q53SF8 | 14.81297 | 14.99180 |
| Splicing factor, proline- and glutamine-rich | Q2I379 | 13.24550 | 14.87215 |
| SPRY domain containing 3, isoform CRA_b | A0A024RAX4 | 19.42803 | 18.80773 |
| Sterile alpha motif domain-containing protein 9-like | Q8IVG5 | 13.75997 | 15.12553 |
| SUMO-specific isopeptidase USPL1 | Q5W0Q7 | 15.11830 | 14.59143 |
| SWI/SNF complex subunit SMARCC2 | F8VXC8 | 14.42036 | 15.65633 |
| Taste receptor type 2 member 46 | P59540 | 12.54436 | 16.86867 |
| T-cell lymphoma invasion and metastasis 2 variant | Q59EY7 | 16.10230 | 16.28703 |
| Testis tissue sperm-binding protein Li 71n | A0A140VKF4 | 15.39276 | 13.73707 |
| Threonine--tRNA ligase 2, cytoplasmic | A2RTX5 | 17.90647 | 17.04920 |
| Thrombospondin type-1 domain-containing protein 7A | A0A087WY08 | 17.01590 | 15.40557 |

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|---|------------------|-----------------------------------|----------------------------------|
| Thyroxine-binding globulin | P05543 | 16.44707 | 18.77900 |
| TNRC4 protein | Q9BTF3 | 16.33397 | 17.22477 |
| Trans-3-hydroxy-L-proline dehydratase | Q96EM0 | 14.12070 | 13.77217 |
| Transmembrane protein 229B | G3V2T8 | 14.82793 | 12.89740 |
| Trichohyalin | Q07283 | 16.68580 | 16.72860 |
| Trinucleotide repeat containing 15, isoform CRA_a | A0A024R4A5 | 13.84173 | 14.66457 |
| TRPC4 protein | Q3KR50 | 17.21833 | 16.11907 |
| Truncated platelet membrane glycoprotein Ib beta | A0A140GX62 | 13.60433 | 14.77530 |
| Truncated RHCE | J9ZZ28 | 18.99867 | 18.10427 |
| TUBGCP6 protein | B2RWN4 | 16.26080 | 16.56540 |
| Tubulin tyrosine ligase-like family, member 1, isoform CRA_a | A0A024R4U6 | 13.11090 | 15.70617 |
| Tumor rejection antigen (Gp96) 1 variant | Q59FC6 | 17.29333 | 18.52753 |
| Tyrosine-protein kinase BAZ1B | Q9UIG0 | 15.19106 | 17.72040 |
| U3 small nucleolar RNA-interacting protein 2 | O43818 | 16.73207 | 14.76257 |
| Uncharacterized protein CAD | Q53SY7 | 17.11370 | 13.54680 |
| Uncharacterized protein DKFZp686A1195 | Q6MZU1 | 15.03130 | 16.42427 |
| Uncharacterized protein DKFZp686L0869 | Q7Z3U5 | 14.45300 | 17.45910 |
| Uncharacterized protein DKFZp686L2367 | Q63HJ5 | 17.28743 | 17.08380 |
| Uncharacterized protein EPAS1 | Q53SM6 | 15.06363 | 14.10163 |
| Uncharacterized protein KIAA1841 | Q6NSI8 | 14.40963 | 15.84723 |
| Unconventional myosin-Va | A0A087WY00 | 17.21363 | 15.56977 |
| Vacuolar protein sorting-associated protein 13B | Q7Z7G8 | 15.46263 | 16.61873 |
| Vacuolar protein sorting-associated protein 13C | Q709C8 | 16.45590 | 18.08480 |
| Vitamin D binding protein protein | Q6LDC6 | 22.42233 | 20.99427 |
| von Willebrand factor A domain-containing protein 5B2 | E9PF42 | 15.77030 | 16.46133 |
| YY1AP-related protein1 | A4PB67 | 15.91973 | 16.14080 |
| Zinc finger and BTB domain-containing protein 46 | Q86UZ6 | 15.81593 | 14.15240 |
| Zinc finger and SCAN domain-containing protein 20 | P17040 | 18.36460 | 16.58563 |
| Zinc finger MYM-type protein 6 | O95789 | 13.46780 | 16.11133 |
| Zinc finger protein 268 | F5H467 | 15.31090 | 16.09290 |
| Zinc finger protein 40 | F5H212 | 15.72953 | 16.81953 |
| Zinc finger protein 42 (Myeloid-specific retinoic acid-responsive), isoform CRA_a | A0A024R4R6 | 17.47743 | 15.90940 |

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|---------------------------------|------------------|-----------------------------------|----------------------------------|
| Zinc finger protein 536 | K7EQN6 | 14.72180 | 16.48410 |
| Zinc finger protein 579 | Q8NAF0 | 15.43530 | 17.13520 |
| zinc finger protein 726P1 | Q15940 | 14.38427 | 15.56697 |
| Zinc finger protein 804A | B9EGN2 | 15.25293 | 16.16153 |
| Zinc finger RNA-binding protein | Q96KR1 | 15.07440 | 16.33907 |
| ZMAT1 protein | A7MD47 | 16.62973 | 16.62247 |

^aValues represent the log2 fold change of precursor signal intensities in particular scramble KKKU-213A compared to siRNA KKKU-213A detected by LC-MS/MS

^bValues represent the log2 fold change of precursor signal intensities in particular scramble KKKU-100 compared to siRNA KKKU-100 detected by LC-MS/MS

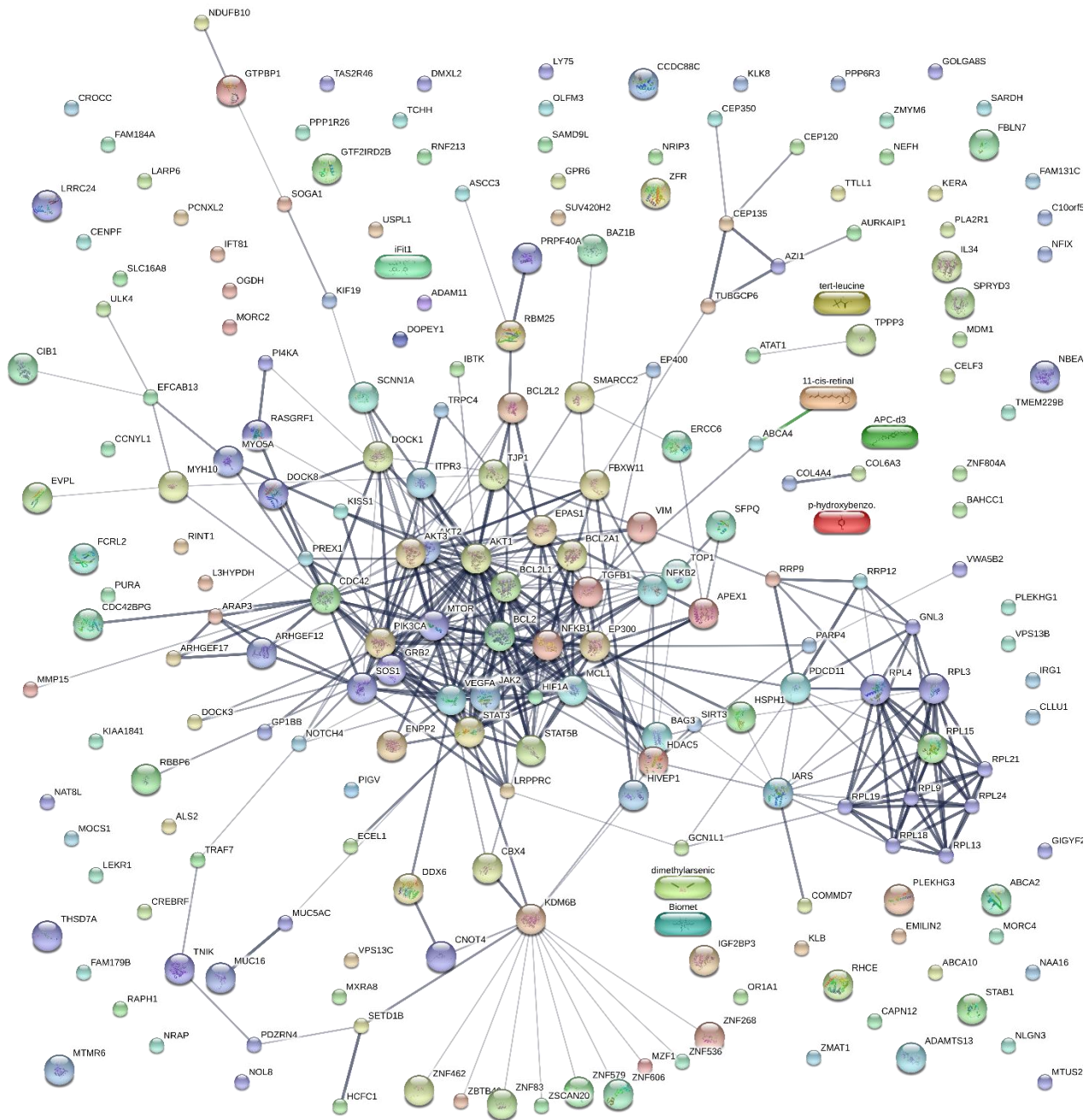


Figure S1. Protein interaction networks of APEX1 and the related proteins of scramble CCA cells.