**Supplementary Table 7.** Genes whose expression in CD44+/CD24– MCF-7/ADR cells compared with CD44+/CD24+ MCF-7/ADR cells was statistically significantly different, with a fold change of greater than 2\*

Gene symbol	Gene name	Location	Type(s)	Fold change
ABCA1	ATP-binding cassette, subfamily A (ABC1), member 1	Plasma membrane	Transporter	5.386
ABCA13	ATP-binding cassette, subfamily A (ABC1), member 13	Unknown	Transporter	-2.957
ABCA5	ATP-binding cassette, subfamily A (ABC1), member 5	Plasma membrane	Transporter	-2.274
ABCD2	ATP-binding cassette, subfamily D (ALD), member 2	Cytoplasm	Transporter	-3.018
ACER1	Alkaline ceramidase 1	Cytoplasm	Enzyme	-2.642
AHR	Aryl hydrocarbon receptor	Nucleus	Ligand-dependent nuclear receptor	-3.006
ANXA1	Annexin A1	Plasma membrane	Other	-2.419
APEX1	APEX nuclease (multifunctional DNA repair enzyme) 1	Nucleus	Enzyme	-2.689
ATOX1	ATX1 antioxidant protein 1 homolog (yeast)	Cytoplasm	Transporter	-2.973
BAG1	BCL2-associated athanogene	Cytoplasm	Other	-3.071
BCL2L1	BCL2-like 1	Cytoplasm	Other	-2.228
BIRC5	Baculoviral IAP repeat-containing 5	Cytoplasm	Other	-2.236
CCNE1	Cyclin E1	Nucleus	Transcription regulator	2.19
CDKN1A	Cyclin-dependent kinase inhibitor 1A (p21, Cip1)	Nucleus	Other	2.338
CFLAR	CASP8 and FADD-like apoptosis regulator	Cytoplasm	Other	-2.007
CIAPIN1	Cytokine induced apoptosis inhibitor 1	Cytoplasm	Other	2.811
CLDN3	Claudin 3	Plasma membrane	Transmembrane receptor	2.555
CLDN5	Claudin 5	Plasma membrane	Other	1.997
CLPTM1L	CLPTM1-like	Unknown	Other	-2.227

DOT1L	DOT1-like, histone H3 methyltransferase (Saccharomyces cerevisiae)	Nucleus	Phosphatase	2.797
FAS	Fas (TNF receptor superfamily, member 6)	Plasma membrane	Transmembrane receptor	2.476
GGT1	γ-Glutamyltransferase 1	Cytoplasm	Enzyme	2.965
GSTM2	Glutathione S-transferase μ2 (muscle)	Cytoplasm	Enzyme	4.358
GSTM4	Glutathione S-transferase µ4	Cytoplasm	Enzyme	4.407
GSTO1	Glutathione S-transferase ω1	Cytoplasm	Enzyme	2.239
HSF1	Heat shock transcription factor 1	Nucleus	Transcription regulator	-2.051
HSPB1	Heat shock 27kda protein 1	Cytoplasm	Other	2.055
HSPH1	Heat shock 105kda/110kda protein 1	Cytoplasm	Other	-2.507
IL6	Interleukin 6 (interferon, β2)	Extracellular space	Cytokine	-2.351
JUN	Jun oncogene	Nucleus	Transcription regulator	2.175
KCNMA1	Potassium large conductance calcium-activated channel, subfamily M, α member 1	Plasma membrane	Ion channel	3.231
LBR	Lamin B receptor	Nucleus	Enzyme	-2.744
MDM2	Mdm2 p53 binding protein homolog (mouse)	Nucleus	Transcription regulator	-4.209
MLH3	MutL homolog 3 (Escherichia coli)	Nucleus	Other	2.218
MMP9	Matrix metallopeptidase 9 (gelatinase B, 92kda gelatinase, 92kda type IV collagenase)	Extracellular space	Peptidase	3.064
MT2A	Metallothionein 2A	Cytoplasm	Other	-2.327
POLB	Polymerase (DNA directed), β	Nucleus	Enzyme	-2.373
RAD23A	RAD23 homolog A (S. cerevisiae)	Nucleus	Other	-2.051
RAD51	RAD51 homolog (reca homolog, <i>E. coli</i> ) ( <i>S. cerevisiae</i> )	Nucleus	Enzyme	-3.171
RPL36 (includes EG:25873)	Ribosomal protein L36	Cytoplasm	Other	-2.647

SIRT2	Sirtuin (silent mating type information regulation 2 homolog) 2 (S. cerevisiae)	Nucleus	Transcription regulator	1.981
SIRT3	Sirtuin (silent mating type information regulation 2 homolog) 3 (S. cerevisiae)	Cytoplasm	Enzyme	2.008
SIRT4	Sirtuin (silent mating type information regulation 2 homolog) 4 (S. cerevisiae)	Cytoplasm	Enzyme	-2.881
SLC16A2	Solute carrier family 16, member 2 (monocarboxylic acid transporter 8)	Plasma membrane	Transporter	-3.999
SLC19A3	Solute carrier family 19, member 3	Plasma membrane	Transporter	2.659
SLC22A1	Solute carrier family 22 (organic cation transporter), member 1	Plasma membrane	Transporter	-2.826
SLC22A4 (includes EG:6583)	Solute carrier family 22 (organic cation/ergothioneine transporter), member 4	Plasma membrane	Transporter	2.621
SLC25A15	Solute carrier family 25 (mitochondrial carrier; ornithine transporter) member 15	Cytoplasm	Transporter	2.437
SLC29A1	Solute carrier family 29 (nucleoside transporters), member 1	Plasma membrane	Transporter	2.127
SLC3A1	Solute carrier family 3 (cystine, dibasic and neutral amino acid transporters, activator of cystine, dibasic and neutral amino acid transport), member 1	Plasma membrane	Transporter	4.425
SLC7A10	Solute carrier family 7, (neutral amino acid transporter, y+ system) member 10	Plasma membrane	Transporter	-2.025
SLC7A11	Solute carrier family 7, (cationic amino acid transporter, y+ system) member 11	Plasma membrane	Transporter	2.264
SLCO4A1	Solute carrier organic anion transporter family, member 4A1	Plasma membrane	Transporter	2.54
SRC	V-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)	Cytoplasm	Kinase	-2.028
STARD4	Star-related lipid transfer (START) domain containing 4	Cytoplasm	Transporter	2.098

TAP1	Transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)	Cytoplasm	Transporter	-2.026
TCEAL4	Transcription elongation factor A (SII)-like 4	Unknown	Other	-2.536
TNF	Tumor necrosis factor (TNF superfamily, member 2)	Extracellular space	Cytokine	2.149
TPRKB	TP53RK binding protein	Unknown	Other	-3.287
TRAF1	TNF receptor-associated factor 1	Cytoplasm	Other	2.343
UBL5	Ubiquitin-like 5	Cytoplasm	Other	-2.528
UGCG	UDP-glucose ceramide glucosyltransferase	Cytoplasm	Enzyme	2.407
XRCC4	X-ray repair complementing defective repair in Chinese hamster cells 4	Nucleus	Other	2.308

<sup>\*</sup>APEX = Apurinic-apyrimidinic endonuclease; ATX1 = antioxidant protein; BCL2 = B-cell CLL/lymphoma; IAP = Inhibitor of apoptosis protein; CASP8 = apoptosis-related cysteine peptidase; FADD = FAS-associated death domain protein; CLPTM1 = cleft lip and palate associated transmembrane protein; DOT1 = disruptor of telomeric silencing; TNF = Tumor necrosis factor; Mdm2 = mouse double minute 2; MutL= *E. coli* DNA mismatch repair gene; START = Star-related lipid transfer domain; MDR/TAP = Multidrug resistant/ transporter associated with antigen processing; TP53RK = TP53 regulating kinase.