ADCY2 HTR7 LPAR3 PTGDR GPR1 GRM8 OXER1 PALM PARY12 SSTR1 GNA14 SSTR5 CORT ADGRD1 -MARCO HTR1B G_protein Coupled receptor signaling pathway, coupled to cyclic nucleotide second messenger PTGDR2 CYSLTR1 **HGF** FZD4 ADCY8 HRH2 MYLK2 TBXA2R RGS2 TNNI3
PR TTR1D FDXN4 WNK3 ADM CHRM2 TTBK1 DNM1, GRPR TNNC2 CNTFR FENT LSAMP CX3CR1 GRIN2D ADRA1B HSPB7 AKAP6 KCNIP4 ¢ACNA2D1/SLC22A5/CACNB3 PTODS PITX3 HOXB8 HTR2A ACPP EXPLOYER KCNH2FXYD1 TNFRSF21 TNFRS NRGN SNAP25
RAG1 NRAS1
CHRNA5 BBS12 GABRA5
CHRNA5 BBS12 GABRA5 B4GALT2 FUOM CHRNA5 BBS12 GABRA5 SLC8A1 KCNG2 CLIC2

CRABP2 GRIA1 BTG2 EGR2 CALB1 CAMK2B P2RY1 ATP2B1 PTGER3 CELF2 POMC CYL

MAFF DEAFT NCOA2 SEZ6L PREX2 FOS TACR1 DRD4 DRD1 KCNMB4 TNNI3K ANK2ATP2B3 CACMB4

COL17A1 FOU2F3 PYY2 LRRN4 SORCS3 THRB FGF13 regulation of system 660448 R1H3 BDKRB2 ACSS1 IL

DACT2 MSX2 EN1 NTRK2 CEND1 THRB PTGS2 GRIN2A PLCE1 IL6ST ACAT2 ERG28 PMEL

SLC4A7 RIPOR2 ZDHHC21 ZFP36 CREBRA MYO15A EGFR SLC6A3 LEPR PTAFR PIK3CG FDX1 TM7SF2 RBP4 SGPP2

ABCB6 REG3G SDINK6 ATP8A2 FDEEL PLCB1 GCN14 SNCA SPHK1 ADH1A BEST TM7SF2 RBP4 SGPP2 / ÆRG28/PMEL/ ERRFI1 PLEB1 GCNT4 SNCA SPHK1 ADH1A-BEST TM7SF2 RBP4 FRAS1 SOSTDC1 EVPL HPSE ADAM28 CLN6 APP WNT5A RNF180 CYP7B1 MAOB PCK1 PCSK9 PCK2 CYP46A1 ACSS2 F12

R2D KRT17 CPD50A LANDAR TCF15 APO Program SUNTABLY COMPOUND METABOLIC PROCESS1 DDT SULT1E1 PLB1 MVK CLOCK SPRR2D KRT17 AQP3 TGM3 —SOX21 SPRR2A LAMA5
HOXC13 CSTA PKP1 SCEL SPINKSkin development STRN COHPOSITIVE DESCRIPTION of Inflammator NESTENSE EBP DUOXA2 ADH1B DHCR7 TLR6 COL1A2 APOC1 APOA1 MVD WCER2 CUBN S100A12 RPS19 PPL SPRR2B PRSS8 SQX9 CTSV LAMC2 DCT SORU1 PTS FOFTE PLDA ELF3 GRIZONIUTAT structure organization4 LCAT CCL18 TNRSF4 DLL1 MET PCSK6 WNT10B FERMT1 COLSA1 CYR61 DSC3 HPN ABCG1 DISP3 ALOXE3 HES1 MADCAM1 COL5A3 TIMP!

KLKB1 PLA2G2A PLA2G7 MAG FASN CX3CL1 DUOXA1 CYP24A1 TLR3 TI

STMN1 CREB3L1 COL2A1 ACAN DDR2 NCAN homoph@ADM3adhesion via plasma membrane adhesion molecules PCDHG8 PCDHG82 S10

ICAM5 ELN HAS2 KDR KLK2 VCAM1 PLXNB3 PCDHGB7 PCDHGA9 PCDHGA9

CTRR1 P4HR MMP9 17444 COL8A2 TILL1 1.55. PITX2 ALOXE3 HES1 MADCAM1 COL5A3 TIMP PLXNB3 PCDHGB7 PCDHG9 PCDHGA5PCDHGB5 CLDN18-AMIGO1 CTRB1 P4HB MMP9 LAMA4 COL8A2 TLL1 LRP1 ALB PCDHGBCDHR1 CDH15PCDHGA5PCDHGB5 CDHA9

PDPN GPM6B COL14A1 EFEMP2 ICAM4 IGSF9B PCDHGBCDHR1 CDH15PCDHGA8CDHA12 CLDN7

MFAP4 SPINT1 MATN3 COL10A1 MELTF PCDHGA12 PCDHGACDHB16PCDH10 CRB3 PCDHGA4 SELP

HSD17B12 CHADL FLFT2 MFAP5 COL4A2 PCDHGA10 PCDHGC4 PCDHGA1 CLDN2 CDH2

ITGA2B KLK4 JAM3 CTRBC CDH40 SOCIAL COL10A1 PCDHGC5 PCDHGA3 PCDHAC2 NECTIN4 PCDHA9 ABI3BP JAM3 CTRB2 CDH18 PCDHGB4 PCDHB8 CD209 FAT2 PCDHGB3