```
EIF3M EIF3K
                                                                         EEF1G
                                                                   EIF3D
                                                                                RPL7A
                                                                          RPS26
GHRHR
           PTGDR •
PTGER1WNT3A KISS1/
                  OXT UTS2
                                                          RPS29CTOME_SELENOAMINO_ACID_METABOLISM
      ・・・・ REACTOINE_SELENGAMINO_ACID
1 DRD5 GRPR HRH3 CCK GPR143 RELITIB科R21E_EUKARYOTIC_TRANSLATION_ELONGATION
VIP_INSL3 TRANSLATION_ELONGATION
 VIPR1 DRD5
        NPW GPR68BXA2KNTSR MLNR ADRA1D
                                                                                            RPL5
                                                                       RPS23
              GPR68BXAZR TO THE ADRB2 COTRANSLATIONAL_PROTEINT PS5 GETING FIF 4 AF MERLANDE CXCL3 FOR ADRATA
WNT10A/OXER1-SAA1 GALR3AS2R4ACKR4
           NPB NMUR2PRM1 PYY DRD2/GPER1
                                                                                    RPL23A
                                                                             EIF3I RPL27
ADRA2B DRD3 RXFP4 CXCL6 CXCR3
HTR1FP2RY4 OPRK1 PDYNGALR2 NPY4R
TAS2R9 ADDRESS OF TAS2R9
                                                                                        EEF2 RPN2
                  RXFP3 PPY GNRH1 CRHR1
                                                                                  SSR<sub>2</sub>
TAS2R9 ADRB1
  TAS2R31 GNRH2 CCR10 CXCR5
                                                                                           ITGB4
                   LPAR3
TAS2R30 TAS2R41
                                                                                                       size
                            HTR1B
                                                                                COL1A1
  REACTOME_CLASS_C_3_METABOTROPIC_GLUTAMATE_PHEROMONE_RECEPTORS
                                                                                            THBS1
 RGSL1
                                                          REACTOME_SYNDECAN_INTERACTIONS
                                                                                               COL5A1
                                                                             PPM1A
                                                                                    TGFB1 COL3A1
                                                                                    SMAD3 ITGAV
                                                                    ARHGEF18
                                                                  TRIM33
                                                                            USP9X
                                                               CBL
                                                                                         SKIL
                                                                             MTMR4
                        REACTOME SIGNALING BY TGF BETA RECEPTOR COMPLEX
                                                                  UBB
                                                                                      NEDD8
                                                    HIPK2
                                                                                 JUNB
                                                                   NCOR2
                                                          TBL1XR1
   MYO9B ARHGEF1 ARHGAP26
                                                                  NCOR1
                                         AGO1
                                                                                     SP1
                                 REACTOME REGULATION OF MECP2 EXPRESSION ASMAD2TFURIN
MYO9A
                                    ●IQGAP1
                                               HDAC2
           ERBIN
                    ARHGEF17
                                                              HDAC3
 ARHGEF12
                                  RHOB
                                            TNRC6A
                                                                  AGO2
ILIPREACTOME_RHOC_GTPASARHGAP5
                                                            AGO3
                ARHGAP35 AKAP13
                                  ARHGDIA TNRC6B
      CAVIN1
                                                 CAMK2A MECP2
       ARHGEF10
                                  FMNL3
                 DLC1 ARHGEF10L
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