ADAMTS1, MMP14, PHYKPL ITGA4 A2M JAM2_ITGAE SERRINE1 VCAM1-ITGA8 MATN3/ITGA9 EMILIN3 FBN1 / LTBP1 II FBLN FMOOD47 M CD44 NID CDH1_ITGA5_ITGB8_tropgrepagell surface interactions
PDGFRA (CREB1) I SOCS6 TGFB2 ADAM19 - ITGA5 ITGB8 TOTO GOLSAT LOXL4 PDGFRA CREB1/IRS1 SPP1 .ITPR1 EFEMP2 BGN TIMP2 COL9A3 FGFBP3 TRIB3 THES4, SH2B3 CAPMADAMTS18 P4HA2 DST ITGA2, ADAM12/ SPRED2FGF9 ANOTSKACB ADAMTS9TLL2 LOXL2 HSPG2 COL9A1 / HBS1 \PIK3R1PDGFRB /GABRG2 HAPINGAMICOLIZAExitatellular matrix organizaten2NTRK2—SPHK1 FBLNMMP24 P3H2 COL 15A1 COL 5A1 TGB1 DNAL4CY FIABRES JAKEGER2 ADAMTS8 MADA POOR FOR TO TO THE PROPERTY OF TH 35 POOLIGEON TO COLSA1 COL 82 CAPNO CTSK LOXLO COLGACOL1424MA1 POGFA 46ABRGAKT3 130 SDC1 MMP9 ADAM1ABI2MRG38PRY1 / IDOA MMP2 CTSV ——SDC1 MMP9 ADAM14BI2NRG38PRY1 IF
CRTAP DDR2 SERPINH1 COL11A1 POGFB ITPR3 SHC DAV1— 177 CRTAP DDR2 SERPINH1 COL11A1 POGEB 11 TO DOCK SERPINH1 POGEB 11 TO DOCK SERPINH COL11A1 POGEB 11 TO DOCK SERPIN SK LTBP2 PRKOA MAPK11 WASFIN3CBRIT1 SHETPROLEDGEC CASK LTBP2 /PRK#A MAPK11 TGFB1 SDC3
Non-integrin membrane-ECE/REGE/Action BOE HBEGF STAT6 SOS1 EXET 12 VRK NRGMAPK 12 FGF 19 TNS3 /NRXN1 SPARC/THEM4: ADAMTS5' SPINT2PRKOEM3 GAB2 EGFR PDE3B CYFIP2 GRB2 ERBBBPRK SHC3 ARHGEF7 GABRB2 PCSK5 DUSP4 FUT1 FLRT3 NRP1 FIAM1 EGEULADCYARI WWOX RPPRR1BFYN ROCK2 PRKCB PIK2CA **\K\T**G \DUSP3 ATP6V0E2 WEGFC GAB1 LYN \ \ KITLG 'DUSP3 ATP6V1G2 GALNT3 APH1B STAM2 AXLBD SPRED1