cn5C vs. cn5P ANKRD46Ap.Arg126= GALDH1A3: p.Leu195=RB55 M.BLAA:/p.Asp323 Tyr. This is the Complete DIVOY mDK200R-Synonymous mutations Functional mutations TCIRG1ENPP2: p.Asp616Asn8G Cancer genes NAVB: p.Gly140Val - KLABCC12 TEX10: p.Alao12Glu ADAMTSLS: p.pro582Ser-e407 FAILS
PIK3CG: p.Arg13LysDENND1A: p.Gly85515 244 PSMD13: pOR6T12 GPR37: p. Ser575Ph SF: p.Trp1045Ter MGRN1: p.Gly195 ppp 1839: npargage INME A: p.Lvs2451; p.Glv2377er A: TBCEL-TECTA: n.Asn1080Asn CRT: p.Glv103: p.Glv2377er A: TBCEL-TECTA: p.Asp1080Asn 132 For 1 HAP1: p.Arg397Cyana PRPF3/p.Glu617Ter Ala516Val SEC14L5: pus 7225. Arg 145Thr. p. FSIP2: p.Glu3691Ter RNF10: p.Ala516Val USH2A: p.Pro2431Ala/ POLR3E: p.Gln626Ter 24: p.Leu1080= FSHR: p.Leu244= ARPC5L: p.Ile15Met LAMA1: p.Gly/1941Ala SLC24A4: p.Ile185Leu MCL1: p.Phe254= DBF4B: p.His183= PCDH15: p.Arg341Leu or 0.4 GALNT17:/p.Glu196Lys GPR171: p.Ala91= CRB2: p.Leu661= OPN4: p.Leu328Met NYAP1: p.Ala444Ser DENND5A: p.Leu349Val TTN: p.Asp20002Glu CNR1: p.Phe381= TTN: p.Ile5832Val COL5A1: p.Gln1185Ter 0.2 0.0

0.2 0.4 0.6 0.8 CCF of cn5C (CIS+AIS)

1.0

0.0