57C1 vs. 57P1

L2HGDH: p.Ala443Thr. OR52E4: p.Thr297Asn
DCc.pATA31D1: p.Val642 = SPRY2: p.Cys218

V65DH: p.Pr6603Thrys RASGRP3: p.Val573 = C16orf90: p.Gl NFF2I 2. n Glv81Ala Synonymous mutations Functional mutations TTN: p.Ala19065Thr C16orf90: p.Glu ACACB: p.Thr2366Met PLA2G4A: p.Ala452 Cancer genes EXOC6B: p.Gln267Leu

SLC30A7: p.Phe159Leu

KCNJ10: p.Ala201: p.Pro921 IGH R: р. Гуг115= CKAP2L: p.Arg284= LSM1: p.Arg120Leu PKN3: p.Glu708Ter HEATR5B: p.Glu1640Gln SLC28A1: p.Cys647= OR10K2: p.Val12= PDE3B: p.Cys35= OR5/1V1: p. Leu203= Primary ### P.I hr461 | P.Pro4288er | C16ørf78: p.As | DCLRE1C: p.His412 | TASOR2: p.Tyr429Cys | Tasor4 | C16ørf78: p.As | C16ørf78: SORBSZ: p.Pro428Ser\_/C16orf78: p.Asp7Glu DPP6: p.Ala63Thr MTUS1: p.Lys1150= | COBL: p.Thr461= / PLEKHG2: p.Gln915ArgfsTer131 / PKP3: p.Ala590= / GREB1L: p.Leu984= KCNQ3: p.Thr838Met TYR: p.Gly191Val ST18: p.Met126Ile TYR: p.Tyr156= MYO18B: p.Pro182Ala 6 Jo 0.4 MYH7B: p.Leu1128= DDX49: p.Val105= 0.2 0.0 0.0 0.2 0.6 8.0 1.0 0.4

CCF of 57C1 (CIS+AIS)