NPHP1 (2q13_NPHP1) -	-0.6	0.1	0.1	0
PMP22 (HNPP) -	-0.1	0.5	-0.1	-0.1
OTUD7A (15q13.3_BP4-BP5) -	-2.1	2.6	2.6	1.8
PRODH (22q11.2_LCRA-D) -	-0.9	0.3	-0.9	-0.1
UQCRC2 (16p12.1) -	-0.3	-0.6	-1	-0.4
NDE1 (16p13.11) -	-0.1	-0.4	-1	0.1
PEX11B (1q21_TAR) -	0.2	-0.6	0.5	0.5
TUFM (16p11.2dist) -	1.3	1.3	0.2	1.4
COX10 (HNPP) -	0	-1	0.1	-0.6
SCARF2 (22q11.2_LCRA-D) -	0.9	0.7	0.1	-0.4
POLR3GL (1q21_TAR) -	1.2	1.2	0	-1.2
MYH11 (16p13.11) -	-0.1	-3.3	-0.9	-1
NCAPH (2q11.2) -	2.3	2.4	1.8	-0.2
PRRT2 (16p11.2prox) -	-0.1	0		-0.8
		-		
ANAPC1 (2q13) -	-2	-3.1	0.2	-1.4
GRAP (SMS) -	-1.2	-1.2	-1.2	-0.3
CD19 (16p11.2dist) -	1.2	1.3	0.1	-1.2
HJV (1q21_TAR) -	-1.1	-1.1	0	-0.4
IGLL1 (22q11.2_LCRD-H) -	-1.1	-1.1	-8.6	-1.1
PIGW (17q12) -	-1.1	-1.1	-1.1	-4.2
	0.6	0.4	-0.4	1.1
SLC25A1 (22q11.2_LCRA-D) -				-1.1
GP1BB (22q11.2_LCRA-D) -	-0.7	-0.7	-0.7	-0.1
RBP3 (10q11.21q11.23) -	1.2	-3.1	1	-1.2
CNNM4 (2q11.2) -	2.1	2.1	0.9	-1.1
FDFT1 (8p23.1) -	-0.3	-1	-1	0
SNAP29 (22q11.2_LCRA-D) -	0.7	0	-0.7	0.8
CDC45 (22q11.2_LCRA-D) -	-1.9	-3.3	-0.3	-1.2
SGCG (13q12.12) -	-1.8	-1.2	0.4	-1
CHAT (10q11.21q11.23) -	-1.9	1.1	-2.9	-1.2
NRROS (3q29) -	-0.8	-0.8	-0.8	-1.1
NSMCE3 (15q11q13_BP3-BP4) -	-0.2	-0.2	-0.2	-0.8
			-	
TFRC (3q29) -	2.4	3	2.5	-1.2
PCYT1A (3q29) -	-1.2	1.4	2.2	-1.1
SACS (13q12.12) -	-2.7	-1.9	-2.7	-0.2
OTOA (16p12.1) -	-0.5	-1.5	-2.9	0.1
TXNRD2 (22q11.2_LCRA-D) -	-0.9	1.6	0.4	-0.2
MIPEP (13q12.12) -	-0.9	-2.3	-3.3	-0.1
PI4KA (22q11.2 LCRA-D) -	-1.9	_4	5	-1.2
LMAN2L (2q11.2) -	-1.8	1.6		-1.7
TBX6 (16p11.2prox) -	0.5	-3.7	-2.5	-0.1
ATP2A1 (16p11.2dist) -	-1.6	-2.7	1.2	-0.9
DYNLT2B (3g29) -	-0.49	3	0.9	-0.65
TANGO2 (22q11.2_LCRA-D) -	-2.3	-0.3	2.3	0.4
B9D1 (SMS) -	-0.9	0.4	0.4	-0.6
ZNHIT3 (17q12) -	1.6	0.4	0.4	1.6
ERCC6 (10q11.21q11.23) -	-1	-3.1	-4.1	-0.3
FAN1 (15q13.3_BP4-BP5) -	-0.9	-2.4	-4	-0.2
LZTR1 (22q11.2_LCRA-D) -	-1.6	-1.9	-2.1	-0.2
MERTK (2q13) -	-0.9	-2.6	1.2	-0.8
TRPM1 (15q13.3_BP4-BP5) -	-0.2	-1.2	-2.3	-1
ABCC6 (16p13.11) -	-1.2	-0.3	-2.8	-0.3
ATPAF2 (SMS) -	0.2	3.1		3.5
RNF168 (3g29) -	-1.8	2.1	2.4	-1.1
NCF1 (7q11.23) -	0	-2.4	3.9	-0.73
NSMCE3 (15q11q13_BP3-BP5) -	1.8	1.8	1.8	-0.85
MDH2 (7q11.23_dist) -	3	1.5	5.5	-1.3
OTUD7A (15q11q13_BP3-BP5) -	-1.8	1.9	1.9	1.7
		3.9		
MAT1A (10q23) -	-1.6		-2.5	-0.59
BBS4 (15q24_BPA-BPC) -	-3	-2.2	0.3	4.9
STRA6 (15q24_BPA-BPC) -	-2.1	-2.6	3.4	-1.2
RP1L1 (8p23.1) -	-1.6	-3.1		-0.91
ALDH3A2 (SMS) -	-3	-1.8	1.4	-0.93
CDHR1 (10q23) -	-1.8	-2.2	4	-1.1
TOP3A (SMS) -	-1.8	-3.4	4.4	-1.2
TNFRSF13B (SMS) -	-2.1	-2	2.4	0.1
MPI (15q24_BPA-BPC) -	-2.6	4.3	3.5	-0.3
POR (7q11.23_dist) -	4.6	3.1	1.5	0.9
FAN1 (15q11q13_BP3-BP5) -	-0.9	-2.4	-4	-0.1
MYO15A (SMS) -	-1.3	-2.1	-3.2	0.2
TRPM1 (15q11q13_BP3-BP5) -	-0.3	-1.3	-2.4	-0.9
HERC2 (PWS/AS_BP1-BP3) -	-1.7	-1.3		0.4
OCA2 (PWS/AS_BP1-BP3) -	-2	-1.1	-1.5	0.8
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	A F.F.			
	AFR	AMR	EAS	EUR

orange: CNV+SNV cyan: SNV+SNV

