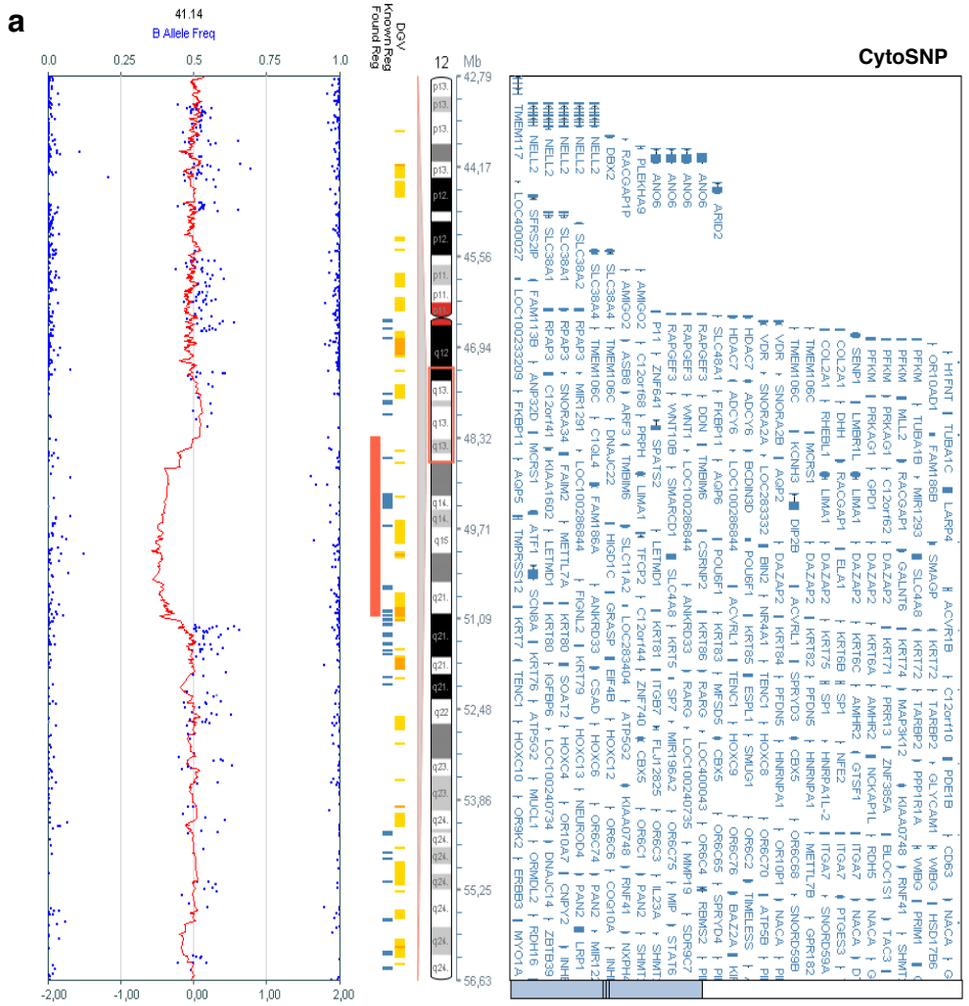
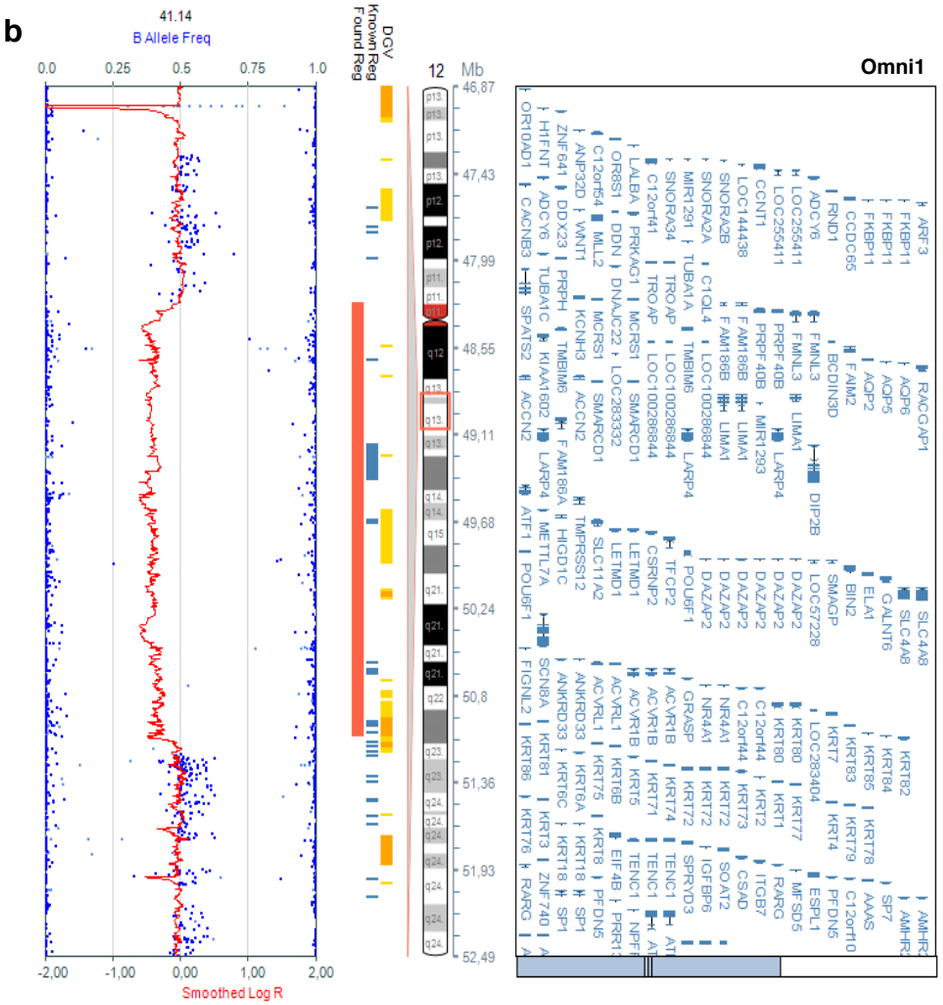
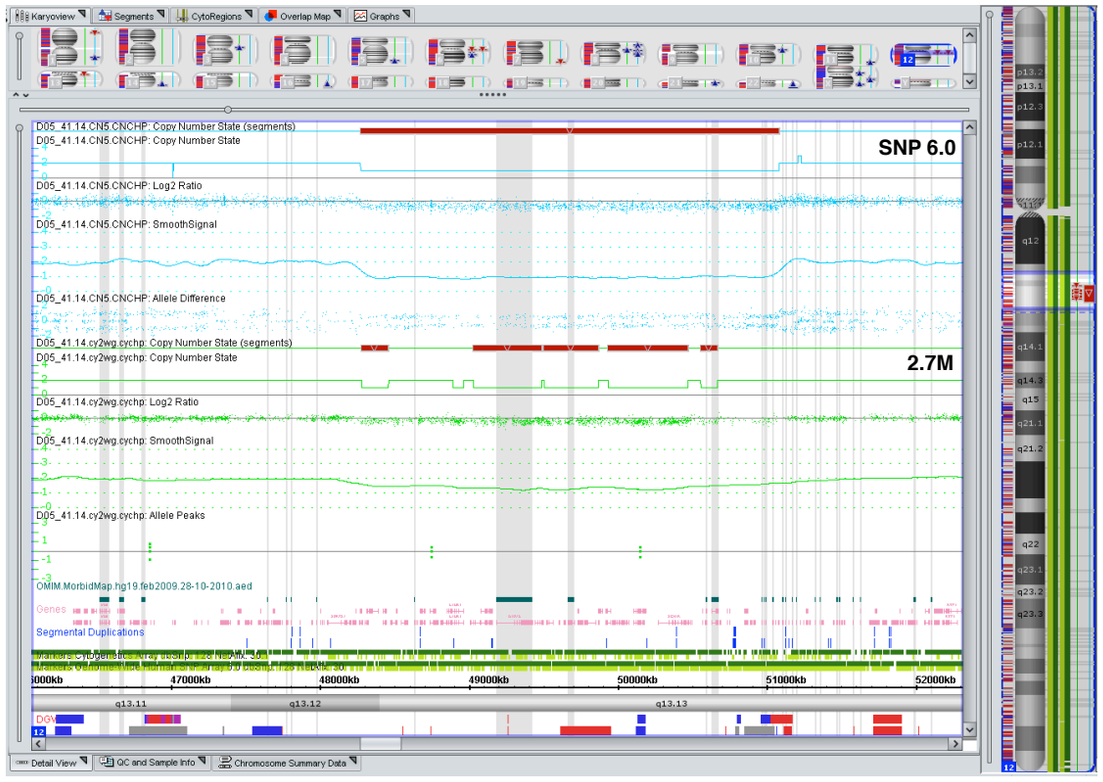
a


CytoSNP
41.14
B Allele Freq
Smoothed Log R
DGV
Known Reg
Found Reg
12
Mb

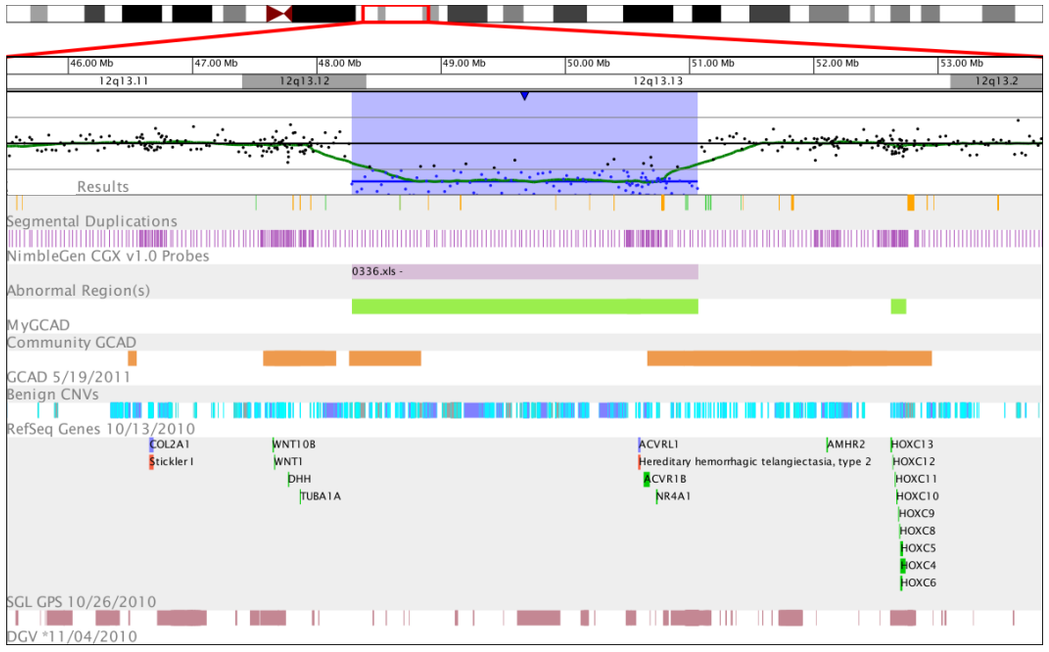
b


Omni1
41.14
B Allele Freq
Smoothed Log R
DGV
Known Reg
Found Reg
12
Mb

c


SNP 6.0
2.7M
D05_41.14.CN5.CNCHP: Copy Number State (segments)
D05_41.14.CN5.CNCHP: Copy Number State
D05_41.14.CN5.CNCHP: Log2 Ratio
D05_41.14.CN5.CNCHP: SmoothSignal
D05_41.14.CN5.CNCHP: Allele Difference
D05_41.14.cy2wg.cychp: Copy Number State (segments)
D05_41.14.cy2wg.cychp: Copy Number State
D05_41.14.cy2wg.cychp: Log2 Ratio
D05_41.14.cy2wg.cychp: SmoothSignal
D05_41.14.cy2wg.cychp: Allele Peaks
OMIM.MorbidMap.hg19.feb2009.28-10-2010.aed
Genes
Segmental Duplications
DGV

d


CGX-12
Results
Segmental Duplications
NimbleGen CGX v1.0 Probes
0336.xls -
Abnormal Region(s)
MyGCAD
Community GCAD
GCAD 5/19/2011
Benign CNVs
RefSeq Genes 10/13/2010
SGL GPS 10/26/2010
DGV *11/04/2010