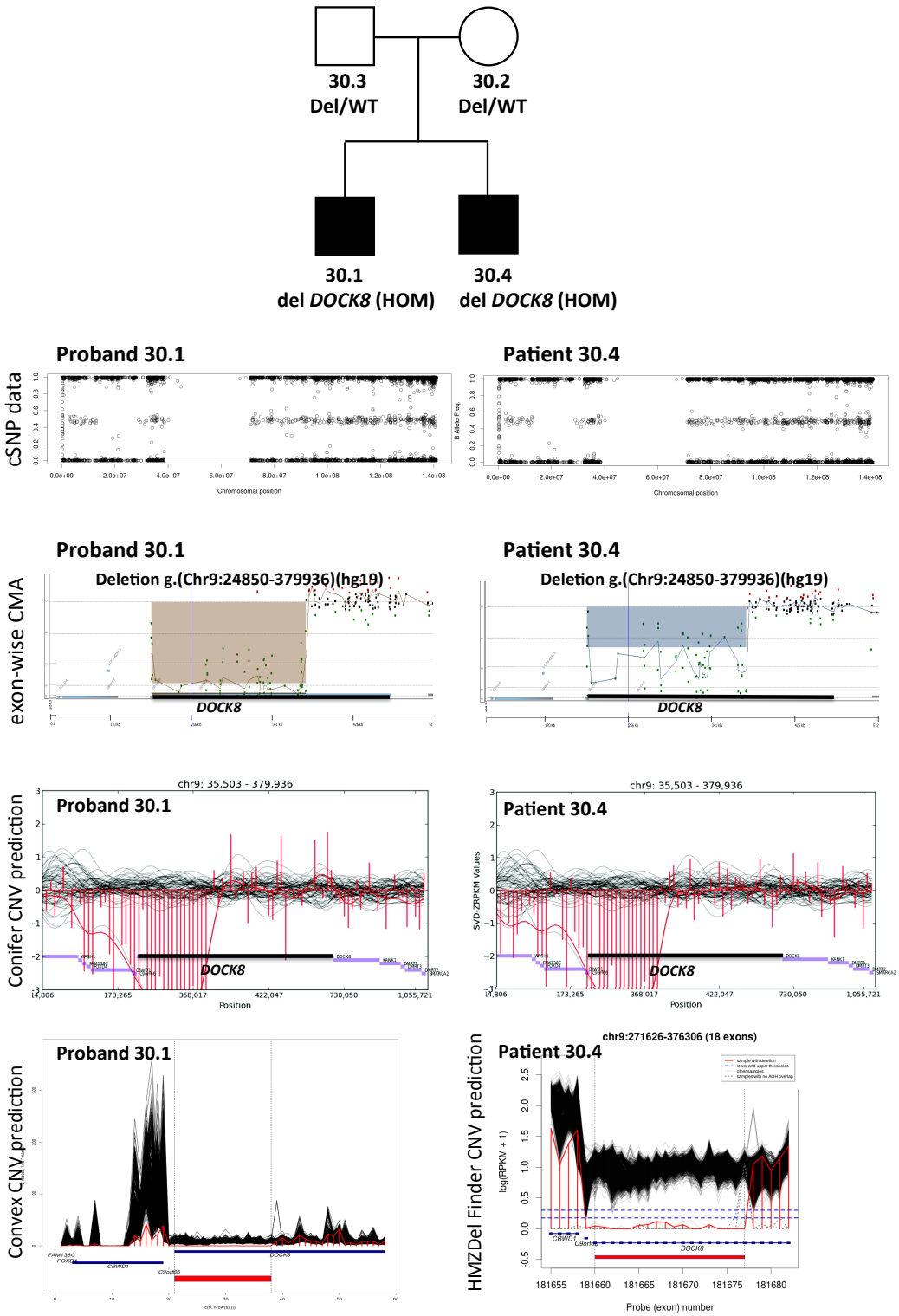
30.3
Del/WT
30.2
Del/WT
30.1
del DOCK8 (HOM)
30.4
del DOCK8 (HOM)
Proband 30.1
Patient 30.4
cSNP data
Chromosomal position
B Allele Freq.
exon-wise CMA
Deletion g.(Chr9:24850-379936)(hg19)
DOCK8
Conifer CNV prediction
chr9: 35,503 - 379,936
Position
SVD-ZRPKM Values
Convex CNV prediction
HMZDel Finder CNV prediction
chr9:271626-376306 (18 exons)
log(RPKM + 1)
Probe (exon) number

Figure E4.