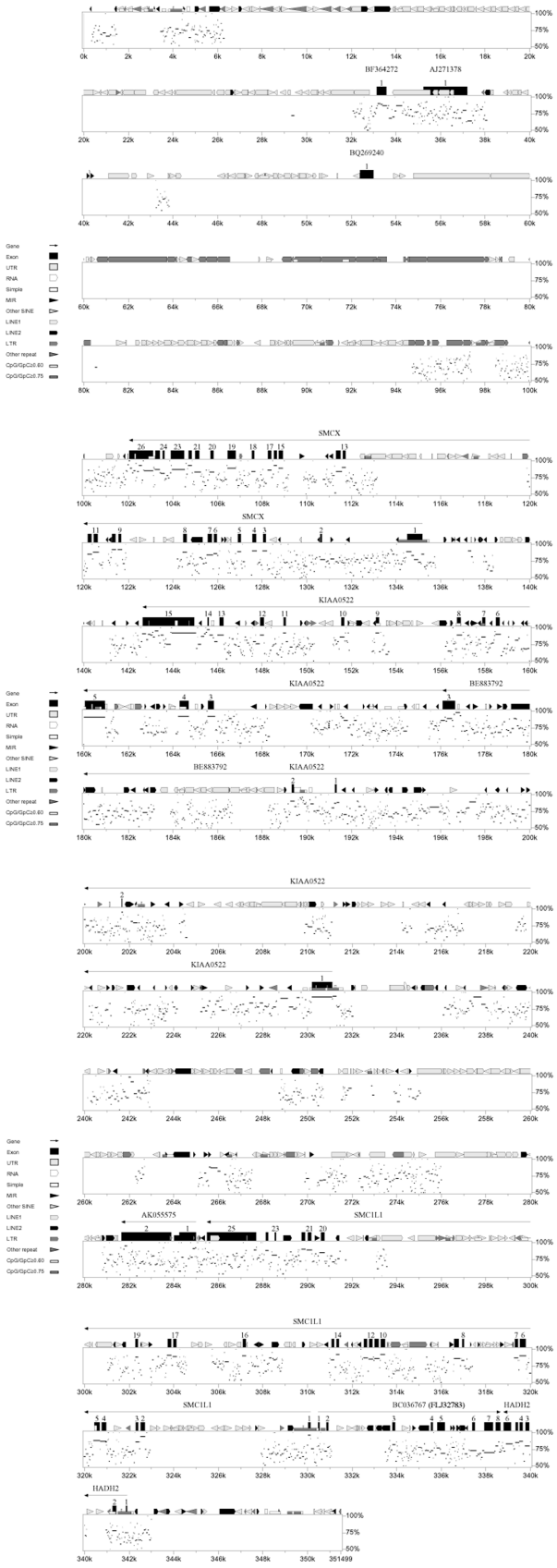
Supplemental Figure S1.



Percent identity plot between the domain of escape from Xp11.2 and the region of conserved synteny in mouse. The human mouse sequences were first analyzed by Repeat Masker (http://ftp.genome.washington.edu/). Percent sequence identity between 50-100% is given. Features of the human sequence including genes/ESTs, exons, CpG content ≥ 0.60 and 0.75, and repetitive elements are shown above the plot.