Supplementary Figure 27: L1HS Insertions and Deletions

Supplementary Figure 27. L1 HS Insertions and Deletions. Sites of structural variation that contained insertions (blue) or deletions (red) of L1 HS elements were identified by experimental characterization of the fosmids or examination of the human genome reference sequence (build35), respectively. The average discordancy for the fosmids mapped to each region is plotted. If we assume L1 HS polymorphisms are 6-8 kb in length, the analysis provides an estimate of the lower-bound sensitivity of our approach.