Figure S1.—CGH plots of the log2 of the ratio of the hybridization signal of genomic DNA to mouse Chr 2 BACs from heterozygous Pax6 deletion mutants relative to the hybridization signal from C3H wildtype. For all four Pax6 deletion mutants analysed a distinct region surrounding the Pax6 Chr 2 region displayed a series of overlapping BACs with log2 ratios lower than the copy number call threshold of -0.3. The Pax6^{12Neu} deletion was estimated to be the longest (~5.8 Mb) and the Pax6^{11Neu} deletion the shortest (~450 kb) of the four deletions analysed.