Supplementary Figure 1. Southern analysis of the RR-HO survivors with a chromosome VI specific probe, RET2

The samples from Figure 2D were subjected to PFGE that was performed using CHEF Mapper XA with a size range of 250-2500 kb using the Auto Algorithm mode. The clone numbers are indicated above. The lanes marked in blue, red, and green indicate type-1, -2, and -3 samples, respectively. The lanes marked in gray and black indicate samples with new fusion(s) of chromosomes VI and III accompanied by Leu+ recombination. See text for more details. L = LS20, Pre, pre-induction conditions; and IND, induced conditions