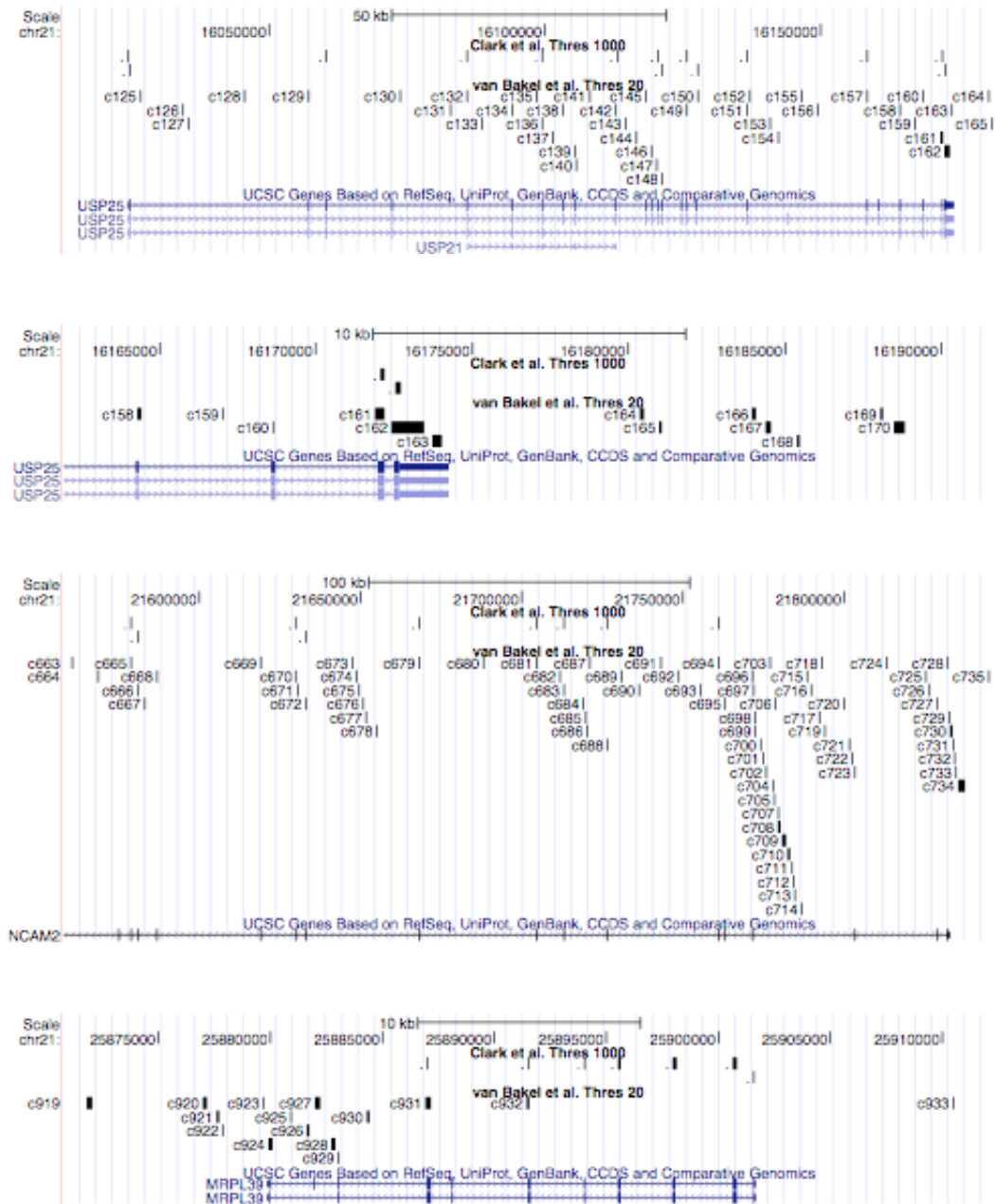## Figure S4

Figure S4: Genome browser screenshots showing annotation and transfrags from the van Bakel *et al*. [1] and the Clark *et al*. (ENCODE) tiling arrays using a threshold that gives similar recall values for both. The van Bakel *et al*. arrays show many more scattered transfrags.

1. van Bakel H, Nislow C, Blencowe BJ, Hughes TR (2010) Most "dark matter" transcripts are associated with known genes. PLoS Biol 8: e1000371.