

SUPPLEMENTAL MATERIAL

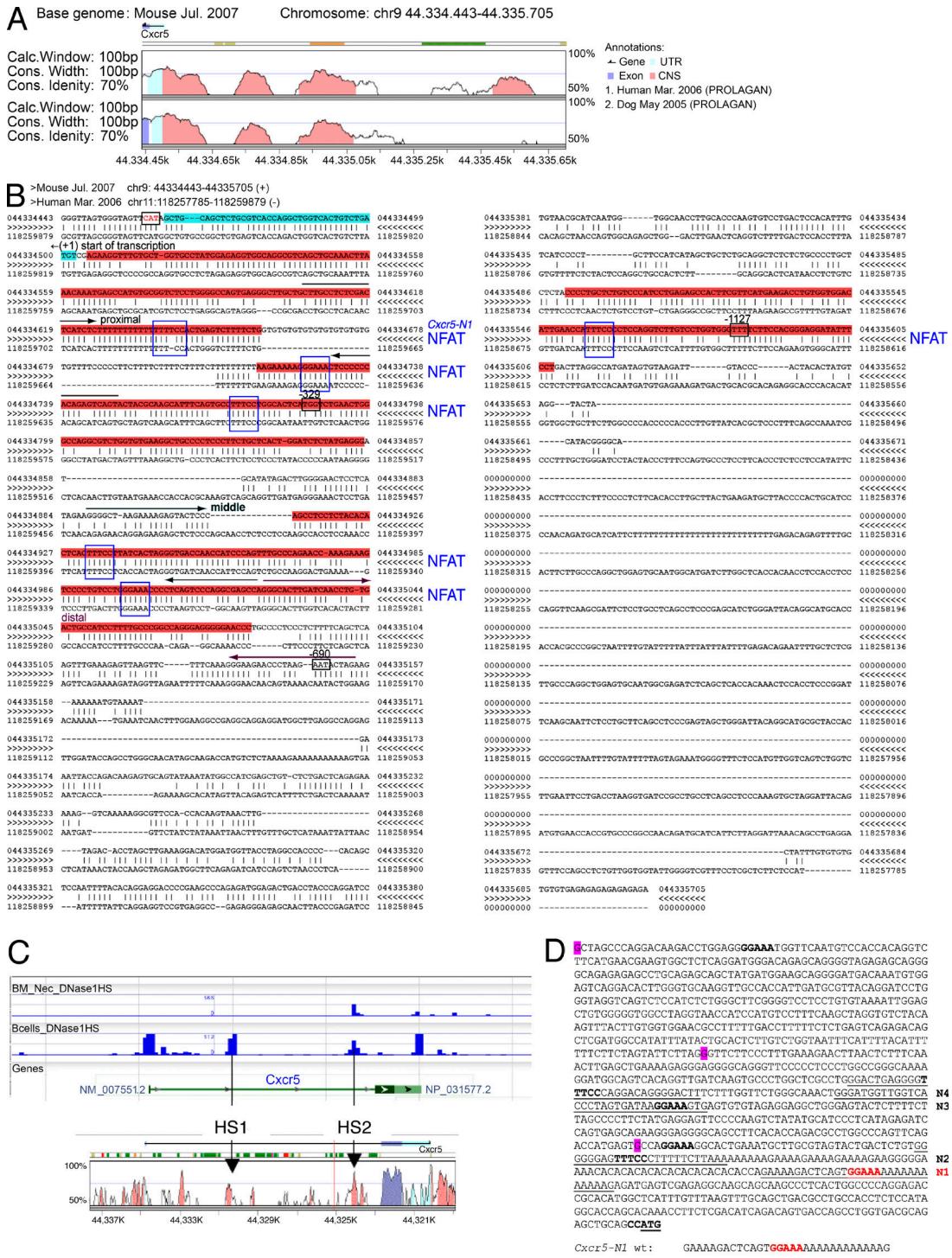
Vaeth et al., <http://www.jem.org/cgi/content/full/jem.20130604/DC1>

Figure S1. Scheme of Cxcr5 promoter region. (A) Alignment of human and dog Cxcr5 promoter reveals conserved noncoding sequences (CNS). (B) DNA sequence alignment of murine and human Cxcr5 promoter shown in A. Potential binding sites of NFAT, constructs used in reporter gene assays, and primers for ChIP experiments are illustrated. (C) Identification of potential regulatory elements (HS2 and HS1) within the Cxcr5 gene using public DNase I mapping data. (D) Sequence of the Cxcr5 promoter; starting nucleotides for luciferase constructs in purple, probes used for EMSA underlined and names given at the side. The GG→AA mutation within Cxcr5-N1 (red) is shown below.