



**Supplementary Figure 1. Organization of the CCR7 gene and location of predicted sterol response elements (SRE)**

To identify potential binding sites for transcription factors that may be important in the regulation of CCR7 gene expression, 6 Kb of the human and mouse CCR7 upstream regulatory regions were analyzed by MatInspector (Bioinformatics. 2005; 21:2933-42). This revealed a conserved sterol responsive element (SRE), in both human and mouse CCR7.