

SUPPELEMENTARY FIGURE

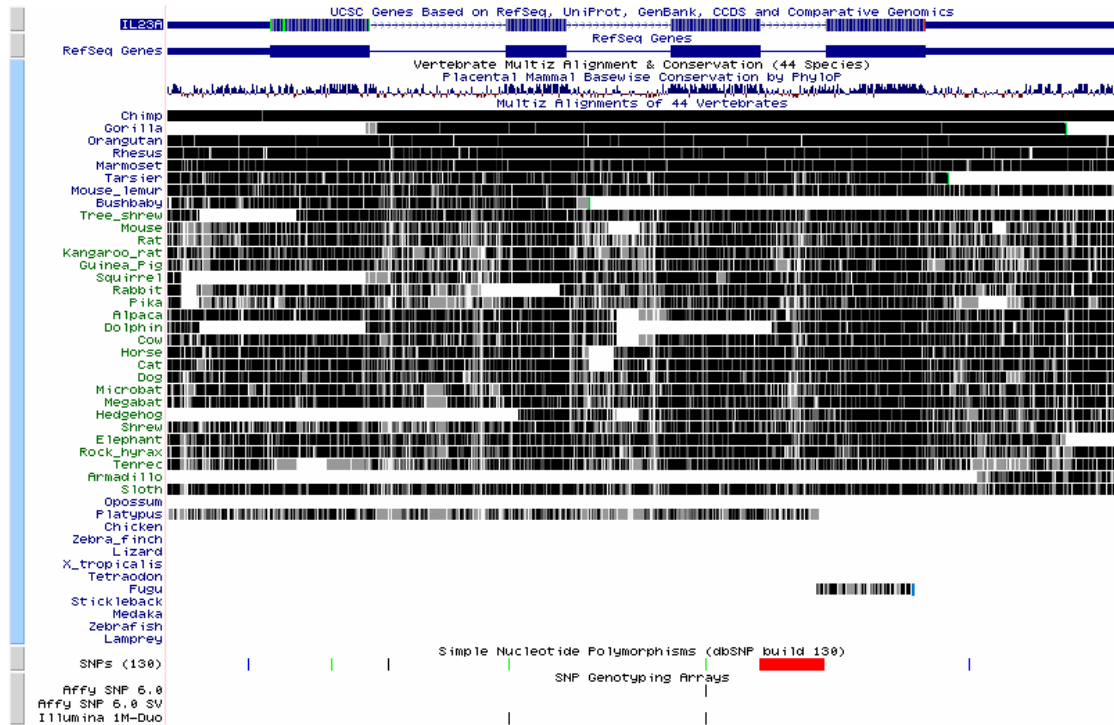


Figure S1. Intra-species sequence comparison of *IL23A*. Within the transcribed region of *IL23A*, the exonic regions are most highly conserved across the species. Eleven of the 44 vertebrates did not reveal a homologue for this gene. The UCSC Genome Browser (<http://genome.ucsc.edu>) was used to determine the sequence similarity of *IL23A* coding regions across 44 vertebrates, including 31 placental mammals, eight of which were primates.