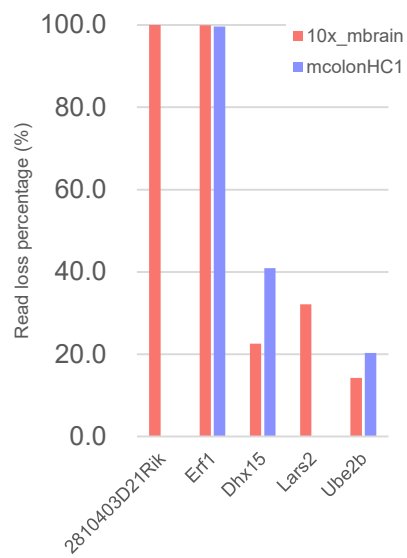




Additional file 2. Read loss percentage of the most misaligned genes in mouse