SFig 1

CTRL KD		
ADAR KD		
CTRL KD 1 Illumina	G	
CTRL KD 2 Illumina		
CTRL KD 3 Illumina	G G G	
	T T G A C C C C	
	UBE2I	

Supplemental Figure 1 | Example of novel A-to-I editing found with nanopore data. IGV shots of nanopore and Illumina data aligned to hg38. There were no reads aligning to UBE2I in the second Illumina CTRL KD replicate.

Supplementary Table 2

Gene Set Name	p-value	FDR q-value
DNA metabolic process	2.75E-13	4.27E-09
DNA repair	2.21E-11	1.72E-07
Cellular response to DNA damage stimulus	1.80E-10	9.34E-07
Cellular response to stress	3.39E-10	1.32E-06
Chromosome	1.45E-09	4.49E-06
Cell cycle	3.27E-09	8.47E-06
Transferase complex	7.38E-09	1.62E-05
Catalytic complex	8.34E-09	1.62E-05
Small molecule metabolic process	6.89E-08	1.19E-04

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Supplementary Table 2 | GO terms associated with haplotype-specific transcripts found in hybrid castaneus x 129 mouse data.