**Figure S4.** AD hybrids show high chromosomal aneuploidy. For each AD hybrid isolate the normalized depth of reads aligned to a combined AD reference is depicted across the chromosomes of the H99 and JEC21 genomes. In some isolates, the loss of a chromosome of one ancestry appears compensated by the gain of an extra copy in the other ancestry, noted by brackets, resulting in homozygosity of some regions of the hybrid genome.