Table S2

SNPs deduced from the sequencing of LdiAmB1000.1 and absent from the sequencing of its Ldi263WT parent.

SNP type	Total[1]	In CDS[1,2]	In CDS non-synonymous[1,2]
Homozygous	18	3	3
Heterozygous	470	123	85
Total	488	126	88

[1] Number of SNPs with ≤ 0.1% wrong call rates (phred scale quality value of ≥ 30).

[2] CDS, Coding sequence