

Table S10. Considered coordinates and location of ancestry informative SNPs for *AMY2B*, *GCKR*, and *PHYH*.

Gene	Coordinates ¹	Expanded Coordinates ²	ai.SNPs ³
<i>GCKR</i>	chr17:24411781- 24887707	chr17:23911781-25387707	chr17.24309172 chr17.24312880 chr17.25295979
<i>PHYH</i>	chr2:25986659-26125955	chr2:25486659-26625955	chr2.25670153
<i>AMY2B</i>	chr6:49348429-50097961	chr6:48848429-50597961	chr6.50074785 chr6.50081582 chr.50089070

¹These coordinates range from the transcription start site of the nearest non-overlapping 5' and 3' flanking genes according to the UCSC genome browser using the CanFam2 build.

²Consistent with Vonholdt et al. (2010), these coordinates expand the previous column by 500kb on each side to include the full potential range of an introgressed haplotype.

³ai.SNPs refer to ancestry informative SNPs. Defined by an absolute value Δ DAF between dogs and wolves greater than two standard deviations from the genome-wide mean.