Title: DNA Elements for Constitutive Androstane Receptor- and Pregnane X Receptor-mediated Regulation of Bovine CYP3A28 Gene

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S2 Fig. Resolution of the two encountered gaps in the bovine *CYP3A28* **promoter region.** The re-sequencing of *CYP3A28* promoter region was able to solve the first gap starting at -1492 bp revealing a sequencing artefact (A), as well as the second gap at base -2988 with a new genomic sequence (B). Two of the five single nucleotides polymorphisms (-2899T>G and -2981_-2979 insA) are here shown circled.