A dual targeted β-defensin and exome sequencing approach to identify, validate and functionally characterise genes associated with bull fertility

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Supplementary tables

Table S1: Summary statistics for β-defensin sequencing

Table S2: Significantly associated SNPs from TS

Table S3: Summary statistics for WES

Table S4: Significantly associated SNPs from WES

Table S5: Gene-ontology analysis of SNPs divergent between fertility groups