

Table S1. The diameter of the cervix and diameter of left and right uterine horns, mean daily milk yield for the first 60 days of lactation and blood concentrations of total cholesterol, triglycerides, non-esterified fatty acids, β -Hydroxybutyrate across all studied groups (Mean $\pm$ SD). P100- pregnant animals up to 100 days postpartum, P200- pregnant between 100 and 200 days postpartum, C- culled (including two cows sold to other farms at day 70 and 92 postpartum, two cows culled at 145 and 173 because of lameness and six not pregnant after 200 days), $\varnothing$ C- cervix diameter, $\varnothing$ LH- left horn diameter, $\varnothing$ RH- right horn diameter, MY60- mean daily milk yield for first 60 days of lactation, TC- total cholesterol, TG- triglyceride, NEFA- non-esterified fatty acids, BHBA- β -Hydroxybutyrate, P⁴-progesterone.

Parameter		P100	P200	C	<5% PMNs	>5% PMNs	<10% PMNs	>10% PMNs
n		11	14	10	20	15	26	9
$\varnothing$ C	mm	32.18 $\pm$ 3.12	34.50 $\pm$ 3.67	32.80 $\pm$ 3.74	33.90 $\pm$ 4.14	32.47 $\pm$ 2.56	33.85 $\pm$ 3.76	31.67 $\pm$ 2.50
$\varnothing$ LH	mm	28.45 $\pm$ 3.36	29.07 $\pm$ 4.71	27.90 $\pm$ 3.73	29.50 $\pm$ 4.17	27.27 $\pm$ 3.39	29.19 $\pm$ 4.08	26.67 $\pm$ 3.08
$\varnothing$ RH	mm	27.55 $\pm$ 3.62	29.29 $\pm$ 4.68	28.40 $\pm$ 4.19	29.85 $\pm$ 4.18	26.67 $\pm$ 3.52	29.11 $\pm$ 4.02	26.67 $\pm$ 4.30
MY60	kg	37.36 $\pm$ 6.36	38.79 $\pm$ 6.36	34.30 $\pm$ 9.65	36.75 $\pm$ 7.63	37.47 $\pm$ 7.70	36.77 $\pm$ 7.17	37.89 $\pm$ 8.98
TC	mmol/L	4.02 $\pm$ 0.93	4.43 $\pm$ 0.75	3.66 $\pm$ 1.10	4.13 $\pm$ 0.97	4.02 $\pm$ 0.96	4.11 $\pm$ 0.89	4.02 $\pm$ 1.16
TG	mmol/L	0.18 $\pm$ 0.031	0.19 $\pm$ 0.022	0.18 $\pm$ 0.015	0.19 $\pm$ 0.025	0.18 $\pm$ 0.019	0.19 $\pm$ 0.024	0.17 $\pm$ 0.019
NEFA	mmol/L	0.63 $\pm$ 0.39	0.55 $\pm$ 0.25	0.54 $\pm$ 0.14	0.59 $\pm$ 0.33	0.54 $\pm$ 0.16	0.59 $\pm$ 0.30	0.52 $\pm$ 0.17
BHB	mmol/L	1.03 $\pm$ 0.33	0.97 $\pm$ 0.29	0.99 $\pm$ 0.17	0.98 $\pm$ 0.26	1.01 $\pm$ 0.29	0.97 $\pm$ 0.24	1.07 $\pm$ 0.35
P ⁴	ng/mL	0.87 $\pm$ 1.14	0.71 $\pm$ 1.05	0.54 $\pm$ 0.10	0.46 $\pm$ 0.69	0.95 $\pm$ 1.21	0.57 $\pm$ 0.85	0.91 $\pm$ 1.23

Table S2. Selected genes, primer forward, and reverse sequences used for RT-qPCR.

Gene	NCBI GenBank Accession number	Primer forward sequence	Fragment size
		Primer reverse sequence	
<i>IL1B</i>	NM_174093.1	AAGGCTCTCCACCTCCTCTC TTTGGGGTCTACTTCCTCCA	186
<i>IL6</i>	NM_173923.2	TCTGGGTTCATCAGGCGAT TGTTTGTGGCTGGAGTGGTT	196
<i>CXCL8</i>	NM_173925.2	GTGCTCTCTTGGCAGCTTT GGTGAAAGGTGTGGAATGT	118
<i>IL17A</i>	NM_001008412.1	GCTCTTGTGAAGGCAGGAGT ATTGCGGTGGAGAGTCCAAG	177
<i>PTGDS</i>	NM_174791.4	ACACTTCACCACCTTTGCCA GCACTTGTCAGTCTTCGGCA	76
<i>PTGS1</i>	NM_001105323	CAACTGCACCATCCCTGAGA GGGGATAAGGTTGGAACGCA	175
<i>PTGS2</i>	NM_174445.2	AATCTTCCAGTCGCAGTAG TTGAGGCAGTGTTGATGAT	376
<i>PTGES</i>	NM_174443.2	GAAGAAGGCTTTTGCCAACCC AAGACCAGGAAGTGCATCCG	198
<i>PRXL2B</i>	NM_001040598.1	GCCAAAGGTGGTGATAAAGTGC GAGCATGCCTCTTCATCGCA	146
<i>INHBA</i>	NM_174363.2	GGACGGAGGGCAGAAATGAA TTCCTGGCTGTGCCTGATTC	80
<i>INHA</i>	NM_174094.4	CGGCTGCAGGTGCCA CTGGGATGGCTGGAACACAT	86
<i>FST</i>	NM_175801.3	CTATGCTAGCGAGTGTGCCA GTGTCTTCCGAAATGGAGTTGC	99
<i>POSTN</i>	NM_001040479.1	CCCCATAACTGTCTACAAGCCA TGACCTTGGTGACCTCTTCTTGC	200
<i>C20RF29</i>	XM_582695.5	CCTTCAAGAGCCCCCTGT GGGTCCTTTTCCAACCTCTCC	64
<i>SLC30A6</i>	NM_001075766.1	TGATGAGGAAACCTAGCCCTGCC TCGGGCTGCTCCAAAAAGCGT	142