

## **Comparison of five different RNA sources to examine the lactating bovine mammary gland transcriptome using RNA-Sequencing**

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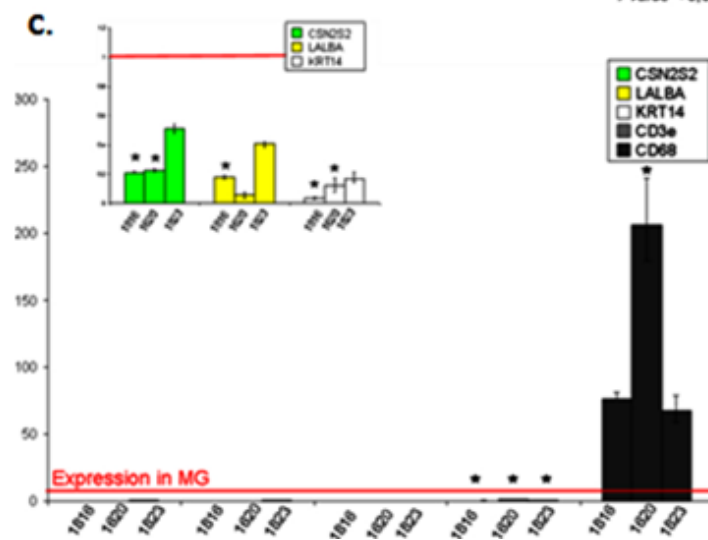
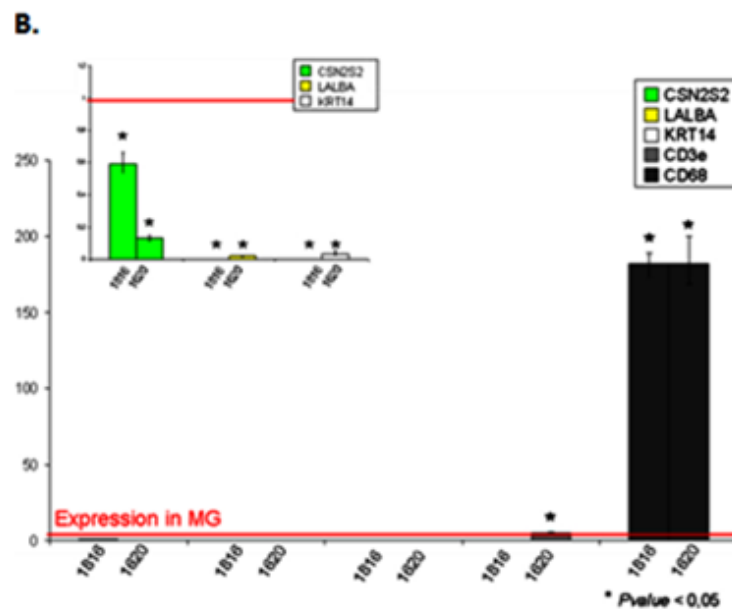
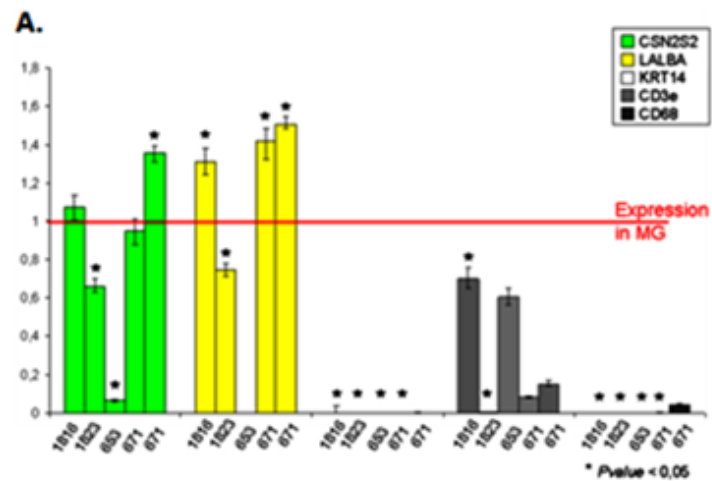
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**Supplementary File 1** Markers of myoepithelial cells, milk epithelial cells and immune cells used to validate the selectivity and specificity of laser capture microdissection by qPCR.

<b>Genes</b>	<b>Primers</b>	<b>Sequence 5'&gt;3'</b>	<b>Amplicon size (nt)</b>
<b>κ-casein (CSN3)</b> <i>GenBank: NM_174294.2</i>	forward reverse	AGG TGC AAT GAT GAA GAG TTT TTT C CCC AAA AAT GGC AGG GTT AA	66
<b>Cluster of differentiation 3 epsilon (CD3e)</b> <i>GenBank: NM_174011.3</i>	forward reverse	ACG CTG TAC CTG AAA GCA AGA AAT ACA CCA GCA GCA GCA AG	118
<b>Cluster of differentiation 68 (CD68)</b> <i>GenBank: NM_001045902.1</i>	forward reverse	GAT CTG CTC TCC CTG AAG CTAC A CAT TGG GAC AAG AGA AAC TTG GT	79
<b>Cluster of differentiation 18 (CD18)</b> <i>GenBank: M81233.1</i>	forward reverse	AGC GAC CTC AGG GAG TAC CA TTA TCG TTG TTC CAC TGG GAC TT	65
<b>Keratin 14 (KRT14)</b> <i>GenBank: NM_001166575.1</i>	forward reverse	CCC AGC TCA GCA TGA AAG C AGC GGC CTT TGG TCT CTT C	57

**Supplementary File 2** Quantification of specific markers for mammary myoepithelial cells (Krt14), lymphocytes (CD3e), neutrophils (CD18), macrophages (CD68 and CD18) and MEC (CSN3) by qPCR.



**Supplementary File 3** Correlations between highly expressed genes ( $\geq 500$  RPKM) of MGT, SC, LCMEC, MFG and mMEC transcriptome in Jersey and Normande cows.

		MGT		SC	LCMEC		MFG		mMEC	
		Jersey	Normande	Jersey	Jersey	Normande	Jersey	Normande	Jersey	Normande
<b>MGT</b>	<b>Jersey</b>	1.00	-	-	-	-	-	-	-	-
	<b>Normande</b>	0.97	1.00	-	-	-	-	-	-	-
<b>SC</b>	<b>Jersey</b>	0.98	0.91	1.00	-	-	-	-	-	-
<b>LCMEC</b>	<b>Jersey</b>	0.99	0.99	0.95	1.00	-	-	-	-	-
	<b>Normande</b>	0.99	0.99	0.95	1.00	1.00	-	-	-	-
<b>MFG</b>	<b>Jersey</b>	0.94	0.85	0.97	0.90	0.89	1.00	--	-	-
	<b>Normande</b>	0.96	0.93	0.96	0.95	0.94	0.96	1.00	-	-
<b>mMEC</b>	<b>Jersey</b>	0.08	0.01	0.16	0.04	0.01	0.36	0.24	1.00	-
	<b>Normande</b>	0.05	0.02	0.09	0.04	0.02	0.19	0.16	0.58	1.00

MGT: mammary gland tissue; SC: milk somatic cells; LCMEC: laser microdissected mammary epithelial cells; MFG: milk fat globules; mMEC: antibody- captured milk mammary epithelial cells.

**Supplementary File 4** Correlations between normally/medium expressed genes (ranged from 500 to 10 RPKM) of MGT, SC, LCMEC, MFG and mMEC transcriptome in Jersey and Normande cows.

		MGT		SC	LCMEC		MFG		mMEC	
		Jersey	Normande	Jersey	Jersey	Normande	Jersey	Normande	Jersey	Normande
<b>MGT</b>	<b>Jersey</b>	1.00	-	-	-	-	-	-	-	-
	<b>Normande</b>	0.91	1.00	-	-	-	-	-	-	-
<b>SC</b>	<b>Jersey</b>	0.95	0.97	1.00	-	-	-	-	-	-
<b>LCMEC</b>	<b>Jersey</b>	0.91	0.91	0.95	1.00	-	-	-	-	-
	<b>Normande</b>	0.84	0.93	0.99	0.95	1.00	-	-	-	-
<b>MFG</b>	<b>Jersey</b>	0.87	0.87	0.81	0.95	0.96	1.00	--	-	-
	<b>Normande</b>	0.96	0.86	0.87	0.90	0.96	0.95	1.00	-	-
<b>mMEC</b>	<b>Jersey</b>	0.40	0.14	0.75	0.36	0.10	0.42	0.12	1.00	-
	<b>Normande</b>	0.35	0.32	0.39	0.34	0.22	0.35	0.34	0.27	1.00

MGT: mammary gland tissue; SC: milk somatic cells; LCMEC: laser microdissected mammary epithelial cells; MFG: milk fat globules; mMEC: antibody- captured milk mammary epithelial cells.

**Supplementary File 5** Correlations between lowly expressed genes (<10 RPKM) of MGT, SC, LCMEC, MFG and mMEC transcriptome in Jersey and Normande cows.

		MGT		SC	LCMEC		MFG		mMEC	
		Jersey	Normande	Jersey	Jersey	Normande	Jersey	Normande	Jersey	Normande
<b>MGT</b>	<b>Jersey</b>	1.00	-	-	-	-	-	-	-	-
	<b>Normande</b>	0.83	1.00	-	-	-	-	-	-	-
<b>SC</b>	<b>Jersey</b>	0.77	0.78	1.00	-	-	-	-	-	-
<b>LCMEC</b>	<b>Jersey</b>	0.75	0.77	0.73	1.00	-	-	-	-	-
	<b>Normande</b>	0.84	0.78	0.87	0.75	1.00	-	-	-	-
<b>MFG</b>	<b>Jersey</b>	0.81	0.82	0.82	0.73	0.73	1.00	--	-	-
	<b>Normande</b>	0.87	0.81	0.83	0.63	0.67	0.76	1.00	-	-
<b>mMEC</b>	<b>Jersey</b>	0.13	0.06	0.81	0.06	0.06	0.46	0.34	1.00	-
	<b>Normande</b>	0.12	0.13	0.21	0.08	0.10	0.12	0.61	0.18	1.00

MGT: mammary gland tissue; SC: milk somatic cells; LCMEC: laser microdissected mammary epithelial cells; MFG: milk fat globules; mMEC: antibody- captured milk mammary epithelial cells.

**Supplementary File 6** Correlation values using the ranking/position of the RPKM gene expression values ranged from the most expressed gene to the lowest expressed gene in mammary gland tissue, SC, LCMEC, MFG and mMEC transcriptome in Jersey and Normande cows.

		MGT		SC	LCMEC		MFG		mMEC	
		Jersey	Normande	Jersey	Jersey	Normande	Jersey	Normande	Jersey	Normande
<b>MGT</b>	<b>Jersey</b>	1.00	-	-	-	-	-	-	-	-
	<b>Normande</b>	0.96	1.00	-	-	-	-	-	-	-
<b>SC</b>	<b>Jersey</b>	0.94	0.94	1.00	-	-	-	-	-	-
<b>LCMEC</b>	<b>Jersey</b>	0.93	0.93	0.92	1.00	-	-	-	-	-
	<b>Normande</b>	0.92	0.94	0.90	0.90	1.00	-	-	-	-
<b>MFG</b>	<b>Jersey</b>	0.94	0.91	0.95	0.93	0.89	1.00	-	-	-
	<b>Normande</b>	0.91	0.91	0.92	0.89	0.94	0.90	1.00	-	-
<b>mMEC</b>	<b>Jersey</b>	0.91	0.82	0.94	0.90	0.80	0.93	0.80	1.00	-
	<b>Normande</b>	0.82	0.76	0.73	0.80	0.79	0.80	0.84	0.69	1.00

MGT: mammary gland tissue; SC: milk somatic cells; LCMEC: laser microdissected mammary epithelial cells; MFG: milk fat globules; mMEC: antibody- captured milk mammary epithelial cells.