

Supplementary information

S1. Number of outliers, minimum, maximum, mean, and standard deviation of the proportions of type I, type IIA and type IIX myofibers in Holstein cows, as estimated using four deconvolution methods (CIBERSORTx, Cellanneal, DeconvR-NNLS, and DeconvR-RLM), with Brahman and Wagyu as reference breeds. The table also includes values obtained from direct biopsies of the *Longissimus dorsi* muscle taken in the same cows.

Myofiber	Methodology	Reference Breed	Outliers	Min (%)	Max (%)	Mean (%)	SD (%)
Type I	Biopsy	Holstein	0	13.98	38.98	23.09	5.61
	Cellanneal	Brahman	0	38.64	72.14	55.14	9.38
		Wagyu	0	31.49	39.05	36.01	1.99
	CIBERSORTx	Brahman	1	12.60	24.85	18.67	3.48
		Wagyu	2	10.99	21.26	16.09	2.91
	DeconvR – NNLS	Brahman	0	29.89	47.12	37.41	3.61
		Wagyu	0	24.92	40.59	29.21	3.24
	DeconvR – RLM	Brahman	1	11.35	26.04	17.39	3.59
		Wagyu	1	10.38	22.96	15.36	3.07
Type IIA	Biopsy	Holstein	0	26.88	69.40	40.73	9.20
	Cellanneal	Brahman	0	1.46	34.89	15.89	10.19
		Wagyu	0	34.44	44.38	38.83	2.11
	CIBERSORTx	Brahman	1	36.20	47.50	42.41	3.22
		Wagyu	2	56.76	68.15	62.40	2.94
	DeconvR – NNLS	Brahman	0	0.00	48.12	19.10	13.83
		Wagyu	0	38.59	72.26	61.43	8.39
	DeconvR – RLM	Brahman	1	33.72	50.02	43.64	4.28
		Wagyu	1	58.29	70.59	64.66	3.27
Type IIX	Biopsy	Holstein	0	9.33	59.14	36.17	9.98
	Cellanneal	Brahman	0	23.91	36.67	28.97	3.23
		Wagyu	0	21.39	31.33	25.15	2.66
	CIBERSORTx	Brahman	1	32.25	43.10	38.93	2.46
		Wagyu	2	17.59	26.13	21.51	1.72
	DeconvR – NNLS	Brahman	0	21.99	64.55	43.49	12.82
		Wagyu	0	0.00	33.42	9.36	9.24
	DeconvR – RLM	Brahman	1	32.71	45.41	38.97	3.42
		Wagyu	1	15.90	27.25	19.97	2.55

Min: Minimum; Max: Maximum; SD: Standard Deviation; NNLS: Non-Negative Least Square; RLM: Robust Linear Model; n: number of cows with results after removing outliers. Different letters indicate statistically significant differences between groups, as determined by the Dunn's test with Bonferroni correction ($p < 0.00139$).