

Association of Missense *MTTP* Gene Polymorphism with Carcass Characteristics and Meat Quality Traits in Pigs

KATARZYNA ROPKA-MOLIK¹, PRZEMYSŁAW PODSTAWSKI²,
KATARZYNA PIÓRKOWSKA¹, MIROSŁAW TYRA³

¹Department of Animal Genomics and Molecular Biology, National Research Institute of Animal Production, Balice, Poland

²University of Agriculture in Cracow, Cracow, Poland

³Departement of Animal Breeding and Genetics, National Research Institute of Animal Production, Balice, Poland

Supplementary Online Material (SOM)

Table S1. Family structure of all pig breeds analyzed

Number of animals	Large White	Landrace	Pulawska	Duroc	Pietrain	Whole population
n of boars/n of pigs per sire	51/4.80	64/4.22	21/3.43	10/3.80	18/3.00	164/4.20
n of sows/n of pigs per dam	158/1.55	161/1.67	47/1.53	29/1.31	37/1.45	435/1.58

Table S2. Association between ENSSSCT00000010052.2:c.2518C>T polymorphism and slaughter traits evaluated in different pig breeds

Slaughter traits	Genotype	Large White (n = 245)	Landrace (n = 273)	Pietrain (n = 54)	Pulawska (n = 72)	Duroc (n = 38)
Carcass yield (%)	CC	77.57 ± 0.34 ^a	76.34 ± 0.20	78.82 ± 0.34	77.30 ± 0.22 ^{a•}	75.61 ± 0.35
	CT	76.90 ± 0.26 ^b	76.32 ± 0.15	78.19 ± 0.22	77.14 ± 0.19 ^{ab}	76.91 ± 0.68
	TT	77.00 ± 0.28 ^{ab}	76.55 ± 0.14	78.68 ± 0.49	76.73 ± 0.25 ^b	—
Weight of loin (kg)	CC	7.70 ± 0.16	7.58 ± 0.10	8.01 ± 0.13 ^{a•}	7.20 ± 0.14	7.36 ± 0.00
	CT	7.81 ± 0.12	7.56 ± 0.07	7.81 ± 0.08 ^{ab}	7.38 ± 0.13	7.96 ± 0.00
	TT	7.83 ± 0.13	7.71 ± 0.07	7.52 ± 0.18 ^b	7.13 ± 0.17	—
Weight of ham without bone and skin (kg)	CC	9.27 ± 0.15	9.19 ± 0.10	10.31 ± 0.12	8.95 ± 0.15	9.34 ± 0.09
	CT	9.17 ± 0.12	9.05 ± 0.07	10.59 ± 0.08	8.77 ± 0.14	9.47 ± 0.29
	TT	9.05 ± 0.13	9.06 ± 0.07	10.39 ± 0.17	8.82 ± 0.18	—
Average backfat thickness (cm)	CC	1.32 ± 0.09 ^b	1.34 ± 0.06	1.15 ± 0.08	1.48 ± 0.08	1.35 ± 0.24
	CT	1.44 ± 0.07 ^b	1.32 ± 0.04	1.11 ± 0.05	1.57 ± 0.07	1.56 ± 0.38
	TT	1.56 ± 0.07 ^a	1.34 ± 0.04	1.07 ± 0.12	1.50 ± 0.09	—
Loin eye area (cm ²)	CC	54.20 ± 1.39	53.77 ± 0.87	67.67 ± 1.69	50.87 ± 1.41	48.80 ± 0.04
	CT	52.26 ± 1.07	53.38 ± 0.66	66.48 ± 1.10	49.69 ± 1.27	49.35 ± 0.15
	TT	53.64 ± 1.17	52.91 ± 0.61	64.37 ± 2.44	50.43 ± 1.64	—
Lean meat percentage	CC	61.05 ± 0.74	61.23 ± 0.50	66.67 ± 0.60	59.45 ± 0.81	62.27 ± 0.43
	CT	60.35 ± 0.57	60.87 ± 0.38	67.63 ± 0.39	58.66 ± 0.73	60.95 ± 1.24
	TT	59.87 ± 0.62	60.86 ± 0.35	66.09 ± 0.87	58.98 ± 0.94	—
Weight of primary cuts (kg)	CC	24.15 ± 0.58	23.80 ± 0.20	26.90 ± 0.24	23.07 ± 0.32	23.80 ± 0.23
	CT	23.81 ± 0.44	23.67 ± 0.15	27.30 ± 0.16	22.78 ± 0.28	24.38 ± 0.52
	TT	24.01 ± 0.48	23.66 ± 0.14	26.66 ± 0.35	22.88 ± 0.37	—

values are presented as Least Squares Means ± standard errors

means with different superscripts differ significantly between genotypes (^{a,b}*P* ≤ 0.05, ^{a,b•}*P* ≤ 0.1 – trends)