# RELATIONSHIP OF INTRAMUSCULAR FAT, FATTY ACIDS AND CONJUGATED LINOLEIC ACIDS WITH TENDERNESS IN PORK

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## Introduction

Intramuscular fat (IMF) that is a primary visual stimulus to customers plays an important role in driving people to purchase meat. It is generally accepted that a higher level of IMF has a positive influence on tenderness, but not been illustrated consistently (Brewer, 2001; Fernandez, 1999a & 1999b; Van Laack, 2001). Conjugated linoleic acids (CLA) has anticancerogenic, antiatherogenic, antidiabetic effects and so on. Its content and division maybe are also relative with meat tenderness. With increasing of unsaturated/ *cis* fatty acids and decreasing of *trans* fatty acids (FA), the meat hardness will reduced correspondingly (Wood, 2003). The couple of C18:0/C18:2 or CLA may be a key factor for pork tenderness.

The aim of this study was to research the relationship between IMF/FA/CLA and pork tenderness. The mathematical model for IMF/FA/CLAlinking pork tenderness was to be built up.

## **Materials and Methods**

*M. longissimus dorsi* (LD) was obtained from nine kinds of crossbred swines (9 groups) at 10-12 rib (n = 66). LD samples were stored at  $3\pm1^{\circ}$ C for 3 d to test their shear force. 1) A piece of LD (3cm thickness) was heated at 80°C in water bath, the middle temperature of meat up to 70°C and kept 10min, then cooled at 2°C for 12 h. 10-12 meat columns from the same piece ( $\oint = 1.27$ cm) were determined shear force (SF)with tenderometer (C-LM3, PRC). IMF(Fortin, 2005), ether-16 FAs (containing a couple of CLA; Aldai, 2006) were analysed by Saxhlet and gas-chromatography, respectively. All of data were treated as Means ± SD with one-way analysis of variance (ANOVA) by SPSS 13.0. Pearson correlation coefficients and regression equations were set up.

#### **Results and Discussion**

*Rrelationship between IMF content and pork tenderness in 9 groups.* IMF content in LD was the range of 1.37% - 3.78% in 9 crossbreds. There was no the significant negative linear relativity between IMF and SF (r = -0.230, P = 0.074) as Fortin (2005) by Pearson correlation coefficient. However, the log<sup>-</sup> curve relativity ("Inverse" mathematical model) was estimated between both of them. Its regression equation was  $Y_{SF} = 0.104 \times 1/X_{IMF-c}$  (R<sup>2</sup> = 0.702, P<0.001\*\*\*, Figure 1) for pork.

*Relationship between FAs and pork tenderness.* 16 FAs were assayed by gas-chromatography in this study. In pork, the composition of FAs as saturated and unsaturated FAs was around half by half. There was significant negative relativity between C14:0, C18:2, C18:3 and C22:6 with SF in pork, respectively (r = -0.31, r = -0.48, r = -0.52, r = -0.47;  $p < 0.01^{**}$ ). C18:1 and C18:2 (*cis9/tans11, trans10, cis12*) were UFA in pork mostly. There was logarithmic correlation coefficient betweent FAs and SF, especially up to very significantly in C18:0/C18:2 (r = 0.619, p < 0.01^{\*\*}, Table 1). Both of their relation was clearly by "Logarithmic" mathematical model:  $Y_{SF} = 13.678 \times lnX_{C18:0/C18:2}$ ,  $R^2 = 0.775$ ,  $p < 0.001^{***}$ ; Figure 2).

## Conclusions

There were signicantly negative Log-relativity between IMF content and pork tenderness, while positive Log-relativity between the ratio of C18:0/C18:2 and pork tenderness by "Inverse" curve model. There was



F Figure 2 Log relation with the ratio of C18:0/C18:2 and SF

C18:0/C18:2

3.50

Table 1 Correlation coefficient between FAs and SF in pork									
	$\mathbf{X}_1$	$X_2$	<b>X</b> <sub>3</sub>	$X_4$	$X_5$	$X_6$	X7	$X_8$	X9
Y	0.095	-0.306*	0.048	-0.088	-0.192	0.057	0.326*	0.307	-0.484**
	X <sub>10</sub>	X11	X <sub>12</sub>	$\mathbf{X}_{13}$	3	X <sub>14</sub>	X15	X <sub>16</sub>	X <sub>6</sub> / X <sub>9</sub>
Y	-0.526**	-0.069	0.191	-0.18	37 0.	124	-0.474**	-0.047	0.619**

Figure 1 Log relation between IMF and SF

Notes: Y-SF, X<sub>1</sub>-C12:0, X<sub>2</sub>-C14:0, X<sub>3</sub>-C16:0, X<sub>4</sub>-C16:1(c9), X<sub>5</sub>-C17:0, X<sub>6</sub>-C18:0, X<sub>7</sub>-C18:1(c9), X<sub>8</sub>-C18:1(t9), X<sub>9</sub>-C18:2(c9,12), X<sub>10</sub>-C18:3(c6,9,12), X<sub>11</sub>-C18:3(c9,12,15), X<sub>12</sub>-C20:0, X<sub>13</sub>-C20:4(c5,8,11,14), X<sub>14</sub>-C22:4 (c5,8,11,14), X<sub>15</sub>-C22:6(c4,7,10,13,16,19), X<sub>16</sub>-C18:2 (*cis9*, *trans11*); \*p < 0.05 and \*\*p < 0.01 meant the difference significantly between crossbreds.

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