

# PLECTIN POSTMORTEM DEGRADATION AND ITS RELATION TO WATER DISTRIBUTION IN FRESH PORK

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## I. OBJECTIVES

Plectin is an intermediate filament (IF)-based versatile cytolinker protein of the plakin family. Plectin links the desmin IF and anchors them to specific cytoskeletal structures and cytoplasmic organelles in skeletal muscle, forming a complex IF network that has a great influence in sustaining and strengthening the integrity of cytoarchitecture. A previous study has shown that plectin was a potential substrate of calpain-1 during the conversion of muscle to meat. Therefore, the current study aimed to evaluate the contribution of plectin postmortem degradation to the changes in water distribution in fresh pork.

## II. MATERIALS AND METHODS

*Longissimus thoracis* muscles (between the 10th and 14th thoracic vertebra with the size of 15 cm × 8 cm × 4 cm) were collected from both sides of 44 carcasses within 1 h postmortem and aged at 4°C for 0 h, 6 h, 12 h, 1 d, 3 d, 7 d, and 13 d under vacuum package. *Longissimus thoracis* samples from one side were used to prepare the whole protein samples of 5 mg/mL for the Western blot analysis of plectin degradation. Samples from the other side were used for low field-nuclear magnetic resonance analysis (LF-NMR) based on the spin-spin transverse relaxation time ( $T_2$ ) analysis. Statistical significance of the differences among individual means was assessed by Duncan's multiple-range tests. Pearson correlation coefficients were obtained to evaluate the correlation between different protein degradation and LF-NMR parameters.

## III. RESULTS

Figure 1 shows that plectin was found to be significantly degraded during postmortem aging. The amount of intact plectin was rapidly reduced at the early postmortem aging ( $P < 0.05$ ) and almost disappeared at day 3. Meanwhile, the degraded plectin under the intact band emerged at 0 h and accumulated fast during the first 3 d aging ( $P < 0.05$ ). However, the amount of the degraded product reduced between day 3 and day 13 ( $P < 0.05$ ), indicating a further degradation of plectin during later stage of postmortem aging. Table 1 shows the correlations between LF-NMR  $T_2$  parameters (water population [P] and transverse relaxation time [T]) and plectin postmortem degradation. The amount of intact plectin during 1-d aging was significantly correlated to  $P_{21}$  at day 1 and 3 ( $P < 0.01$ ). There were significantly positive correlations between plectin intensity within 1 d and  $P_{22}$  at 1 d and 3 d ( $P < 0.01$ ). Moreover, plectin content at 0 h was significantly correlated to  $T_{21}$  and  $T_{22}$  at 1 d ( $P < 0.01$ ). As  $P_{21}$  and  $P_{22}$  represent the intra- and extra-myofibrillar water, respectively, these results indicate that plectin might contribute to the water distribution during condition including the decrease of intra-myofibrillar water and the increase of extra-myofibrillar water early postmortem. As  $T_{21}$  and  $T_{22}$  reflect the changes in intra- and extra-myofibrillar space, the results above imply the contribution of plectin to the structural changes including the decrease of intra-myofibrillar space and the space between myofibrils and sarcolemma early postmortem.

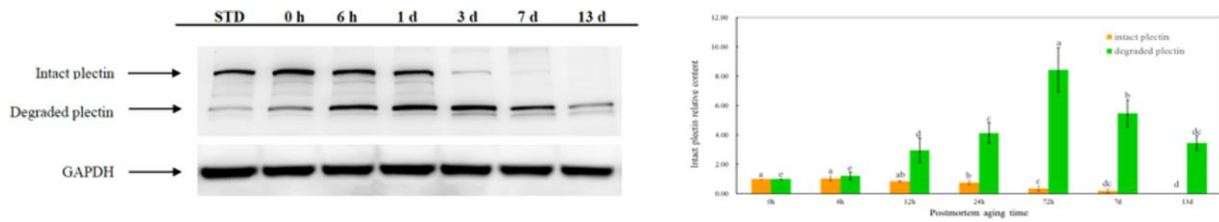


Figure 1. Western blot analysis of plectin expression in postmortem pork LT muscle. Note: 50  $\mu$ g of total protein samples was loaded per lane. The standard was a sample with a stable band in the trial test and was loaded onto each gel

Table 1 Pearson correlation coefficients and levels of significance for the correlations between plectin degradation and LF-NMR T<sub>21</sub>, T<sub>22</sub>, P<sub>21</sub>, P<sub>22</sub>.

Intact plectin	P <sub>21</sub>				P <sub>22</sub>				T <sub>21</sub>				T <sub>22</sub>			
	1d	3d	7d	13d	1d	3d	7d	13d	1d	3d	7d	13d	1d	3d	7d	13d
0h	-0.549**	-0.500**	-0.421**	-0.002	0.576**	0.544**	0.551**	0.386**	-0.445**	-0.119	0.031	-0.264	-0.415**	0.026	-0.253	-0.338*
6h	-0.404**	-0.394**	-0.051	0.117	0.428**	0.406**	0.269	0.275	-0.172	0.133	0.022	-0.239	-0.322*	0.029	0.195	-0.187
1d	-0.486**	-0.380*	-0.246	0.070	0.480**	0.414**	0.409**	0.310*	-0.230	0.142	0.058	0.238	-0.350*	0.052	0.221	-0.198
3d	-0.549**	-0.338*	-0.354*	0.043	0.566**	0.425**	0.421**	0.118	-0.431**	0.182	0.034	0.145	-0.466**	0.116	0.140	-0.054

- Note: \*\*: Correlation is significant at the 0.01 level. \*: Correlation is significant at the 0.05 level. (n=44).

#### IV. CONCLUSION

Plectin degradation was found to be negatively correlated to P<sub>21</sub>, T<sub>21</sub>, and T<sub>22</sub> and positively correlated to P<sub>22</sub>. The correlations were established between plectin content at the earlier aging time and the population of T<sub>2</sub> water at the later days, indicating the potential contribution of plectin degradation to the changes of water distribution during postmortem aging.

Keywords: plectin, pork, postmortem aging, water-holding capacity