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Journal Prevention

- 1 Cleavage of FNDC5 and insights into its maturation process

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48 Abstract

49 FNDC5 corresponds to an irisin precursor that increases with exercise. Studies suggest that irisin 50 mediates beneficial effects in adipose tissues, skeletal muscle, bone, and brain. However, the cleavage 51 and maturation processes of FNDC5 have not been clearly identified. This study aimed to show that the 52 signal peptide and transmembrane domain of FNDC5 were associated with the secretion of its 53 ectodomain. Localization studies identified the signal peptide that was responsible for endoplasmic 54 reticulum targeting activity of nascent FNDC5 and showed that the FNDC5 ectodomain corresponding to 55 irisin could be transported across the membrane by a transmembrane domain. Analysis of cleavage 56 constructs revealed that the ectodomain of FNDC5 could be cleaved from its signal peptide and 57 transmembrane attachment. Genetic ablation of the signal peptide cleavage site blocked N-glycosylation 58 of FNDC5. Identification of the FNDC5 maturation process should facilitate our understanding of irisin 59 secretion.

60 Introduction

61 FNDC5 (fibronectin domain-containing protein 5) was initially discovered and characterized by two 62 research groups with a focus on the fibronectin type III repeat domain and the peroxisomal targeting 63 motif, respectively (Ferrer-Martinez et al., 2002; Teufel et al., 2002). In examining the metabolic benefits 64 of exercise that are expressed by muscle in response to PGC-1α activation, irisin has been identified as a 65 myokine that is produced after cleavage of the ectodomain portion of the type I membrane protein, 66 FNDC5 (Bostrom et al., 2012).

Following the initial description of irisin, many authors have tried to identify its physiological role. Studies suggest that irisin mediates certain beneficial effects in adipose tissues, skeletal muscle, bone, and brain (Colaianni et al., 2015; Reza et al., 2017; Wrann et al., 2013; Xiong et al., 2018). In particular, irisin has been shown to mediate the thermogenesis of white adipose tissue (Bostrom et al., 2012; Zhang et al., 2014). Since the peptide was first described, interest in its therapeutic potential has increased and been linked to a wide range of therapeutic effects, including improved metabolism, energy homeostasis, cardiovascular health, and cognition (Leal et al., 2018; Young et al., 2019).

74 The reported circulating concentrations of irisin seem to differ dramatically in human serum or 75 plasma, which have raised doubts about the validity of the commercial enzyme-linked immunosorbent 76 assay (ELISA) kits used to detect circulating irisin and even about the existence of irisin in humans 77 (Crujeiras et al., 2015; Polyzos et al., 2015). Undervaluation data obtained from tandem mass 78 spectrometry reveal that irisin circulates at a level at or above the levels quantitated for many other 79 important biological hormones, such as leptin and insulin (Jedrychowski et al., 2015). These findings 80 confirm the actual existence of irisin in humans. Information regarding different FNDC5 protein 81 fragments, the presence of mature or precursor forms of irisin, and glycosylation of the protein probably 82 contribute to immunoassay quality and irisin detection accuracy (Erickson, 2013; Perakakis et al., 2017).

83 The irisin sequence was described in 2012 by Bostrom et al. Using mass spectrometry, the irisin 84 peptide (KDEVTMKE) was found in the medium of the N-terminus of FNDC5 (without the signal 85 peptide) fused with the C-terminus of the FC-domain of IgG construct (Bostrom et al., 2012). The peptide 86 sequence was consistent with amino acids 133-140 of FNDC5 according to UniProt entry Q8K4Z2. The 87 peptide that was shed into the media might have been only a small portion of the irisin. The protein could 88 be detected in serum, but cleavage site(s) and exact size both remain a matter of debate (Sanchis-Gomar 89 et al., 2015). Therefore, the formation sequence of the irisin precursor is needed to better understand how 90 secreted irisin is formed. In addition, validating whether irisin is actively cleaved from FNDC5 might 91 explain the differences in circulating concentrations of irisin between acute and chronic training and 92 reveal additional stimulators of irisin secretion. The data obtained in this study identify the cleavage and 93 maturation process of FNDC5 by fusing different FNDC5 protein fragments with DsRed2 and/or EGFP

94 and should facilitate an improved detection of irisin in physiological concentrations.

95 Materials and Methods

96 Bioinformatic identification of FNDC5

97 The signal peptide sequence of FNDC5 was analyzed using SignalP-software 98 (http://www.cbs.dtu.dk/services/SignalP/) (Petersen et al., 2011). Sequences with positive C, Y, S, and 99 D-scores were considered positive for a signal sequence. The transmembrane domain of FNDC5 was 100 analyzed in the Phobius web interface (http://phobius.cgb.ki.se) (Kall et al., 2004). Phobius provides a list 101 of the location of the predicted transmembrane helices. If the whole sequence is labeled as cytoplasmic or

102 non-cytoplasmic, the prediction is that it contains no membrane helices.

103 Plasmid construction

To express FNDC5 and its different domains as EGFP or DsRed2 tagged fusion proteins, the DNA 104 105 sequences coding FNDC5 and its different domains were cloned into pEGFP-C1, pEGFP-N1, or 106 pDsRed2-C1 vectors (Clontech, USA), respectively, to produce the following: N-terminus of FNDC5 107 fused with EGFP (pEGFP-C1-FNDC5); C-terminus of FNDC5 fused with EGFP (pEGFP-N1-FNDC5); 108 N-terminus of signal peptide fused with EGFP (pEGFP-C1-SP); N-terminus of signal peptide fused with 109 DsRed2 (pDsRed2-C1-SP); N-terminus of transmembrane-cytoplasmic (C) domains fused with EGFP 110 (pEGFP-C1-TM-C); and N-terminus of irisin-transmembrane-cytoplasmic domains fused with EGFP 111 cleavage, (pEGFP-C1-irisin-TM-C). То analyze FNDC5 plasmid called a 112 pDsRed2-C1-FNDC5-EGFP-N1 was constructed. Potential ectodomain-cleavage-deficient mutants 113 (pEGFP-N1-FNDC5deletion¹³⁹KE¹⁴⁰) were constructed via site-directed mutagenesis of 114 pEGFP-N1-FNDC5 using the QuikChange strategy with high-fidelity Pfu DNA polymerase (TaKaRa, 115 China) and verified by sequencing. Signal-peptide-cleavage-deficient mutants (pEGFP-N1-FNDC5 116 deletion²⁸AD²⁹), N-terminus of irisin fused with EGFP (pEGFP-C1-Irisin), and N-terminus of 117 transmembrane domains (TM) fused with EGFP (pEGFP-C1-TM) have been previously described (Nie 118 and Liu, 2017). pDsRed2-C1-EGFP-N1 and pDsRed2-C1-NLS-EGFP-N1 were constructed as EGFP 119 and DsRed2 fusion control plasmids.

120 Cell transfection and fluorescence microscopy

Human HeLa and 293T cells were cultured in DMEM medium (Gibco, USA) supplemented with 10% fetal calf serum, 100 U of penicillin, and 100 μ g/mL streptomycin. Cells were grown at 37°C in a humidifier with a 5% CO₂ atmosphere. Cells were transfected with the indicated plasmids using LipofectamineTM 2000 transfection reagent (Thermo Fisher Scientific, USA) according to the manufacturer's protocol. The fluorescent pattern was viewed using a Nikon AIR laser-scanning confocal microscope.

127 Antibodies and reagents

128The following antibodies and reagents were used: EGFP (sc-8334, Santa Cruz Biotechnology, 1:1000,
v/v), irisin (ab174833, Abcam, 1:1000, v/v), α-tubulin (ab15246, Abcam, 1:100, v/v), FITC labeled goat
anti-rabbit IgG (ab6717, Abcam, 1:1000, v/v), HRP labeled goat anti-rabbit IgG (ab6721, Abcam,
1:10000, v/v), PNGase F (P0704S, New England Biolabs), RIPA Lysis Buffer (20-188, Millipore),
Protease Inhibitor Cocktail (P8340, Sigma-Aldrich), TBST buffer (CW0043, Cwbiotech, China), BCA
Protein Assay Kit (23225, Thermo Fisher Scientific), ECL Plus Western Blotting Substrate (32132,
Thermo Fisher Scientific), and Prestained Protein Ladder (26616, Thermo Fisher Scientific).

135 Western Blotting analysis

136 Whole cell lysates were prepared by separately adding 1 mL of RIPA Lysis Buffer and 5 μ L Protease

137 Inhibitor Cocktail to each of the groups. After a 30-minute incubation on ice, samples were centrifuged at

138 12,000 rpm for 15 minutes at 4°C to pellet cellular material, and supernatants were assessed for protein 139 concentrations. Protein concentrations were assessed using the BCA Protein Assay Kit according to the 140 manufacturer's protocol. Western blotting was performed as follows: whole cell lysates in RIPA lysis 141 buffer were mixed with 5× SDS-PAGE sample buffer. Proteins were subjected to SDS-PAGE with a 12% 142 acrylamide gel. The gel-separated proteins were transferred to PVDF membranes, incubated with 143 primary antibodies overnight at 4°C, and probed with HRP labeled secondary antibodies for 1 hour at 144 room temperature. The PVDF membranes with protein-antibody complexes were washed three times 145 using TBST buffer. The proteins on the PVDF membranes were visualized with Thermo Scientific Pierce 146 ECL Plus Western Blotting Substrate on a Tanon 5200 (Tanon, China) detection system. The experiment 147 was repeated at least three times.

148 **Results**

149 Identification of potential transport signals within the FNDC5 protein

As shown in Figure 1A, FNDC5 in mice contains 209 amino acids with a 28 amino acid signal peptide

N-terminal, followed by an ectodomain domain, a transmembrane domain spanning 152-170 amino acids, and a cytoplasmic domain segment. Using SignalP 4.1 server, we found that FNDC5 contains an

152 actus, and a cytoplastice domain segment. Using Signal 4.1 server, we round that TTDCS contains an 153 N-terminal signal peptide sequence and that the cleavage site is located between ²⁸A and ²⁹D (Figure 1B).

Phobius analysis revealed that FNDC5 contained transmembrane helices. The N-terminal sequence of the

155 transmembrane helices was the non-cytoplasmic portion and the C-terminal sequence of the

156 transmembrane helices was the cytoplasmic portion (Figure 1C).

157 Localization of constructs containing transport signals

158 Constructs containing transport signals were prepared (Figure 2B) and verified by sequencing and 159 western blotting analysis (Figure 2D) and subcellular localization was evaluated. As shown in Figure 2A, 160 DsRed2 labeled signal peptide (DsRed2-SP fusion protein) was observed in a vesicular-like compartment around the nucleus in FITC labelled a-tubulin HeLa cells. To determine the specific vesicular 161 162 compartment in which the signal peptide was localized, we co-transfected EGFP labeled signal peptide 163 (EGFP-SP fusion protein) and pDsRed2-ER vector (an ER labeling marker). The co-localization of 164 EGFP-SP with DsRed2-ER indicates that the signal peptide is an ER transport signal. To identify the 165 transmembrane domains responsible for the transmembrane localization of the ectodomain, we 166 constructed combinatorial fragments of FNDC5 domains without a signal peptide. As shown in Figure 167 2C, EGFP-TM was localized in the cytomembrane. EGFP-irisin-TM-C behaved similarly to 168 EGFP-TM-C with respect to transmembrane localization.

169 FNDC5 cleavage

170 We generated non-cleavage cytoplasm localization control by fusing DsRed2 with EGFP, and 171 non-cleavage nucleus localization control by fusing nuclear localization signals (NLS) with 172 DsRed2-EGFP. As shown in Figure 3A, the yellow fluorescence of DsRed2-EGFP was evenly 173 distributed throughout the cytoplasm, and the yellow fluorescence of DsRed2-NLS-EGFP was evenly 174 distributed throughout the cell nucleus in HeLa cells. Exploration of the fluorescent pattern of 175 DsRed2-FNDC5-EGFP showed that the red and green fluorescence were slightly merged (Figure 3B). 176 The red fluorescence of the DsRed2-FNDC5-EGFP protein was observed in a vesicular-like 177 compartment. These results suggested that FNDC5 occurs during proteolytic cleavage.

In view of the proteolytic cleavage of FNDC5, we compared DsRed2-FNDC5-EGFP constructs with
 constructs containing various transport signals in the lysates of 293T cells by western blotting. In cells
 transfected with the pDsRed2-C1-FNDC5R-EGFP-N1 plasmid (Figure 3C, lane 5), a 30 kDa cleavage
 product was observed, which was in agreement with the signal peptide cleavage form (DsRed2-SP) of

DsRed2-FNDC5-EGFP. Meanwhile, we observed a 50 kDa cleavage product (Irisin-TM-C-EGFP),
which was similar to the EGFP-Irisin-TM-C construct (Figure 3C, lane 3) and the signal peptide cleavage
product of FNDC5-EGFP construct (Figure 3C, lane 6). In addition, a 36 kDa cleavage product
(TM-C-EGFP) was observed, which was in agreement with the EGFP-TM-C construct (Figure 3C, lane
and the signal peptide and irisin cleaved product of FNDC5-EGFP construct (Figure 3C, lane 6).

187 Effect of cleavage-deficient mutation on N-glycosylation of FNDC5

188 To investigate the role of the signal peptide cleavage and transmembrane domain cleavage during 189 FNDC5 maturation, a signal-peptide-cleavage-deficient mutant by alanine and aspartate acid deletion (deletion ${}^{28}AD^{29}$) and a potential ectodomain-cleavage-deficient mutant by lysine and glutamic acid deletion (deletion ${}^{139}KE^{140}$) were forcibly expressed in 293T cells. PNGase F was used to remove the 190 191 entire N-glycan structure. FNDC5-EGFP (Figure 4A lane 1) and FNDC5-EGFP-deletion¹³⁹KE¹⁴⁰ (Figure 192 193 4A lane 4) proteins displayed a highly heterogeneous molecular weight pattern. The heterogeneous 194 pattern changed after PNGase F treatment (Figure 4B lanes 1 and 4), and the latter band was in agreement 195 with the predicted molecular weight of FNDC5-EGFP protein. However, EGFP-FNDC5 (Figure 4A lane 196 2) and FNDC5-EGFP-deletion²⁸AD²⁹ (Figure 4A lane 3) proteins displayed little or no glycosylation 197 pattern compared to that of the FNDC5-EGFP protein. Moreover, the pattern was slightly decreased by 198 PNGase F treatment (Figure 4B lanes 2 and 3). Meanwhile, the same result was obtained using 199 anti-FNDC5 antibody detection (Figure 4C and Figure 4D).

200 Discussion

Irisin is a myokine secreted into circulation by skeletal muscles under certain physiological conditions, such as aerobic training (Figure 5). Furthermore, the localization of irisin sequences alone (EGFP-irisin)

has been previously described (Nie and Liu, 2017). We have shown that FNDC5 contains an N-terminal

signal sequence (Figure 1B) and that the signal peptide of FNDC5 can result in the retention or retrieval of nascent FNDC5 in the ER (Figure 2A). The signal peptide contains two arginines (11R and 15R) in the

205 of hascent FNDC5 in the ER (Figure 2A). The signal peptide contains two arginines (FIR and FSR) in the 206 middle, and an aromatic amino acid, tryptophan (8W and 17W), at both ends (Figure 1A) that possibly 207 conforms to the arginine-based ER localization consensus motif $\Phi/\Psi/RRXR$ (Michelsen et al., 2005; Nie 208 et al., 2016) (where Φ/Ψ denotes an aromatic or bulky hydrophobic residue and X represents any amino 209 acid).

210 Fluorescence analysis of the cleavage construct (pDsRed2-C1-FNDC5-EGFP-N1) revealed that the 211 signal peptide and transmembrane domain cleavage products were partially merged (Figure 3B). In the 212 lysates of 293T cells transfected with the cleavage construct plasmid, the 30 kDa fragment observed 213 corresponded to the signal peptide cleavage form (DsRed2-SP) of DsRed2-FNDC5-EGFP. In addition, a 214 53 kDa cleavage product (irisin-TM-C-EGFP) wherein the DsRed2-signal peptide was cleaved was 215 observed (Figure 3C). These results suggested that the presence of the signal peptide maintained ER 216 retention or retrieval of FNDC5 until it was masked as a result of signal sequence cleavage. The findings 217 also suggest that FNDC5 maturation includes at least two proteolytic cleavages: one between the signal 218 peptide and ectodomain, and the other between the ectodomain and transmembrane domain.

Nascent secretory protein synthesis begins in the ER, which is charged with the tasks of correctly integrating nascent proteins and ensuring correct post-translational modification and folding (Barlowe and Miller, 2013; Dancourt and Barlowe, 2010). The removal of the signal peptide cleavage site (deletion ²⁸AD²⁹) has a significant effect on the N-glycosylation of FNDC5, which is mostly retained in the ER [14]. In addition, the removal of a potential ectodomain cleavage site (deletion ¹³⁹KE¹⁴⁰) was observed, similar to that of wild type FNDC5 (Figure 4). The experiments suggested that signal peptide sequence cleavage of nascent FNDC5 occurs before downstream N-glycosylation tasks take place.

Once secretory protein FNDC5 has completed its signal peptide cleavage and N-glycosylation modification, it becomes capable of forward traffic. We have examined the possible contribution of each of the transmembrane containing segments of FNDC5 to ectodomain transmembrane localization, which

was achieved by chimera constructs (EGFP-TM, EGFP-TM-C, and EGFP-irisin-TM-C). The
 EGFP-irisin-TM-C construct behaves similarly to EGFP-TM-C with respect to transmembrane
 localization and is independent of irisin sequences (Figure 2C). Therefore, the ability of the FNDC5
 transmembrane domain to act as an inserted membrane transport signal is a striking feature.

233 Crucial data obtained from co-immunoprecipitation study revealed that aV family integrin 234 complexes are irisin receptors in osteocyte (Kim et al., 2019). The findings re-kindled interest in irisin as 235 a candidate target in the treatment of osteoporosis and raised a concern that its use for the browning of 236 adipose tissue, might also lead to osteoporosis (Anastasilakis et al., 2019; Farmer, 2019). Moreover, the 237 research progress of irisin in heart disease, Alzheimer's disease, and dihydromyricetin stimulation of 238 irisin secretion has made encouraging progress (Li et al., 2018; Lourenco et al., 2019; Zhou et al., 2015). 239 Our current data allowed us to clearly identify the cleavage between the signal peptide and ectodomain, 240 and the other between the ectodomain and transmembrane domain. Furthermore, based on the results 241 shown in our Supporting Information, a 35 kDa cleavage fragment (TM-C-EGFP) was observed in lysates of 293T cells transfected with the pEGFP-N1-FNDC5deletion¹³⁹KE¹⁴⁰ plasmid. We favor the 242 243 possibility that the precise proteolytic cleavage site between the ectodomain and transmembrane domain position does not occur merely between ¹⁴⁰E and ¹⁴¹M. 244

In view of the findings reported here, the proteolytic cleavages of FNDC5 are evident. FNDC5 maturation includes at least two proteolytic cleavages: one between the signal peptide and ectodomain, and the other between the ectodomain and transmembrane domain. Our findings provide experimental evidence regarding the functioning of the irisin and FNDC5 with the aspect of cleavage and maturation. Further studies are needed to explore the precise ectodomain cleavage sites and the molecular mechanism of cleavage regulation. Moreover, studies on the mechanisms involved in the proteolytic cleavages would allow us to assess stimulators of irisin secretion.

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Conflicts of Interest: The authors declare that they have no conflicts of interest with the contents of this article.

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Legends



Figure 1. Diagram of the structural domain sequences of FNDC5. (A) Schematic drawing of FNDC5. The signal peptide is boxed in red. The fibronectin type III (FNIII) domain is underlined in yellow. The transmembrane domain is boxed in green. The cytoplasmic domain is underlined in blue. (B) Result of FNDC5 signal peptide analysis using SignalP-software. (C) Result of FNDC5 transmembrane domain analysis using the Phobius web interface.



Figure 2. Identification of construct containing transport signal localizations. (A) Representative images of FNDC5 signal peptide intracellular localization. pEGFP-C1-SP and pDsRed2-C1-SP plasmids were transfected into HeLa cells. pDsRed2-ER vector was used as an ER labeling marker (Clontech, USA). α-tubulin was detected using an FITC labeled antibody. Nuclei were stained with DAPI. (B) Schematic representation of the construction of different plasmids. (C) Representative images of intracellular localization of FNDC5 transmembrane domain and construction. pEGFP-C1-TM, pEGFP-C1-TM-C, pEGFP-C1-irisin, and pEGFP-C1-irisin-TM-C plasmids were transfected into HeLa cells. pEYFP-Mem vector (Clontech, USA) was used as aplasma membrane target marker. Nuclei were stained with DAPI. (D) Western blotting analysis was carried out using EGFP antibodies on protein extracts collected from 293T cells transfected with pEGFP-C1-irisin (Lane 1), pEGFP-C1-SP (Lane 2), pEGFP-C1-TM (Lane 3), pEGFP-C1-TM-C (Lane 4), pEGFP-C1-irisin (Lane 5), and pEGFP-C1-irisin-TM-C (Lane 6) plasmids.

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Figure 3. Analysis of FNDC5 cleavage. (A) Representative images of intracellular localization of DsRed2-EGFP and DsRed2-NLS-EGFP proteins. pDsRed2-C1-EGFP-N1 plasmid was used as a non-cleavage cytoplasm localization control. pDsRed2-C1-NLS-EGFP-N1 plasmid was used as a non-cleavage nucleus localization control. Representative images of DsRed2-FNDC5-EGFP protein localization. pDsRed2-C1-FNDC5-EGFP-N1 plasmid was transfected into HeLa cells. (C) Western blotting analysis was carried out using EGFP antibodies on protein extracts collected from 293T cells transfected with pEGFP-C1 (Lane 1), pEGFP-C1-TM-C (Lane 2), pEGFP-C1-irisin-TM-C (Lane 3), pEGFP-C1-FNDC5 (Lane 4), pDsRed2-C1-FNDC5-EGFP-N1 (Lane 5), and pEGFP-N1-FNDC5 (Lane 6) plasmids.



Figure 4. Effect of mutation on FNDC5 N-glycosylation. 293T cells were transfected with pEGFP-N1-FNDC5 (Lane 1), pEGFP-C1-FNDC5 (Lane 2), pEGFP-N1-FNDC5-deletion²⁸AD²⁹ (Lane 3), and pEGFP-N1-FNDC5-deletion¹³⁹KE¹⁴⁰ (Lane 4) plasmids. Protein lysates (20 μ g protein per lane) were immunoblotted after treatment (+) or non-treatment (-) with PNGase F. Western blotting analysis was carried out using EGFP and FNDC5 antibodies.

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Figure 5. Model of FNDC5 maturation and cleavage. Nascent FNDC5 proteins are translated in the ER. Once the FNDC5 protein is ready for forward traffic, signal peptide sequence cleavage activity operates to ensure that FNDC5 is not recognized by the retrieval or retention machinery before the N-glycosylated FNDC5 is fully folded and assembled. Ectodomain cleavage occurs on the cell plasma membrane. Subsequent to ectodomain cleavage, irisin is generated. The ectodomain cleavage of FNDC5 must be tightly regulated, as the permanent release of irisin is incompatible with a state of health. In addition, it is an area lacking information regarding the precise ectodomain cleavage site, and furthermore, the molecular mechanism of FNDC5 cleavage regulation.

Highlights

- The signal peptide responsible for ER targeting of FNDC5 was identified
- FNDC5 ectodomain was cleaved from its signal peptide and transmembrane • attachment
- Genetic ablation of the signal peptide cleavage site blocked FNDC5 ٠ N-glycosylation