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Role of MuscleBlind-Like proteins in the regulation of expression of CaV61 isoforms in adult skeletal muscle

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Voltage-gated calcium channels (CaVs or VGCCs) are major regulators of calcium-related cellular functions. In skeletal muscle, though the essential component of the pore channel is the CaVa1 subunit, the CaVB1 subunit is an essential subunit guaranteeing CaV fine-tuning activity.

CaVβ1-E and CaVβ1-D are two different isoforms of CaVβ1 protein in skeletal muscle, expressed during embryogenesis and in healthy innervated adult muscle, respectively. Importantly, our recent study demonstrated that the embryonic CaVB1-E isoform expression increases after a nerve damage in adult skeletal muscle and enables the expression of GDF5 (Growth Differentiation Factor 5) to counteract excessive muscle wasting (*Traoré et al. 2019*). However, the mechanisms leading to the increase in CaVβ1-E expression are unknown to date.

Our RNAseq data analysis in innervated versus denervated muscles revealed MuscleBlind-Like (MBNL) proteins as potential candidates regulating CaVB1 expression in skeletal muscle. Interestingly, in a human model of Dystrophy Myotonic 1 (DM1), the sequestration of MBNLs in toxic nuclear aggregates is related to an impaired splicing of CaVB1 transcript (CACNB1) (Arandel et al. 2017).

Here, we evaluate the effect of a modulation of MBNLs protein levels on the expression of CaVB1 isoforms in both *in vitro* and *in vivo* systems as well as in pathological mouse models of DM1.





Mbnl1 modulates the expression of CaV_{β1} isoforms in vivo



Variations of Cacnb1 isoforms at transcriptional level

RT-PCR triplex Cacnb1 Ex2-3					
Ctl	OE shMbpl1 CoVR1E				
And Personnel in Concession, Name	Cacnb1-E				
	Cacnb1-D				



• Variations of CaVβ1 isoforms at protein level



Mbnl3 modulates the expression of CaVβ1 isoforms *in vivo*



A downregulation of Mbnl1 induces an increase of the expression of CaV β 1-E in mouse skeletal muscle through a direct or indirect mechanism.

Cacnb1-D seems to be expressed after the activation of an alternative promoter

	Mouse	e Cacnb1 g	gene	
primer_fw	4 5 6	7A 7B 8	9 10 11	12 13 // 14
Ex ATG				primer_rv

• Nanopore sequencing on mouse and human muscles



Mouse						
	TA inn	TA den	GAS inn	GAS den		
Ex_ATG	100,00	100,00	100,00	100,00	100,00	
11 <mark>0</mark> 0011111111110000	94,58	92,01	89,96	82,83	89,84	
11 <mark>0</mark> 0011100000011001	0,00	0,00	0,00	11,19	2,80	
11 <mark>0</mark> 0011110111111001	0,44	2,84	5,37	1,37	2,50	
11 <mark>0</mark> 0011111101111001	2,12	1,90	1,68	1,58	1,82	
11 <mark>0</mark> 0011101111111001	0,76	0,81	0,70	0,69	0,74	
11 <mark>0</mark> 0011111011111001	0,49	0,73	0,66	0,73	0,65	
10 <mark>0</mark> 001111111111001	0,53	0,59	0,55	0,45	0,53	
11 <mark>0</mark> 001111111101001	0,42	0,34	0,32	0,39	0,37	
11 <mark>0</mark> 0001111111111001	0,18	0,19	0,17	0,18	0,18	
11 <mark>0</mark> 0011111110111001	0,15	0,11	0,15	0,19	0,15	
11 <mark>0</mark> 0010111111111001	0,18	0,13	0,13	0,09	0,13	
11 <mark>0</mark> 0011111001111001	0,13	0,14	0,14	0,10	0,13	
11 <mark>0</mark> 0011110101111001	0,02	0,04	0,08	0,08	0,05	
11 <mark>0</mark> 001111111110001	0,02	0,05	0,09	0,03	0,05	
11 <mark>0</mark> 001111111111111	0,00	0,12	0,01	0,00	0,03	
1100011100000001001	0.00	0.00	0.00	0.10	0.03	

<u>Human</u>					
	BC1	BC2	BC3	BC4	
Ex_ATG	100,00	100,00	100,00	100,00	100,00
110 <mark>0</mark> 001011011110110101	90,85	93,86	88,45	88,20	84,60
110 <mark>0</mark> 001011010110110101	0,26	0,35	0,36	0,32	6,61
110 <mark>0</mark> 001011011100110101	2,08	2,20	1,94	2,04	1,85
100 <mark>0</mark> 000000000000000000000000000000000	1,58	0,23	0,71	2,36	0,74
110 <mark>0</mark> 001011001110110101	0,81	0,85	0,67	0,61	0,71
110 <mark>0</mark> 001011011010110101	0,68	0,73	0,72	0,62	0,63
110 <mark>0</mark> 000000000000000000000000000000000	0,01	0,01	2,61	1,70	0,51
11C <mark>0</mark> 00101000000000000001	0,01	0,02	1,45	0,00	0,50
110 <mark>0</mark> 001011011110110001	0,30	0,35	0,40	0,31	0,38
100 <mark>0</mark> 001011011110110101	0,31	0,34	0,48	0,37	0,33
110 <mark>0</mark> 001010001110110101	0,03	0,08	0,03	0,06	0,31
110 <mark>0</mark> 001011010110110001	0,01	0,00	0,01	0,01	0,31
110000000000000000000000000000000000000	0,66	0,01	0,00	1,50	0,25
1100001011010000000001	0,01	0,00	0,14	0,00	0,19
1100001011010100110101	0,01	0,02	0,02	0,02	0,16
	i				

The « Ex_ATG1 » intronic region is never asociated with an initiation of transcription in Ex1.

 \rightarrow The expression of *Cacnb1* isoforms could be transcriptionally regulated, with the existence of an alternative promoter.

shMbnl1+2 mouse model of DM1 :

0 = Exon absent / 1 = Exon present

MBNLs impacts *Cacnb1* mRNA stability

Luciferase-Cacnb1D 3'UTR Luciferase-*Cacnb1E 3'UTR*

Cacnb1-E increases in DM1 pathological models

HSA mouse model of DM1:





CONCLUSIONS & PERSPECTIVES

- Mbnl1 negatively regulates Mbnl3 \bullet
- A downregulation of Mbnl1, and the subsequent increase of Mbnl3, leads to increased CaVβ1-E and decreased CaVβ1D expression levels *in vivo*
- The expression of *Cacnb1-D* seems to be under the control of an alternative promoter
- *Cacnb1* transcripts stability is modulated by MBNLs *in vitro* through their 3'UTR •
- Mouse models of DM1 are associated with an increased *Cacnb1-E* expression
- Studying the potential alternative promoter controlling *Cacnb1-D* expression •
- Characterization of the splicing events occurring at *Cacnb1* Ex2-3 and Ex13-14 \bullet
- Studying a potential cross-regulation of CaVβ1-D on CaVβ1-E expression \bullet
- Deciphering the role of CaV β 1-E in DM1 pathophysiology \bullet







