

## **Supplemental Material to:**

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**Modulation of distinct isoforms of L-type calcium channels by Gq-coupled receptors in *Xenopus oocytes*:  
Antagonistic effects of G $\beta$  $\gamma$  and protein kinase C**

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**<http://www.landesbioscience.com/journals/channels/article/22016/>**

Channel composition	<i>n</i>	<i>G</i> <sub>max</sub> $\mu S$	<i>V</i> <sub>rev</sub> <i>mV</i>	<i>V</i> <sub>o</sub> <i>mV</i>	<i>K</i> <sub>o</sub> <i>mV</i>
$\alpha_{1C}$ -LNT, $\alpha_2\delta$	14	6.3 $\pm$ 1	66.5 $\pm$ 2	9.7 $\pm$ 1	8.4 $\pm$ 0.3
$\alpha_{1C}$ -LNT, $\alpha_2\delta$ , $\beta_{2b}$	12	40.2 $\pm$ 8.4	63.1 $\pm$ 1.8	-4.8 $\pm$ 1.1	6.7 $\pm$ 0.3
$\alpha_{1C}$ -LNT,+9a, $\alpha_2\delta$	14	7.8 $\pm$ 1.5	72.7 $\pm$ 6.2	10.6 $\pm$ 0.4	8.4 $\pm$ 0.2
$\alpha_{1C}$ -LNT,+9a, $\alpha_2\delta$ , $\beta_{2b}$	20	41.0 $\pm$ 5.6	62.8 $\pm$ 1.6	-4.8 $\pm$ 0.8	6.8 $\pm$ 0.2
$\alpha_{1C}$ -SNT, $\alpha_2\delta$	7	20.68 $\pm$ 5.5	65.2 $\pm$ 1.9	5.1 $\pm$ 0.3	8.2 $\pm$ 0.5
$\alpha_{1C}$ -SNT, $\alpha_2\delta$ , $\beta_{2b}$	6	51.4 $\pm$ 11.9	61.2 $\pm$ 1	-11.6 $\pm$ 1.8	6.3 $\pm$ 0.5
$\alpha_{1C}$ -SNT,+9a, $\alpha_2\delta$	8	37.2 $\pm$ 13.4	74.8 $\pm$ 6.8	2.6 $\pm$ 0.6	8.7 $\pm$ 0.8
$\alpha_{1C}$ -SNT,+9a, $\alpha_2\delta$ , $\beta_{2b}$	12	43.6 $\pm$ 9.9	60.3 $\pm$ 0.5	-9.8 $\pm$ 1.4	6.5 $\pm$ 0.2

Voltage	isoform	$\tau_{slow}$ <i>ms</i>	<i>A</i> <sub>slow</sub>	$\tau_{fast}$ <i>ms</i>	<i>A</i> <sub>fast</sub>	<i>C</i>
0mV	$\alpha_{1C}$ -LNT (n=5)	2571 $\pm$ 52	0.42 $\pm$ 0.01	844 $\pm$ 46	0.37 $\pm$ 0.03	0.21 $\pm$ 0.02
	$\alpha_{1C}$ -LNT,+9a (n=6)	4138 $\pm$ 707	0.35 $\pm$ 0.03	987 $\pm$ 66	0.52 $\pm$ 0.08	0.12 $\pm$ 0.06
20mV	$\alpha_{1C}$ -LNT (n=5)	2056 $\pm$ 95	0.46 $\pm$ 0.02	634 $\pm$ 33	0.44 $\pm$ 0.03	0.103 $\pm$ 0.01
	$\alpha_{1C}$ -LNT,+9a (n=6)	2672 $\pm$ 225	0.46 $\pm$ 0.02	619 $\pm$ 62	0.44 $\pm$ 0.03	0.103 $\pm$ 0.02
40mV	$\alpha_{1C}$ -LNT (5)	2148 $\pm$ 152	0.51 $\pm$ 0.02	490 $\pm$ 43	0.45 $\pm$ 0.03	0.04 $\pm$ 0.02
	$\alpha_{1C}$ -LNT,+9a (n=6)	2577 $\pm$ 258	0.55 $\pm$ 0.01	377 $\pm$ 68	0.39 $\pm$ 0.03	0.06 $\pm$ 0.03

Weiss 2012. Supplemental Table 1

MLRALVQPATPAYQPLPSHLSAETE	<b><u>ST</u></b> CKGTWHEAQLNHFYISPGG	<b><u>S</u></b> NY	50
<b><u>G</u></b> SPRPAHANMNANAAAGLAPEHIP	<b><u>T</u></b> PGAALSWQAAIDAARQAKLMGSAGN		100
AT <b><u>I</u></b> <b><u>STV</u></b> <b><u>SS</u></b> TQRKRQQYGKPKKQG	<b><u>S</u></b> TTATRPPRALLCLTLKNPIRRAC <b><u>I</u></b>		150
VEWKPF $\bar{E}$ III			160

Weiss 2012. Supplemental Figure 1

**A**

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Rabbit SM,  $\alpha_{1C_b}$       gaggcactccagcgggcttgcatgccagaagaaaggaagtttgcttggttagtcact 1450
|||||  |||||  |||  ||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||
Human chromosome 12 gaggcactccggcgggcatgcttgatcagaagaaaggaagtttgcttggttagtcact 2510936

Rabbit SM,  $\alpha_{1C_b}$       ccacagagacccatg 1472
|||||  |||||  |||||
Human chromosome 12 ccacagaaacccatg 2510951
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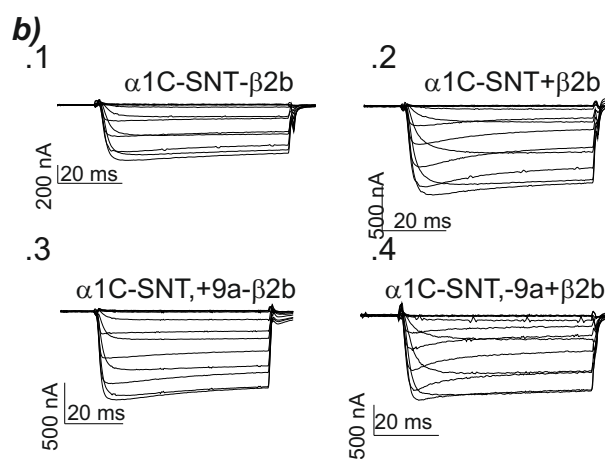
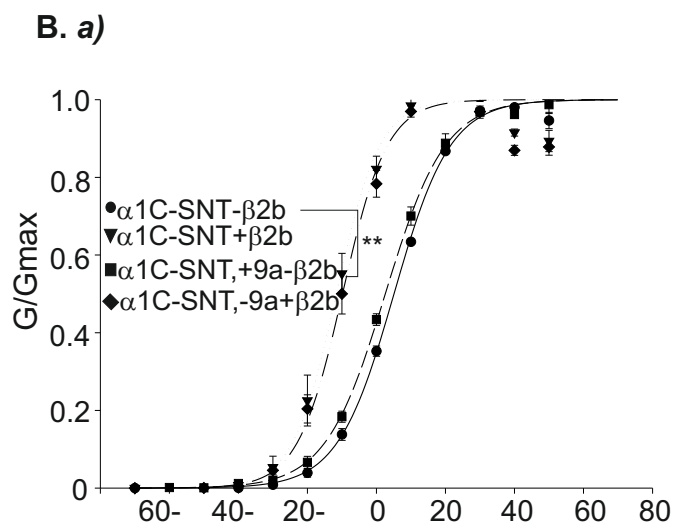
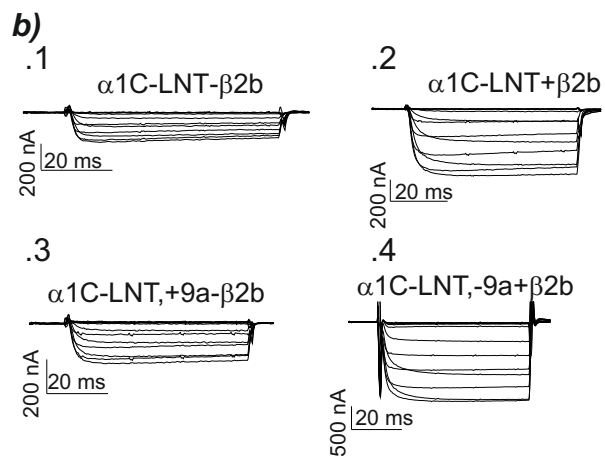
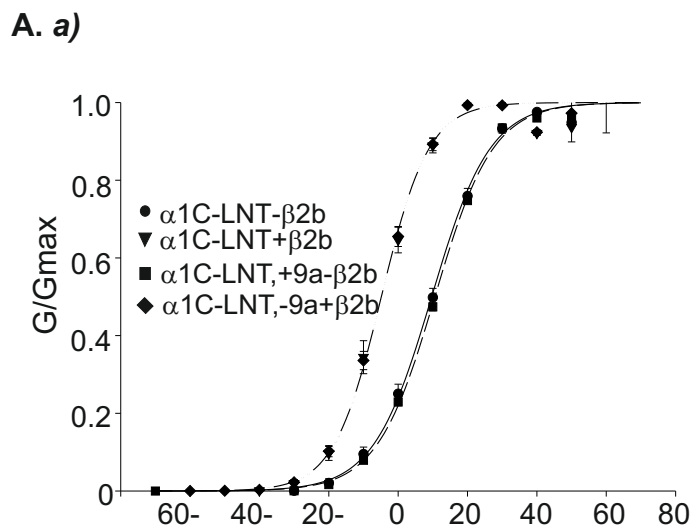
**B**

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 $\alpha_{1C}$ , NT-rabbit long    GEFSKEREKAKARGDFQKLREKQQLEEDLKGYLDWITQAEDIDPENEDEGMDEEKPRN-- 493
 $\alpha_{1C,77}$ , NT-human short GEFSKEREKAKARGDFQKLREKQQLEEDLKGYLDWITQAEDIDPENEDEGMDEEKPRN-- 463
 $\alpha_{1C_b}$ , rabbit SM      GEFSKEREKAKARGDFQKLREKQQLEEDLKGYLDWITQAEDIDPENEDEGMDEEKPRNRG 465
*****

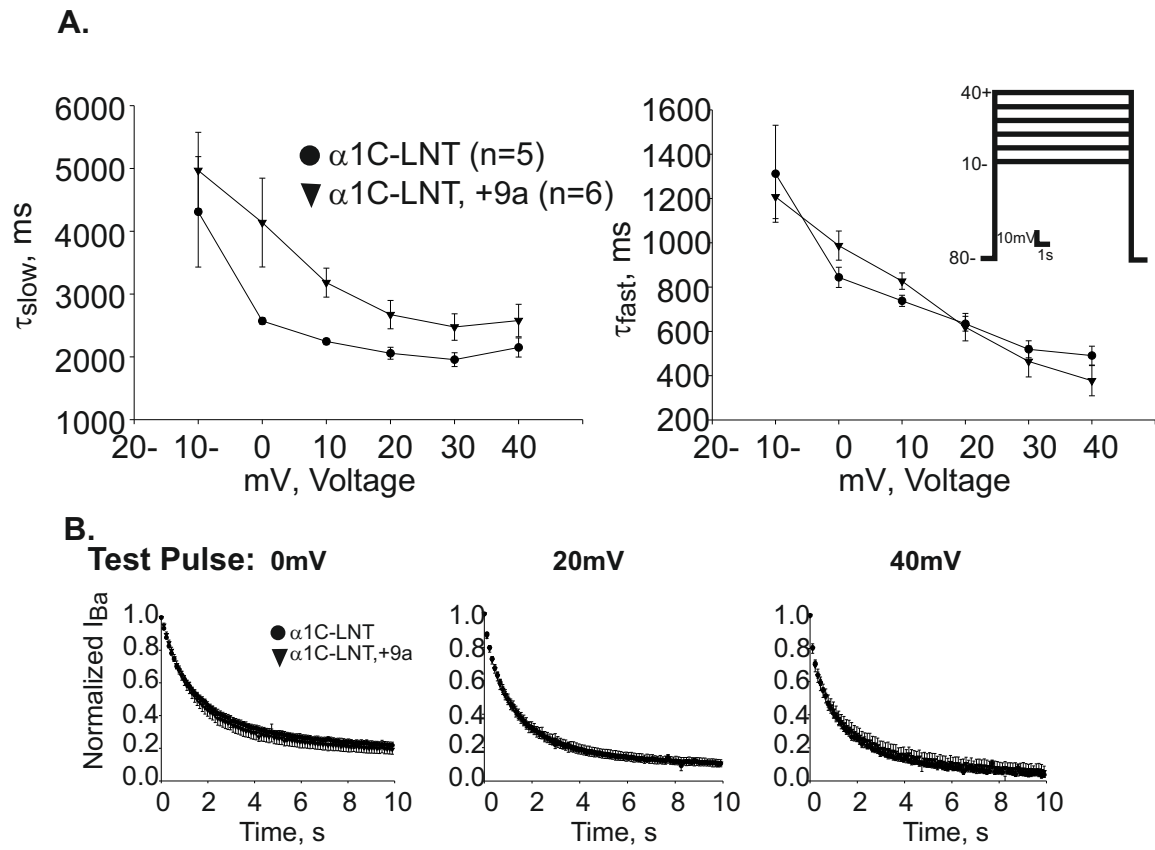
 $\alpha_{1C}$ , NT-rabbit long    -----MSMPTSETESVNTENVAGGDIEGENCGARLAHRISKS 530
 $\alpha_{1C,77}$ , NT-human short -----MSMPTSETESVNTENVAGGDIEGENCGARLAHRISKS 500
 $\alpha_{1C_b}$ , rabbit SM      TPAGLHAQKKGKFAWFHSTETHVSMPTSETESVNTENVAGGDIEGENCGARLAHRISKS 525
                               :*****

 $\alpha_{1C}$ , NT-rabbit long    KFSRYWRRWNRFCRRKCRAAVKSN 554
 $\alpha_{1C,77}$ , NT-human short KFSRYWRRWNRFCRRKCRAAVKSN 524
 $\alpha_{1C_b}$ , rabbit SM      KFSRYWRRWNRFCRRKCRAAVKSN 549
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Weiss 2012. Supplemental Figure 2



Weiss 2012. Supplemental Figure 3



Weiss 2012. Supplemental Figure 4

## Supplemental Material

**Supplemental Table 1.** Top. Activation parameters of four  $\alpha_{1C}$  isoforms. Bottom. Inactivation parameters of two  $\alpha_{1C}$ -LNT isoforms.

**Supplemental Figure 1. The amino acid composition of rabbit long-NT.** Ser/Thr mutations are underlined: mutant 1 (mut I), single underline; mutant 2 (mut II), double underline.

**Supplemental Figure 2. A 75 nucleotide long insertion in L1 of  $\alpha_{1C}$ .** *A.* Nucleotide alignment of the L1 insertion in rabbit and the part of human chromosome 12 encoding  $\alpha_{1C}$ . *B.* a.a. alignment of the entire L1 reveals a highly homologous sequence among the different species and isoforms. Identical a.a. are marked by asterisk, gaps are marked by dashes. Only short-NT human L1 is shown, as this region is identical in human long-NT isoform.

**Supplemental Figure 3.  $\alpha_{1C,+9a}$  constructs have similar activation parameters as  $\alpha_{1C,-9a}$  constructs.** Oocytes were injected with one of four different  $\alpha_{1C}$  isoforms,  $\alpha_{2\delta}$  with or without  $\beta_{2b}$ . G-V curves of the +9a isoforms vs. their controls reveal no significant changes in activation kinetics, when expressed with or without the auxiliary  $\beta_{2b}$  subunit. *Aa)* G-V curves of long-NT isoforms. *b)* representative traces of two LNT isoforms.  $I_{Ba}$  measured at 10 mV steps from -70 mV to +50 mV. *Ba)* G-V curves of short-NT isoforms. *b)* representative traces of two SNT isoforms.  $I_{Ba}$  measured at 10 mV steps from -70 mV to +50 mV. \*\*,  $p < 0.01$  by t-test; brackets indicate the two groups with a statistical significance effect.

**Supplemental Figure 4.  $\alpha_{1C,+9a}$  constructs have similar inactivation kinetics as  $\alpha_{1C,-9a}$  constructs.** *A.* Oocytes were injected with  $\alpha_{1C}$ -LNT  $\pm 9a$ ,  $\alpha_{2\delta}$ ,  $\beta_{2b}$ . The inactivation curve was fitted with two time constants,  $\tau_{slow}$  and  $\tau_{fast}$  (see Experimental Procedures for details). There was no significant difference between the two isoforms. *Indent:* The inactivation protocol consists of 6, 10s,

incrementing voltage steps (from -10mV to +40mV). B. Representative normalized Ba<sup>2+</sup> currents at 3 selected test pulses. Each point is mean±S.E.M.  $\alpha_{1C,-9a}$  n=5;  $\alpha_{1C,+9a}$  n=6.