



# Disruption of asparagine-linked glycosylation to rescue and alter gating of the $\text{Na}_v1.5\text{-Na}^+$ channel

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## Abstract

SCN5A gene encodes the voltage-gated sodium channel  $\text{Na}_v1.5$  which is composed of a pore-forming  $\alpha$  subunit of the channel. Asparagine (N)-linked glycosylation is one of the common post-translational modifications in proteins. The aim of this study was to investigate impact of N-linked glycosylation disruption on the  $\text{Na}^+$  channel, and the mechanism by which glycosylation regulates the current density and gating properties of the  $\text{Na}^+$  channel. The  $\text{Na}_v1.5\text{-Na}^+$  channel isoform ( $\alpha$  submit) derived from human was stably expressed in human embryonic kidney (HEK)-293 cells (Nav1.5-HEK cell). We applied the whole-cell patch-clamp technique to study the impact of N-linked glycosylation disruption in Nav1.5-HEK cell. Inhibition of the N-glycosylation with tunicamycin caused a significant increase of  $\text{Na}_v1.5$  channel current ( $I_{\text{Na}}$ ) when applied for 24 h. Tunicamycin shifted the steady-state inactivation curve to the hyperpolarization direction, whereas the activation curve was unaffected. Recovery from inactivation was prolonged, while the fast phase ( $\tau_{\text{fast}}$ ) and the slow phase ( $\tau_{\text{slow}}$ ) of the current decay was unaffected by tunicamycin.  $I_{\text{Na}}$  was unaffected by tunicamycin in the presence of a proteasome inhibitor MG132 [N-[(phenylmethoxy)carbonyl]-L-leucyl-N-[(1S)-1-formyl-3-methylbutyl]-L-leucinamide], while it was significantly increased by tunicamycin in the presence of a lysosome inhibitor butyl methacrylate (BMA). These findings suggest that N-glycosylation disruption rescues the  $\text{Na}_v1.5$  channel possibly through the alteration of ubiquitin–proteasome activity, and changes gating properties of the  $\text{Na}_v1.5$  channel by modulating glycan milieu of the channel protein.

**Keywords** Glycosylation ·  $\text{Na}_v1.5$  channel · Tunicamycin · MG132 · Ubiquitin–proteasome pathway

## Abbreviations

N-linked	Asparagine-linked	TEA-Cl	Tetraethylammonium chloride
HEK cells	Human embryonic kidney	TEA-OH	Tetraethylammonium hydroxide
MG132	N-[(phenylmethoxy)carbonyl]-L-leucyl-N-[(1S)-1-formyl-3-methylbutyl]-L-leucinamide	HEPES	4-(2-Hydroxyethyl)-1-piperazineethanesulfonic acid
BMA	Butyl methacrylate	G	Conductance
DMEM	Dulbecco's modified Eagle's medium	I-V curve	Current–voltage curve
$I_{\text{Na}}$	Voltage-gated sodium channel current	ER	Endoplasmic reticulum
LQT3	Long QT syndrome type 3	FOXO1	Forkhead box protein O1

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## Introduction

In cardiovascular system many ion channels regulate transmembrane currents that make much contribution to excitation in cardiomyocytes [1–7]. Among them the voltage-gated  $\text{Na}^+$  channel current ( $I_{\text{Na}}$ ) plays a critical and significant role for the fast upstroke of the action potentials and is essential for proper conduction of the electrical impulse. Voltage-gated sodium channels are composed of a pore-forming  $\alpha$  subunit and auxiliary  $\beta$  subunits. SCN5A gene encodes