

Effects of retigabine and the novel M-current activator BHV-7000 on epilepsy-associated *KCNQ2* variants

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ABSTRACT

Rationale: The voltage-gated potassium channels *KCNQ2* (K_v7.2; Q2) and *KCNQ3* (K_v7.3; Q3), encoded by *KCNQ2* and *KCNQ3*, respectively, co-assemble to form a channel complex (Q2/Q3) responsible for generating the M-current that regulates neuronal excitability. *KCNQ2* pathogenic variants identified in children with developmental and epileptic encephalopathy (DEE) most often exhibit loss-of-function with dominant-negative effects. Pharmacological potentiation of M-current with retigabine (RET) or newer investigational agents such as BHV-7000 (BHV; formally KB-3061) is a potential therapy for this condition. Here, we compared the effects of RET with BHV on wild-type (WT) Q2/Q3 and channels comprised of DEE-associated *KCNQ2* variants.

Methods: We expressed Q2/Q3 channels in Chinese hamster ovary (CHO) cells and recorded whole-cell currents using automated planar patch clamp first in the absence then presence of vehicle (DMSO) or M-current activator (RET or BHV). Experiments were conducted using cells transfected with an equal mixture of WT and variant *KCNQ2* along with WT *KCNQ3* to generate heteromultimeric channel complexes that recapitulated the heterozygous state. Specific channel activity was determined by applying the M-current blocker XE-991 (10 μM) at the end of experiments, and only XE-991-sensitive currents were analyzed.

Results: At 3 μM, both compounds induced significant hyperpolarizing shifts in the voltage-dependence of activation (delta V_{1/2}) of WT channels, but the effect was greater for BHV (RET = -18.7±1.7, n=13; BHV = -32.1±1.4, n=8). Both compounds boosted current amplitude to a similar degree (relative to the no drug condition: RET = 7.0±0.9 fold increase; BHV = 6.2±0.9 fold increase) when measured during -10 mV test pulses. The half-maximal effective concentration (EC₅₀) for delta V_{1/2} of WT channels were RET = 4.9 μM; BHV = 1.4 μM. We also assessed the effects of RET and BHV on channels incorporating *KCNQ2* variants for which clinical responses to retigabine treatment were reported. BHV (3 μM) induced greater hyperpolarizing shifts in activation V_{1/2} than RET (3 μM).

Conclusions: The M-current activators RET and BHV exhibited effects on WT Q2/Q3 channels and restored current for channels containing DEE-associated pathogenic *KCNQ2* variants. We are testing several additional variants and determining concentration-response relationships for a more complete understanding of these effects. Delineating the heterogeneity of M-current activator responsiveness of pathogenic variants may enable better deployment of precision pharmacotherapies for *KCNQ2*-DEE.

Background

KCNQ2 pathogenic variants identified in children with developmental and epileptic encephalopathy (DEE) most often exhibit loss-of-function with dominant-negative effects. Pharmacological potentiation of M-current is a potential therapy for this condition. Our previous results (Vanoye et al, 2022, JCI Insight) demonstrated genotype-dependent differences in the response of *KCNQ2* variants to retigabine, a proposed precision therapy for *KCNQ2* developmental and epileptic encephalopathy. In this study we investigated whether the newer investigational agent BHV-7000 (BHV; formally KB-3061) exhibits genotype-dependent differences.

Methods

Cell Culture: Chinese hamster ovary cells (CHO-K1) stably expressing *KCNQ3*-WT (CHO-Q3) were grown in F-12 nutrient mixture medium supplemented with fetal bovine serum, penicillin and streptomycin with hygromycin selection at 37°C in 5% CO₂.

Molecular Biology: Variants were introduced into human *KCNQ2* using Quikchange mutagenesis (Agilent technologies). *KCNQ2* variants were expressed from plasmid pIRES2_KCNQ2_EGFP, whereas *KCNQ2*-WT was expressed from plasmid pIRES2_KCNQ2_CyOFP. These plasmids included green or orange fluorescent proteins, respectively, as transfection markers. The *KCNQ2* reading frame of all constructs was sequenced completely.

Electroporations: Wild type plus *KCNQ2* variant cDNAs were transiently co-transfected into CHO_KCNQ3 cells using the MaxCyte STX system.

Flow Cytometry: Transfection efficiency was evaluated by flow cytometry (CytoFLEX, Beckman Coulter) using a 488 nm laser and filters for green fluorescence (FITC, *KCNQ2*_variants coupled to EGFP expression), and orange fluorescence (PEA, *KCNQ2*_WT couple to CyOFP expression).

Electrophysiology: Automated planar patch clamp recording was performed on the Nanion SyncroPatch 768 PE platform. External solution contained (in mM): 140 NaCl, 4 KCl, 2.0 CaCl₂, 1 MgCl₂, 10 HEPES, 5 glucose, pH 7.4. The composition of the internal solution was (in mM): 60 KF, 50 KCl, 10 NaCl, 20 EGTA, 10 HEPES, 5 mM MgATP, pH 7.2. Retigabine (SIGMA) and BHV-7000 (Biohaven) were added from 10 mM stock solutions dissolved in DMSO. DMSO volume was constant for all compound concentrations (0-30 μM; 3 μl/ml). Whole-cell currents were recorded at room temperature from holding potential of -80 mV using 1000 ms depolarizing pulses from -80 to +40 mV (in 10 mV steps), followed by a 250 ms step to 0 mV to analyze tail currents. Number of cells (n) is given on the figure legends.

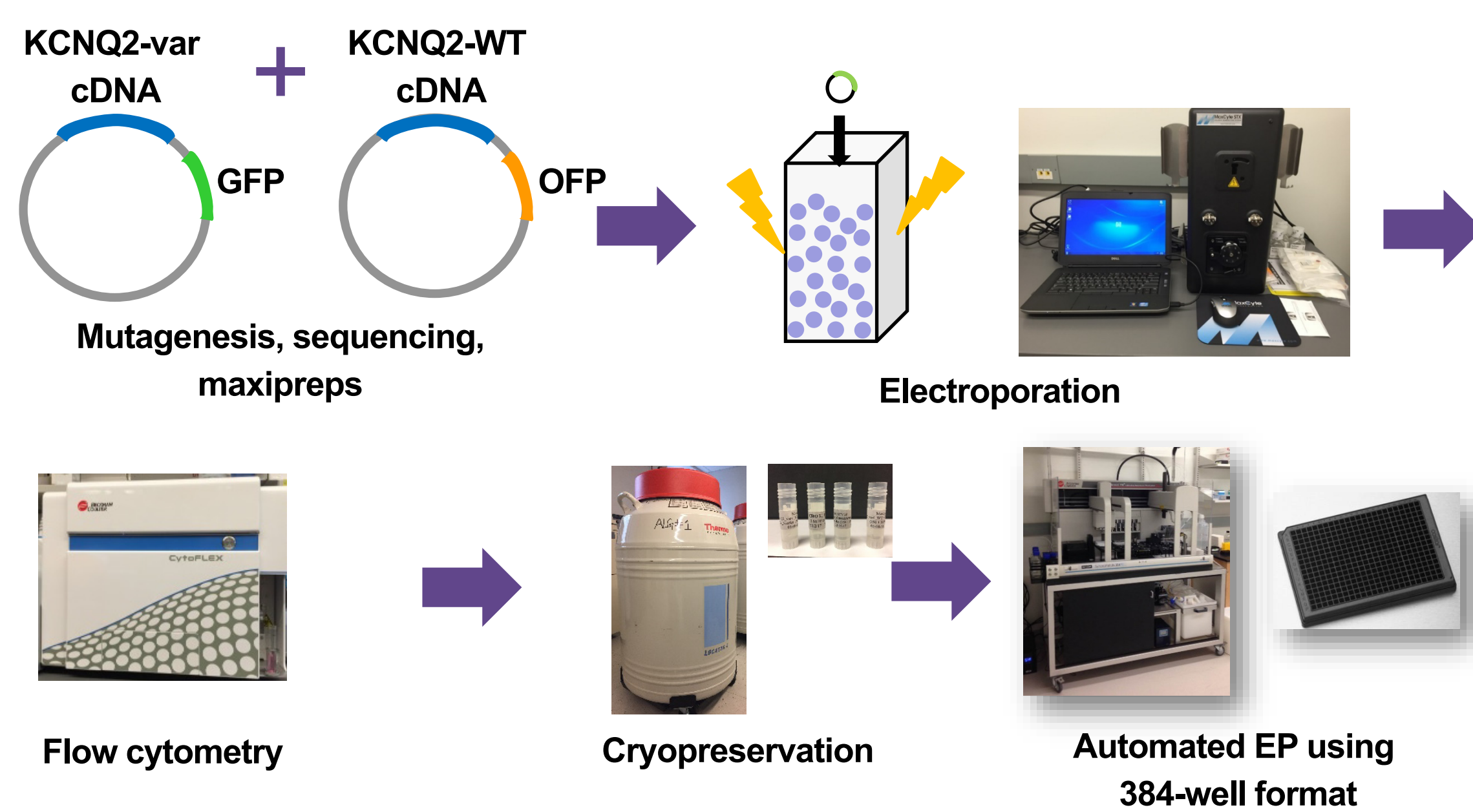


Fig. 1 – Experimental work-flow for high throughput functional studies of *KCNQ2* variants expressed in CHO-K1 cells stably expressing human *KCNQ3* (CHO-Q3 cells).

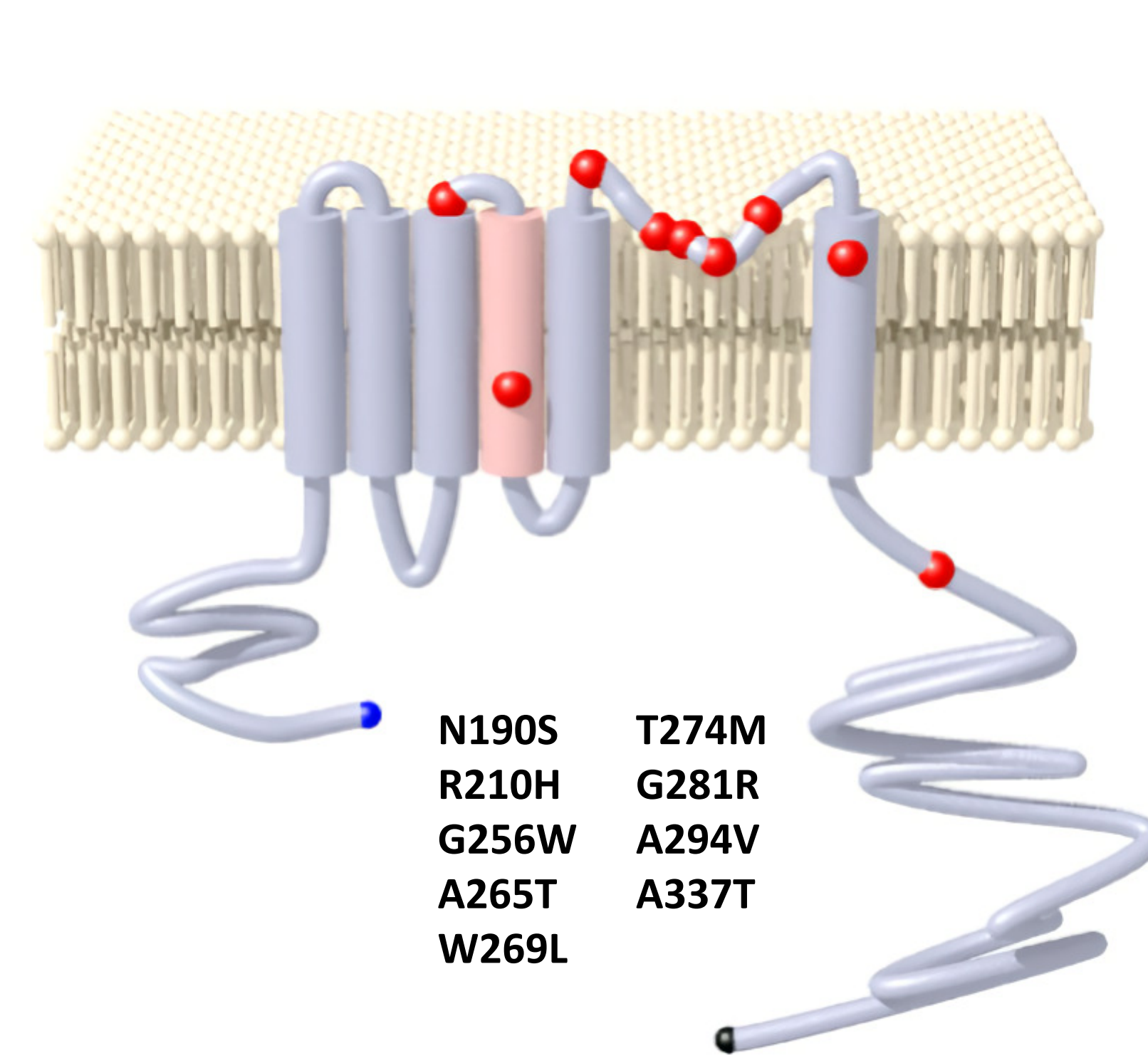


Fig. 2 – Variants analyzed in this study and location in the *KCNQ2* protein.

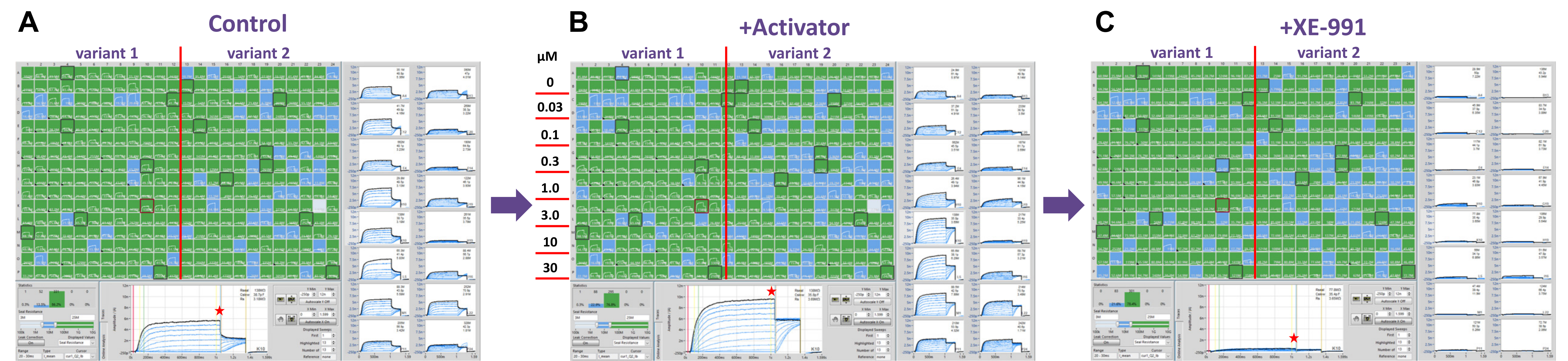
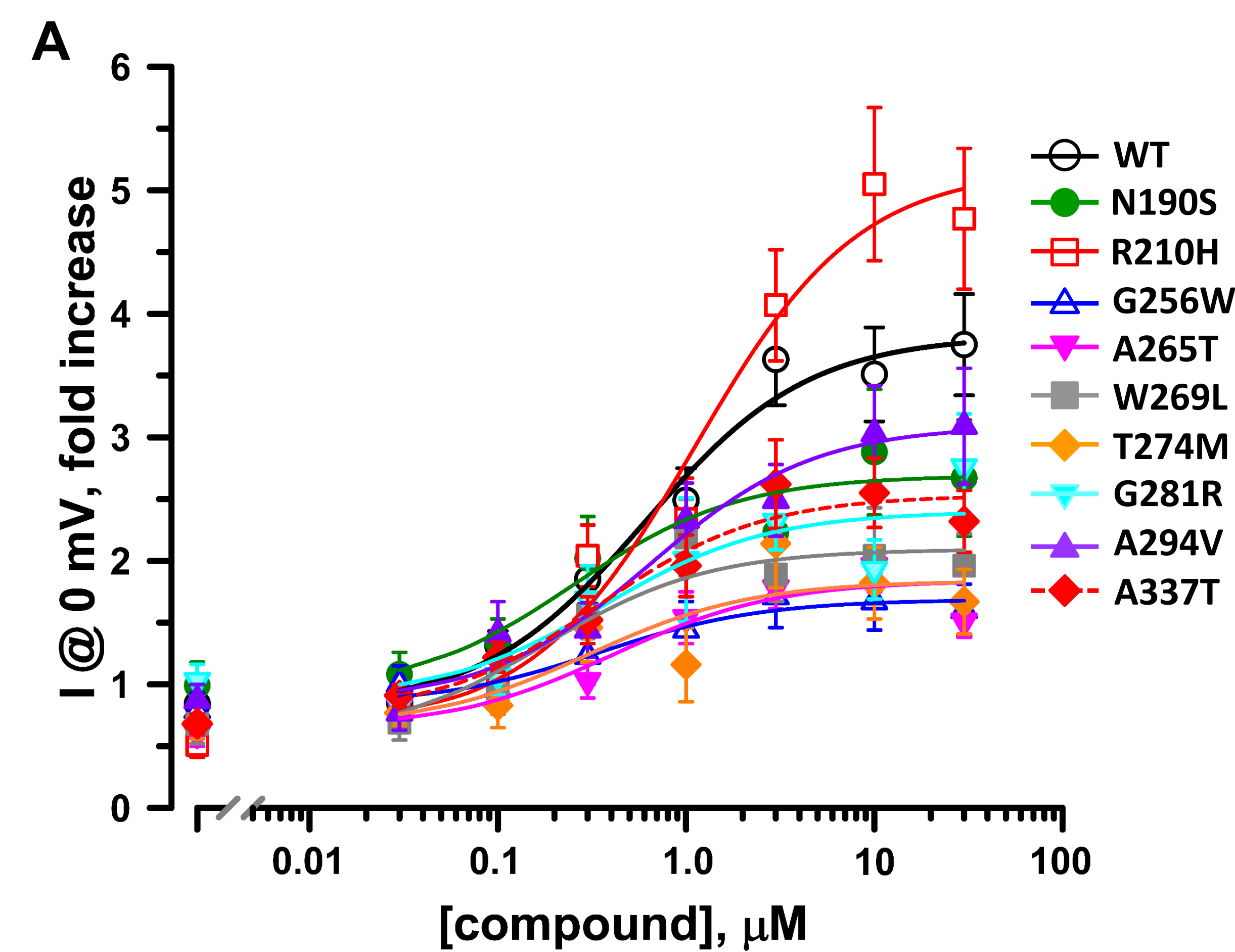


Fig. 3 Whole-cell recording protocol and use of XE-991 to isolate *KCNQ2* + *KCNQ3* current.

Screen-shots of whole-cell current recordings from CHO-Q3 cells transiently expressing *KCNQ2* variants (2 variants, 12 columns per variant, 192 data points per variant) recorded under Control conditions (A), following addition of 0-30 μM Activator (24 data points per concentration). (B), and after the addition of 10 μM XE-991 (C) Currents recorded in the presence of XE-991 were digitally subtracted from the Control and +Activator data. Only the XE-991-sensitive currents were analyzed. Whole-cell currents were measured at the end (998 ms) of a 1000 ms long voltage step (*).

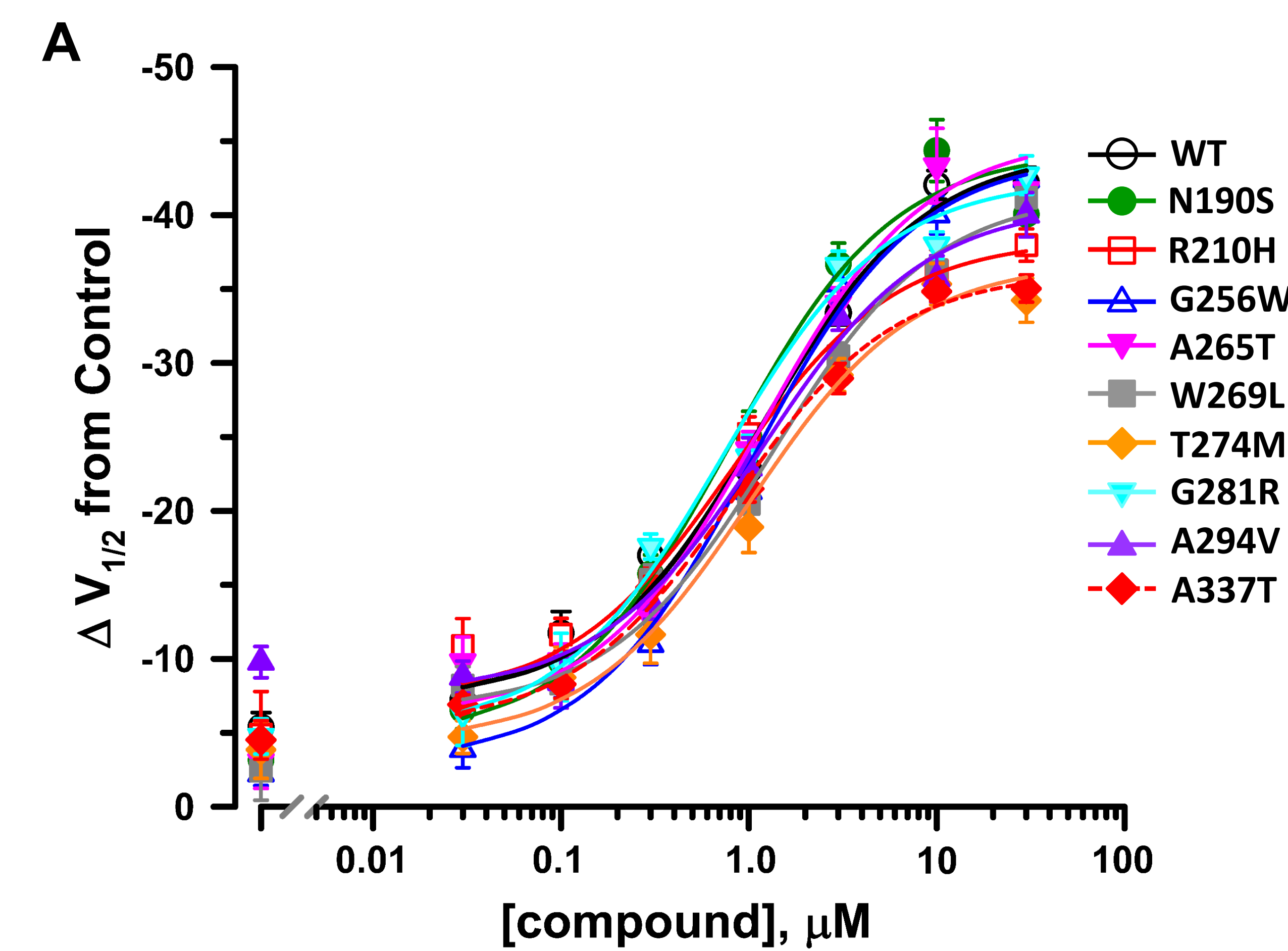


Channel variant	EC ₅₀ , μM	Maximum fold increase	Fold increase @ EC ₅₀
WT	0.6 ± 0.2	3.8 ± 0.2	2.3
N198S	0.2 ± 0.2	2.7 ± 0.2	1.8
R210H	1.1 ± 0.4	5.2 ± 0.3	2.9
G256W	0.3 ± 0.3	1.7 ± 0.1	1.3
A265T	0.4 ± 0.2	1.8 ± 0.1	1.2
W269L	0.2 ± 0.1	2.1 ± 0.3	1.3
T274M	0.3 ± 0.3	1.8 ± 0.2	1.2
G281R	0.4 ± 0.3	2.4 ± 0.2	1.6
A294V	0.6 ± 0.4	3.1 ± 0.2	2.0
A337T	0.3 ± 0.2	2.5 ± 0.1	1.6

Fig. 4 – Concentration response for BHV-7000 induced changes in whole-cell current density recorded at 0 mV

A. Average concentration response curves obtained from *KCNQ2* wild type and variants expressed as heterozygous channels (+*KCNQ2*-WT) exposed to various BHV-7000 concentrations. B. List of parameters from fitted data. N = 12-23 per concentration.

+Activator whole-cell currents were normalized to whole-cell currents recorded under Control conditions. Normalized data were fit with the equation $Y = \text{Bottom} + X * (\text{Top-Bottom}) / (EC_{50} + X)$.



Channel variant	EC ₅₀ , μM	Max. ΔV _{1/2} , mV	ΔV _{1/2} @ EC ₅₀ , mV
WT	1.2 ± 0.2	-44.4 ± 1.0	-25.9
N198S	0.8 ± 0.1	-44.4 ± 1.4	-24.1
R210H	0.8 ± 0.2	-38.4 ± 1.0	-22.8
G256W	1.1 ± 0.1	-44.1 ± 0.9	-23.6
A265T	1.2 ± 0.2	-45.3 ± 1.5	-25.6
W269L	1.3 ± 0.2	-41.5 ± 1.2	-24.0
T274M	1.0 ± 0.2	-36.9 ± 1.3	-20.6
G281R	0.7 ± 0.1	-42.5 ± 1.0	-23.8
A294V	1.2 ± 0.2	-40.7 ± 1.3	-24.2
A337T	0.8 ± 0.1	-36.2 ± 0.7	-20.8

Fig. 5 – Concentration response for BHV-7000 induced hyperpolarization of activation voltage-dependence

A. Average concentration response curves obtained from *KCNQ2* wild type and variants expressed as heterozygous channels (+*KCNQ2*-WT) exposed to various BHV-7000 concentrations. B. List of parameters from fitted data. N = 6-20 per concentration.

Voltage-dependence of activation values under Control and +Activator conditions were derived from tail currents normalized to maximal tail current amplitude and expressed as a function of the preceding voltages. Data were fit to a Boltzmann function $I(V) = \text{Bottom} + (\text{Top-Bottom}) / (1 + \exp((V_{1/2} - V) / \text{slope}))$.

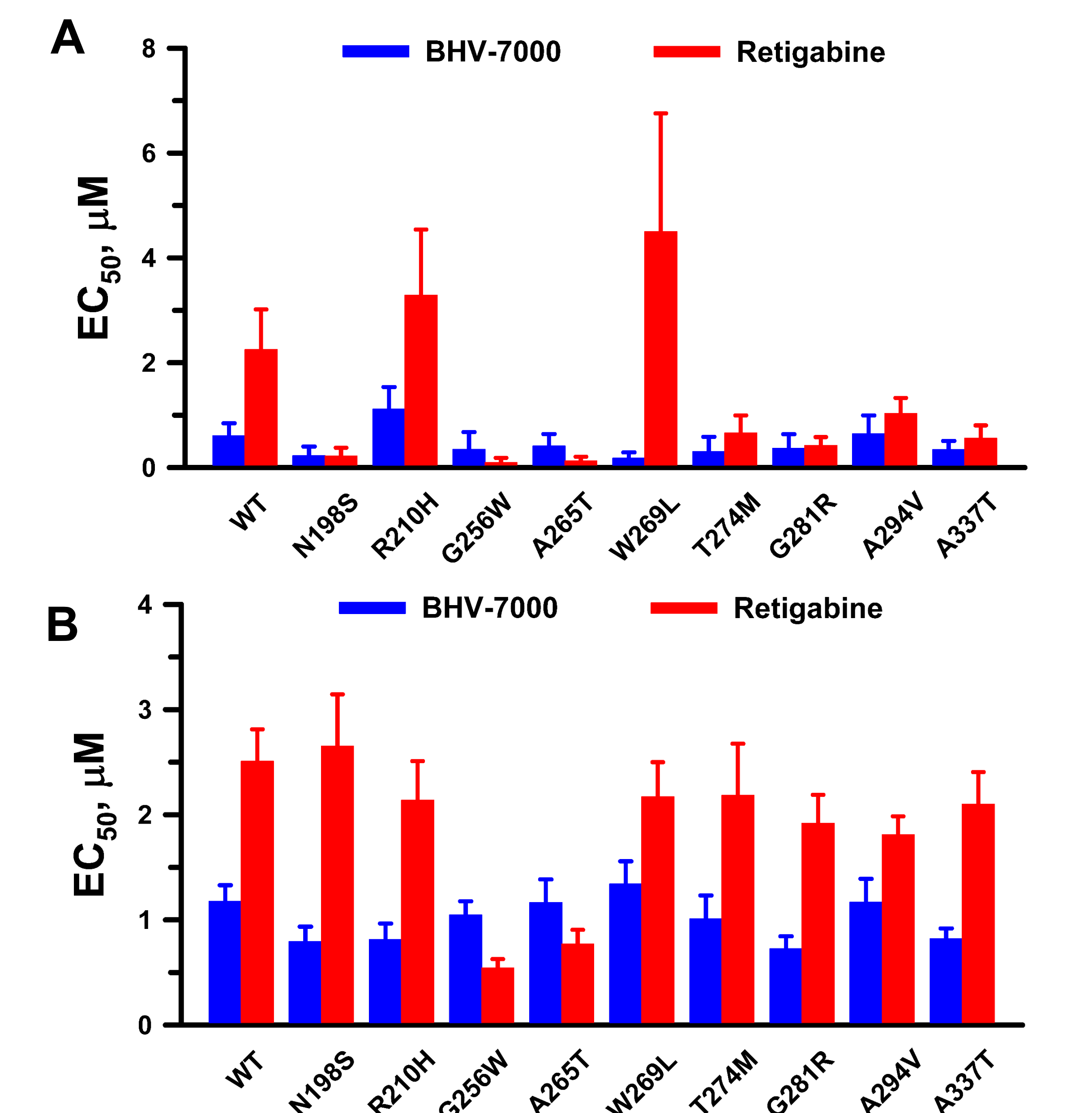


Fig. 6 – EC₅₀ values calculated for (A) changes in current density recorded at 0 mV, and (B) activation V_{1/2}.

EC₅₀ values were derived from concentration response curves obtained from *KCNQ2* wild type and variants channels expressed as heterozygous channels (+*KCNQ2*-WT) exposed to various BHV-7000 and retigabine concentrations. N = 5-23 per concentration.

Summary

1. Pharmacological potentiation of variant *KCNQ2* channels expressed in the heterozygous state can be analyzed by automated patch clamp.
2. Automated patch clamp analysis of variant *KCNQ2* channels reveal that *KCNQ2* channels exhibit dissimilar responses to activator exposure.
3. EC₅₀ values calculated for BHV-7000 and retigabine-induced increase in current density and hyperpolarization of the voltage-dependence of activation show larger heterogeneity for retigabine and more potent modulation for BHV-7000.

Conclusion

Pharmacological potentiation of pathogenic *KCNQ2* channels is variant dependent. Further understanding of variant-specific responsiveness may enable better deployment of targeted therapies.

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