Supplementary Table 10. Statistics: Analysis of Variance of tetanic mechanical properties of TTX-Sensitive and -Resistant Skeletal Muscles from *Thamnophis sirtalis* and *Thamnophis atratus* (pertaining to Figures 3 and Supp. Fig. 4).

Test:	Kruskal-Wallis (Non-Parametric ANOVA)			
Variable	Baseline Force (N g ⁻¹) by Genotype	F _{max} (N g ⁻¹) by Genotype	dFdt _{max} (N g ⁻¹ s ⁻¹) by Genotype	dFdt _{min} (N g ⁻¹ s ⁻¹) by Genotype
chi-squared	25.986	20.107	19.895	18.059
df	3	3	3	3
p-value	9.6 x 10 ⁻⁶	0.0002	0.0002	0.0004
Observations	52	52	52	52
Variable	Time to 10% F _{max} (s) by Genotype	Time to dFdt _{max} (s) by Genotype	Time to dFdt _{max} (s) by Genotype	Time to $dFdt_{min}$ (s) by Genotype
chi-squared	28.828	20.451	20.457	7.3948
df	3	3	3	3
p-value	2.43 x 10 ⁻⁶	0.0001	0.0001	0.0603
Observations	52	52	52	52

Definitions:

Baseline Force (N g⁻¹)

F_{max} (N g⁻¹)

Time to 10% F_{max} (s)

Time to 50% F_{max} relaxation (s)

Contraction Duration from 10% maximal contraction to 50% post-maximal relaxation (s)

 $dFdt_{max}$ (N $g^{-1} s^{-1}$)

 $dFdt_{min}$ (N g^{-1} s^{-1})

Time to dFdt_{max} (s)

Time to dFdt_{min} (s)

Contaction Duration from dFdt_{max} to dFdt_{min} (s)

Optimal baseline tension on the muscles in Newtons/gram-tissue

Maimum force produced by each phasic contraction in Newtons/gram

Time to develop 10% of maximal force from stimulus onset (seconds)

Time required to half-maximally relax from stimulus onset (seconds)

(Time to 50% relaxation) - (Time to 10% contraction) (seconds)

Positive peak of first derivative (in Newtons/gram-tissue per second)

Negative peak of first derivative (in Newtons/gram-tissue per second)

Time from stimulus onset to peak rate of force development (seconds)

Time from stimulus onset to peak rate of force relaxation (seconds)

(Time to dFdtmin) - (Time to dFdtmax) (seconds)

^{*}Note: WTa = Ancestral Thamnophis atratus, EPN = TTX resistant Thamnophis atratus; WTs = Ancestral Thamnophis sirtalis, LVNV = TTX resistant Thamnophis sirtalis

^{**}Note: Shaded values represent adjusted p-values that fall below a predetermined significance level (α =0.05)