

## ***Abcc8* (Sulfonylurea Receptor-1) Impact on Brain Atrophy after Traumatic Brain Injury Varies by Sex**

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### **Abstract**

Females have been understudied in pre-clinical and clinical traumatic brain injury (TBI), despite distinct biology and worse clinical outcomes versus males. Sulfonylurea receptor 1 (SUR1) inhibition has shown promising results in predominantly male TBI. A phase II trial is ongoing. We investigated whether SUR1 inhibition effects on contusional TBI differ by sex given that this may inform clinical trial design and/or interpretation. We studied the moderating effects of sex on post-injury brain tissue loss in 142 male and female ATP-binding cassette transporter subfamily C member 8 (*Abcc8*) wild-type, heterozygote, and knockout mice (12–15 weeks). Monkey fibroblast-like cells and mouse brain endothelium-derived cells were used for *in vitro* studies. Mice were injured with controlled cortical impact and euthanized 21 days post-injury to assess contusion, brain, and hemisphere volumes (vs. genotype- and sex-matched naïves). *Abcc8* knockout mice had smaller contusion volumes ( $p=0.012$ ) and larger normalized contralateral (right) hemisphere volumes (nRHV;  $p=0.03$ ) after injury versus wild type. This was moderated by sex: Contusions were smaller ( $p=0.020$ ), nRHV was higher ( $p=0.001$ ), and there was less global atrophy ( $p=0.003$ ) in male, but not female, knockout versus wild-type mice after TBI. Less atrophy occurred in males for each copy of *Abcc8* lost ( $p=0.023$ – $0.002$ , all outcomes). *In vitro*, sex-determining region Y (SRY) stimulated *Abcc8* promoter activity and increased *Abcc8* expression. Loss of *Abcc8* strongly protected against post-traumatic cerebral atrophy in male, but not female, mice. This may partly be mediated by SRY on the Y-chromosome. Sex differences may have important implications for ongoing and future trials of SUR1 blockade.

**Keywords:** atrophy; brain injuries; central nervous system; genotypic intracranial hemorrhage; sex; mice; traumatic

### **Introduction**

Multiple pre-clinical and early clinical studies suggest that inhibiting sulfonylurea receptor 1 (SUR1) may be neuroprotective after traumatic brain injury (TBI).<sup>1–4</sup> Based on encouraging results in preliminary studies demonstrating reduced cerebral edema and post-traumatic contusion growth,<sup>1–9</sup> a phase II trial is currently evaluating the efficacy of glyburide, a SUR1 inhibitor, in contusional TBI (ASTRAL, NCT03954041), and a phase III trial is ongoing in stroke (NCT2869453). However, previous TBI ther-

apies have reached similar points of translation only to fail in phase III trials.<sup>10–12</sup> Identifying subgroups likely to respond to a novel therapy may both improve the probability of demonstrating efficacy and increase effectiveness if adopted into clinical practice.

Male bias is increasingly recognized in TBI research and neuroscience research as a whole.<sup>13</sup> Although epidemiological studies report that females make up 30–40% of hospitalized TBI patients,<sup>14–17</sup> they frequently comprise only 20% of patients enrolled into phase III

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therapeutic trials.<sup>11,12,18–21</sup> A recent scoping review of >150 clinical reports by Gupte and colleagues identified that although males are ~40% more likely to suffer TBI, most studies report worse clinical outcomes in females.<sup>22</sup> They further identified a discontinuity between clinical and pre-clinical studies, given that the latter identify poorer outcomes in males.<sup>22</sup> Females are additionally grossly under-represented in both pre-clinical and clinical studies—a phenomenon that is only partly attributable to disease epidemiology.<sup>22</sup>

Sex is also a major source of biological heterogeneity; it influences molecular drivers of secondary injury and functional outcomes after TBI.<sup>23,24</sup> Sex-based differences in SUR1 inhibition have been reported for non-TBI neurological diseases.<sup>25,26</sup> However, pre-clinical SUR1-TBI reports have been limited to males,<sup>1,2,7,8</sup> and females have been under-represented in clinical SUR1-related reports.<sup>3,4,27,28</sup> Sex-based differences in response to SUR1 inhibition post-TBI could have important implications for both design and analysis of current and future clinical trials, as well as bedside prognosis and treatment.<sup>29</sup>

In humans, biological sexual dimorphism is driven largely by the sex-determining region Y (SRY) gene on the Y chromosome.<sup>30</sup> A previous study demonstrated that SRY forms a transcriptional complex with specificity protein 1 (Sp1), an activator of ATP-binding cassette transporter subfamily C member 8 (*Abcc8*; SUR1) transcription,<sup>31</sup> which synergistically activates transcription of the monoamine oxidase A gene in neural tissue.<sup>32</sup> A similar mechanism could contribute to sex differences in *Abcc8* post-traumatic expression.

We hypothesized that sex moderates effects of genetic SUR1 inhibition on contusion volume and brain atrophy after contusional TBI in mice, and that the SRY gene contributes to these differences.

## Methods

### Mice

The University of Pittsburgh Institutional Animal Care and Use Committee approved the experiments. One hundred forty-two adult (12- to 15-week-old) male and female mice with global *Abcc8* knocked out on a C57/BL6J background were used with heterozygote and wild-type littermates.<sup>33</sup> Mice were given food and water *ad libitum*, group housed (12-h day-night cycles) until injury, and single housed thereafter.

### Injury

Mice underwent a controlled cortical impact (CCI) model of moderate-severe TBI.<sup>2</sup> Briefly, mice were anesthetized with 4% isoflurane in a 2:1 N<sub>2</sub>O/O<sub>2</sub> gas mixture and placed in a stereotactic frame (Kopf, Tujunga, CA). A left scalp incision was made. A dental drill was used to make a 5-mm craniotomy over the left parietal cortex, and the bone flap was removed. CCI was performed

with a pneumatic impactor (Bimba, Monee, IL) with a flat 3-mm tipped impounder on the left parietal cortex (5 m/s, depth = 1.2 mm, dwell time = 50–60 ms). Mice recovered in a temperature-controlled chamber before returning to their cage. Naïve mice were euthanized at 12–15 weeks without injury.

### Histological volumetric analyses

Contusion volume (CV) and brain hemisphere (right hemisphere volume [RHV] and left hemisphere volume [LHV]) volumes were measured as previously described (Supplementary Methods).<sup>34</sup> Total brain volume (TBV) was also calculated (RHV + LHV). Normalized values for each parameter in injured mice were calculated as a percentage of the corresponding mean hemisphere or mean total brain volume for sex- and genotype-matched naïve mice (e.g., normalized CV<sub>male,KO</sub> [nCV<sub>male,KO</sub>] = CV<sub>injured,male,KO</sub>/LHV<sub>naive,male,KO</sub>). We normalized to naïve volumes, rather than right (contralateral) hemisphere volume of the same mouse, because right hemisphere atrophy after injury was disproportionate across genotype and sex (Table 1). Volume measurements were performed by technicians blinded to genotype and sex.

### Cell culture

COS-7 cells were cultured as previously described (Supplementary Methods).<sup>35</sup> Control cell lines and cell lines expressing SRY were developed by transfecting bEnd.3 cells (Supplementary Methods). Transfections were performed by using Lipofectamine 2000 (Invitrogen, Carlsbad, CA), unless otherwise indicated. Colonies that survived in a medium containing 0.2 mg/mL of hygromycin B were selected, and SRY expression was confirmed by immunoblot analysis.

### Luciferase promoter assay

SRY-stimulated activity of the promoter region of *Abcc8* was determined using luciferase reporter plasmids (Supplementary Methods).<sup>36</sup> COS-7 cells were transfected with luciferase reporter plasmids containing rat *Abcc8* or *Fra1* promoter, and expression was compared with cells transfected with chicken  $\beta$ -actin promoter-driven luciferase. Photinus luciferase reporter plasmids (pGL3-basic; Promega, Madison, WI), containing the rat *Abcc8* promoter (–578 to +80) or rat FOS-like antigen 1 (*Fra1*) promoter (–878 to +61), were transfected into COS-7 cells using Lipofectamine 2000 (Invitrogen). *Fra1* was used as a positive control for SRY-driven expression.<sup>37</sup> Cotransfection of pRL-CMV, Renilla luciferase expression plasmid was a control for transfection efficiency. After transfection, cells were maintained in basal conditions for 24 h, then transfected with either rat SRY, human SRY, or empty vector. Photinus and Renilla luciferases activity in extracts of transfected cells were measured using the Dual Luciferase Reporter

**Table 1. Sex, Genotype, and Injury Effects on Total Brain and Hemispheric Volumes**

	Naïve (mm <sup>3</sup> , mean ± SD)			Injured (mm <sup>3</sup> , mean ± SD)			Normalized (% , mean ± SD)		
	Wildtype	Knockout	p value	Wild type	Knockout	p value	Wild type	Knockout	p value
Total brain volume									
<i>Stratified by sex</i>									
Male	262.1 ± 18.1	239.7 ± 19.4	<b>0.011*</b>	216.4 ± 15.2	218.2 ± 10.9	0.70	82.4 ± 5.8	91.0 ± 4.5	<b>0.001*</b>
Female	257.7 ± 17.6	250.4 ± 11.7	0.25	232.1 ± 11.6	221.1 ± 9.6	<b>0.02</b>	90.1 ± 4.6	88.3 ± 3.7	0.32
Base model (all mice): $p_{\text{genotype}} = \mathbf{0.004}^\dagger$ , $p_{\text{sex}} = \mathbf{0.037}$ , $p_{\text{injury}} < \mathbf{0.001}^\dagger$							Base model (all mice): $p_{\text{genotype}} = 0.06$ , $p_{\text{sex}} = 0.32$		
Interaction model (all mice): $p_{\text{interaction}} = \mathbf{0.023}^\dagger$							Interaction model (all mice): $p_{\text{interaction}} = \mathbf{0.001}^\dagger$		
Ipsilateral (left) hemisphere volume									
<i>Stratified by sex</i>									
Male	130.4 ± 9.1	118.5 ± 10.7	<b>0.012*</b>	95.9 ± 6.9	96.1 ± 4.1	0.53	73.5 ± 5.3	81.1 ± 3.4	<b>0.0004*</b>
Female	127.6 ± 8.7	123.6 ± 5.9	0.20	105.1 ± 4.2	99.3 ± 5.3	<b>0.008</b>	82.4 ± 3.3	80.3 ± 4.2	0.21
Base model (all mice): $p_{\text{genotype}} = \mathbf{0.001}^\dagger$ , $p_{\text{sex}} = \mathbf{0.014}^\dagger$ , $p_{\text{injury}} < \mathbf{0.0001}^\dagger$							Base model (all mice): $p_{\text{genotype}} = \mathbf{0.047}^\dagger$ , $p_{\text{sex}} = \mathbf{0.008}^{\text{at}}$		
Interaction model (all mice): $p_{\text{interaction}} = \mathbf{0.022}^\dagger$							Interaction model (all mice): $p_{\text{interaction}} < \mathbf{0.0001}^\dagger$		
Contralateral (right) hemisphere volume									
<i>Stratified by sex</i>									
Male	131.7 ± 9.1	121.2 ± 9.0	<b>0.013</b>	120.2 ± 9.8	122.0 ± 8.1	0.62	91.2 ± 7.4	100.1 ± 6.7	<b>0.004*</b>
Female	130.1 ± 9.1	126.8 ± 5.9	0.31	127.0 ± 5.8	121.8 ± 7.2	0.07	97.6 ± 4.5	96.0 ± 5.6	0.47
Base model (all mice): $p_{\text{genotype}} = \mathbf{0.019}^\dagger$ , $p_{\text{sex}} = 0.11$ , $p_{\text{injury}} = \mathbf{0.009}^\dagger$							Base model (all mice): $p_{\text{genotype}} = \mathbf{0.046}^\dagger$ , $p_{\text{sex}} = 0.70$		
Interaction model (all mice): $p_{\text{interaction}} = \mathbf{0.037}^\dagger$							Interaction model (all mice): $p_{\text{interaction}} = \mathbf{0.004}^\dagger$		

\*Represents statistical significance in sex- and injury-stratified models at  $p = 0.0125$ . <sup>†</sup>Represents statistical significance in base or interaction model at  $p = 0.05$ .

SD, standard deviation.

Assay System (Promega). Values of luciferase activity with SRY were divided by those from empty vector transfection to obtain fold-induction.<sup>38</sup>

**Electrophoretic mobility shift assay**

We measured the effect of SRY on the  $K_d$  value of binding of Sp1 to the 22-bp-long cis-regulatory element by electrophoretic mobility shift assay (EMSA; Supplementary Methods).<sup>32</sup> To measure DNA binding activity of Sp1 to the putative binding sites in the *Abcc8* promoter region, we used a biotinylated double-stranded DNA oligonucleotide probe containing the Sp1 binding sites (5'-GTGGGGGCGGGGCGGGGCGGGCCT-3'). Lysates from COS-7 cells transfected with two expression vectors driving expression of SRY and Sp1 or Sp1 and control empty vector were incubated with the biotinylated DNA fragments for 20 min at 25°C. To examine the DNA binding activity of Sp1 in high salt concentration, additional sodium chloride was added to the reaction mixture as indicated. Biotinylated DNA oligonucleotides on the membrane were detected by a chemiluminescent nucleic acid detection module kit (ThermoFisherScientific, Waltham, MA), and the images were processed by using commercial imaging software (FUJIFILM Life Science, Cambridge, MA).

**RNA isolation and real-time polymerase chain reaction**

Determination of the relative abundance of *Abcc8* messenger RNA (mRNA) in bEnd.3 cells transfected with either empty vector or SRY was performed as previously described (Supplementary Methods).<sup>31</sup> Total RNA was extracted using TRIzol Reagent (Invitrogen). RNA was

further purified with Amplification Grade DNase I (Invitrogen). Complementary DNA was synthesized from 1 µg of total RNA of each sample using SuperScript III Reverse Transcriptase (Invitrogen). Abundance of *Abcc8* mRNA in samples was determined by real-time PCR (ABI PRISM 7300; Applied Biosystems, Carlsbad, CA) and normalized to *S18*. *Fral* and vascular endothelial growth factor (*Vegf*) were used as positive and negative controls, respectively, of SRY-driven expression.

**Statistical analysis**

This study was powered (0.8) to detect a difference in CV between wild-type and knockout mice in analyses stratified by sex with  $n = 12$  mice per group for each sex and genotype (Supplementary Methods). An additional 12 heterozygous mice were used for secondary analyses of how the number of alleles knocked out affects outcome. An equal number of naïve mice of the same sex and genotypes and similar age (12–15 weeks) were tested. Data are presented as mean ± standard deviation (SD). Associations between sex, genotype, and histological lesions were assessed with multi-variable linear regression models (Supplementary Methods). Primary models included two-level genotypes (wild-type and knockout), sex, and injury status (for LHV, RHV, and TBV) or only injured mice (for CV). Secondary analyses for the association between number of alleles knocked out and histological outcome (including heterozygous mice) were performed using a Wilcoxon-type non-parametric test of trend across genotypes.<sup>39</sup>

Use of regression models was *a priori* selected over multiple independent *t*-tests. Our large sample size of

142 mice allowed for the use of regression models including sex and genotype as covariates to address the research questions of interest without model overfitting. Alpha was set at 0.05 for initial regression Wald tests for genotype, 0.025 for *post hoc* analyses stratified by sex, and 0.0125 for *post hoc* analyses stratified by both sex and injury (Bonferroni). All analysis was performed with Stata software (version 16.1; StataCorp LP, College Station, TX).

## Results

One hundred forty mice were included in the final analysis: Of 72 injured mice, 2 died after injury (1 female wild type, 1 male heterozygote), resulting in  $n=11-12$  per group. For naïve mice, breeding limitations resulted in only 10 obtainable male wild types, for a total of 70 naïve mice analyzed. Males weighed more than females ( $31.5 \pm 3.0$  vs.  $23.2 \pm 2.5$  g;  $p < 0.001$ ). Body weight did not differ by genotype in either sex.

### Effect of sex on neuropathology

In wild-type mice, males and females had similar absolute contusion, hemisphere, and brain volumes post-injury (all  $p > 0.05$ ). However, wild-type females were protected from post-injury atrophy versus wild-type males: They had higher post-injury nTBV ( $90.1 \pm 3.7\%$  vs.  $82.4 \pm 5.8\%$ ;  $p = 0.001$ ), nRHV ( $97.6 \pm 4.4\%$  vs.  $91.2 \pm 7.4\%$ ;  $p = 0.02$ ), and nLHV ( $82.4 \pm 3.3\%$  vs.  $73.5 \pm 5.3\%$ ;  $p = 0.0001$ ).

### Sex, genotype, and total brain volume

Male, but not female, *Abcc8* knockout mice were protected from post-injury atrophy/TBV loss with increased protection for each *Abcc8* allele knocked out (Fig. 1A; Table 1).

In all mice, a genotype-by-sex-by-injury interaction suggests that relationships between TBV and

these variables are interdependent ( $p_{\text{interaction}} = 0.023$ ;  $p_{\text{interaction\_normalized}} = 0.001$ ). In uninjured mice, male, but not female, *Abcc8* knockouts had smaller brains/lower TBV versus wild types ( $p = 0.011$ ; Table 1). However, male *Abcc8* knockout mice were also protected from post-injury whole-brain atrophy (higher nTBV) versus wild type ( $p = 0.001$ ; Table 1; Fig. 1A-ii). This protection was almost 2-fold—total atrophy in male wild-type mice was 17.6% of naïve TBV, whereas it was only 8.9% in male knockout mice (100% nTBV). Knockout did not protect females. In the secondary analyses including heterozygotes, there was increased post-injury sparing of nTBV in males for each copy of *Abcc8* lost ( $p = 0.002$ ; Fig. 1A-ii). Taken together, these results suggest that the role of *Abcc8* in determining TBV both before and after injury depends on the number of alleles knocked out.

### Sex, genotype, and hemisphere volumes

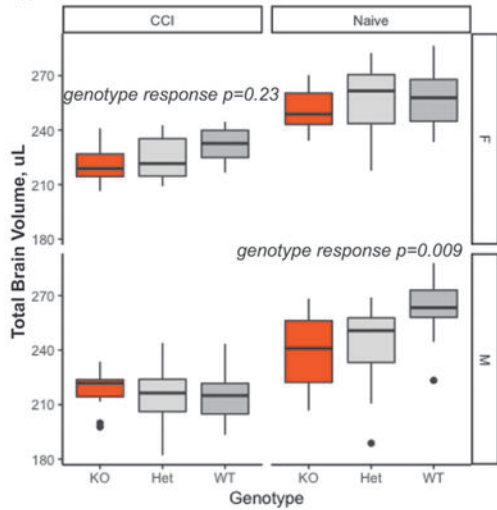
Male, but not female, *Abcc8* knockouts were protected from global post-injury atrophy in *both* left (injured) and right (contralateral) hemispheres (Table 1). Again, the effect was dependent on the number of alleles knocked out (Fig. 1B,C).

In the left (ipsilateral) hemisphere, a genotype-by-sex-by-injury interaction indicates that the association between LHV and these three variables are interdependent ( $p_{\text{interaction}} = 0.022$ ,  $p_{\text{interaction\_normalized}} < 0.0001$ ; Table 1). As with TBV, the association between genotype, sex, and LHV differed before versus after injury: In naïves, LHV was smaller in male knockouts versus wild types, but similar across genotypes in females (Table 1; Fig. 1B-i). After injury, LHV was lower in female knockouts versus wild types ( $p = 0.008$ ), but not in males (Table 1). Post-injury benefit of *Abcc8* knockout was exclusively observed in males: nLHV was higher in male knockouts

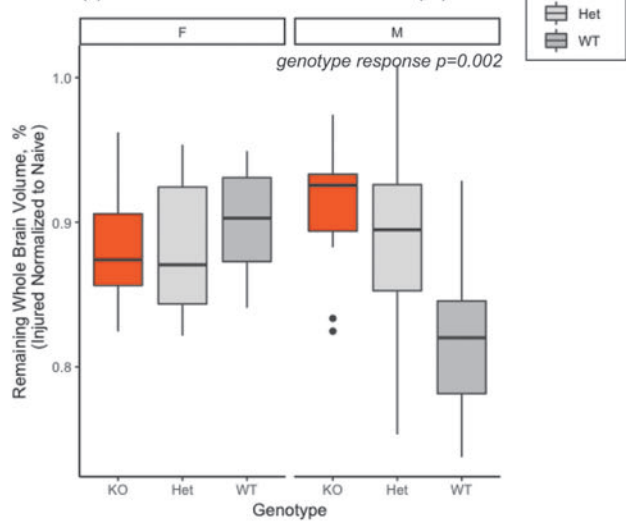
**FIG. 1.** Genotype effects of *Abcc8* knockout on total brain and hemispheric volumes differ by sex. **(A-i)** There was a significantly lower total brain volume for each copy of *Abcc8* lost (wild type: dark gray; heterozygote: light gray; knockout: red) in male naïve mice ( $p < 0.0125$ ), **(A-ii)** whereas after injury there was less brain volume lost after injury for each copy of *Abcc8* knocked out (wild type: dark gray; heterozygote: light gray; knockout: red;  $p < 0.025$ ). **(B-i)** There was a significant decrease in left hemisphere volume for each copy of *Abcc8* lost in naïve male mice (wild type: dark gray; heterozygote: light gray; knockout: red) in male naïve mice ( $p < 0.0125$ ), **(B-ii)** whereas after injury there was less diffuse hemisphere loss after injury for each copy of *Abcc8* knocked out (wild type: dark gray; heterozygote: light gray; knockout: red;  $p < 0.025$ ). **(C-i)** There was no significant effect of sequential *Abcc8* allele knockout on right hemisphere volume in naïve mice (wild type: dark gray; heterozygote: light gray; knockout: red;  $p > 0.0125$ ), **(C-ii)** whereas after injury there was less diffuse hemisphere loss after injury for each copy of *Abcc8* knocked out in male mice (wild type: dark gray; heterozygote: light gray; knockout: red;  $p < 0.025$ ). *Abcc8*, ATP-binding cassette transporter subfamily C member 8 (SUR1); CCI, controlled cortical impact; Het, heterozygote; KO, knockout; WT, wild type.

### A Genotype Response of *Abcc8* in Total Brain Volume (Both Hemispheres) by Sex

(i) Absolute Volume in Naive vs CCI

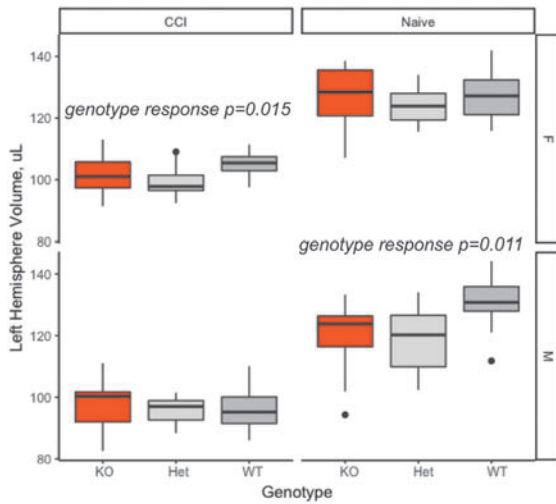


(ii) CCI Volume Normalized to Naive (%)

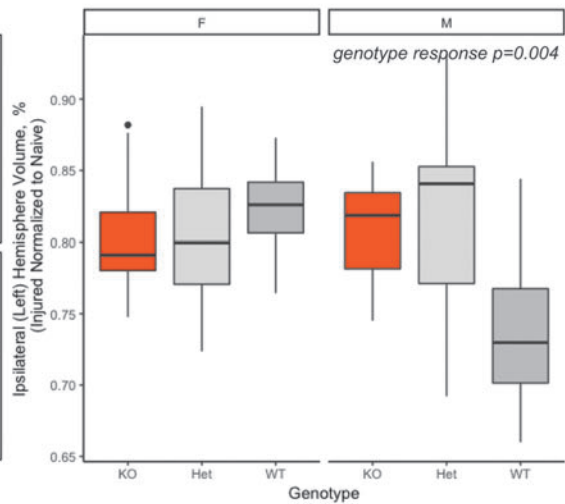


### B Genotype Response of *Abcc8* in Ipsilateral (Left) Hemisphere Volume by Sex

(i) Absolute Volume in Naive vs CCI

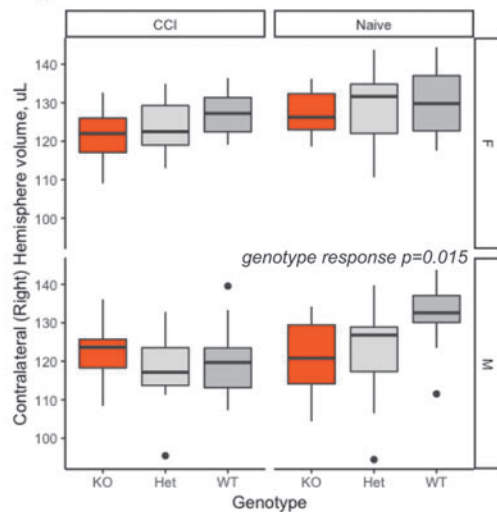


(ii) CCI Volume Normalized to Naive (%)

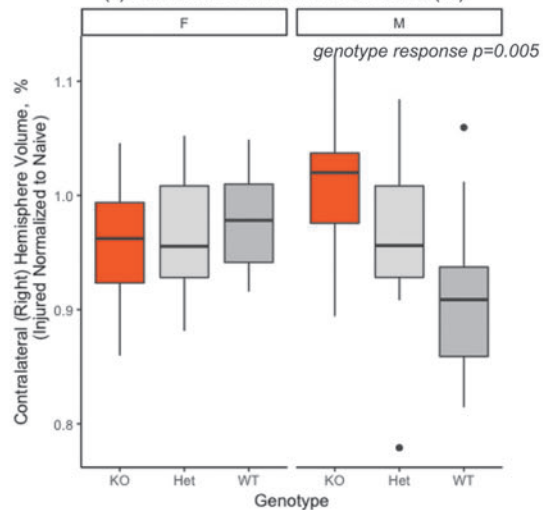


### C Genotype Response of *Abcc8* in Contralateral (Right) Hemisphere Volume by Sex

(i) Absolute Volume in Naive vs CCI



(ii) CCI Volume Normalized to Naive (%)



versus wild types ( $p=0.0004$ ), but similar in females across genotypes, denoting a male-specific protection (Table 1; Fig. 1B-ii). When heterozygotes were included, there was a significant loss of LHV in male naïve mice ( $p=0.011$ ) and increased protection against atrophy in males after injury for each *Abcc8* allele lost ( $p=0.004$ ; Fig. 1B-ii).

In the contralateral (right) hemisphere, a genotype-by-sex-by-injury interaction model again suggested an interdependence between these three variables in their effects on RHV ( $p_{\text{interaction}}=0.037$ ;  $p_{\text{interaction\_normalized}}=0.004$ ). In naïves, male knockouts had lower RHV ( $p=0.013$ ), though it was not statistically significant after adjustment for multiple comparisons. This was not noted in females (Fig. 1C-i). nRHV was markedly higher in male *Abcc8* knockouts versus male wild types, with essentially no post-injury volume loss (Fig. 1C-ii). nRHV did not differ by genotype in females. The decrease in RHV in naïve males was dependent on the number of alleles knocked out ( $p=0.015$ ; Fig. 1C-i), and nRHV loss decreased (i.e., less right hemisphere atrophy) in injured males for each copy of *Abcc8* lost ( $p=0.005$ ; Fig. 1C-ii).

### Sex, genotype, and contusion volume

CV was lower in knockout versus wild-type mice (Table 2, base model). *Abcc8* knockout was protective in male ( $17.7 \pm 3.7$  vs.  $15.9 \pm 3.6$  mm<sup>3</sup>;  $p=0.02$ ), but not female, mice (Fig. 2B-i). There was no genotype-by-sex interaction (Table 2). There was no difference in nCV between wild types and knockouts of either sex (Fig. 2B-ii).

### Sex-determining region Y-mediated ATP-binding cassette transporter subfamily C member 8 expression

In COS-7 cells luciferase activity driven by the *Abcc8* promoter was markedly increased (up to 3.5-fold) in response to SRY coexpression (Fig. 3A). Serial deletion of the *Abcc8* promoter identified a 22-bp cis-regulatory element (−100/−79) that when deleted caused a marked reduction in basal *Abcc8* expression and completely abrogated SRY-mediated stimulation of *Abcc8* expression (Fig. 3B). Sequence analysis of this region revealed no binding sites for SRY, but three binding sites for the transcription factor, Sp1. This suggests that SRY likely activates *Abcc8* transcription indirectly rather than direct DNA binding. In an EMSA, coexpression of SRY with Sp1 increased the affinity of Sp1 to its binding sites ~2- to 3-fold and stabilized the Sp1/DNA complex versus Sp1 alone across a wide range of NaCl concentrations (Fig. 3C). This suggests that SRY may increase Sp1-mediated *Abcc8* transcription. When the effect of SRY on *Abcc8* expression was evaluated in two independent cell lines originating from brain microvascular endo-

thelium (i.e., overexpression of SRY in bEnd.3 cells), an ~6-fold increase in *Abcc8* mRNA was observed versus empty vector (Fig. 3D).

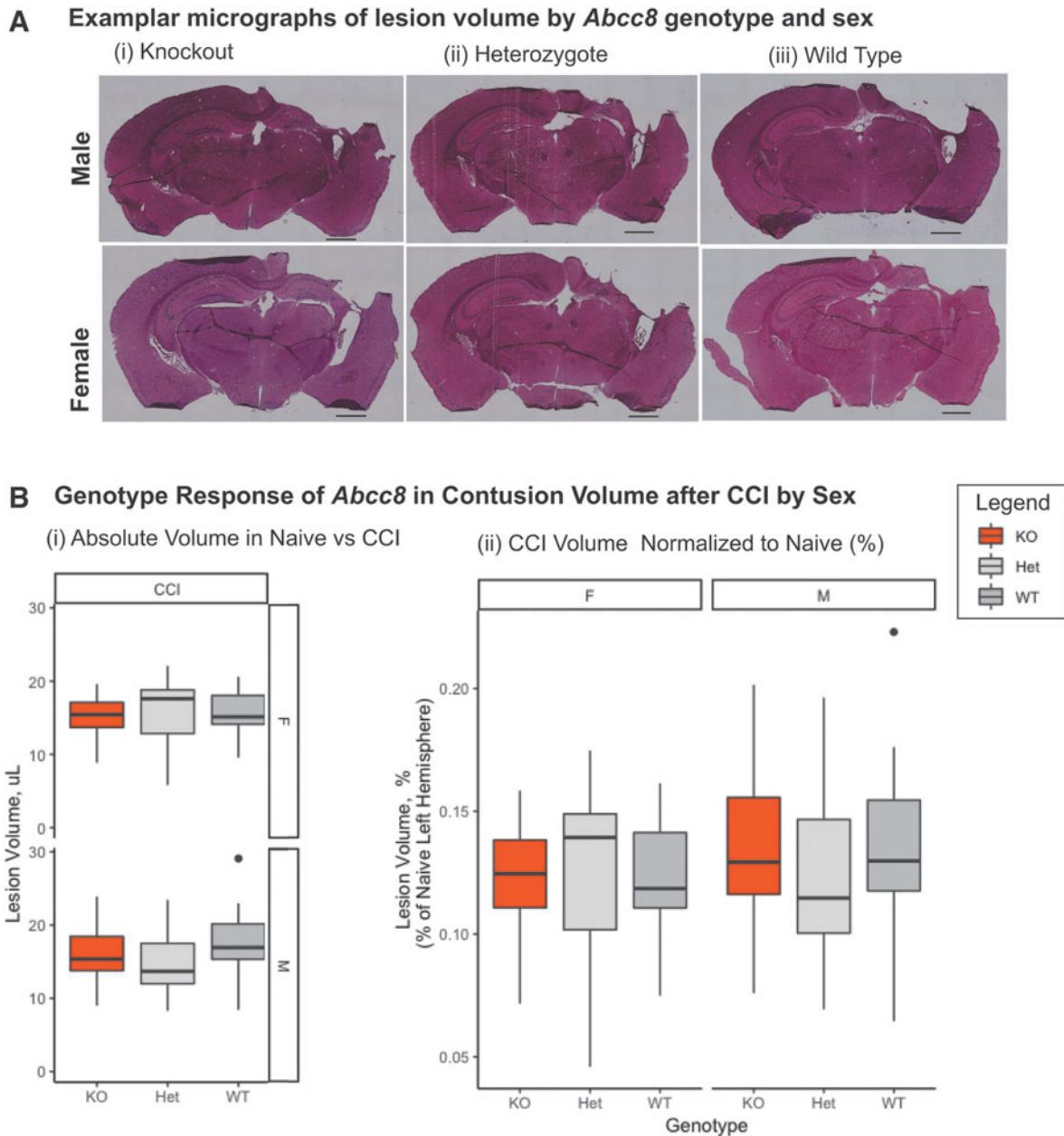
### Discussion

In this study, genetic knockout of *Abcc8* (SUR1) yielded smaller contusions and conferred a robust protection from brain tissue atrophy after experimental TBI in male, but not female, mice. Protection appeared dependent on the number of alleles knocked out. Our *in vitro* studies suggest that this sex difference may partly be related to Y-chromosome gene SRY-mediated stimulation of *Abcc8* transcription by Sp1. To our knowledge, this is the first study to identify sex differences associated with SUR1 inhibition/absence after TBI.

### Sex and traumatic brain injury

Supported by both clinical and pre-clinical studies, it is widely acknowledged that biological sex may moderate pathophysiology and outcome after TBI.<sup>22,40–44</sup> Effects are complex and vary based on mechanism, severity, and measured outcome.<sup>22</sup> Nevertheless, females have been understudied in pre-clinical and clinical TBI, despite distinct biology and worse clinical outcomes versus males.<sup>22</sup> Previous studies, specifically in the CCI model, have found inconsistent effects of sex on CV<sup>23,40,45</sup>. Some report smaller lesions in females<sup>23,40</sup> whereas others identify no difference.<sup>45</sup> Although we found no impact of sex on 21-day CV, total post-traumatic tissue loss was lower in wild-type females versus males.

Sex-based differences after TBI may be mediated by both hormonal and non-hormonal mechanisms.<sup>46</sup> Several studies have implicated disruption of normal sex hormone regulation as a key prognostic factor after TBI.<sup>47,48</sup> Similarly, pre-clinical data suggest that ovariectomized rats may have phenotypic similarities to male rats in development of secondary injury cascades and response to therapy.<sup>49–51</sup> There is also increasing evidence that estrous cycle phase may be associated with outcome after TBI, with better outcomes in the follicular phase (potentially dependent on follicular phase progesterone level).<sup>52–55</sup> The sex hormone, progesterone, has been extensively studied as a neuroprotectant, although randomized clinical trials were ultimately negative.<sup>10,23,56–58</sup> However, controversy over dosing and other details of those trials remain. Non-hormonal mechanisms that may mediate sex differences after TBI include sex-based differences in the activation of neuronal death pathways,<sup>49,59</sup> differences in excitotoxicity,<sup>60,61</sup> different antioxidant defense mechanisms,<sup>59</sup> differences in autophagy,<sup>62</sup> and differential expression of neurotrophic factors.<sup>63</sup> There may be additional complex differences related to both sex and gender in mechanism of injury, circumstances surrounding injury, and medical comorbidities.<sup>64</sup>



**FIG. 2.** Genotype effects of *Abcc8* knockout on contusion volume after injury differ by genotype and sex. **(A)** Representative photomicrographs of post-injury hemispheres from median mouse in each group. Top row from left to right: male knockout, male heterozygote, and male wild type. Second row from left to right: female knockout, female heterozygote, and female knockout. Scale bar is 1000  $\mu$ m. **(B)** There was no relationship between number of alleles knocked out and contusion volume for either sex, either before **(i)** or after **(ii)** normalization. *Abcc8*, ATP-binding cassette transporter subfamily C member 8; CCI, controlled cortical impact; Het, heterozygote; KO, knockout; WT, wild type.

**Sex, sulfonylurea receptor 1, and traumatic brain injury**

*Abcc8* knockout in male, but not female, mice reduced CV in our study. A novel ~2-fold reduction in global brain tissue atrophy was also noted in male, but not female, *Abcc8* knockouts after CCI (8.9% vs.

17.6%). This protective effect increased for each additional *Abcc8* allele knocked out.

Although these sex-based differences have not been reported, CV reduction with *Abcc8* knockout is consistent with earlier literature of pharmacological SUR1 inhibition. An early pre-clinical study of glyburide in TBI

**Table 2. Sex and Genotype Effects on Contusion Volume**

	Injured (mm <sup>3</sup> , mean ± SD)			Normalized (% , mean ± SD)		
	Wildtype	Knockout	p value	Wildtype	Knockout	p value
Contusion volume						
Stratified by sex						
Male	17.7 ± 3.7	15.9 ± 3.6	<b>0.020*</b>	13.6 ± 4.1	13.4 ± 3.4	0.91
Female	15.8 ± 3.5	15.2 ± 3.0	0.73	12.4 ± 2.8	12.3 ± 2.4	0.96
Base model (all mice):				Base model (all mice):		
$p_{\text{genotype}} = \mathbf{0.012}^\dagger$ , $p_{\text{sex}} = 0.45$				$p_{\text{genotype}} = 0.25$ , $p_{\text{sex}} = 0.64$		
Interaction model (all mice):				Interaction model (all mice):		
$p_{\text{interaction}} = 0.26$				$p_{\text{interaction}} = 0.95$		

\*Represents statistical significance in sex and injury stratified models at  $p = 0.025$ .

†Represents statistical significance in base or interaction model at  $p = 0.05$ .

SD, standard deviation.

reported an ~50% reduction in 24-h CV.<sup>8</sup> A more modest reduction (~30%) persisted at 21 days in a blinded, rigorous multi-center pre-clinical study in male rat CCI recently reported by the Operation Brain Trauma Therapy (OBTT) consortium.<sup>7</sup> Of the 12 drugs tested to date by OBTT, only glyburide decreased 21-day CV after CCI.<sup>7</sup> Surprisingly, reductions in male CV observed with glyburide were larger than those noted in our study of genetic SUR1 knockout,<sup>1,7</sup> albeit in rats versus mice and at different post-injury time points. Nonetheless, this suggests complex effects of constitutive, complete, and global SUR1 knockout or, potentially, off-target beneficial effects of glyburide. SUR1 overexpression has also been observed in predominantly male (81%) human traumatic contusions, the corresponding injury pattern to CCI.<sup>28</sup> Two placebo-controlled randomized trials (with ~87% and 72% male patients, respectively) have demonstrated reduced contusion expansion with glyburide.<sup>3,4</sup> The ongoing multi-center phase II ASTRAL trial of glyburide is specifically enrolling patients with contusional TBI (NCT03954041).

Although the protective effect observed in the contusion was expected, the robust protection against diffuse atrophy has not previously been described. It is consistent with a small randomized clinical trial demonstrating improved functional outcome in diffuse axonal injury<sup>27</sup>—of note, this study had only 2 females (out of 40 patients). Contralateral damage after CCI has been previously described and is associated with neuroinflammation or Wallerian degeneration, among other mechanisms.<sup>65,66</sup> How SUR1 contributes to extracontusional brain atrophy remains to be defined. However, a greater global tissue sparing effect by knockout vs. glyburide therapy might reflect glyburide's limited BBB permeability.

A pre-clinical study of trigeminal neuralgia reported sex-based differences in SUR1-mediated pathology: SUR1 activation by selective-agonist diazoxide attenuated capsaicin-induced mechanical hypersensitivity in male, but not female, rats.<sup>26</sup> Despite this, mechanisms underlying effects of sex on SUR1-mediated secondary

injury are unknown. Our *in vitro* findings suggest that the SRY gene may stimulate Sp1-mediated *Abcc8* expression. If *Abcc8* expression is greater in males, then this could contribute to preferential protection of SUR1 knockout in males. Surprisingly, in naïve male mice, *Abcc8* knockouts had lower brain volumes. Although the neurodevelopmental importance of SUR1 remains undefined, our data suggest a teleological role for SUR1 in normal central nervous system (CNS) development (particularly in males) either directly or indirectly by systemic effects on metabolic function and growth.<sup>25</sup>

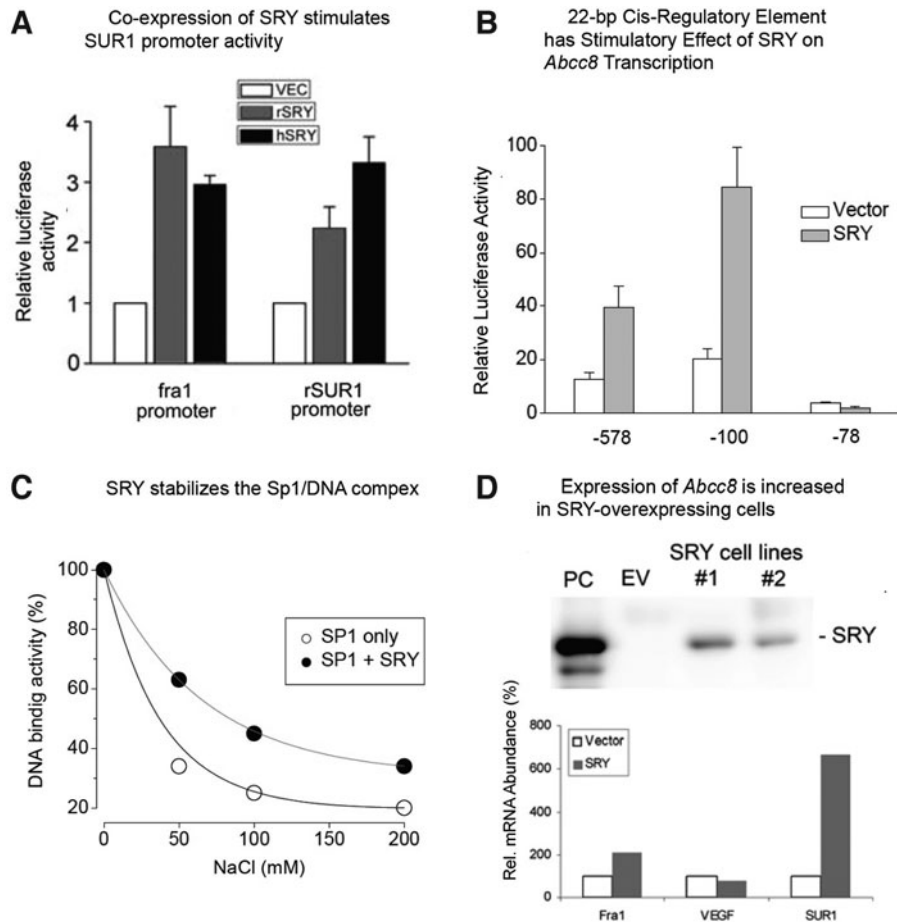
Research on SUR1 in TBI has been exclusively (pre-clinical) or predominantly (clinical) conducted in males.<sup>1-4,7,8,27,28</sup> This is suboptimal for a variety of reasons, including potential biological sex-based differences in the role of this pathway post-injury with ensuing disparate responses to targeted inhibition. Although the strong neuropathological protection in males is highly encouraging for ongoing and future trials of SUR1 inhibition in contusional TBI, our finding that females were not protected may be important to optimize and inform trial design/analysis and future targeted therapy. Further work evaluating mechanistic differences in this pathway between the sexes, as well as comparing global versus conditional genetic inhibition versus pharmacological inhibition, may valuably identify potential off-target effects as well as the role of dose/timing in males versus females.

### Strengths and limitations

This is one of the largest pre-clinical studies of sex-based differences in TBI, with findings that potentially inform ongoing and future clinical trials of SUR1 inhibition. Our large sample size allowed for a detailed analysis of both sex and genotype effects and inclusion of naïve mice of both sexes, and all three genotypes allowed for appropriate normalization of baseline tissue volumes and assessment of non-injury-related effects of SUR1, which have not previously been reported. Our injury model specifically mimics human contusional TBI and is relevant to the population being enrolled in ASTRAL, and both the model and the neuropathological outcomes are well established. Blinding research technicians to sex and genotype decreased risk of bias.

Our study also has several limitations. Although genetic knockout allowed us to assess complete SUR1 inhibition, this does not necessarily reflect pharmacological inhibition, which may produce incomplete inhibition and/or off-target effects. Pharmacological inhibition can also be modulated with regard to dose and timing. Surprisingly, our findings suggest that genetic inhibition may underestimate potential efficacy, at least on CV.<sup>1,7</sup> Global *Abcc8* knockout has consequences outside the CNS, particularly on glucose metabolism and the cardiovascular system<sup>25,67</sup>; this study does not parse out CNS





**FIG. 3.** Sex-determining region Y (SRY) expression may stabilize Sp1 (specificity protein 1) and increase *Abcc8* expression in males. **(A)** Coexpression of SRY stimulates SUR1 promoter activity. A luciferase reporter construct containing an ~4-kb promoter region of rat *Abcc8* was cotransfected into COS-7 cells along with plasmids directing expression of rat SRY (gray), human SRY (black), or vector alone (white). A luciferase reporter construct containing a promoter region of the rat FOS-like antigen 1 (*Fra1*) gene, which is regulated by SRY, was used as a positive control. **(B)** A 22-bp-long cis-regulatory element (–100/–79) plays a critical role in the stimulatory effect of SRY on transcription of *Abcc8*. A luciferase reporter construct containing –578/+80, –100/+80, or –78/+80 of the promoter region of the rat *Abcc8* gene was transfected together with empty vector (Vector) or expression plasmid with SRY (SRY), and luciferase activities were measured. Each luciferase activity is represented as relative to the luciferase activity driven by a  $\beta$ -actin promoter ( $n=5$ ). **(C)** SRY stabilizes the Sp1/DNA complex. The biotinylated 22-bp-long cis-regulatory element (–100/–79 region) was used as a probe in an electrophoretic mobility shift assay. DNA binding activities of lysate from COS7 cells transfected with Sp1 plasmid alone or Sp1 and SRY expression plasmids together were compared at the indicated salt concentrations. **(D-i)** Stable expression of SRY was validated in two independent cell lines (#1 and #2; PC indicates positive control). **(D-ii)** Comparison of mRNA abundance for *Fra1* (positive control), vascular endothelial growth factor (VEGF; negative control), and *SUR1* in control (white) and SRY-overexpressing (gray) bEnd.3 cells. *Abcc8*, ATP-binding cassette transporter subfamily C member 8.

versus non-CNS effects. Global *Abcc8* knockouts also indiscriminately inhibit all SUR1-regulated channels, including constitutively expressed potassium channels in the CNS.<sup>68</sup> Additionally, our study is limited by not having tracked the estrous phase of female mice, with increasing pre-clinical and clinical evidence that the estrous cycle and endogenous female sex hormones may play an important role in modulating secondary injury after TBI.<sup>52–55,69,70</sup>

Our approach, however, mimicked clinical practice where injury can occur at any point in the cycle, and resultant endocrine dysregulation may impede clinical assessment of cycle phase after severe injury.<sup>48</sup> Nevertheless, the impact of estrous phase on response to this pathway should be assessed in future studies, particularly those testing glibenclamide therapy, given its use in current clinical trials. Further, investigating the impact of the estrous phase on this pathway in future work may help identify novel and potentially intervenable targets involved in lesion volume and brain atrophy after TBI. Given the large sample size required to appropriately power the study and normalize appropriately across both sex and genotype, our specific focus on neurohistopathological effects, and a desire to promptly generate findings on this important issue (particularly in the context of the radiographical end-point of the ongoing ASTRAL study), behavioral testing was beyond the current scope and was not performed. However, behavioral performance is a key metric for therapy translatability, and previous studies have demonstrated sex-specific responses to therapy in behavioral outcomes.<sup>71–74</sup>

Follow-up pre-clinical work investigating effects of sex differences in this pathway on behavioral outcome may importantly guide upcoming phase 3 clinical trial design to evaluate potentially disparate effects of SUR1 inhibition in male and female patients (sample-size calculations, analysis plans, etc.). Similarly, our study is limited by studying mechanisms only *in vitro*, and not assessing whether *Abcc8* expression was increased *in vivo* in these mice. These studies will be critical for a better understanding of the potential translational significance of the observed sex-based differences and should be performed in the future.

## Conclusion

*Abcc8* knockout reduced contusion volume and protected against global brain-tissue atrophy after experimental moderate-severe TBI: benefit was restricted to males, with increasing protection for each additional *Abcc8* allele knocked out. Studies exploring sex-dependent effects of pharmacological SUR1 inhibition are warranted—these could have important implications for analysis of the ongoing ASTRAL trial evaluating contusion expansion, as well as design of future studies testing SUR1 inhibition in TBI, and beyond.

## Authors' Contributions

S.T.: Conception and design of the study, acquisition and analysis of data, drafting a significant portion of the manuscript, substantive revision of manuscript, and final approval before submission. B.E.Z.: Conception and design of the study, acquisition and analysis of data, drafting a significant portion of the manuscript, substantive revision of manuscript, and final approval before submission. P.M.K.: Conception and design of the study, drafting a significant portion of the manuscript, substantive revision of the manuscript, and final approval prior to submission. J.M.S.: Conception and design of the study, acquisition and analysis of data, substantive revision of the manuscript, and final approval before submission. V.G.: Conception and design of the study, acquisition and analysis of data, substantive revision of the manuscript, and final approval before submission. M.S.K.: Conception and design of the study, acquisition and analysis of data, substantive revision of the manuscript, and final approval before submission. S.K.W.: Conception and design of the study, acquisition and analysis of data, drafting a significant portion of the manuscript, substantive revision of the manuscript, and final approval before submission. R.S.B.: Drafting a significant portion of the manuscript, substantive revision of the manuscript, and final approval before submission. K.J.F.: Acquisition and analysis of data, substantive revision of the manuscript, and final approval before submission. V.A.V.: Acquisition and analysis of data, substantive revision of the manuscript, and final approval before submission. R.M.J.: Conception and design of the study, acquisition and analysis of data, drafting a significant portion of the manuscript, substantive revision of the manuscript, and final approval before submission.

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## Author Disclosure Statement

R.M.J. is a paid consultant for and on the advisory board for Biogen. J.M.S. holds a US patent (7,285,574), “A novel non-selective cation channel in neural cells and methods for treating brain swelling.” J.M.S. is a member of the board of directors for and holds shares in Remedy Pharmaceuticals and is a paid consultant for Biogen.

## Supplementary Material

Supplementary Methods

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