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RYANODINE RECEPTOR TYPE 1 (RYR1) POSSESSING MALIGNANT HYPERTHERMIA MUTATION R615C EXHIBITS HEIGHTENED SENSITIVITY TO DYSREGULATION BY NONCOPLANAR 2,2',3,5',6-PENTACHLOROBIPHENYL (PCB 95)

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Abstract

Malignant hyperthermia (MH) susceptibility is conferred by inheriting one of >60 missense mutations within the highly regulated microsomal Ca^{2+} channel known as ryanodine receptor type 1 (RyR1). Although MH susceptible patients lack overt clinical signs, a potentially lethal MH syndrome can be triggered by exposure to halogenated alkane anesthetics. This study compares how non-coplanar 2,2',3,5',6-pentachlorobiphenyl (PCB 95), a congener identified in environmental and human samples, alters the binding properties of [^3H]ryanodine to ryanodine receptor type 1 (RyR1) *in vitro*. Junctional sarcoplasmic reticulum (SR) was isolated from skeletal muscle dissected from wild type pigs ($\text{w}_t\text{RyR1}$) and pigs homozygous for MH mutation R615C ($\text{MH}\text{RyR1}$), a mutation also found in humans. Although the level of $\text{w}_t\text{RyR1}$ and $\text{MH}\text{RyR1}$ expression is the same, $\text{MH}\text{RyR1}$ shows heightened sensitivity to activation and altered regulation by physiological cations. We report here that $\text{MH}\text{RyR1}$ shows more pronounced activation by Ca^{2+} , and is less sensitive to channel inhibition by Ca^{2+} and Mg^{2+} , compared to $\text{w}_t\text{RyR1}$. In a buffer containing 100nM free Ca^{2+} , conditions typically found in resting cells, PCB 95 (50–1000nM) enhances the activity of $\text{MH}\text{RyR1}$ whereas it has no detectable effect on $\text{w}_t\text{RyR1}$. PCB 95 (2 μM) decreases channel inhibition by Mg^{2+} to a greater extent in $\text{MH}\text{RyR1}$ (IC_{50} increased 9-fold) compared to $\text{w}_t\text{RyR1}$ (IC_{50} increased by 2.5-fold). PCB95 reduces inhibition by Ca^{2+} 2-fold more with $\text{MH}\text{RyR1}$ than $\text{w}_t\text{RyR1}$. Our data suggest that non-coplanar PCBs are more potent and efficacious toward $\text{MH}\text{RyR1}$, and have more profound effects on its cation regulation. Considering the important roles of Ca^{2+} and Mg^{2+} in regulating Ca^{2+} signals involving RyR channels, these data provide the first mechanistic evidence that a genetic mutation known to confer susceptibility to pharmacological agents also enhances sensitivity to an environmental contaminant.

Keywords

Ryanodine receptor; calcium; malignant hyperthermia; polychlorinated biphenyls; gene-environment interaction.

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Introduction

Ryanodine receptors (RyRs) are large molecular weight proteins that assemble as tetramers and function as high conductance calcium channels that release Ca^{2+} that is stored within the sarcoplasmic/endoplasmic reticulum (SR/ER) of most cells. The large cytoplasmic domain of RyRs physically and/or functionally associate with plasma membrane proteins affording mechanisms that tightly couple extracellular stimuli into localized intracellular calcium signals that amplify and integrate cellular signaling events such as excitation-contraction (E-C), excitation-transcriptional, and excitation-secretion coupling (Meissner 2002; Fill and Copello, 2002; Wamhoff et al., 2006). Moreover, RyRs have been shown to regulate energy metabolism by mitochondria (Hajnoczky et al., 2002). There are three genetic isoforms of RyRs: RyR1, which is responsible for skeletal muscle E-C coupling; RyR2, which is responsible for cardiac muscle E-C coupling; and RyR3, originally characterized in the central nervous system (Pessah et al., 1985; Takeshima et al., 1989; Marks et al., 1989; Zorzato et al., 1990; Nakai et al., 1990; Otsu et al., 1990; Giannini et al., 1995; Hakamata et al., 1992). In recent years, it has become clear that all three RyR isoforms are widely expressed in both excitable and non-excitable cells.

Dysfunction of RyR1 and its closely related cardiac isoform, RyR2, are known to contribute to heritable diseases of muscle. Missense mutations within three “hot spots” of the *ryr1* gene have been linked to malignant hyperthermia (MH) susceptibility, central core disease (CCD), and multi-minicore disease (MmD) (MacLennan et al., 1990; Du et al., 2004; Avila et al., 2001). Interestingly, analogous mutations within the *ryr2* gene have been linked to cardiac arrhythmogenesis, including arrhythmogenic right ventricular dysplasia type 2 (ARVD2), catecholaminergic polymorphic ventricular tachycardia (CPVT) and familial polymorphic ventricular tachycardia (FPVT) (Bers and Perez-Reyes, 1999; Marks et al., 2002; Wang et al., 2004).

MH is an inherited, life-threatening disorder of skeletal muscle in human and animals possessing point mutations within RyR1. The prevalence of the MH genotype in humans is 1 out of 20,000 anesthetized adults (MacLennan and Phillips, 1992). It should be noted though that the incidence of MH mutations in the human population is likely to be much higher since only a fraction of the population undergoes general anesthesia with triggering agents and susceptible individuals do not always trigger with a single anesthetic. Patients expressing RyR1 with an MH mutation (MH RyR1) show no overt clinical signs of disease. However when exposed to halogenated inhalation anesthetics and/or depolarizing blockers of skeletal muscle contraction, individuals can undergo a fulminant MH episode characterized by a striking increase in end tidal carbon dioxide, intense production of heat, tachycardia, unstable hemodynamics, lactic acidosis, and skeletal muscle rigidity (Gronert GA, 1980; Gronert and Antognini, 1994). Thus, MH is a true pharmacogenetic disorder that remains sub-clinical until triggered by a chemical or physical stressor. If symptoms of a clinical MH episode are not promptly treated with administration of the muscle relaxant dantrolene, the fulminant episode is lethal. Dantrolene is therefore the drug of choice to abrogate clinical MH. The mechanism by which dantrolene exerts its relaxing action on muscle contractility is by preventing RyR1 channels from assuming long-lived open states (Paul-Pletzer et al., 2001; Kobayashi et al., 2005). The fulminant MH syndrome results from the uncontrollable release of SR-stored Ca^{2+} through MH RyR1 (MacLennan and Phillips, 1992). A number of studies have shown that MH RyR1 is hypersensitive to Ca^{2+} channel activation and/or hyposensitive to channel inhibition (Lopez et al., 2005; Yang et al., 2003; Louis et al., 2001; Ording et al., 1997; Sei et al., 2002; Wappler et al., 1999 & 2003; Wehner et al., 2002).

Although MH susceptible patients can be successfully identified using the *In Vitro Contracture Test* (IVCT) (European Malignant Hyperthermia Group, 1984), the underlying mechanisms

that lead to a fulminant MH episode and its severity in patients are still poorly understood (Loke and MacLennan, 1998). Although >60 mutations have been linked to MH susceptibility, only about 50% of MH episodes have been linked to RyR1 mutations. Furthermore, patients with a known MH mutation may not go into a fulminant MH episode on the first or even subsequent exposures to triggering agents (e.g. anesthetics), and they exhibit different degrees of severity in response to the same triggering agent among patients in the same susceptible family (Loke and MacLennan, 1998). This variability in triggering the clinical MH phenotype could reflect the influence of yet unidentified physiological and/or environmental modifiers that allosterically sensitize the RyR1 channel complex toward activation. To examine this hypothesis, we tested if a representative non-coplanar polychlorinated biphenyl, 2,2',3,5',6-pentachlorobiphenyl (PCB 95), differentially influences $_{wt}$ RyR1 and $_{MH}$ RyR1 function *in vitro*.

Previous studies from our laboratory have shown that ortho-substituted PCB congeners potentially affect the mobilization of Ca^{2+} from skeletal SR and brain ER by shifting RyR1 sensitivity to Ca^{2+} and Mg^{2+} through a highly selective and structurally specific interaction with the FKBP12 (FK506-binding protein)/RyR1 complex (Wong and Pessah, 1996 & 1997). The most active PCB congeners toward RyR1 include many of those found at highest concentrations in human tissues including PCB 153 (2,2',4,4',5,5'-hexachloro), 170 (2,2',3,3',4,4',5-heptachloro) and 180 (2,2',3,4,4',5,5'-heptachloro) (Pessah et al., 2006). In the present study, we use PCB 95, one of the most potent RyR1 sensitizers identified to date (Pessah et al., 2006), to test the hypothesis that $_{MH}$ RyR1 is more highly susceptible than its wild type counterpart to dysregulation by non-coplanar PCBs. In addition to its potent RyR1 activity, PCB 95 has been selected for these initial studies since it has been recently reported to constitute 3–4% of the total PCBs found in contaminated San Francisco Bay sediments (Hwang et al., 2006), and has also been detected in several human tissues when analytically resolved (Covaci et al., 2002; Chu et al., 2003; DeCaprio et al., 2005; Jursa et al., 2006). Thus PCB 95 may be more frequently found in human and animal tissues than previously thought since it is often not resolved from PCB 66 (2,3',4,4'-tetrachloro) by GC. For example 4.2% of the total PCB burden in Fox River fish was PCB59+66 (Kostyniak et al., 2005). In a recent study of serum samples drawn from a Native American population, PCB 95 contributed an average of 1% of total serum PCBs detected (DeCaprio et al., 2005). Furthermore, analysis of age-dependent congener profile revealed that PCB 95 were higher (> 2% of total PCBs measured) among the youngest 5% of the study population (age range 18–24) (DeCaprio et al., 2005).

The human $_{MH}$ RyR1 mutation R614C occurs in about 2% of MH susceptible patients, and the homologous mutation (R615C RyR1) occurs in the “Pietrain” pig line. The Pietrain is homozygous for the R615C RyR1 mutation, and was selected for growth and leanness traits; *i.e.*, faster-growing pigs at heavier slaughter weights while providing high quality leaner meat (Band et al., 2005). In the present study, junctional SR membrane vesicles are isolated from skeletal muscle dissected from R615C RyR1 homozygote MH susceptible Pietrain pigs and their wild type breed-matched counterparts. We report for the first time that $_{MH}$ RyR1 exhibits significantly higher sensitivity to a non-coplanar PCB than does wild type RyR1.

Materials and Methods

Materials

—[3H]Ryanodine ([3H]Ry; 60–90 Ci/mmol; >99% pure) was purchased from Perkin-Elmer New England Nuclear (Wilmington, DE). PCB 95 (>99% pure) was purchased from Ultra Scientific (North Kensington, RI). Chemiluminescence chemicals were purchased from PerkinElmer Life Science Products. Unlabeled ryanodine (>99% by HPLC) and leupeptin were purchased from Calbiochem (San Diego, CA). Monoclonal 34C–(RyR1)-1° antibody was obtained from Developmental Studies Hybridoma Bank (Iowa City, IA). Polyvinylidene

difluoride microporous membranes were purchased from Millipore (Bedford, MA). Phenylmethylsulfonylfluoride (PMSF), imidazole, and A-5906-(RyR1)-2° antibody for RyR1 were purchased from Sigma (Saint Louis, MO). All other reagents were of the highest purity commercially available.

Membrane Preparations

— Porcine skeletal muscle junctional SR (JSR) membrane vesicles were isolated from flash-frozen longissimus dorsi muscle from 3 “Pietrain” (R615C-RyR1 homozygous) MH susceptible pigs and 3 “Near Pietrain” MH normal (control) pigs that are wildtype at the *ryr1* locus (Band et al., 2005). The pigs were raised at the University of Minnesota as a standing herd and their genotype assessed by PCR. Pigs were slaughtered by approved methods, and skeletal muscle dissected and frozen immediately. Frozen skeletal muscle was a generous gift of Dr. Bradley R. Fruen, University of Minnesota, and shipped frozen to UC Davis. The junctional SR (JSR) membrane vesicles were prepared from the frozen muscle by a modification of the method of Saito et al., (Saito et al., 1984). Briefly, 110 g of muscle was homogenized in 5-fold (w/v) ice-cold buffer consisting of 5mM imidazole, 300 mM sucrose, 100 µg/ml PMSF, and 10 µg/ml leupeptin, pH 7.4. The homogenate was centrifuged at 8,000 × g for 10 min. The resulting supernatant was collected and centrifuged at 110,000 × g for 1 hr. The crude microsomal pellets were suspended in homogenization buffer with a glass Dounce homogenizer and heavy SR vesicles from this solution further purified by centrifugation at 70,000 × g on a sucrose gradient for 15 hr at 4°C. Junctional SR vesicles enriched in RyR1 Ca²⁺ channel complexes were collected from the 38–45% sucrose interface and centrifuged at 110,000 × g for 1 hr. The pellet was collected and suspended in storage buffer containing 10% sucrose, aliquoted in microfuge tubes, and rapidly frozen in liquid nitrogen and stored at –80°C until needed. Protein concentrations were measured by the method of Lowry using BSA as a standard (Lowry et al., 1951). Porcine JSR membranes were shown enriched in functional RyR1 channels and active Ca²⁺ transport properties (Fill et al., 1990; Mickelson et al., 1988; Ta and Pessah, in preparation).

Western Blot Analysis

—Proteins were denatured in 1:1 in a solution containing 10% mercaptoethanol and 10% sucrose-HEPES buffer for 15 min at 80 °C before being loaded onto a 3–10% gradient sodium dodecyl sulfate-polyacrylamide gel and subjected to electrophoresis at 200 V for 45 min at 4° C. The size-separated proteins were then transferred slowly onto polyvinylidene difluoride microporous membranes using an electroblotter for 16 hr at 30 V followed by a rapid transfer for 1 hr at 150 V. The transferred proteins were then incubated in TTBS buffer (25 mM Tris base, 137 mM NaCl, 0.05% Tween 20, pH 7.4) containing 5% nonfat dry milk at ambient temperature for 30 min. The blot was then probed with 34C primary antibody (1:200 dilution) diluted 200 times in TTBS buffer plus 1% bovine serum albumin at 24 °C. After 1 hr, the blot was rinsed in TTBS buffer three times and then incubated with the secondary antibody (goat anti-rabbit IgG at a 1:20,000 dilution) for 1 hr at 24 °C. After a final rinse step with TTBS, enhanced chemiluminescence techniques were used to visualize the immunoblots.

Measurement of [³H]Ry Binding

—Since [³H]Ry binds with high affinity to an open state of the RyR channel, occupancy of [³H]Ry-binding sites provides a convenient biochemical indicator of modulation of the open probability of the channel (i.e. channel activity) by exogenously added ligands (Pessah et al., 1985, 1987 and 1991). Equilibrium measurements of specific high-affinity [³H]Ry binding and incubation conditions were determined according to the method developed by Pessah et al. (Pessah et al., 1985; Pessah et al., 1987; Needleman and Hamilton, 1997). To determine maximum binding level of _{WT}RyR and _{MH}RyR, an unlabeled ryanodine titration binding

experiment shown in Fig. 1A was performed in high salt buffer containing 1 M KCl, 15 mM NaCl, 20 mM HEPES, 100 μ M CaCl₂, 1 nM [³H]Ry with various concentration of unlabeled ryanodine, incubated for 3 hr at 37°C. The influence of PCB 95 concentration on [³H]Ry binding to RyR1 (Fig. 2) was determined by incubating the SR vesicles in physiological intracellular salt binding buffer (140 mM KCl, 15 mM NaCl, 20 mM HEPES, 100 nM CaCl₂, 1 nM [³H]Ry at pH 7.4) and was incubated for 20 hr at 24°C. Similarly, the effect of PCB 95 on the modulation of ryanodine binding by Ca²⁺ and Mg²⁺ was determined by obtaining the Ca²⁺ activation curves (Fig. 3) and Mg²⁺ inhibition curves (Fig. 4) in the presence of various PCB 95 concentrations added from a 100x stock in dimethylsulfoxide before initiating the binding by addition of radioligand. Free concentration of Ca²⁺ (50nM to 50 μ M) was adjusted by titrating EGTA based on calculations determined using the Bound and Determined (B.A.D.) software program (Brooks and Storey, 1992). Nonspecific binding was assessed by incubating SR vesicles with 1000-fold excess unlabeled ryanodine. The binding reactions were filtered through GF/B glass-fiber filters. The filters were washed twice with 3 ml of ice-cold harvest buffer containing 20 mM Tris, 250 mM KCl, 15 mM NaCl, 50 μ M CaCl₂, pH 7.1 and were soaked in 5 ml of scintillation cocktail overnight. The radioactivity was measured using a scintillation counter (Beckman, model LS 6000IC). Specific [³H]Ry binding was determined by subtracting nonspecific binding from total binding and was greater than 90% under all buffer conditions, and unaffected by PCB 95.

Data Analysis

— The ORIGIN™ computer software (Microcal Software, Inc., Northampton, MA) was employed to analyze [³H]Ry binding data. Non-linear regression for one-site binding (Fig. 1A) and Scatchard analysis (Fig. 1B) were utilized to determine maximum receptor activation (B_{max}) and dissociation constant (K_d) for w_t RyR1 and M_H RyR1. Analysis of the effect of PCB 95 on [³H]Ry receptor binding parameters was performed using non-linear curve fitting of data across the entire dose-response curve using a sigmoidal equation to calculate the concentration needed to obtain 50% of the maximum receptor activity (EC_{50} or IC_{50}). One-way ANOVA was utilized to compare B_{max} , K_d , protein expression, EC_{50} or IC_{50} values for w_t RyR1 and M_H RyR1 to test if a population difference between the two groups were significant and to report P values. Experiments were performed in triplicates and the data are presented as mean $\pm$ standard error of the replicated experiments on muscle prepared from three animals per genotype.

Results

In order to determine if junctional microsomal membrane vesicles isolated from skeletal muscle possessing RyR1 missense mutation R615C were quantitatively different compared to preparations isolated in an identical manner from wild type animals, [³H]Ry receptor binding affinity and density were compared. Fig. 1A shows equilibrium binding curves measured with 25 μ g of junctional microsomal protein from either wild type or MH susceptible preparations. These determinations were made in the presence of high salt which was shown to maximize the binding of [³H]Ry to the high affinity site on RyR1 (1 M KCl, 100 μ M Ca²⁺, 20 mM HEPES, 15 mM NaCl) (Pessah et al., 1987; Pessah and Zimanyi 1991a). [³H]Ry was titrated from 0.25 to 129 nM and the reactions were quenched by rapid filtration after 3 hr incubation at 37°C. Scatchard plot analysis showed that both w_t RyR1 and M_H RyR1 preparations possessed the similar [³H]Ry binding-site density and affinity (B_{max} of 3.5 ± 0.2 and 3.0 ± 0.1 pmol/mg protein ($P < 0.288$), K_d of 2.9 ± 0.3 and 2.1 ± 0.2 nM ($P < 0.039$), respectively, $n = 6$, Fig. 1B). The level of w_t RyR1 and M_H RyR1 expression in the two preparations was confirmed by western blot analysis (Fig. 1C) and densitometric analysis of western blots did not reveal any significant difference in the level of protein expression between the two preparations ($P < 0.85$, Fig. 1D).

Previous studies have shown that MH missense mutations resulted in hypersensitivity of RyR1 channels to channel activation by calcium, 4-chloro-*m*-cresol, and caffeine (Ording et al., 1997; Sei et al., 2002; Wappler et al., 1999 & 2003; Wehner et al., 2002; Yang 2003). In addition, a recent study by Lopez et al. showed that expression of several human MH mutations in myotubes result in persistent elevation of resting cytoplasmic Ca^{2+} (Lopez et al., 2005). In the present study, we compared the effect of PCB 95 on MH RyR1 function with that of wt RyR1 under conditions that mimic the cytoplasmic Ca^{2+} concentration in resting cells. These experiments were performed under more physiological salt conditions (140 mM KCl, 15 mM NaCl, 20 mM HEPES, 1 nM [^3H]Ry, 100 nM Ca^{2+} , pH 7.4), and incubation was allowed to proceed for 20 hr at 24°C. Under this assay condition, PCB 95 increased the amount of [^3H]Ry bound in a PCB 95 dose-dependent manner, and reflects the activating actions of PCB 95 on overall channel activity (Wong and Pessah, 1996). The EC_{50} for PCB 95 activation of wt RyR1 was >3-fold higher than that of MH RyR1 (EC_{50} s of $3.6 \pm 0.2 \mu\text{M}$ vs. $1.1 \pm 0.1 \mu\text{M}$, respectively, $n = 6$; Fig. 2). Furthermore, Fig. 2, *inset*, highlights the more profound effect of lower concentrations ($\leq 1000 \text{ nM}$) of PCB 95 on activation of MH RyR1 , concentrations that had little detectable effect on wt RyR1 .

RyR1 endogenous modulators such as Ca^{2+} and Mg^{2+} play an important role in regulation of the channel function to ensure the proper cellular signaling (Meissner, 2002). We therefore examined if these physiological cations differentially regulated wt RyR1 and MH RyR1 in the presence and absence of 250 nM PCB 95. Fig. 3A shows the dose-response for Ca^{2+} activation of [^3H]Ry binding and Fig. 3B shows the Logit-log analysis for [^3H]Ry binding in the presence and absence of 250 nM PCB 95. In the absence of 250 nM PCB 95, Fig. 3A shows that EC_{50} s for wt RyR1 and MH RyR1 are significantly different (170.8 ± 2.5 and $217.0 \pm 5.9 \text{ nM}$, respectively, P -value < 0.05) and Logit-log analysis in Fig. 3B shows that MH RyR1 exhibited a 3-fold steeper Logit slope compared with that of wt RyR1 (1.5 ± 0.8 and 0.5 ± 0.2 , respectively, $n = 6$, Table 1A). Similarly, in the presence of 250 nM PCB 95, EC_{50} s for wt RyR1 and MH RyR1 are also significantly different (148.6 ± 7.6 and $171.5 \pm 0.7 \text{ nM}$, respectively, P -value < 0.05) and the Logit slopes for both MH RyR1 and wt RyR1 are steeper compared to their respective controls lacking PCB 95 (3.3 ± 1.1 and 2.5 ± 0.9 , respectively, $n = 6$, Table 1A).

It is important to note that specifically at $1 \mu\text{M}$ [Ca^{2+}] $_{\text{free}}$ (shown in Fig. 3A), MH RyR1 attained a 4-fold higher absolute amount of optimal [^3H]Ry binding compared to wt RyR1 (0.25 ± 0.01 vs. $0.065 \pm 0.011 \text{ pmol/mg protein}$). Interestingly, the presence of PCB 95 increased the binding level of wt RyR1 preparations to that observed with MH RyR1 preparations free of PCB95, whereas PCB 95 further increased optimal binding of MH RyR1 2.5-fold (to $0.62 \pm 0.01 \text{ pmol/mg protein}$). Thus PCB 95 increased the level of [^3H]Ry binding in MH RyR1 preparations by nearly 10-fold compared to wt RyR1 preparations free of PCB 95 at $1 \mu\text{M}$ [Ca^{2+}] $_{\text{free}}$, reflecting its much higher potency toward MH RyR1 . Fig. 3D summarizes the EC_{50} s for Ca^{2+} activation of [^3H]Ry binding in the presence of several PCB 95 concentrations. As PCB 95 concentration is increased, both MH RyR1 and wt RyR1 became more sensitive to activation by Ca^{2+} ; however, the impact on MH RyR1 was more significant than that of wt RyR1 . Although the EC_{50} s for wt RyR1 and MH RyR1 were significantly different from one another in the absence of PCB 95, they diverge significantly as PCB95 concentration is increased (* $P < 0.05$ for PCB 95 $< 250 \text{ nM}$, and ** $P < 0.001$ for PCB 95 $> 250 \text{ nM}$).

We next examined the effect of PCB 95 on the ability of Mg^{2+} to inhibit [^3H]Ry binding to wt RyR1 and MH RyR1 in buffer containing optimal ($100 \mu\text{M}$) Ca^{2+} , mimicking the local Ca^{2+} concentration attained at the cytoplasmic face of RyRs during physiological activation (Perez et al., 1999; Niggli E, 1999; Jaggar et al., 2000). As shown in Fig. 4A, the presence of $2 \mu\text{M}$ PCB 95 significantly increases the amount of [^3H]Ry specific binding to both MH RyR1 and wt RyR1 preparations. Furthermore, in the presence of PCB 95, RyRs became hyp insensitive to inhibition by Mg^{2+} . The Logit slope for Mg^{2+} inhibition of MH RyR1 was

much steeper compared to that of w_t RyR1 (-0.76 ± 0.96 vs. -0.21 ± 0.05 , Table 1B). However, in the presence of $2 \mu\text{M}$ PCB 95, the Logit slopes for both M_H RyR1 and w_t RyR1 were similar (-0.754 ± 0.15 and -0.89 ± 0.091 , respectively, Table 1B.) Importantly, the action of PCB 95 on Mg^{2+} inhibition is significantly more profound on M_H RyR1 than on w_t RyR1 (Fig. 4C). In the presence of $2 \mu\text{M}$ PCB 95, the IC_{50} s for inhibition of w_t RyR1 and M_H RyR1 increased to $737 \pm 59 \mu\text{M}$ ($n = 6$) and $2659 \pm 112 \mu\text{M}$ ($n = 6$), respectively, Table 1B. Fig. 4D summarizes the IC_{50} s for magnesium inhibition in the presence of several PCB 95 concentrations. As PCB 95 concentration increased, RyRs became increasingly hyposensitive to Mg^{2+} inhibition for both M_H RyR1 and w_t RyR1; however, the impact was significantly greater on M_H RyR1 (** $P < 0.001$) than that of w_t RyR1.

Ca^{2+} above $100 \mu\text{M}$ is also known to inhibit RyR1 in a dose-dependent manner (Pessah et al., 1985; 1987). We therefore studied the effect of PCB 95 on the inhibitory potency of Ca^{2+} using equilibrium $[^3\text{H}]\text{Ry}$ binding analysis under the same buffer conditions described above (graphs are not shown). In the absence of PCB 95, Ca^{2+} inhibited both w_t RyR1 and M_H RyR1 with similar potencies (IC_{50} s are $110.7 \pm 10.6 \mu\text{M}$ and $133.9 \pm 26.5 \mu\text{M}$, Table 1C). In the presence of $2 \mu\text{M}$ PCB 95, however, the inhibitory potencies for w_t RyR1 were significantly greater ($P < 0.001$) from that of M_H RyR1 ($1.2 \pm 0.1 \text{ mM}$ and $2.6 \pm 0.2 \text{ mM}$, respectively).

Discussion

MH susceptible individuals remain sub-clinical until triggered by exposure to general anesthetic agents such as halothane, enflurane, isoflurane, desflurane, and sevoflurane, or a depolarizing muscle blocker such as succinylcholine (Gronert and Antognini, 1994). Considering that MH susceptibility is conferred by any one of more than 60 point mutations in RyR1 and requires an environmental trigger to elicit a clinical episode, we tested if M_H RyR1 has a particularly heightened sensitivity to a non-coplanar PCB of environmental relevance, PCB 95. The rationale for testing this hypothesis is based on the fact that MH susceptible humans may not trigger on the first or even after several exposures to triggering agents (e.g. such as inhalation anesthetics), and they exhibit different degrees of severity in response to the same triggering agent among patients in the same susceptible family (Loke and MacLennan, 1998). This variability in phenotype could reflect the influence of exposures to one or more environmental modifiers that allosterically sensitize the RyR1 channel complex toward become more active (*i.e.*, hyperactive). We first chose to examine the possible influence of non-coplanar PCB 95 on M_H RyR1 since: (1) its mechanism of interaction with w_t RyR1 from skeletal and cardiac muscle (Wong and Pessah, 1996; 1997) and brain (Wong et al., 1997) are well characterized, (2) its actions on primary neurons and PC12 cells suggest RyR-mediated mechanisms are involved in its cellular toxicity (Wong et al 2001; Gafni et al., 2004), (3) the availability of breed-matched muscle tissue from MH susceptible and MH normal pigs, and (4) PCB 95, when resolved by gas chromatography, contributes 4–5% of the PCB burden in San Francisco Bay sediments (Hwang et al., 2006), and is commonly detected in human serum (DeCaprio et al., 2005; Jursa et al., 2006), adipose (Covaci et al., 2002), and liver, kidney, muscle, and brain (Chu et al., 2003).

In the present study, we compared the influences of PCB 95, a non-coplanar PCB, on the regulation of M_H RyR1 and w_t RyR1 using $[^3\text{H}]\text{Ry}$ binding analysis, a reliable indicator of channel conformation, and a sensitive and reliable reporter of how physiological (Pessah et al., 1987; Pessah and Zimanyi, 1991b; Fessenden et al., 2000; Feng et al., 2002) and xenobiotic modifiers influence the channel's activity (Buck et al., 1992; Buck et al., 1999; Chen et al., 1999; Feng et al., 1999; Chen et al., 2003). Consistent with previous studies (Mickelson et al., 1988), we found no significant difference in RyR1 protein expression between junctional SR preparations isolated from MH susceptible and wild type muscle, indicating that the mutation does not cause alterations in RyR1 protein transcription, translation and targeting. Therefore,

measurable alteration in aspects of channel regulation of MH RyR1 compared to w_t RyR1 must be due primarily to abnormal regulation of the channel complex possessing the R615C mutation (Yang et al., 2003). In the present study, we found PCB 95 differentially altered the properties MH RyR1 and w_t RyR1, especially regulation conferred by physiologically important cations.

MH RyR1 is more profoundly influenced by PCB 95 than w_t RyR1 in several respects. PCB 95 dose response curves shown in Fig. 2 reveal that MH RyR1 possessing point mutation R615C were significantly more sensitive to activation by PCB 95. In the presence of low PCB 95 concentration (<250 nM), EC_{50} values for activation by Ca^{2+} for w_t RyR1 and MH RyR1 were not different from one another, suggesting that higher PCB 95 concentrations can produce the same level of allosteric activation of w_t RyR1 seen with MH RyR1, and that indeed the sensitizing action of PCB 95 on both genotypes were saturable. However, PCB 95 (<250 nM) significantly enhanced [3H]Ry binding to MH RyR1 but not w_t RyR1, reflecting the higher sensitivity of MH RyR1 channels to activation by PCB 95. Voltage clamp studies of single channels reconstituted in bilayer lipid membranes currently underway in our lab have verified that MH RyR1 respond with a significantly higher open probability than w_t RyR1 when exposed to nanomolar PCB 95 (Pessah et al., in preparation). In the present study, as the concentration of PCB 95 was increased, EC_{50} s for activation of [3H]Ry binding with w_t RyR1 and MH RyR1 were affected; however, the shift in the dose-response curve was more profound with MH RyR1. Therefore, PCB 95 enhanced the sensitivity for achieving a Ca^{2+} -induced open conformation of the channel, measured as an increased binding of [3H]Ry. In this respect, the Ca^{2+} sensitivity of MH RyR1 was more dramatically shifted than w_t RyR1. Interestingly, Gronert et al. reported that physiological or emotional stress, anxiety, excitement, or sudden changes of environmental temperature can also initiate MH in some patients (Gronert et al., 1980). The present results raise a compelling possibility, that exposure to low levels of RyR-active PCBs such as PCB 95 could further sensitize MH susceptible individuals to the adverse effects of general anesthetic agents, depolarizing muscle blockers, or general environmental stresses such as temperature or emotional state to a greater extent than MH normal individuals. Since Ca^{2+} regulation of RyRs is essential for proper function of muscle cells, RyR inhibition by mM of Ca^{2+} and Mg^{2+} act as a protective mechanism to ensure intracellular calcium homeostasis.

The massive increase in cytosolic Ca^{2+} concentration seen in a triggered MH patient is due to inability of cations (Ca^{2+} and Mg^{2+}) to inhibit RyR1 from releasing calcium, especially when intracellular Ca^{2+} concentrations reached an abnormally high level (Lopez et al., 2005; Yang et al., 2003; Louis et al., 2001; Ording et al., 1997; Sei et al., 2002; Wappler et al., 1999 & 2003; Wehner et al., 2002). In the present study, we found that channel inhibition by Ca^{2+} and Mg^{2+} was not affected by low concentration of PCB 95 (<250 nM); however, at a concentration of PCB 95 ($\geq 1 \mu M$), RyRs became hyposensitive to channel inhibition by these cations. IC_{50} for Ca^{2+} inhibition of w_t RyR1 increased by 10-fold compared with its control, whereas IC_{50} for MH RyR1 increased by 20-fold relative to its respective control (without $2 \mu M$ PCB 95). IC_{50} for Mg^{2+} inhibition of w_t RyR1 increased by 2.5-fold compared with its control, whereas IC_{50} for MH RyR1 increased by 9-fold relative to their respective control (without $2 \mu M$ PCB 95). In the presence of $2 \mu M$ PCB 95, IC_{50} values for MH RyR1 are 2-fold higher for calcium inhibition and more than 3-fold higher for magnesium inhibition compared to those of w_t RyR1. Therefore, in the presence of PCB 95, physiological feedback inhibition of the channel open state by Ca^{2+} and Mg^{2+} are greatly affected in both genotypes, but more profoundly influenced in the presence of an MH susceptibility point mutation.

Missense mutations on *ryr1* gene are known to cause about 50% of the clinical MH cases reported (Gronert et al., 2004). However, the fact that *ryr1* mutations are expressed in cells other than muscle raises the possibility that exposure to RyR1-active PCBs and possibly other persistent organic pollutants may elicit a broader toxicological impact in individuals carrying

these mutations. RyR1s are expressed within the central nervous system (Ledbetter et al., 1994; Martin et al., 1998) and in specific immune cell such as B lymphocytes (Sei et al., 1999) and dendritic cells (O'Connell et al., 2002; Goth et al., 2006) where they contribute essential aspects of function. In fact, human MH mutations have been recently shown to increase the sensitivity of B lymphocytes to xenobiotic challenges (Ducreux et al., 2004; Sei et al., 2002). The present findings raise the intriguing question if individuals possess RyR mutations would be at higher risk of adverse outcomes resulting from exposure to non-coplanar PCBs and possibly other persistent organic pollutants.

If the higher sensitivity of MH RyR1 point mutation R615C to dysregulation by PCB 95 shown here extends to their influence on the integrity of RyR-mediated Ca^{2+} signaling could influence functions in a number of organ systems.

The more pronounced effects of PCB 95 on $[^3\text{H}]\text{Ry}$ receptor binding to MH RyR1 compared to that of WT RyR1 is likely to extend to complex mixtures of non-coplanar PCBs known to predominate in several human tissues, including PCBs 153, 170, and 180 (Pessah et al., 2006). The results therefore raise the possibility that individuals who possess MH mutations within RyR1 and have been exposed to non-coplanar PCBs, may be more susceptible their adverse influences, and possibly amplify those of triggering agents. Recent development of knock-in mice that express human MH RyR1 mutations (Chelu et al., 2006; Yang et al., 2006) will permit direct testing of this hypothesis *in vivo*.

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Figure 1.

Determination of protein expression and binding affinity for w_t RyR1 and MH RyR1.

(A) Non-linear regression for [3H]Ry binding at equilibrium measured in high salt buffer (1 M KCl, 20 mM HEPES, 15 mM NaCl, 1 nM [3H]Ry, 3 hrs incubation at 37°C) in the presence of unlabeled ryanodine at various concentrations was employed to determine B_{max} and K_{-1} . (B) Scatchard plots for both w_t RyR1 and MH RyR1. (C) Western Blots of w_t RyR and MH RyR. JSR lane refers to junctional sarcoplasmic reticulum prep obtained from rabbit and is used as a MW marker ($M_r = 565,000$). (D) Densitometric analysis of Western blots. Data are mean $\pm$ standard error of $n = 6$. Non-linear regression and Scatchard plots were obtained as described in Methods.

Figure 2.

PCB 95 enhances $[^3\text{H}]\text{Ry}$ binding to porcine wtRyR1 and MH-RyR1 . The influence of PCB 95 on $[^3\text{H}]\text{Ry}$ binding was tested in the presence of 100 nM free Ca^{2+} in the assay buffer (mimicking the intracellular resting state) as described in Methods. *Inset* shows the higher potency of PCB 95 $< 1\mu\text{M}$ on MH-RyR1 compared with wtRyR1 . Data are mean $\pm$ standard error of $n = 6$. EC_{50}s were obtained by fitting data to a Sigmoidal curve as described in Methods.

Figure 3.

PCB 95 more significantly sensitizes M_H RyR1 to Ca^{2+} activation compared to w_t RyR1.

Calcium activation of [3 H]Ry binding to porcine junctional SR vesicles enriched in w_t RyR1 and M_H RyR1 were performed in a common binding buffer (140 mM KCl, 20 mM HEPES, 15 mM NaCl, 100 μ M $CaCl_2$), 1 nM [3 H]Ry, pH 7.4, 20 hours incubation at 24°C. The free [Ca^{2+}] was adjusted by adding EGTA to the assay buffer based on calculations made with the B.A.D. program. (A) [3 H]Ry specific binding for w_t RyR1 and M_H RyR1 in the presence of 250 nM PCB 95. (B) Logit-log analysis showing the shift in the slope and half-activation constant EC_{50} for Ca^{2+} in the presence of 250 nM PCB 95. (C) Normalized Ca^{2+} activation curves for w_t RyR1 and M_H RyR1 for data in (A) highlighting the rightward shifts in the Ca^{2+} activation in the presence of PCB 95. (D) Summary of EC_{50} values for Ca^{2+} activation measured in the presence of various concentrations of PCB 95. *Statistical analysis:* ANOVA one-way, P-values: * ≤ 0.05 , ** ≤ 0.001 . Data are mean $\pm$ standard error of n = 6.

Figure 4. PCB 95 attenuates the inhibitory effects of Mg^{2+} to a greater extent in MH RyR1 than WT RyR1 . Mg^{2+} inhibition of $[\text{H}^3]\text{Ry}$ binding was measured in an assay buffer containing 140 mM KCl, 20 mM HEPES, 15 mM NaCl, 100 μM CaCl_2 , 1 nM $[\text{H}^3]\text{Ry}$, pH 7.4 and incubated for 20 hours at 24°C. (A) $[\text{H}^3]\text{Ry}$ specific binding for WT RyR1 and MH RyR1 in the presence and absence of 2 μM PCB 95. (B) Logit-log analysis of the data shown in (A). (C) Normalized inhibition curves for Mg^{2+} in the presence and absence of 2 μM PCB 95 highlighting the leftward shifts in inhibition for WT RyR1 and MH RyR1 . (D) Summary of IC_{50} values for Mg^{2+} in the presence of various concentrations of PCB 95. *Statistical analysis:* ANOVA one-way comparison, ** P-values ≤ 0.001 . Data are mean $\pm$ standard error of $n = 6$.

Table 1

(A) EC₅₀s (nM) for activation of wtRyR1 and MH₁RyR1 by Ca²⁺ with and without 250 nM PCB 95. (B) and (C) IC₅₀s (μM) for inhibition of RyR1s by Mg²⁺ and Ca²⁺, respectively, with and without 2 μM PCB 95. EC₅₀s and IC₅₀s were obtained by Sigmodal curve fit. Slopes and R² values were obtained by for Logit-log analysis.

(A)		Calcium Activation	Logit Analysis	
		EC ₅₀ (nM)	Slope	R ²
wtRyR1		170.8 ± 2.5	0.52 ± 0.18	0.85
MH ₁ RyR1		216.9 ± 5.9	1.46 ± 0.81	0.79
wtRyR1 + 250 nM PCB95		148.7 ± 7.6	2.52 ± 0.92	0.85
MH ₁ RyR1 + 250 nM PCB95		171.5 ± 0.7	3.27 ± 1.06	0.87

(B)		Magnesium Inhibition	Logit Analysis	
		IC ₅₀ (μM)	Slope	R ²
wtRyR1		300 ± 82	− 0.22 ± 0.05	0.87
MH ₁ RyR1		287 ± 62	− 0.76 ± 0.09	0.96
wtRyR1 + 2 μM PCB95		737 ± 59	− 0.89 ± 0.09	0.98
MH ₁ RyR1 + 2 μM PCB95		2659 ± 11	− 0.75 ± 0.15	0.89

(C)		Calcium Inhibition	Logit Analysis	
		IC ₅₀ (μM)	Slope	R ²
wtRyR1		110.7 ± 10.6	− 0.16 ± 0.03	0.93
MH ₁ RyR1		133.9 ± 26.5	− 0.12 ± 0.02	0.99
wtRyR1 + 2 μM PCB95		1201.4 ± 111	− 0.79 ± 0.14	0.93
MH ₁ RyR1 + 2 μM PCB95		2560.8 ± 180	− 1.01 ± 0.09	0.98