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Persistence of Reduced Aggression in Vasopressin 1b Receptor Knockout Mice on a More “Wild” Background

Heather K. Caldwell^{1,*} and W. Scott Young III²

¹Department of Biological Sciences and the School of Biomedical Sciences, Kent State University, Kent, OH 44242

²Section on Neural Gene Expression, NIMH, NIH, DHHS, Bethesda, MD 20892

Abstract

It has been previously reported that vasopressin 1b receptor knockout mice (Avpr1b^{-/-}) have reduced levels of aggressive behavior compared to wildtype littermates. However, as the background of the mice were always a mixture of 129/SvJ and C57BL/6, we wanted to determine if the phenotype persisted when our laboratory line was crossed with a wild-derived sub-species of house mice. To this end, we crossed our Avpr1b^{-/-} mice with *Mus musculus castaneus*, one of few sub-species that will breed with laboratory strains. Subsequent F₂ offspring, which were approximately 50% *Mus musculus castaneus*, were tested in a resident-intruder behavioral test to assess aggressive behavior. We found that even on this more “wild” background, Avpr1b^{-/-} continued to demonstrate longer attack latencies and fewer attacks in a resident-intruder test than wildtype littermates. These findings are consistent with previous reports of reduced aggressive behavior in Avpr1b^{-/-} mice and show that the deficit does persist on a different background strain. Further, these findings confirm the importance of the Avpr1b to normal displays of social forms of aggressive behavior.

Keywords

Avpr1b receptor; *M.m. castaneus*; hippocampus

Introduction

The nonapeptide arginine vasopressin (Avp) has been consistently implicated in the regulation of aggressive behavior across species [1-12]. Currently there are two identified receptor subtypes that are centrally expressed: the vasopressin 1a receptor (Avpr1a) [13,14] and the vasopressin 1b receptor (Avpr1b) [15,16]. The Avpr1a has a wide distribution [14,17,18] and has been implicated in the regulation of a variety of behaviors, including aggressive behavior [5,9,19-26]. The Avpr1b, on the other hand, appears to have a somewhat more restricted distribution, but has also been implicated in the regulation of aggression [1,10,11,15,27-30]. Unfortunately, compared to work on the Avpr1a, there is a lack of knowledge about how the Avpr1b regulates aggressive behavior.

One of the hurdles to examining the contribution of the Avpr1b in the regulation of aggressive behavior has been the deficiency of commercially available pharmacological tools. To address this, Wersinger and colleagues generated a mouse line with a genetic disruption of the Avpr1b [10]. The Avpr1b knockout (-/-) mice have reduced levels of social forms of aggressive

*To whom correspondence should be addressed: Heather K. Caldwell, Kent State University, PO Box 5190, 121 Cunningham Hall, Kent, OH 44242. hcaldwel@kent.edu.

behavior and mildly impaired social recognition. Specifically, $Avpr1b^{-/-}$ male mice have longer attack latencies and fewer attacks toward an intruder compared to wildtype controls in neutral arena and resident-intruder tests [10,11]. In a reversed resident-intruder test, where the experimental animals are the intruders, $Avpr1b^{-/-}$ mice will display defensive postures but do not initiate many defensive attacks [11]. While male $Avpr1b^{-/-}$ mice demonstrate deficits in offensive and defensive aggression, female $Avpr1b^{-/-}$ mice have deficits in maternal aggression [11]. So, while there are deficits in forms of aggression that have a “social” component, there is not a global deficit in aggressive behavior as $Avpr1b^{-/-}$ mice have normal predatory aggression, as measured by the time to attack a cricket [11].

One ongoing issue when using laboratory strains of mice is the consistency in the phenotype of the null mutant when it is on a different background. To explore this we examined whether the reduced aggressive behavior that is observed in $Avpr1b^{-/-}$ mice would endure on a more “wild” background. However, getting laboratory strains to mate with more “wild” mice can often be challenging. Fortunately we had access to a population of wild-derived *Mus musculus castaneus* (*M.m. castaneus*) mice that have been reported to have high levels of aggressive behavior and are one of the few wild-derived subspecies that will breed with laboratory strains [31]. We then crossed our $Avpr1b$ line with *M.m. castaneus* to generate a new line that represented a genetic mixture somewhere between the two and measured aggressive behavior.

Methods

Targeted Disruption of the Vasopressin 1b Receptor Gene and PCR Analysis

The generation of the $Avpr1b^{-/-}$ mouse line has been previously described [10]. The offspring were genotyped at weaning using PCR analysis of DNA isolated from tail clips as previously described [10,11]. All experimental procedures were approved by the National Institute of Mental Health Animal Care and Use Committee, and followed the NIH guidelines “Using Animals in Intramural Research.”

Generation of the “wild” line of $Avpr1b$ mice

The new “wild” line of mice was generated by breeding $Avpr1b^{-/-}$ females, on a mixed background of C57BL/6J and 129X1/SvJ generated by W.S.Y [10], with male *M.m. castaneus* mice to generate $Avpr1b$ heterozygous (+/-) mice. The $Avpr1b^{+/-}$ mice were then bred with one another to generate F₂ $Avpr1b^{+/+}$ and $Avpr1b^{-/-}$ mice that were on average 50% *M.m. castaneus*. However, it should be noted that it is likely that the flanking regions of the disrupted $Avpr1b$ gene would be likely to come from the 129X1/SvJ embryonic stem cell background. These animals were then tested in a resident-intruder behavioral test as described below. It was observed that the new “wild” F₂ generation were much more “reactive” than were the original line; they tended to jump out of the cage more and were highly aggressive when handled. While the observations were not systematic, this increase in aggression likely reflects heightened defensive aggression rather than offensive aggression (H.K.C., personal observations).

Animals

The *M.m. castaneus* were provided to us by Dr. Christine Kozak in the Viral Biology Section of NIAID. These mice were received in approximately 1998 from a colleague at Roswell Park who had derived them from wild-trapped mice. They have been randomly-bred since in NIH animal facilities [32-34].

Stimulus “intruders” were 10-week old Balb/c males purchased from NCI-Frederick. Different intruders were used to test the “original” and the “wild” lines of mice. Intruders were group-housed over the course of the experiment. Whether the subject “residents” were of the

“original” line or the more “wild” line, within each line the Avpr1b^{+/+} and Avpr1b^{-/-} mice were of comparable ages (“original” line - Avpr1b^{+/+}: 98.13 ± 8.53 versus Avpr1b^{-/-}: 94.88 ± 7.92 days of age; “wild” line- Avpr1b^{+/+}: 104.42 ± 11.55 versus Avpr1b^{-/-}: 118.57 ± 9.58 days of age) at the time of testing. Residents were initially group-housed in single-sex cages. All animals were housed in a 12L:12D light cycle with food and water available *ad libitum*.

Resident-Intruder Test

Two different resident-intruder tests were completed at two different times on the two lines of mice. The “original” line was tested in March of 2007 while the “wild” line was tested in April 2007. Aside from the dates of testing all other aspects, including housing in the same animal facility, were identical.

Resident males (“original” line: Avpr1b^{+/+} (n=8) and Avpr1b^{-/-} (n=8) ; “wild” line: Avpr1b^{+/+} (n=7) and Avpr1b^{-/-} (n=7)) were singly housed for at least 14 days prior to testing.. Testing was conducted during the dark phase of the light:dark cycle approximately 1 hour after lights out. The test was initiated when an intruder was added to the home cage of the resident male. If no aggressive behavior was observed in the first 5 minutes, a latency of 300s was recorded and the test ended. Otherwise, the test lasted 2 minutes after an attack was first observed. For each resident the latencies to attack as well as the attack frequencies were scored by an observer blind to the genotypes. Subjects were given 3 tests with 3 days between each test. Intruders were only used once each day and residents were never tested with the same intruder.

Statistical Analysis

The data collected across days were analyzed using a repeated measures analysis of variance (ANOVA), with genotype as the between-subjects factor and day as the within-subjects factor. The cumulative attack latencies and attack frequencies were also calculated and compared between groups using a one-way ANOVA. Only animals that attacked were included in the statistical analyses, this resulted in one Avpr1b^{+/+} animal being excluded. Additionally, the latency to attack and attack frequency on day 3 of testing for another Avpr1b^{+/+} mouse was not videotaped due to experimenter error, so the data for this animal was only included in the cumulative attack latency.

To compare the levels of aggressive behavior of the new “wild” line of Avpr1b^{-/-} mice with the previous “original” line, a repeated measures ANOVA was used with genotype and line as the between-subjects factor and day as the within-subjects factor. For all analyses, a p value of < 0.05 was considered statistically significant.

Results

The attack latencies of the “wild” line showed a main effect of day ($F_{(2,20)}=7.71$, $p=0.003$) and of genotype ($F_{(1,10)}=12.79$, $p=0.005$) but no interaction (Figure 1A). Attack latencies shortened from day to day with repeated testing and Avpr1b^{-/-} mice had longer attack latencies compared to Avpr1b^{+/+} mice. In the analysis of attack frequencies across days there was a main effect of day ($F_{(2,20)}=8.28$, $p=0.002$) and of genotype ($F_{(1,10)}=1268.01$, $p=0.036$) but no interaction (Figure 2A). There was the expected increase in attack frequency from day to day with repeated testing and Avpr1b^{-/-} mice displayed fewer attacks than Avpr1b^{+/+} mice. These genotypic differences in attack latency and attack frequency were reflected in statistically significant differences between the groups in cumulative attack latency ($F_{(1,10)}=5.87$, $p=0.038$) (Figure 1B) and in cumulative attack frequency ($F_{(1,12)}=5.12$, $p=0.045$) (Figure 2B).

When the “wild” line was statistically compared to the “original” line there were no differences between the two strains in either the latency to attack or the attack frequency. There were, however, the expected main effects of day and of genotype on the latency to attack (day: $F_{(2,46)}=11.21$, $p=0.001$; genotype: $F_{(1,23)}=20.77$, $p=0.001$) and attack frequency (day: $F_{(2,46)}=16.04$, $p=0.001$; genotype: $F_{(2,23)}=9.03$, $p=0.006$) (Table 1).

Discussion

In the current study we have demonstrated that, on a different background strain, absence of a functional *Avpr1b* gene results in significant reductions in aggressive behavior compared to wildtype controls. These results provide compelling evidence that the role of the *Avpr1b* is conserved within sub-species and likely across species as these results are consistent with studies in Syrian hamsters and mice that found that oral administration of an *Avpr1b* antagonist results in reduced aggression [1,35].

A study that described *M.m. castaneus* as being more aggressive than C57BL6 mice was one of the precipitants of the current study [31]. However, the line generated here that were approximately 50% *M.m. castaneus* do not appear to have higher levels of aggression compared to the “original” line; though these mice were highly reactive compared to the “original” line (described above). Even with the lack of heightened aggression in this “wild” line compared to the “original” line, the persistence of the phenotype is the most critical observation. It is likely that if these mice were further back-crossed into the *M.m. castaneus* sub-species they would show increases in their baseline aggressive behavior.

As the evidence supporting a critical role of the *Avpr1b* in the regulation of aggressive behavior mounts, one issue that still remains is where in the brain *Avp* is acting via the *Avpr1b*. While highly expressed in the pituitary, the *Avpr1b* mRNA is also prominently expressed within the CA2 pyramidal neurons of the hippocampus [28,30]. A recent publication examining changes in blood oxygen levels (BOLD), although focused on the role of the *Avpr1a*, showed that the male rat hippocampus has increased activity in the presence of a female mate and intruder male [Imaging the Neural Circuitry and Chemical Control of Aggressive Motivation. Ferris CF, Stolberg T, Kulkarni P, Murugavel M, Blanchard R, Blanchard DC, Febo M, Brevard M, Simon NG. BMC Neurosci. 2008 Nov 13;9(1):111. [Epub ahead of print]]. It should be noted that *in situ* hybridization histochemistry for the *Avpr1b*, reveals only two other areas that express *Avpr1b* mRNA, the paraventricular nucleus and the anterior amygdala. Within these two nuclei there were relatively few labeled neurons [28]. Detection of *Avpr1b* protein has remained elusive but the abundance of message within CA2 hippocampus is intriguing. We have hypothesized that the role of the *Avpr1b* within the CA2 field may be to help in the formation of memories that are accessory olfactory-based [28,36]. If this is the case, then the *Avpr1b* may be helping to encode the social context and perhaps even stimulate the retrieval of a previous social memory. Therefore, the reduced aggressive behavior caused by a null mutation of the *Avpr1b* gene is a part of a larger, more global deficit in response to social stimuli. The presence of the *Avpr1b* within the CA2 field of hippocampus across mouse, rat, and human suggests that whatever its role may be, its location appears to be evolutionarily conserved [28]. Future work will focus on the contribution of the CA2 field of hippocampus to the behavioral phenotype observed in *Avpr1b* knockout mice.

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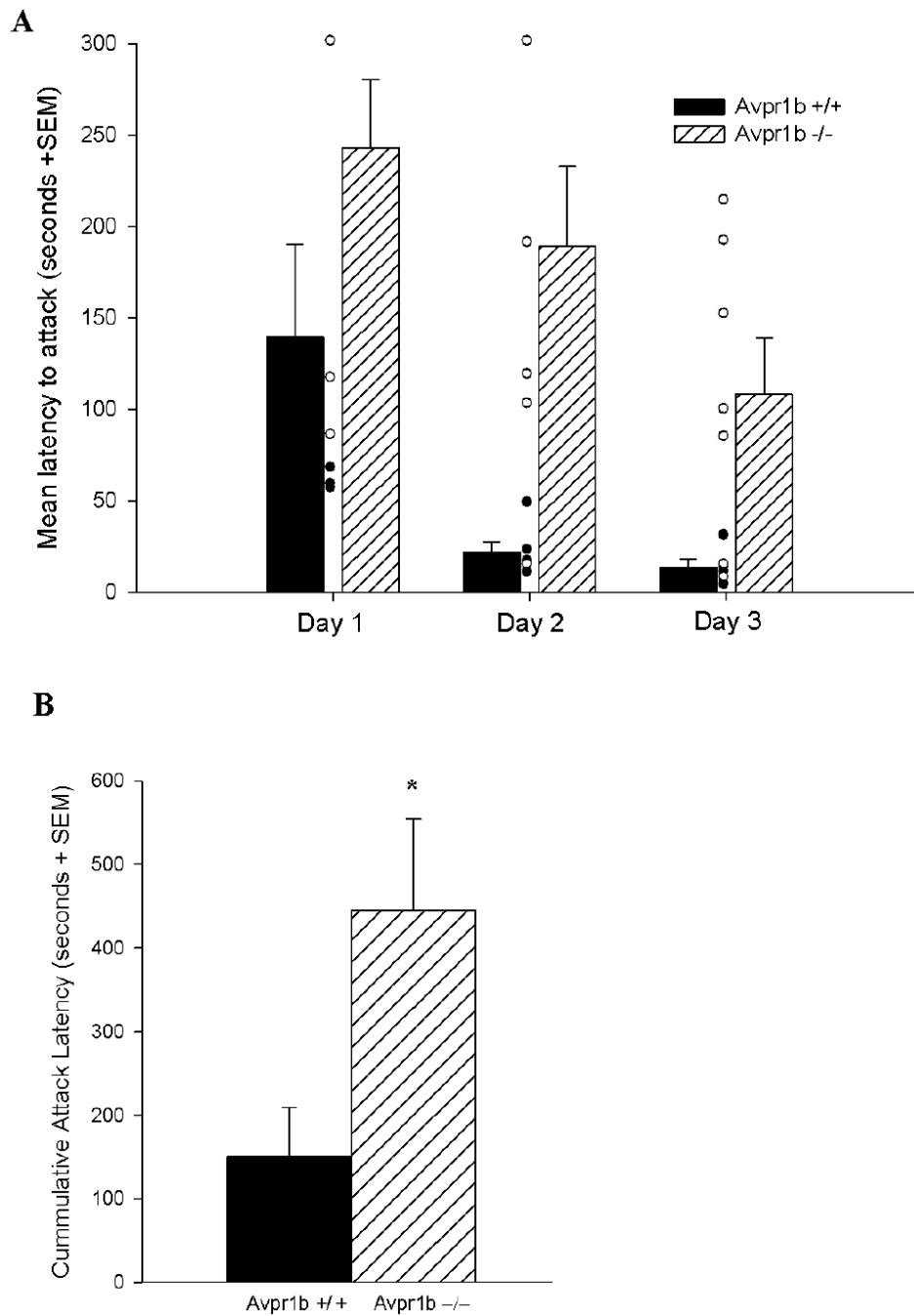


Figure 1. Attack latencies (A) across test days (including individual data points) and (B) cumulatively for Avpr1b wildtype (+/+) and knockout (-/-) mice on a more “wild” background. In (A) there were main effects of day and of genotype, but no interaction. Avpr1b^{-/-} mice had longer attack latencies compared to Avpr1b^{+/+} mice. In (B) Avpr1b^{-/-} mice had longer cumulative attack latencies than Avpr1b^{+/+} mice. (Mean ± SEM)

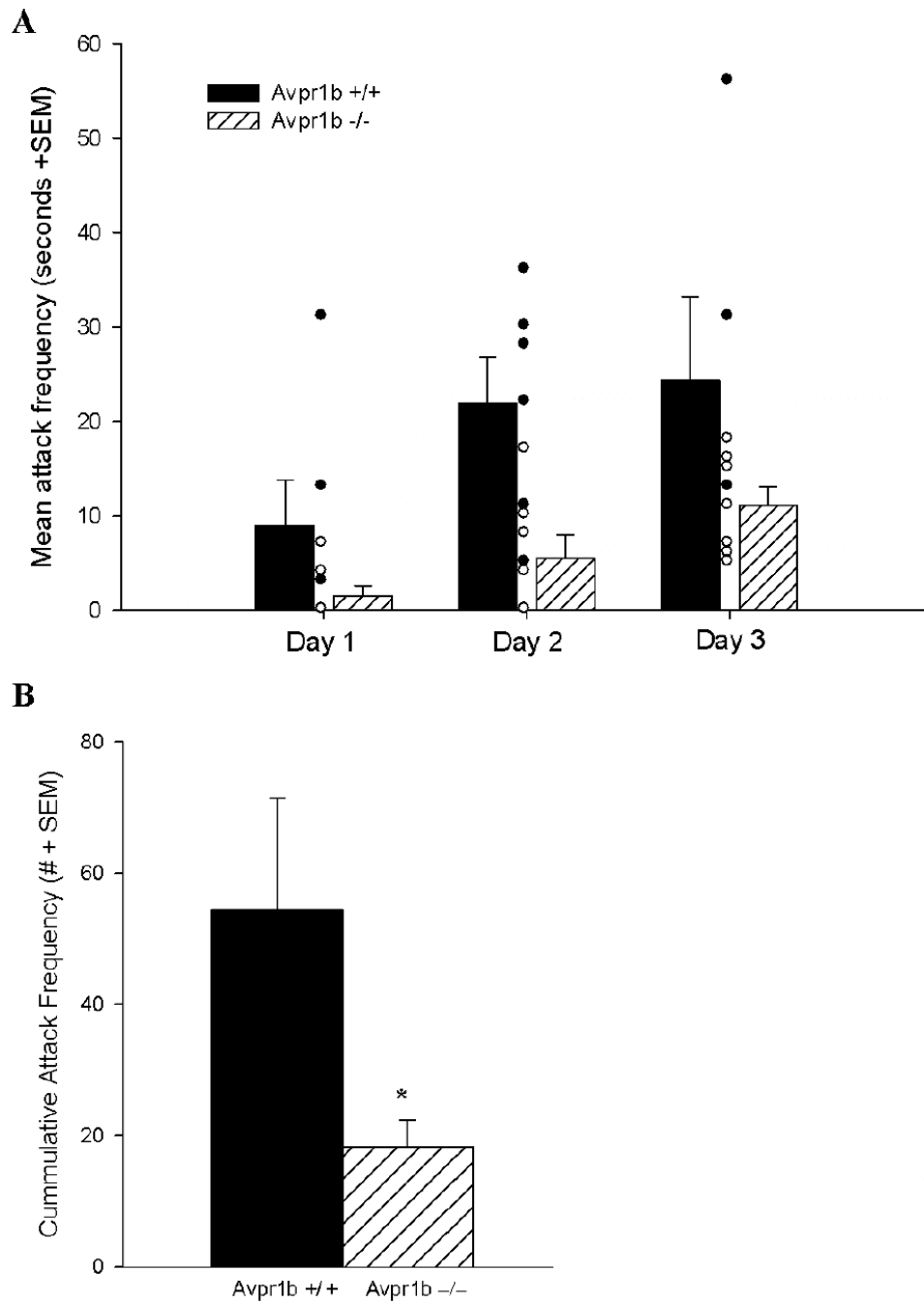


Figure 2. Attack frequencies (A) across test days (including individual data points) and (B) cumulatively for Avpr1b wildtype (+/+) and knockout (-/-) mice on a more “wild” background. In (A) there were main effects of day and of genotype, but no interaction. Avpr1b^{-/-} mice had fewer attacks compared to Avpr1b^{+/+} mice. In (B) Avpr1b^{-/-} mice displayed fewer cumulative attacks than Avpr1b^{+/+} mice. (Mean ± SEM)

Table 1

	Attack Latency (Mean \pm SEM)			Attack Frequency (Mean \pm SEM)		
	Day 1	Day 2	Day 3	Day 1	Day 2	Day 3
Avpr1b^{+/+}	107.40 \pm 48.19	16.80 \pm 2.33	13.80 \pm 4.44	10.80 \pm 5.50	19.20 \pm 4.85	24.40 \pm 8.84
“wild” line	119.50 \pm 30.34	50.88 \pm 18.36	56.13 \pm 35.56	9.13 \pm 2.15	9.38 \pm 2.27	16.38 \pm 3.09
Avpr1b^{-/-}	243.00 \pm 36.95	189.14 \pm 43.70	108.43 \pm 30.65	1.57 \pm 1.07	5.57 \pm 2.45	11.14 \pm 1.99
“original” line	249.00 \pm 24.95	182.25 \pm 43.45	192.13 \pm 45.65	2.13 \pm 1.04	5.63 \pm 1.94	10.88 \pm 4.41