

G Proteins $G\alpha_{i1/3}$ Are Critical Targets for *Bordetella pertussis* Toxin-Induced Vasoactive Amine Sensitization

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Pertussis toxin (PTX) is an AB₅-type exotoxin produced by the bacterium *Bordetella pertussis*, the causative agent of whooping cough. *In vivo* intoxication with PTX elicits a variety of immunologic and inflammatory responses, including vasoactive amine sensitization (VAAS) to histamine (HA), serotonin (5-HT), and bradykinin (BDK). Previously, by using a forward genetic approach, we identified the HA H₁ receptor (*Hrh1/H₁R*) as the gene in mice that controls differential susceptibility to *B. pertussis* PTX-induced HA sensitization (Bphs). Here we show, by using inbred strains of mice, F₁ hybrids, and segregating populations, that, unlike Bphs, PTX-induced 5-HT sensitivity (Bpss) and BDK sensitivity (Bpbs) are recessive traits and are separately controlled by multiple loci unlinked to 5-HT and BDK receptors, respectively. Furthermore, we found that PTX sensitizes mice to HA independently of Toll-like receptor 4, a purported receptor for PTX, and that the VAAS properties of PTX are not dependent upon endothelial caveolae or endothelial nitric oxide synthase. Finally, by using mice deficient in individual $G\alpha_{i/o}$ G-protein subunits, we demonstrate that $G\alpha_{i1}$ and $G\alpha_{i3}$ are the critical *in vivo* targets of ADP-ribosylation underlying VAAS elicited by PTX exposure.

Infection with the bacterium *Bordetella pertussis* can lead to whooping cough, usually typified by paroxysmal coughing and, in severe cases, oropharyngeal tissue swelling, hypertension, pneumothorax, and long-lasting aftereffects (1). The principal active agent in *B. pertussis*, pertussis toxin (PTX), is an AB₅-type secreted exotoxin (2–4). The A (active) subunit catalyzes the ADP-ribosylation and thereby impairment of α -subunit signaling in $G\alpha_{i/o}$ -linked heterotrimeric guanine nucleotide regulatory protein (G-protein) complexes (5). This covalent modification prevents the $G\alpha\beta\gamma$ complex from associating with G protein-coupled receptors (GPCRs) on the cell membrane (6). The B (binding) pentamer is thought to bind cell surface receptors on a variety of mammalian cells and facilitate the cellular entry of the A subunit (7). PTX elicits diverse physiological responses *in vivo*, including leukocytosis and altered glucose regulation (8), increased blood-brain barrier (BBB) permeability, and systemic vasoactive sensitization (VAAS) to biogenic amines (9). PTX is also used as an ancillary adjuvant to approximate an infectious environmental influence in animal models of tissue-specific autoimmune disease (10). Genetically, one of PTX's physiological effects, VAAS to histamine (HA), is controlled by a single autosomal dominant locus (*Bphs*, for *B. pertussis*-induced HA sensitization) and has been used as an intermediate phenotype for genetic studies of organ-specific autoimmune diseases (11, 12).

Although several cell types have the capacity to produce HA, mast cells are thought to be a main source during tissue inflammation (13). Mast cells also produce large amounts of serotonin (5-HT) and bradykinin (BDK), which, together with HA and other lipid and glycosylated mediators, are stored in granules. During inflammation, soluble and cell surface molecules can act as secretagogues to stimulate degranulation and release of factors from mast cells, which can affect vascular permeability (14), leading to antigen leakage from or inappropriate leukocyte access to tissues (13).

We previously studied VAAS in response to HA by using a forward genetic approach to identify the HA H₁ receptor (*Hrh1/H₁R*) as the gene underlying Bphs (11, 12), a subphenotype for

tissue-specific autoimmune diseases elicited with the aid of PTX. A T cell-specific role for H₁R in disease susceptibility was found (15), but a bone marrow chimeric approach revealed that expression of H₁R in the nonhematopoietic compartment was responsible for Bphs (16), prompting the closer look at the mechanism of Bphs that we took in this study.

During the positional cloning of *Bphs*, we observed that SJL/J mice were also susceptible to PTX-induced 5-HT sensitization (Bpss, for *B. pertussis*-induced 5-HT sensitization) while C3H.*Bphs*^{SJL/J} mice were not, thereby excluding *Hrh1* as a candidate for Bpss (17). These results indicate that the phenotypic variation in responsiveness to Bphs and Bpss reflects genetic control of distinct intermediate phenotypes rather than allelic variation in genes controlling overall susceptibility to VAAS. To address this possibility and to gain insight into the genetic control of VAAS to other vasoactive amines, we challenged a panel of PTX-sensitized inbred, F₁, and N2 backcross mice with 5-HT and BDK and studied them for Bpss and Bpbs (*B. pertussis*-induced BDK sensitization), respectively. Unlike *Bphs/Hrh1*, which is dominant, we show that *Bpss* and *Bpbs* are recessive traits and are controlled by separate complex genetic mechanisms unlinked to 5-HT or BDK receptors.

It is known that PTX-induced VAAS to HA (VAASH) requires intoxication with active holotoxin (18) in a mechanism that presumably involves ADP-ribosylation of the $G\alpha_{i/o}$ class of G pro-

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TABLE 1 Bphs, Bpss, and Bpbs of inbred mice^a

| Strain | Bpss | | | | | | | | | Bpbs | | | | | |
|-------------------------|-------------------------|---------------------|---------------------------|--------------------|------------|-----------|------------|------------|--------------------------|------------------|------------------|----------|----------|----------|------------------|
| | Bphs | | | PTX + 5-HT at: | | | | | | LD ₅₀ | PTX + Bdk at: | | | | |
| | Carrier + HA (25 mg/kg) | PTX + HA (25 mg/kg) | Carrier + 5-HT (50 mg/kg) | 50 mg/kg | 12.5 mg/kg | 3.1 mg/kg | 0.78 mg/kg | 0.39 mg/kg | Carrier + Bdk (80 mg/kg) | | 80 mg/kg | 40 mg/kg | 20 mg/kg | 10 mg/kg | LD ₅₀ |
| SM/J | 0/4 | 4/4 ^b | 0/4 | 4/4 ^b | 5/5 | 5/6 | 3/6 | 0/4 | 1.0 | 0/4 | 4/4 ^b | 2/4 | 0/4 | 0/4 | 40.0 |
| SJL/J | 0/4 | 4/4 ^b | 0/4 | 12/12 ^c | 12/12 | 6/11 | 0/8 | | 2.6 | 0/4 | 4/4 ^b | 4/4 | 1/4 | 0/4 | 22.5 |
| AKR/J | 0/4 | 0/4 | 0/4 | 12/12 ^c | 10/10 | 8/17 | 0/8 | | 3.2 | 0/4 | 4/4 ^b | 3/4 | 0/4 | 0/4 | 35.6 |
| MRL/MpJ | 0/4 | 0/4 | 0/4 | 4/4 ^b | 6/6 | 4/8 | 0/4 | | 2.9 | 0/4 | 2/4 | 1/4 | 0/4 | 0/4 | |
| LG/J | 0/4 | 4/4 ^b | 0/4 | 4/4 ^b | 7/8 | 0/4 | | | 5.4 | 0/4 | 2/4 | 0/4 | 0/4 | 0/4 | |
| RFM/UnNCr | 0/4 | 4/4 ^b | 0/4 | 6/6 ^d | 0/6 | 0/4 | | | 26.2 | | | | | | |
| PL/J | 0/4 | 4/4 ^b | 0/4 | 6/6 ^d | 0/8 | 0/4 | | | 29.8 | | | | | | |
| DBA/1J | 0/4 | 4/4 ^b | 0/4 | 5/5 ^d | 0/6 | 0/4 | | | 28.6 | | | | | | |
| SWR/J | 0/4 | 4/4 ^b | 0/4 | 6/6 ^d | 0/4 | 0/4 | | | 22.0 | | | | | | |
| C57BL/6J | 0/4 | 4/4 ^b | 0/4 | 9/15 | 0/14 | 0/8 | | | 0/4 | 4/4 ^b | 4/4 | 0/4 | 0/4 | 0/4 | 28.3 |
| DBA/2J | 0/4 | 4/4 ^b | 0/4 | 4/15 | 0/14 | 0/4 | | | 0/4 | 4/4 ^b | 2/4 | 0/4 | 0/4 | 0/4 | 40.0 |
| A/J | 0/4 | 4/4 ^b | 0/4 | 2/15 | 0/12 | 0/4 | | | 0/4 | 4/4 ^b | 2/4 | 0/4 | 0/4 | 0/4 | 40.0 |
| A/HeJ | 0/4 | 4/4 ^b | 0/4 | 3/6 | 0/6 | 0/4 | | | 0/4 | 4/4 ^b | 4/4 | 2/4 | 0/4 | 0/4 | 20.0 |
| NOD/LtJ | 0/4 | 4/4 ^b | 0/4 | 2/6 | 0/8 | 0/4 | | | | | | | | | |
| FVB/NCr | 0/4 | 4/4 ^b | 0/4 | 2/6 | 0/8 | 0/4 | | | | | | | | | |
| C57BL/10J | 0/4 | 4/4 ^b | 0/4 | 1/6 | 0/4 | 0/4 | | | 0/4 | 2/4 | 0/4 | 0/4 | 0/4 | 0/4 | |
| NZB/BlnJ | 0/4 | 4/4 ^b | 0/4 | 3/8 | 0/6 | 0/4 | | | 0/4 | 4/4 ^b | 3/4 | 2/4 | 0/4 | 0/4 | 22.1 |
| NZW/LacJ | 0/4 | 4/4 ^b | 0/4 | 2/9 | 0/6 | 0/4 | | | 0/4 | 4/4 ^b | 4/4 | 2/4 | 0/4 | 0/4 | 20.0 |
| 129X1/SvJ | 0/4 | 4/4 ^b | 0/4 | 6/10 | 5/14 | 1/5 | 0/4 | | 0/4 | 4/4 ^b | 3/4 | 2/4 | 0/4 | 0/4 | 22.1 |
| BALB/cJ | 0/4 | 4/4 ^b | 1/4 | 3/10 | 6/17 | 0/5 | 0/4 | | 0/4 | 4/4 ^b | 4/4 | 2/4 | 0/4 | 0/4 | 20.0 |
| BALB/cByJ | 0/4 | 4/4 ^b | 0/4 | 0/9 | 0/6 | 0/4 | | | 0/4 | 4/4 ^b | 2/4 | 0/4 | 0/4 | 0/4 | 40.0 |
| CBA/JCr | 0/4 | 0/4 | 0/4 | 0/8 | 0/8 | 0/4 | | | | | | | | | |
| B10.D2n | 0/4 | 4/4 ^b | 0/4 | 0/9 | 0/6 | 0/4 | | | 0/4 | 2/4 | 0/4 | 0/4 | 0/4 | 0/4 | |
| B10.S/SgSnJ | 0/4 | 4/4 ^b | 0/4 | 1/10 | 0/6 | 0/4 | | | 0/4 | 1/4 | 0/5 | 0/5 | 0/5 | 0/5 | |
| C3H/HeJ | 0/4 | 0/4 | 0/4 | 4/14 | 4/15 | 0/4 | | | 0/4 | 4/4 ^b | 4/4 | 1/4 | 0/4 | 0/4 | 22.5 |
| C3H/HeN | 0/4 | 0/4 | 0/4 | 0/4 | 0/4 | 0/4 | | | 0/4 | 4/4 ^b | 4/4 | 2/4 | 0/4 | 0/4 | 20.0 |
| C3H.Bphs ^{SJL} | 0/4 | 4/4 ^b | 0/4 | 0/4 | 0/4 | 0/4 | | | 0/4 | 4/4 ^b | 4/4 | 2/4 | 0/4 | 0/4 | 20.0 |

^a Animals were sensitized with 200 ng PTX by i.v. injection and challenged 3 days later by i.v. injection of HA (25 mg/kg), 5-HT (0.39 to 50 mg/kg), or BDK (10 to 80 mg/kg). Three or four mice were examined at a given dose. Dosages are based on mg/kg (dry weight) free base or peptide. Deaths were recorded at 30 min, and the results are expressed as the number dead/number studied. LD₅₀s of 5-HT and HA were calculated with GraphPad Prism. Contrasts for combined phenotypes: SM = SJL = AKR = MRL; SM < LG = SJL = AKR = MRL; LG < RFM = PL = DBA/1 = SWR. The significance of differences between carrier- and PTX-treated groups was determined by Fisher's exact test. The significance of differences in LD₅₀s among and between the strains was determined by the extra sum of squares *F* test. Bpss overall *F* = 15.9, *P* < 0.0001; Bpbs overall *F* = 1.3, *P* = 0.2.

^b *P* = 0.05.

^c *P* < 0.0001.

^d *P* < 0.01.

teins (19). However, there are several $G\alpha_{i/o}$ proteins and splice variants but the specific *in vivo* target(s) associated with PTX-induced VAAS has remained unclear for nearly 30 years. We show here by using knockout (KO) mice that $G\alpha_{i1}$ and $G\alpha_{i3}$ are the specific targets of ADP-ribosylation required for PTX-induced VAASH.

MATERIALS AND METHODS

Ethics statement. All experimental procedures were approved by the University of Vermont Institutional Animal Care and Use Committee (approval number 10-020) in accordance with relevant institutional guidelines and the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health.

Mice. Most of the inbred strains in Table 1 were purchased from The Jackson Laboratory, except for RFM/UnNCr, FVB/NCr, CBA/JCr, and C3H/HeN mice, which were purchased from the National Cancer Institute/Charles River Laboratories. F₁ hybrid strains and N2 backcross mice, as in Tables 2 and 3, respectively, were generated in-house by using parental strains purchased from Jackson. *Hrh1*-deficient mice (B6.129P-*Hrh1*^{tm1Wat}) (20), C3H.SJL-*Bphs*^S congenic mice (11), were maintained in-house at the University of Vermont. *Bdkrb2*-deficient mice (B6.129S7-*Bdkrb2*^{tm1JhJ}) (21), *Nos3*-deficient mice (B6.129P2-*Nos3*^{tm1UncJ}) (22), caveolin-1-deficient mice (B6.Cg-*Cav1*^{tm1MlsJ}) (23), *Tlr4* mutant C57BL/10ScNJ mice (24), and control C57BL/10J mice were purchased

from Jackson. PKC α -deficient (B6.129-*Prkca*^{tm1Jmk}) (25), PKC β -deficient (129/Sv-*Prkcb*^{tm1Tara}) (26), PKC γ -deficient (B6.129P2-*Prkcg*^{tm1Stl}) (27), PKC δ -deficient (129/SV \times Ola-*Prkcd*^{tm1Qxt}) (5), PKC ϵ -deficient (129/Sv \times C57BL/6-*Prkce*^{tm1Ltg}) (28), and PKC ζ -deficient (129/Sv-*Prkcz*^{tm1mos}) mice (29) were a kind gift from Michael Leitges, Max Planck Institute for Experimental Endocrinology, Hannover, Germany.

$G\alpha_o$ -deficient mice (129SvEv.129SvEv-*Gnao1*^{tm1LBI}) (30), $G\alpha_{i1}$ -deficient mice (129SvOla.129SvEv-*Gnao1.1*^{tm1LBI}) (31), $G\alpha_{i2}$ -deficient mice (129SvOla.129SvEv-*Gnao1.2*^{tm1LBI}) (31), $G\alpha_{i1}$ -deficient mice (129Sv.129SvEv-*Gnai1*^{tm1Drs}) (32), $G\alpha_{i2}$ -deficient mice (B6.129SvEv-*Gnai2*^{tm1LBI}) (33), $G\alpha_{i3}$ -deficient mice (129Sv.129SvEv-*Gnai3*^{tm1LBI}) (31), and $G\alpha_{i1/3}$ doubly deficient mice (129SvEv-*Gnao1*^{tm1LBI} \times 129SvEv-*Gnai3*^{tm1LBI}) (31) were obtained from Lutz Birnbaumer (National Institute of Environmental Health Sciences, Research Triangle Park, NC) and maintained as wild-type (WT, +/+) mice, heterozygotes (+/-), or homozygotes (-/-) on 129SvEv ($G\alpha_o$, $G\alpha_{i1}$, $G\alpha_{i2}$, $G\alpha_{i3}$, $G\alpha_{i1} \times G\alpha_{i3}$), CD1 outbred ($G\alpha_o$), FVB ($G\alpha_{i2}$, 10th backcross), C57BL/6 ($G\alpha_{i2}$, 8th backcross), and crossbred C57BL/6:129SvEv 50:50 ($G\alpha_{i1}$, $G\alpha_{i2}$, $G\alpha_{i3}$) backgrounds. All animals were provided normal mouse chow and water *ad libitum*.

Vasoactive amine sensitization and challenge. Bpss was carried out as described previously (11, 16). Bpss and Bpbs followed a similar protocol. Briefly, 200 ng of PTX (List Biological Laboratories, Campbell, CA) in phosphate-buffered saline was administered intravenously (i.v.) by tail

TABLE 2 Inheritance of Bphs, Bpss, and Bpbs in F₁ hybrid strains^a

| Phenotype and strain | Classification | No. of mice with phenotype/total (P value) | | | Contrasts |
|----------------------------|---|--|-----------------------------|----------------------------|--|
| | | Group 1 | Group 2 | Group 3 | |
| Bphs | | | | | |
| SJL/J | Bphs ⁺ Bpss ⁺ Bpbs ⁺ | 6/6 ^b | 6/6 ^c | 5/6 ^d | |
| C3H/HeJ | Bphs ⁻ | 0/6 ^b | 0/6 ^c | 0/6 ^d | |
| (C3H × SJL)F ₁ | | 6/6 ^b (0.0001) | 6/6 ^c (0.0001) | 4/6 ^d (0.0094) | SJL/J = (C3H × SJL)F ₁ > C3H/HeJ |
| C57BL/6J | Bphs ⁺ Bpss ⁺ | 6/6 ^b | 6/6 ^c | 1/6 ^d | |
| CBA/J | Bphs ⁻ | 0/6 ^b | 0/6 ^c | 0/6 ^d | |
| (B6 × CBA)F ₁ | | 6/6 ^b (0.0001) | 6/6 ^c (0.0001) | 2/6 ^d (0.3) | C57BL/6J = (B6 × CBA)F ₁ > CBA/J |
| Bpss | | | | | |
| SJL/J | Bphs ⁺ Bpss ⁺ Bpbs ⁺ | 12/12 ^e | 12/12 ^f | 6/11 ^g | |
| PL/J (control) | Bpss ⁺ | 6/6 ^e | 0/8 ^f | 0/4 ^g | |
| (PL × SJL)F ₁ | | 8/9 ^e (0.35) | 1/9 ^f (<0.0001) | 0/5 ^g (0.03) | SJL/J > PL/J = (PL × SJL)F ₁ |
| SJL/J | | 12/12 ^e | 12/12 ^f | 6/11 ^g | |
| BALB/cJ | Bpss ⁻ | 3/10 ^e | 6/17 ^f | 0/5 ^g | |
| (BALB × SJL)F ₁ | | 6/9 ^e (0.001) | 4/8 ^f (0.002) | 0/5 ^g (0.02) | SJL/J > BALB/cJ = (BALB × SJL)F ₁ |
| AKR/J | Bphs ⁺ Bpss ⁺ Bpbs ⁺ | 12/12 ^e | 10/10 ^f | 8/17 ^g | |
| DBA/2J | Bpss ⁻ | 4/15 ^e | 0/14 ^f | 0/4 ^g | |
| (AK × DBA)F ₁ | | 4/8 ^e (0.0003) | 0/8 ^f (<0.0001) | 0/5 ^g (0.05) | AKR/J > DBA/2J = (AK × DBA)F ₁ |
| SJL | | 10/10 ^e | 10/10 ^f | 6/10 ^g | |
| B10.S | Bpss ⁻ | 1/10 ^e | 1/10 ^f | 0/10 ^g | |
| (B10 × SJL)F ₁ | | 1/10 ^e (<0.0001) | 0/10 ^f (<0.0001) | 0/10 ^g (0.0006) | SJL/J > B10.S = (B10 × SJL)F ₁ |
| Bpbs | | | | | |
| SJL/J | Bphs ⁺ Bpss ⁺ Bpbs ⁺ | 6/6 ^h | 6/6 ⁱ | 4/6 ^j | |
| B10.S | Bpss ⁻ | 1/6 ^h | 0/6 ⁱ | 0/6 ^j | |
| (B10 × SJL)F ₁ | | 2/6 ^h (0.009) | 1/6 ⁱ (0.0007) | 0/5 ^j (0.008) | SJL/J > B10.S = (B10 × SJL)F ₁ |

^a F₁ hybrids were generated by using the indicated parental strains. Mice were then challenged for Bphs, Bpss, or Bpbs at the indicated doses. Significance of difference in susceptibility between parental strains and F₁ hybrids was determined using Fisher's exact test.

^b HA challenge dose, 25 mg/kg free base.

^c HA challenge dose, 12.5 mg/kg free base.

^d HA challenge dose, 3.1 mg/kg free base.

^e 5-HT challenge dose, 50 mg/kg free base.

^f 5-HT challenge dose, 12.5 mg/kg free base.

^g 5-HT challenge dose, 3.1 mg/kg free base.

^h BDK challenge dose, 80 mg/kg peptide.

ⁱ BDK challenge dose, 40 mg/kg peptide.

^j BDK challenge dose, 20 mg/kg peptide.

vein injection in a volume of 0.1 ml on day 0. On day 3, mice were challenged i.v. with HA dihydrochloride, 5-HT hydrochloride, or BDK acetate (all from Sigma, dosed at milligrams per kilogram of body weight by using the free amine molecular weight). Deaths were recorded at 30 min post-challenge and are shown as the number of animals affected over the total.

TABLE 3 Bphs, Bpss, and Bpbs in N2 backcross mice^a

| Phenotype and strain | No. of mice | | P value | |
|-----------------------------|-------------|------------|---------|---------|
| | Affected | Unaffected | 1 locus | 2 loci |
| Bpss | | | | |
| SJL | 10 | 0 | | |
| (B10.S × SJL)F ₁ | 0 | 10 | | |
| (B10.S × SJL) × B10.S N2 | 20 | 80 | 0.50 | 0.005 |
| Bpbs | | | | |
| SJL | 10 | 0 | | |
| (B10.S × SJL)F ₁ | 0 | 10 | | |
| (B10.S × SJL) × B10.S N2 | 28 | 72 | 0.75 | <0.0001 |

^a SJL, (B10.S × SJL)F₁, and (B10.S × SJL) × B10.S N2 mice were challenged for Bpss (25 mg/kg 5-HT) and Bpbs (40 mg/kg BDK) as in Table 1. Results were tested by chi-square goodness of fit for both one- and two-locus recessive models of inheritance.

Microsatellite genotyping. We performed microsatellite typing as previously described, with liver DNA using informative microsatellite markers that distinguished B10.S and SJL/J mice and are linked to all known 5-HT receptors and the *Bdkrb1/2* locus (34, 35).

RESULTS

Distinct genetic control of Bphs, Bpss, and Bpbs. To further extend our hypothesis that distinct genetic mechanisms control susceptibility to PTX-induced sensitization to individual biogenic amines, we examined 27 different inbred mouse strains for Bphs, Bpss, or Bpbs. We identified two strains that were sensitive to a challenge with all three amines (SJL/J and SM/J), which is consistent with previous reports showing the SJL/J strain is sensitive to both HA and 5-HT challenges (17). Several strains were sensitive to only two amines, but we also observed strains that were mono-sensitive. For example, C57BL10/J mice were susceptible only to Bphs, C3H/HeJ or C3H/HeN mice were susceptible only to Bpbs, and MRL/MpJ mice were susceptible only to Bpss (see Table S1 in the supplemental material). Furthermore, even within a given phenotype, we observed variability in the 50% lethal doses (LD₅₀s). The LD₅₀ of Bpss for SM/J mice was 1 mg/kg 5-HT, whereas that for PL/J mice was 30 mg/kg (Table 1). Bpbs exhibited

LD₅₀s of 20 to 40 mg/kg. The dose of HA used as a control for all strains (25 mg/kg) is approximately 2 LD₅₀s for susceptible strains and readily discriminates Bphs from Bphs-resistant strains (17). The observance of all possible amine sensitivity combinations across an inexhaustive survey of inbred mouse strains indicates that the phenotypic variation in susceptibility to VAAS reflects the genetic control of distinct intermediate phenotypes rather than allelic variation in genes controlling overall susceptibility to intoxication or VAAS.

Susceptibility to Bphs is controlled by *Hrh1*, and this phenotype is inherited in an autosomal dominant fashion (11). We expanded these studies to investigate the inheritance of Bpss and Bpbs by phenotyping (susceptible × resistant)F₁ hybrid mice. (C3H/HeJ × SJL/J)F₁ mice were Bphs, consistent with previous studies (12). Sudweeks et al. observed that (SJL/J × CBA/J)F₁ mice were also Bphs (12). Here we extended this finding by phenotyping (C57BL/6J × CBA/J)F₁ mice for Bphs and observed the same fully penetrant dominant inheritance (Table 2). These data are also consistent with genotype analysis of *Hrh1*, since both SJL/J and C57BL/6J mice harbor the Bphs (*Hrh1*^s) allele while C3H/HeJ and CBA/J mice harbor the Bphs resistance *Hrh1*^r allele (11).

Bpss and Bpbs are two-gene recessive traits. To assess *Bpbs* and *Bpss* inheritance, we generated (susceptible × resistant)F₁ hybrid mice on the basis of the phenotype data in Table 1. For both *Bpss* and *Bpbs*, we used SJL/J mice as the prototypical susceptible strain (SJL/J mice were susceptible to all three vasoactive amines, Bphs⁺ Bpss⁺ Bpbs⁺; Table 1). To rule out strain-specific contributions to the Bpss phenotype, we also employed an F₁ hybrid generated by using AKR/J mice as the susceptible strain (like SJL/J mice, AKR/J mice are Bpss⁺ Bpbs⁺). As a positive control for interstrain crosses in Bpss, we assessed (SJL/J × PL/J)F₁ mice (PL/J is Bpss⁺), and these mice indeed remained Bpss⁺ and were statistically indistinguishable from either parental strain (Table 2). By analyzing the Bpss phenotype of (SJL/J × BALB/cJ), (SJL/J × B10.S), and (AKR/J × DBA/2J)F₁ hybrids, we found that Bpss was recessive since the F₁ mice displayed the phenotype of the Bpss-resistant parental strain (Table 2). We observed a similar recessive inheritance pattern for Bpbs, where (SJL/J × B10.S)F₁ mice were Bpbs resistant, like the B10.S parental strain (Table 2). Collectively, these data again confirm that the phenotypic variation in susceptibility to VAAS reflects the genetic control of responsiveness to individual VAAs rather than susceptibility and resistance to intoxication in general.

The fact that Bphs was reported to be controlled by a single autosomal dominant gene (36) was confirmed when we identified *Hrh1* as the gene controlling Bphs (11). In order to determine the number of loci controlling Bpss and Bpbs, we generated (B10.S × SJL/J) × B10.S N2 backcross mice in sufficient numbers to be able to statistically evaluate by chi-square test both one- and two-locus models of genetic control. The data for both Bpss and Bpbs are consistent with a two-gene model of recessive inheritance (Table 3).

Lack of linkage of Bpss and Bpbs to 5-HT or BDK receptors, respectively. Given that Bphs was identified as the product of *Hrh1* (11), we wondered whether Bpss and Bpbs could be similarly controlled by 5-HT and BDK receptor polymorphism. To test this hypothesis, we determined whether or not Bpss or Bpbs was linked to any of the known 5-HT or BDK receptors, respectively. To this end, (B10.S × SJL/J) × B10.S N2 backcross mice were phenotyped for Bpss or Bpbs and genotyped by using informative microsatellite markers linked to the 14 5-HT genes encoding func-

tional receptors on chromosomes 1, 4, 5, 9, 13, 14, 16, 18, 19, and X and to *Bdkrb1/2* on chromosome 12. Four additional putative 5-HT receptor sequences (*Htr1da*, *Htr1db*, *Htr1ea*, and *Htr1eb*) have been deposited in PubMed Gene. *Htr1da* and *Htr1db* were based on partial *Htr1d* sequence data by Weydert et al. suggesting the possible existence of two *Htr1d* genes on mouse chromosome 4 (37). However, *Htr1da* was shown not to encode a unique functional receptor, and the record for *Htr1db* has been replaced with that for *Htr1d*, which we analyzed. With respect to *Htr1ea* and *Htr1eb*, the situation is similar; *Htr1ea* is actually *Htr1e* and *Htr1eb* is *Htr1f*, both of which we analyzed. Therefore, our analysis covered all known functional 5-HT receptor genes. By chi-square analysis, we did not observe significant linkage of *Bpss* to any of the known 5-HT receptors, nor did we observe linkage of *Bpbs* to *Bdkrb1/2* (Table 4). However, we did observe that *Bdkrb2*-deficient mice (21) were completely protected from Bpss (Table 5), indicating that while not genetically linked to *Bpbs*, *Bdkrb2* signaling is required to elicit PTX-induced sensitization to BDK.

eNOS and caveolin-1 in Bphs. To further investigate the molecular mechanism by which PTX intoxication elicits HA sensitization, we employed genetically modified mice lacking key individual components thought to be involved in PTX activity or the signal propagation involved in HA-mediated vasoregulation. Vascular endothelium expresses the enzyme nitric oxide synthase (eNOS, the protein encoded by *Nos3*), which catalyzes the formation of nitric oxide from L-arginine. This pathway plays a major role in the regulation of vascular tone and blood pressure through the generation of cyclic GMP (cGMP) (38–40). eNOS activation occurs in response to elevation of Ca²⁺ concentrations in the cell (17, 41). Activation of H₁R increases intracellular Ca²⁺ levels as one of the proximal signaling events (42, 43). In addition, H₁R activation has been shown to elevate cGMP levels *in vivo* (44). To address whether *Nos3* is required for Bphs, we sensitized WT C57BL/6J mice and *Nos3*-deficient mice (B6.129 background) (22) with PTX and challenged them with HA. *Nos3*-deficient mice were just as Bphs as WT mice (Table 6), suggesting that VAAS following PTX intoxication is not dependent on *Nos3*. In endothelial cells (ECs), eNOS is highly expressed in caveolae, which are specialized membrane-associated organelles important for the barrier and vasoregulatory functions of these cells. To test whether other features of caveolae distinct from eNOS are important in the regulation of VAASH, we tested the Bphs response of caveolin-1-deficient mice. Since cross regulation of caveolin-1 and eNOS function has been shown (45), we also investigated Bphs in caveolin-1 (*Cav1*) and *Nos3* doubly deficient mice. In contrast to control *Hrh1*-deficient mice, both *Cav1*^{-/-} and *Cav1*^{-/-} *Nos3*^{-/-} mice were Bphs (Table 7). Together, these results indicate that neither the formation of caveolae nor activation of eNOS in ECs is a critical regulator of VAASH.

Individual PKC isoforms do not mediate Bphs. In most mammalian tissues, H₁R induces the activation of phospholipase C (PLC) via PTX-resistant Gα_{q/11}-proteins to generate inositol-1,4,5-triphosphate (IP₃) and diacylglycerol (DAG) from phosphatidylinositol-4,5-bisphosphate (43). IP₃ leads to increased intracellular Ca²⁺, and DAG leads to activation of specific isoforms of protein kinase C (PKC). We therefore examined the requirements for activation of PKC downstream of H₁R in eliciting Bphs. There are at least 12 PKC isoforms (46), and we surveyed selected gene-deficient mice representing representative members of the different isozyme subclasses for their roles in susceptibility to Bphs. We

TABLE 4 Bpss and Bpbs are not linked to 5-HT and BDK receptors, respectively^a

| Receptor | Chr ^b | Distance (cM ^c) | Size (Mbp) | Marker | Distance (cM) | Size (Mbp) | No. of mice | | | | | | χ ² value | P value |
|-----------------------|------------------|-----------------------------|------------|-----------------|---------------|------------|------------------|----|------------------|-------------------|----|--------------------|----------------------|----------------------|
| | | | | | | | Affected | | | Unaffected | | | | |
| | | | | | | | S | H | B | S | H | B | | |
| Bpss | | | | | | | | | | | | | | |
| <i>Htr2b</i> | 1 | 52 | 88 | <i>D1Mit10</i> | 56.6 | 93 | 3 | 12 | 2 | 22 | 14 | 39 | 4.8 | 0.4 |
| <i>Htr5b</i> | | 63 | 123 | | | | | | | | | | | |
| <i>Htr6</i> | 4 | 65 | 139 | <i>D4Mit203</i> | 60 | 129 | 2 | 12 | 6 | 25 | 34 | 17 | 4.9 | 0.4 |
| <i>Htr1d</i> | | 66 | 136 | | | | | | | | | | | |
| <i>Htr5a</i> | 5 | 15 | 28 | <i>D5Mit148</i> | 18 | 32 | 6 | 9 | 3 | 16 | 46 | 14 | 4.5 | 0.5 |
| <i>Htr3a</i> | 9 | 29 | 49 | <i>D9Mit105</i> | 35 | 63 | 9 | 6 | 3 | 21 | 34 | 19 | 6.6 | 0.3 |
| <i>Htr3b</i> | | 29 | 49 | | | | | | | | | | | |
| <i>Htr1b</i> | | 46 | 82 | | | | | | | | | | | |
| <i>Htr1a</i> | 13 | 58 | 106 | <i>D13Mit75</i> | 59 | 107 | 3 | 11 | 6 | 20 | 37 | 18 | 1.2 | 0.9 |
| <i>Htr2a</i> | 14 | 42 | 75 | <i>D14Mit71</i> | 44 | 97 | 6 | 11 | 3 | 17 | 36 | 23 | 2.3 | 0.8 |
| <i>Htr1f</i> | 16 | 39 | 65 | <i>D16Mit14</i> | 34 | 51 | 4 | 9 | 7 | 19 | 32 | 25 | 3.9 | 0.6 |
| <i>Htr4</i> | 18 | 37 | 62 | <i>D18Mit81</i> | 41 | 67 | 4 | 11 | 5 | 19 | 39 | 18 | 0.4 | 1 |
| <i>Htr7</i> | 19 | 33 | 36 | <i>D19Mit88</i> | 34 | 37 | 6 | 8 | 6 | 20 | 34 | 22 | 1.8 | 0.9 |
| <i>Htr2c</i> | X | 66 | 143 | <i>DXMit36</i> | 63 | 143 | 2/4 ^d | 5 | 3/6 ^d | 5/22 ^d | 16 | 12/27 ^d | 3.2/0.1 ^d | 0.7/0.8 ^d |
| Bpbs, <i>Bdkrb1/2</i> | | | | | | | | | | | | | | |
| | 12 | 52 | 107 | <i>D12Mit7</i> | 50 | 104 | 7 | 11 | 10 | 14 | 36 | 15 | 3.4 | 0.6 |

^a (B10.S × SJL/J) × B10.S N2 backcross mice were phenotyped for Bpss or Bpbs as described for Table 3 (25 mg/kg 5-HT and 40 mg/kg BDK, respectively). B, B10.S; H, heterozygous; S, SJL/J (at indicated position with chromosome location for the gene and closest marker indicated). Genotype frequency differences among affected and unaffected BCl mice were evaluated by chi-square analysis in two-by-two contingency tables. Significance was tested by Fisher's exact test.

^b Chr, chromosome.

^c cM, centimorgans.

^d Female/male values.

selected three “conventional” DAG/Ca²⁺-dependent isozymes (PKC-α, -β, and -γ), two DAG-dependent and Ca²⁺-independent “novel” class members (PKC-δ and -ε), and one DAG- and Ca²⁺-independent “atypical” member (PKC-ζ) for PTX sensitization and subsequent HA challenge. All of these gene-deficient mice are on the B6 background, but all carry 129 strain DNA because of gene targeting in strain 129 embryonic stem cells. However, we found no difference in Bpbs in the background control strains over a 16-fold HA dose range (Table 8). As expected, H₁RKO mice were Bpbs resistance, again demonstrating the critical role of *Hrh1* in eliciting Bpbs. Lastly, we observed that all B6.129 PKC KO mice were susceptible to Bpbs (Table 8). Thus, irrespective of the mode of activation or tissue expression patterns, neither PKC-α, -β, -γ, -δ, -ε, nor -ζ individually appears to control Bpbs.

Bpbs is not controlled by *Tlr4*. Experiments with murine splenocytes using radiolabeled PTX and lipopolysaccharide (LPS) showed that both compounds bind to Toll-like receptor 4 (47). In 2004, Kerfoot et al. reported that induction of experimental autoimmune encephalomyelitis (EAE) with PTX as an ancillary adju-

vant and PTX-mediated lymphocyte rolling and adhesion were diminished in the absence of *Tlr4* (48). In conjunction with a recent report showing that commercial PTX preparations can contain LPS/endotoxin (49), we found that our source of PTX contained approximately 40 endotoxin units (EU) per PTX sensitization inoculum. This concentration is equivalent to 0.001 EU/kg of body weight (data not shown), which is far below the 5-EU/kg cutoff in preclinical animal models (50). In light of these considerations, we tested the hypothesis that the VAAS activity of PTX in the Bpbs model was dependent on *Tlr4*. For these experiments, we sensitized WT C57BL/10J mice and C57BL/10ScNJ mice with active holotoxin and challenged them with HA 3 days later. C57BL/10ScNJ mice have a deletion of the *Tlr4* gene that results in the absence of both mRNA and protein (*Tlr4*^{lps-del}) and thus is defective in response to LPS stimulation (24). *Tlr4*^{lps-del} differs from the *Tlr4*^{Lps-d} mutation of C3H/HeJ mice, a point mutation that causes an amino acid substitution, resulting in a hypomorphic *Tlr4* gene product (24). Studying the responsiveness of C57BL/10ScNJ mice therefore precludes the possibility that PTX can still mediate its effects through a non-LPS-responsive allele. We found that both C57BL/10J and C57BL/10ScNJ mice were

TABLE 5 Bpbs in WT and *Bdkrb2*-deficient mice^a

| Strain | No. of mice dead/total after challenge with BDK at: | | | | χ ² value, <i>df</i> | P value |
|---|---|----------|----------|----------|---------------------------------|---------|
| | 80 mg/kg | 40 mg/kg | 20 mg/kg | 10 mg/kg | | |
| C57BL/6J | 3/3 | 3/3 | 1/3 | 0/3 | | |
| B6.129S7- <i>Bdkrb2</i> ^{tm1Jfl/J} | 0/4 | 0/4 | 0/4 | 0/4 | 26.0, 7 | 0.0005 |

^a WT or *Bdkrb2*-deficient mice were sensitized with 200 ng PTX by i.v. injection and challenged 3 days later by i.v. injection of BDK. Significance was determined by Fisher's exact test.

TABLE 6 Bpbs is not mediated by eNOS^a

| Strain | No. of mice dead/total after challenge with HA at: | | |
|--|--|------------|------------|
| | 50 mg/kg | 12.5 mg/kg | 3.13 mg/kg |
| C57BL/6J | 4/4 | 4/4 | 2/4 |
| B6.129.P2- <i>Nos3</i> ^{tm,i/Unc/J} | 4/4 | 4/4 | 1/4 |

^a WT and *Nos3*-deficient mice were analyzed for Bpbs 3 days after the i.v. injection of 200 ng PTX. HA dosages are based on mg/kg (dry weight) free base. Deaths were recorded at 30 min, and the results expressed as the number dead/number studied.

TABLE 7 Disruption of endothelial caveolae does not affect Bphs^a

| Strain | No. of mice dead/total (% of total) after challenge with HA at: | | | |
|---|---|------------|------------|-------------|
| | 25 mg/kg | 12.5 mg/kg | 6.25 mg/kg | 3.125 mg/kg |
| WT | 2/2 (100) | 2/2 (100) | 2/2 (100) | 0/2 (0) |
| <i>Cav1</i> ^{-/-} | 2/2 (100) | 2/2 (100) | 3/3 (100) | 2/5 (40) |
| <i>Cav1</i> ^{-/-} × <i>Nos3</i> ^{-/-} | 2/2 (100) | 2/2 (100) | 3/3 (100) | 1/4 (25) |
| <i>Hrh1</i> ^{-/-} | 0/3 | 0/3 | 0/3 | 0/3 |

^a WT, *Cav1*-deficient, *Cav1/Nos3* doubly deficient, and *Hrh1*-deficient mice were analyzed for Bphs. Deaths were recorded at 30 min, and the results expressed as the number dead/number studied.

Bphs (Table 9). These data are also consistent with our previous study showing that PTX-induced angiogenesis of brain microvascular ECs was not affected by the lack of *Tlr4* signaling (51). Thus, our data suggest that while certain defined effects of PTX, such as leukocyte recruitment, may require *Tlr4*, the HA-sensitizing activity of PTX does not.

Gα_{i1/3} are critical PTX targets for elicitation of VAASH. The ADP-ribosylating activity of PTX is a well-established mechanism of action whereby PTX inhibits the function of Gα_{i/o} proteins (52) and is required to elicit the myriad physiological responses following *in vivo* intoxication, including VAAS and the immune potentiating effects in EAE (18). Conjugation of ADP-ribose to a key cysteine residue in the C terminus of Gα_{i/o} (but not Gα_s, Gα_q, or Gα₁₁) proteins prevents the association of the Gα_{i/o}Gβγ complex with the GPCR (5, 26). Therefore, PTX is a powerful tool to discriminate whether a given GPCR signals through Gα_{i/o}-dependent or Gα_{i/o}-independent mechanisms (19). However, there are several Gα_i proteins (Gα_{i1}, Gα_{i2}, Gα_{i3}, and Gα_z) and Gα_o proteins (Gα_{o1} and Gα_{o2}, splice variant products of the *Gnao* gene) (28). To determine which Gα_{i/o} proteins are targeted by PTX for HA sensitization, we directly assessed the responses of mice genetically deficient in specific Gα_i or Gα_o proteins to HA challenge. We did not test Gα_z since this protein lacks the key cysteine for PTX-mediated ADP-ribosylation (29) and is therefore not affected by PTX. Our hypothesis was that genetic deficiency in one or more Gα_{i/o} proteins would mimic HA hypersensitivity; i.e., such mice would be Bphs without prior PTX sensitization.

We first addressed the role of Gα_o proteins as targets for ADP-ribosylation in VAASH. Gα_{o1} and Gα_{o2} are different gene prod-

TABLE 8 Bphs in PKC KO mice^a

| Strain | No. of mice dead/total after challenge with HA at: | | |
|------------------------------------|--|------------|------------|
| | 50 mg/kg | 12.5 mg/kg | 3.13 mg/kg |
| 129XIIISvJ | 10/10 | 10/10 | 4/6 |
| C57BL/6J | 6/6 | 7/8 | 4/6 |
| B6129PF ₂ /J | 4/4 | 4/4 | 5/6 |
| <i>Hrh1</i> ^{-/-} | 0/4 | 0/4 | 0/4 |
| <i>Prkca</i> (PKCα) ^{-/-} | 3/3 | 3/3 | 3/3 |
| <i>Prkcb</i> (PKCβ) ^{-/-} | 3/3 | 3/3 | 3/3 |
| <i>Prkcc</i> (PKCγ) ^{-/-} | 6/6 | 8/8 | 9/9 |
| <i>Prkcd</i> (PKCδ) ^{-/-} | 4/4 | 3/3 | 3/3 |
| <i>Prkce</i> (PKCε) ^{-/-} | 11/11 | 3/3 | 3/3 |
| <i>Prkcz</i> (PKCζ) ^{-/-} | 4/4 | 3/3 | 1/2 |

^a WT controls and PKC-α, -β, -γ, -δ, -ε, and -ζ singly deficient mice were analyzed for Bphs. Deaths were recorded at 30 min, and the results expressed as the number dead/number studied.

TABLE 9 Bphs in *Tlr4* mutant mice^a

| Strain | No. of mice dead/total after challenge with HA at: | | |
|--------------|--|------------|-------------|
| | 50 mg/kg | 12.5 mg/kg | 3.125 mg/kg |
| C57BL/10J | 5/5 | 6/6 | 1/6 |
| C57BL/10ScNJ | 5/5 | 6/6 | 0/6 |

^a C57BL/10J (*Tlr4*^{+/+}) and C57BL/10ScNJ (*Tlr4*-deficient) mice were analyzed for Bphs. Deaths were recorded at 30 min, and the results expressed as the number dead/number studied.

ucts because of alternative exon usage in the *Gnao* gene (30). *Gnao*-deficient mice lack both Gα_{o1} and Gα_{o2} splice products because of disruption of *Gnao* exon 6, which is common to both gene products (30). Neither *Gnao*-deficient mice nor mice specifically lacking either Gα_{o1} or Gα_{o2} were spontaneously sensitive to HA (Fig. 1). These results demonstrated that the HA-sensitizing activity of PTX was not due to its ADP-ribosylation of Gα_o proteins.

Next we addressed the putative role of Gα_i proteins as targets for PTX. Gα_{i1}, Gα_{i2}, and Gα_{i3} are encoded by distinct genes, *Gnai1*, *Gnai2*, and *Gnai3*, respectively (29), and are widely expressed (28). Compared to WT mice, Gα_{i1}^{-/-} (*P* = 0.0002) and Gα_{i3}^{-/-} (*P* = 0.008) mice were significantly more sensitive to HA, whereas Gα_{i2}^{-/-} mice were not (see Table S2 in the supplemental material). We also analyzed the response of Gα_{i1}^{-/-} Gα_{i3}^{-/-} double-KO mice and found them to also be significantly more susceptible to spontaneous HA sensitivity than WT mice are (*P* = 0.001) (see Table S2). Additionally, if one directly compares the susceptibility to spontaneous sensitivity to HA among the various KO strains, there is a significant overall effect of genotype (*P* < 0.0001) with no significant difference among *Gnao*^{-/-}, Gα_{o1}^{-/-}, Gα_{o2}^{-/-}, and Gα_{i2}^{-/-} mice (*P* = 0.5) or among Gα_{i1}^{-/-}, Gα_{i3}^{-/-}, and Gα_{i1}^{-/-} Gα_{i3}^{-/-} mice (*P* = 0.3); however, there is a highly significant difference in susceptibility to spontaneous HA sensitivity between the two groups (*P* < 0.0001). Taken together, these results demonstrate that Gα_{i1} and Gα_{i3} are clear targets of ADP-ribosylation associated with PTX-induced VAASH.

DISCUSSION

We have focused on understanding the genetic basis for VAAS elicited by PTX sensitization. Changes in vascular permeability elicited during inflammation may promote antigen leakage and contribute to pathogen dissemination. Dysregulated vascular per-

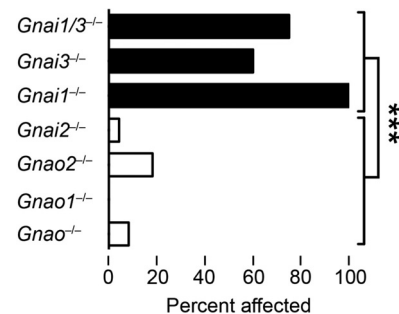


FIG 1 Loss of Gα_{i1} or Gα_{i3} mimics PTX to elicit HA sensitization. Mice deficient in Gα_{i/o} subunits were analyzed for HA sensitivity in the absence of PTX exposure. Mice received 100 mg/kg HA *i.v.*, and deaths were recorded within 30 min. Statistical significance was determined by one-way ANOVA using 8 to 23 mice per group. ***, *P* < 0.001.

meability may allow for inappropriate tissue access to immune cell surveillance, leading to tissue damage (13). In this regard, PTX has long been known to alter vascular permeability through its ability to promote VAAS (53). The vasoactive amines HA, 5-HT, and BDK have all been shown to influence vascular permeability, in particular, the BBB (16, 53, 54). Here we have shown that after PTX sensitization, (i) hypersensitivity to HA, 5-HT, or BDK (Bphs, Bpss, and Bpbs, respectively) is uniquely genetically controlled; (ii) unlike the genetic control of *Bphs*, that of *Bpss* and *Bpbs* and is not linked to any of the known 5-HT or BDK receptors; (iii) Bpbs is mediated by *Bdkrb2*; and (iv) G $\alpha_{11/3}$ proteins are the physiological *in vivo* targets of ADP-ribosylation by PTX associated with VAASH.

Under inflammatory conditions, vascular permeability is modulated by activated leukocytes acting on the vascular unit either through secreted or cell surface factors (13, 54). Mast cells are classically known to produce and store HA, 5-HT, and other amine/lipid/glycosylated mediators (55). More recently, mast cells have been shown to also produce BDK (14). Mast cells perfuse tissues and rapidly respond to external local stimuli which may act as secretagogues to stimulate the release of stored granular contents, including vasoactive amines. In fact, the critical role of mast cell-derived products in the control of vascular permeability was recently shown (14). Concomitant with the relaxed vascular permeability promoted by mast cell degranulation, neutrophils may gain tissue access to promote damage and promote subsequent recruitment of adaptive immune cells (13). From these data, one may then posit that environmental exposure may affect mast cells or the target vasculature to influence vascular permeability.

Mast cell number, function (56, 57), and susceptibility to VAAS vary across different genetic backgrounds (58, 59). Although Bphs was mapped by using the SJL/J-derived *Hrh1* allele (11, 12), up to 75% of the available inbred mouse strains harbor this same allele (11; R. Noubade, unpublished data). We and others have indeed observed that SJL/J mice were also susceptible to Bpss (17, 53), though Bpss was not mediated by *Hrh1* (17). Like HA and 5-HT, a BDK challenge after PTX intoxication also leads to death due to acute hypovolemic and hypotensive shock. Given these phenotypic similarities and the extensive genetic variation across inbred laboratory mouse strains, we tested whether a conserved genetic mechanism controls general VAAS by PTX. To address this possibility, we examined responses to HA, 5-HT, and BDK in a panel of inbred mice. Among these strains, we observed the full spectrum of VAAS to different individual mediators, indicating that the phenotypic variation in responsiveness to PTX reflects the genetic control of distinct intermediate phenotypes rather than allelic variation in genes controlling overall susceptibility to PTX intoxication.

Furthermore, using F₁ hybrid crosses, we found that Bpss and Bpbs are autosomal recessive, while Bphs is autosomal dominant (12). The susceptibility pattern for Bpss and Bpbs in (SJL/J \times B10.S) \times SJL/J N2 crosses was consistent with a two-gene model of genetic control. Unlike *Bphs*, which showed genetic linkage to a key HA receptor, *Hrh1* (11), for both *Bpss* and *Bpbs*, susceptibility was not linked to genes encoding the cognate ligand receptors. Furthermore, although *Bpbs* was not linked to *Bdkrb1/2*, this phenotype was abrogated in *Bdkrb2*-deficient mice, indicating that *Bdkrb2* is nonetheless required for *Bpbs*. Taken together, our results confirm and extend the finding that the phenotypic variation

in susceptibility to VAAS following PTX sensitization reflects the unique genetic control of specific intermediate phenotypes, i.e., Bphs, Bpss, and Bpbs, rather than susceptibility and resistance to intoxication in general.

Previously, we investigated the cell type requirements for H₁R expression in Bphs. Using bone marrow chimeras, we found that the nonhematopoietic compartment determined susceptibility to Bphs (16). We found that EC-specific expression of H₁R was not sufficient to mediate Bphs (16). Here we extended our analysis of ECs in VAASH. Inhibition of G α_i proteins by PTX has been shown to increase eNOS (*Nos3*) expression (60), yet *Nos3*-deficient mice were not protected from Bphs. Similarly, disruption of *caveolin-1* either alone or in conjunction with *Nos3* disruption did not affect Bphs. Another possibility that we have not analyzed here is that VAASH is due to PTX-mediated inhibition of sphingosine-1-phosphate (S1P) receptor signaling in EC. S1P regulates barrier function, and mice deficient in S1P in plasma were more susceptible to HA-induced vascular leakage (61). Nonetheless, administration of PTX to mice expressing EC-specific H₁R did not render them sensitive to VAASH (16), which argues against this notion. Taken together, these results did not formally rule out the possibility that *Hrh1* on EC is required but suggested that coexpression of H₁R in another component of the vascular unit may be required for Bphs. Given the roles for H₁R in vascular smooth muscle and neural function (62, 63), we hypothesize that H₁R signaling in non-ECs of the vascular unit may be an important component of Bphs (16). Indeed, in the central nervous system (CNS), astrocytic end feet projecting into the EC/smooth muscle layers that wrap cerebral arterioles are important in relaying neural signals that control arterial tone in the brain (64).

Our results illustrate a central role for G α protein activation and GPCRs in vasoregulation. The mast cell mediators HA, 5-HT, and BDK all promote vascular leakage/edema (65), and all signal through GPCRs. Nonetheless, diversity in molecular responses is controlled by receptor specificity and the activation of different effector G α proteins (28). Forty-five years ago, Bergman and Munoz introduced the concept of counterbalanced signals controlling VAAS, with a special emphasis on the interplay between HA and the adrenergic-receptor (AR) pathways (66). These authors also suggested that PTX might sensitize to HA by targeting AR pathways. More recent observations are also consistent with the existence of opposing G α protein effects in VAAS. Many ARs are coupled to G α_i proteins (α_{2A} , α_{2B} , α_{2C}) (67), whereas key receptors for the vasoactive amines (H₁R, 5-HT₂, and the B₂ BDK receptor) are all linked to G $\alpha_{q/11}$ (28). It was recently shown that the anaphylactic shock response is controlled by endothelial G $\alpha_{q/11}$ signaling (68). Our data suggest that inactivation of G $\alpha_{11/3}$ proteins by PTX is a key destabilizing effect that may allow unabated signaling through G $\alpha_{q/11}$ or G α_s -linked GPCRs such as those for HA, 5-HT, and BDK.

PTX catalyzes the ADP-ribosylation of G $\alpha_{i/o}$ proteins associated in the G $\alpha/\beta/\gamma$ heterotrimer and prevents the interaction of the trimeric signaling complex with GPCRs (6). In mice, there are eight G $\alpha_{i/o}$ genes (*Gnai1* to -3, *Gnao*, *Gnaz*, and *Gnat1* to -3) which, because of alternative splice forms of *Gnao*, encode at least 10 different protein products (28). PTX has been generally considered a broad-spectrum G $\alpha_{i/o}$ inhibitor with only limited insight into either G $\alpha_{i/o}$ specificity or specific function. Here we have identified G α_{11} and G α_{13} as the targets of ADP-ribosylation associated with Bphs. Mice lacking G α_{11} and G α_{13} were spontaneously

susceptible to Bphs, a response normally observed in intact mice only after intoxication with active PTX holotoxin. It is possible that PTX targets other than $G\alpha_{i1/3}$ contribute to Bphs. This would be detected only if we treated $G\alpha_{i1/3}$ -deficient mice with PTX and observed 100% penetrance. We did not formally address this possibility because we are not aware of any WT or KO mouse strain besides $G\alpha_{i1}$ - or $G\alpha_{i1/3}$ -deficient mice that exhibit spontaneous HA sensitivity at 100% penetrance (17, 36). Therefore, we interpret our present results to be due to incomplete penetrance possibly due in part to HA dosing (17, 36). Furthermore, given that Bphs was seen in all other $G\alpha_{i/o}$ -deficient mice, our results show the specificity of PTX for $G\alpha_{i1/3}$ with regard to HA sensitization.

We hypothesize that H_1R signaling (which uses $G\alpha_{q/11}$) in the vascular unit is counterbalanced by $G\alpha_{i1/3}$ -dependent signals to maintain proper vascular tone and permeability. The mice used here were ubiquitously deficient in $G\alpha_{i1/3}$, precluding the assignment of $G\alpha_{i1/3}$ function to a specific cell type. The GPCR(s) that links to $G\alpha_{i1/3}$ in the control of Bphs is also not known, although ARs remain a possibility. Since the identification of *Hrh1* as *Bphs* in 2002 (11), more has been learned about the other three HA receptors, two of which— H_3R and H_4R —are linked to $G\alpha_{i/o}$ and are therefore sensitive to PTX (63). H_4R expression is restricted to the hematopoietic system, but our finding that Bphs is controlled by the nonhematopoietic system (16) rules out this receptor as a $G\alpha_{i/o}$ -linked candidate for PTX inhibition in Bphs. Expression of H_3R , however, is found in the CNS and in the autonomic nervous system (69) and has been found to regulate BBB permeability and susceptibility to both CNS autoimmune disease (70) and CNS infection in the case of cerebral malaria (71). Accordingly, another potential explanation for the physiological Bphs response could involve PTX-mediated inhibition of H_3R to destabilize baseline vascular permeability and allow HA to signal unabated through another of its receptors (H_1R) to induce acute vasodysregulation.

Collectively, our data reveal that $G\alpha_{i1/3}$ proteins play an important role in the sensing of PTX and the regulation of vascular permeability in response to vasoactive amines. While H_1R signaling in the nonhematopoietic compartment appears to ultimately control Bphs, the hematopoietic compartment is likely to serve as a source of vasoactive amines, for example, with mast cells as major producers of HA. Thus, it is important to understand $G\alpha$ function in specific cell types at the interface of vasculature and tissue. Under inflammatory conditions, $G\alpha_{i2}$ expression in ECs, but not in leukocytes, is important in the extravasation of leukocytes into inflamed lung tissue (72). PTX exposure strongly impairs immune cell migration due to inhibition of $G\alpha_{i/o}$ -linked chemokine receptor signaling (73). Deletion of $G\alpha_{i2}$ affects T and B cell homing under steady-state conditions (33, 74, 75) and leads to T cell hyperreactivity (76). However, not all functions of $G\alpha_{i2}$ are unique to this molecule, since competition for CXCR3-mediated signaling between $G\alpha_{i2}$ and $G\alpha_{i3}$ occurs, where $G\alpha_{i3}$ may compete for and quench $G\alpha_{i2}$ -mediated chemotaxis in *in vitro* migration assays (77). Extravasation of immune cells into tissues under inflammatory conditions involves a combination of chemokines (78, 79) and hence may involve different $G\alpha_i$ members. Our data linking PTX to $G\alpha_{i1/3}$ function and the ability of PTX to enhance the development of inflammatory disease (10) implicate $G\alpha_{i1}$ - and $G\alpha_{i3}$ -linked GPCRs in this process.

Our data may also give insight into the molecular mechanisms discriminating immunity to *B. pertussis* provided through exposure to whole-cell pertussis (wP) via infection or vaccination or via

vaccination with the acellular pertussis (aP) vaccine (which replaced wP in 1997). aP refers to a vaccine in which PTX is inactivated by chemical fixation or by genetic means. This modification decreases the incidence of side effects, but data showing that aP may exhibit less durability and protection than wP are mounting (80). Taken with our data, it is possible that the $G\alpha_{i1/3}$ -linked GPCRs inhibited by PTX are critical for durable immunity to *B. pertussis*. Indeed, production of inflammatory cytokines was increased in stimulated splenocytes from $G\alpha_{i1/3}$ -deficient mice while that of regulatory cytokines was decreased (81), lending support to the notion that $G\alpha_{i1/3}$ damping may contribute to a robust systemic immune response. Identification of these critical GPCRs may lead to strategies for their blockade in order to maximize the benefits of aP while increasing vaccine durability. The hypersensitivity to vasoactive amines released during the course of inflammation, as we have modeled here with Bphs, may be an indicator of such an effective strategy.

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