

Vector Priming Reduces the Immunogenicity of *Salmonella*-Based Vaccines in *Nramp1*^{+/+} Mice

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The present studies in *Nramp1*^{-/-} BALB/c and *Nramp1*^{+/+} CBA mice question the significance of this genotype as a determinant of the level of gut colonization following oral administration of naturally attenuated or highly virulent *Salmonella* strains. In line with previous results in BALB/c hosts, vector priming of CBA mice with *Salmonella enterica* serovar Stanley was found to significantly compromise the immunogenicity of a recombinant construct expressing a foreign pilus protein.

The construction of *Salmonella*-based multivalent vaccines, which can simultaneously confer protection against typhoid and other infectious diseases, would expedite mass immunization programs in developing countries. This strategy is threatened, however, by the possible adverse consequences of the environmental priming of potential vaccinees. Experimental studies have yielded conflicting data concerning the impact of deliberate priming with the vector strain on the immunogenicity of recombinant *Salmonella* vaccines. Two studies in which *aroA* *Salmonella enterica* serovar Dublin vectors were used failed to detect any reduction in responses to foreign antigen delivered orally or systemically to vector-primed hosts (2, 19). Studies in three other laboratories employed *aroA* *S. enterica* serovar Typhimurium (11, 14) or the naturally attenuated *S. enterica* serovar Stanley (1, 18) as vectors and concluded that prior exposure to the vector significantly compromises both serum and intestinal antibody responses to a range of foreign antigens. The factors that determine the impact of vector priming remain unclear but include the nature of the foreign antigen and the *Salmonella* vector (18).

To date, all studies of the significance of vector priming have been performed in BALB/c mice, which have the *Nramp1*^{-/-} genotype and are highly susceptible to *Salmonella* infection. The *Nramp1*-associated difference in the capacity to control the growth of systemically administered virulent serovar Typhimurium is dramatic, with 50% lethal doses typically 1,000-fold higher in resistant *Nramp1*^{+/+} strains (10, 13). The wild-type *Nramp1* protein is primarily expressed in macrophages within the spleen and liver (8, 17), where it is recruited to the membrane of *Salmonella*-containing phagosomes. *Nramp1* functions as a divalent metal ion transporter, suggesting that its key function is to reduce cation availability within the phagosome, but a role in promoting phagosome maturation has also been proposed (3, 7).

The relevance of *Nramp1* in the present context derives from Eisenstein's suggestion (5) that *Nramp1*^{+/+} mice might

be more appropriate hosts for studies of *Salmonella* infection, which are designed to assist the development of typhoid vaccines. This was based on a consideration of the relative significance of humoral and cell-mediated responses in immunity to *Salmonella*. Whereas cell-mediated immunity is critical for the protection of *Nramp1*^{-/-} hosts, antibodies induced by inactivated vaccines are sufficient to protect *Nramp1*^{+/+} mice and humans from (low-dose) challenge with virulent *Salmonella* (5).

The aims of this study were twofold. First, we sought to ascertain whether the *Nramp1* genotype influences the extent to which orally administered *Salmonella* can colonize the gut-associated lymphoid tissue (GALT). Since the systemic spread of attenuated *S. enterica* serovar Typhi vaccines is deemed undesirable (16), the immunogenicity of recombinant vaccines based upon such vectors will depend on their capacity to persist within the GALT. The second objective was to determine whether vector priming compromises the immunogenicity of recombinant *Salmonella* vaccines in *Nramp1*^{+/+} mice. This also has clear implications for the development of human vaccines.

***Salmonella* infection in *Nramp1*^{-/-} and *Nramp1*^{+/+} hosts.** Serovar Stanley carries the same O-4,5,12 lipopolysaccharide (LPS) as serovar Typhimurium but is naturally attenuated for mice; after oral administration, it shows prolonged GALT colonization (1) but is unable to spread systemically (data not shown). Groups of BALB/c (*Nramp1*^{-/-}) and CBA (*Nramp1*^{+/+}) mice were orally dosed with 1.1×10^9 CFU of serovar Stanley, and the levels of GALT colonization were compared by determining Peyer's patch burdens at various intervals, as done previously (18). Similar numbers of bacteria were recovered from the two strains at each of the first three time points, with mean burdens of ca. 10^4 CFU on days 5 through 8 (Fig. 1A). Thereafter, significantly fewer bacteria were recovered from the *Nramp1*^{+/+} hosts. These data suggest that innate immunity in the early period is similarly effective in the two strains, with the CBA mice subsequently benefiting from more effective major histocompatibility complex-linked acquired immune responses.

The similarity of the early colonization profiles was unexpected and prompted an experiment to confirm the *Nramp1* status of the mice being used. Groups of CBA and BALB/c mice were injected intraperitoneally with 40 CFU of the highly

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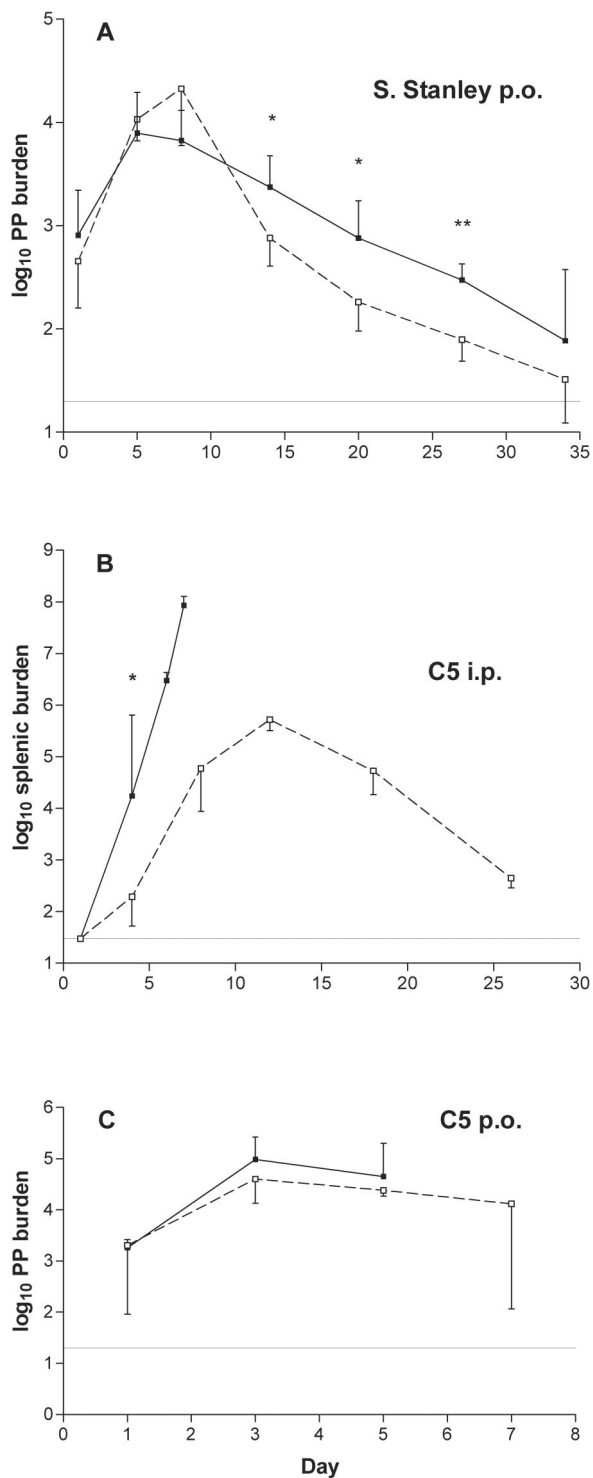


FIG. 1. Comparative growth of *Salmonella* strains in BALB/c and CBA mice. Animals were dosed orally (p.o. [A and C]) or intraperitoneally (i.p. [B]) with serovar Stanley (A) or serovar Typhimurium strain C5 (B and C), as described in the text. Bacterial burdens in Peyer's patches (PP [A and C]) or spleens (B) of BALB/c (■, solid lines) and CBA (□, broken lines) mice were determined by plating homogenates on XLD medium. Graphs show log₁₀ recoveries (geometric mean ± standard deviation; n = 4 at each time point in each experiment), with broken lines showing limits of detection. Student's *t* test (two-tailed) was used to assess the significance of the observed differences (*, *P* < 0.05; **, *P* < 0.01).

virulent C5 strain of serovar Typhimurium. Spleens were removed at various time points, and bacterial loads were determined by plating homogenates on XLD medium (Oxoid). BALB/c mice were unable to control C5 growth in the spleen, and only 4 (of 13) mice were still alive on day 7; these animals had splenic burdens of ca. 10⁸ CFU by this time. In contrast, bacterial loads in the spleens of CBA mice were significantly lower on day 4 (*P* < 0.05) and reached a peak burden of <10⁶ on day 12 (Fig. 1B). Thereafter, the infection was cleared and no mouse of this strain died. This experiment confirmed the presumed difference in the *Nramp1* status of our animals.

The benefit conferred by *Nramp1*, in terms of restricting the systemic growth of *Salmonella*, is less apparent with strains of lower virulence or with *aroA*-attenuated vectors (6, 10, 15). It was therefore of interest to ascertain whether an *Nramp1*-related difference in GALT colonization might be apparent if mice were orally infected with the highly virulent C5 strain. Again, there was no difference in the Peyer's patch colonization profiles seen in BALB/c or CBA mice (dose of 1.1 × 10⁸ bacteria [Fig. 1C]). The last four BALB/c mice died as a result of infection before day 7, whereas the remaining CBA mice showed no visible signs of illness at this time, consistent with their capacity to restrict systemic growth.

Significance of vector priming: relevance of host *Nramp1* status. Groups of BALB/c and CBA mice were orally dosed with 1.1 × 10⁹ CFU of serovar Stanley on day -70, with additional mice kept aside as controls. Ten weeks after primary infection, all animals were dosed with 1.3 × 10⁹ CFU of serovar Stanley-K88, a recombinant expressing the *Escherichia coli* pilus protein K88 (1). Serum and fecal pellet samples were collected at intervals for the determination of anti-K88 and anti-LPS antibody responses by enzyme-linked immunosorbent assay (ELISA), as detailed elsewhere (18).

The efficacy of vector priming was illustrated by the secondary serum (Fig. 2B) and mucosal (Fig. 2D) anti-LPS responses observed in both CBA and BALB/c mice. Control and vector-primed BALB/c mice developed serum anti-K88 (Fig. 2A) and anti-LPS (Fig. 2B) responses comparable to those observed previously (1, 18). Naïve animals displayed strong, consistent immunoglobulin G (IgG) responses to K88, with comparatively weak and delayed responses to vector LPS. Vector priming completely prevented the development of the former and resulted in secondary responses to LPS. A similar pattern of responses was seen in control and vector-primed CBA mice; again, the serum IgG anti-K88 response was completely inhibited as a consequence of prior exposure to the vector strain (Fig. 2A). Of interest was the finding that the primary serum response to K88 was significantly lower in CBA than in BALB/c mice (*P* < 0.01 on days 28 and 69 by one-way analysis of variance with Bonferroni's posttest).

The state of hyporesponsiveness to K88 extended to the mucosal IgA responses in both mouse strains (Fig. 2C), and in BALB/c mice, the impact of vector priming was highly significant (*P* < 0.01 on days 14, 28, 46, and 69). However, the mucosal anti-K88 response in control CBA mice was only modest and consequently, the differences between the (corrected) antibody titers of control and vector-primed mice did not achieve statistical significance. As with the serum IgG response, the primary intestinal IgA response to K88 was sig-

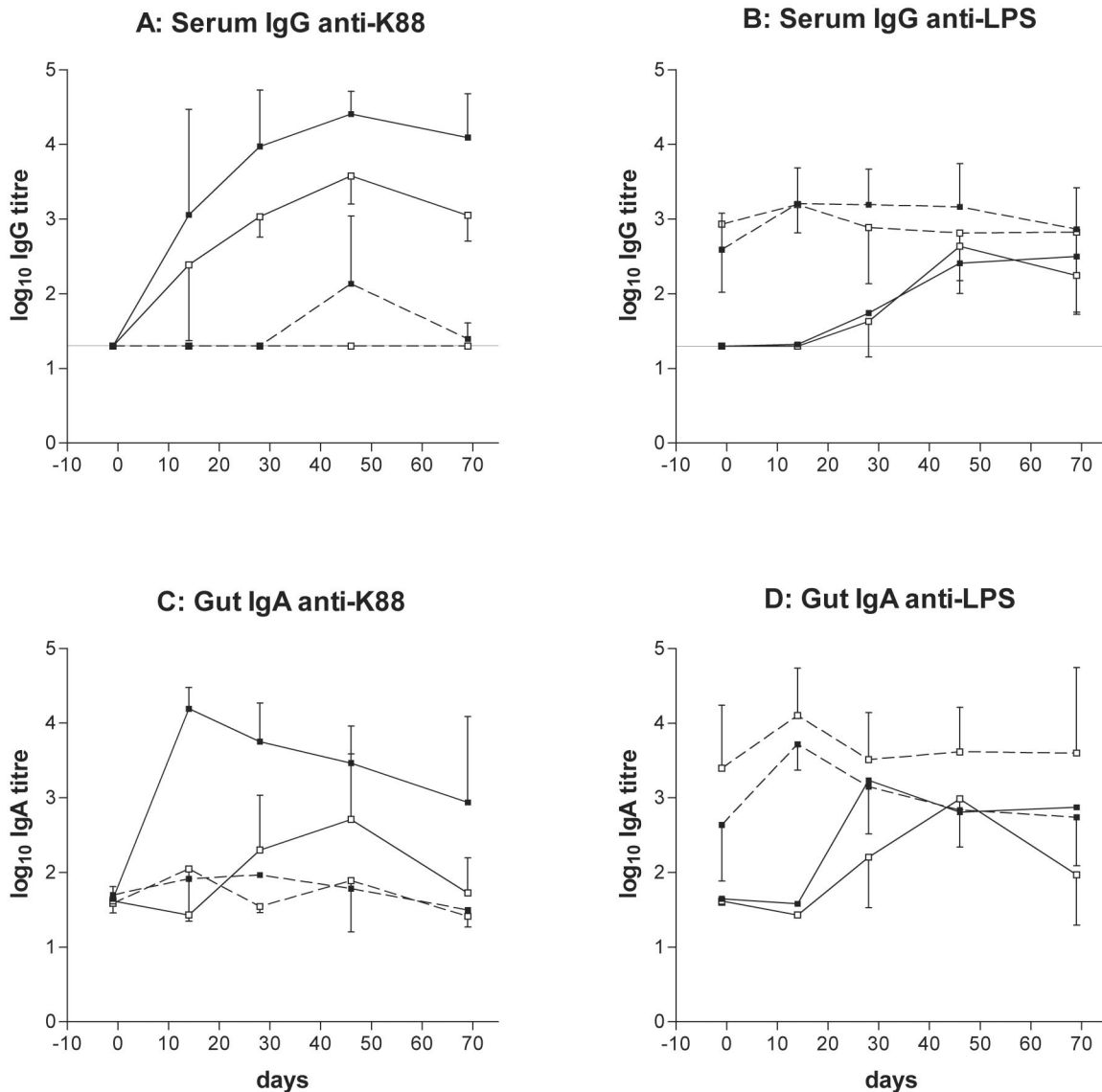


FIG. 2. Impact of vector priming in *Nrampl*^{+/+} mice. Control and vector-primed BALB/c (■) and CBA (□) animals were dosed with serovar Stanley-K88 on day 0. Blood and fecal pellet samples were collected at intervals, and serum IgG (A and B) and gut IgA responses (C and D) were determined by ELISA. The latter are expressed as antibody titer per milligram of IgA to correct for varying immunoglobulin content. Graphs show log₁₀ anti-K88 (A and C) and anti-LPS (B and D) ELISA titers (geometric mean ± standard deviation; *n* = 5 at each time point) in control (solid lines) and vector-primed (broken lines) animals. Broken lines in panels A and B show the limit of detection of serum ELISAs at 1 in 20.

nificantly weaker in CBA than in BALB/c mice on days 14, 28 ($P < 0.01$), and 69 ($P < 0.05$).

IgG subclass analysis of serum anti-LPS responses. Sera collected in the previous experiment were analyzed to ascertain the IgG1:IgG2a subclass ratios of antibodies to LPS in BALB/c and CBA mice. Samples collected 49 and 69 days after primary infection with serovar Stanley and at the same intervals after immunization (of 20-week-old controls) with serovar Stanley-K88 were tested. In all cases, the CBA sera showed a slightly higher proportion of IgG1 antibodies, and at three of the four time points, this difference was significant at the 5% level (Table 1).

Our failure to detect any significant difference in the GALT colonization profiles of attenuated or virulent *Salmonella* strains in mice of different *Nrampl* genotypes suggests that the

complement of resident and migratory phagocytic cells within the infected gut fails to exert any *Nrampl*-mediated control of bacterial growth. This contrasts with the well-recognized and dramatic impact of the *Nrampl* genotype on the fate of systemically administered *Salmonella*. Evidently, the advantage conferred by the *Nrampl*^{+/+} genotype is expressed only systemically, with even *Nrampl*^{+/+} hosts experiencing colonization of the GALT, presumably in deference to the paramount need to restrict systemic growth. Consistent with this notion, *Salmonella* strains capable of prolonged GALT colonization but unable to translocate to organs of the reticuloendothelial system are markedly attenuated compared with wild-type strains such as serovar Typhimurium C5. Examples include mutant strains cured of the virulence plasmid (9) and naturally attenuated strains such as serovar Stanley.

TABLE 1. IgG subclass ratios of serum anti-LPS responses in CBA and BALB/c mice^a

<i>S. enterica</i> strain and sampling time point ^b	CBA	BALB/c
Serovar Stanley		
D49	1.04 ± 0.26	0.81 ± 0.16
D69	1.22 ± 0.26	*0.87 ± 0.10
Serovar Stanley-K88		
D49	1.30 ± 0.38	*0.87 ± 0.23
D69	1.06 ± 0.17	*0.62 ± 0.21

^a Serum IgG anti-LPS responses of CBA and BALB/c mice (from the experiment shown in Fig. 2) were dissected into IgG₁ and IgG_{2a} responses. Data show ratios of IgG₁:IgG_{2a} ELISA titers (geometric mean ± standard deviation; *n* = 5), and an asterisk denotes significance at the 5% level, as determined by the two-tailed *t* test.

^b D, day.

The immunogenicity of recombinant *Salmonella* vaccines designed for use in man will be dependent upon the capacity of the vector strain to colonize the GALT, as the systemic spread of vaccine organisms is undesirable. In this context, the lack of any detectable Nramp1-related impact on GALT colonization by *Salmonella* is encouraging, if the *Nramp1*^{+/+} host indeed represents a more appropriate model for vaccine development (5). The present findings will need to be confirmed in experiments with *Nramp1* congenic mouse strains, however.

BALB/c and CBA mice displayed similar anti-LPS responses following oral infection with serovar Stanley or serovar Stanley-K88, consistent with the similarity of their GALT colonization profiles. However, the significantly different subclass ratios of the serum IgG responses suggest a different Th1/Th2 bias in the two strains. Moreover, antibody responses to the foreign antigen were significantly lower in the *Nramp1*^{+/+} hosts, although this could reflect the involvement of *H-2* genes. The latter might determine responsiveness to a particular foreign antigen (6) and/or effect the more rapid clearance of the vaccine construct from the GALT (Fig. 1A). Further studies with *Nramp1* or *H-2* congenic mice would clarify this issue, but the significant difference in the anti-K88 responses seen here confirms the value of including outbred mice when assessing the immunogenicity of recombinant vaccines (6).

A recent study by Soo et al. (15) used congenic mouse strains to test the impact of Nramp1 on responses to foreign antigens delivered by recombinant *Salmonella*. In this instance, total antibody responses (to a fragment of tetanus toxin) were similar in *Nramp1*^{+/+} and *Nramp1*^{-/-} mice although IgE and cytokine responses were suggestive of a Th1 bias in the former and a Th2 bias in the latter. Whether *Salmonella* infection elicits Nramp1-associated differences in the patterns of cytokine production remains controversial, however (4, 12).

A key finding of the present study was that preexisting antivector immunity compromises the immunogenicity of the serovar Stanley-K88 construct in *Nramp1*^{+/+} mice. This observation extends our previous findings in BALB/c mice (1, 18) to hosts that might better reflect the responses of human vaccine recipients (5). This study therefore supports other recent data (18) indicating that vector priming could prejudice the use of

recombinant *Salmonella* vaccines in regions where these bacteria may be endemic. Further studies will be needed to elucidate the mechanism of hyporesponsiveness seen in vector-primed hosts before compensating strategies can be designed and evaluated.

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