

# Complement-producing maternal microchimeric cells override infection susceptibility in complement-deficient murine offspring

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1 **Complement-producing maternal microchimeric cells override infection**  
2 **susceptibility in complement-deficient murine offspring**

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15 **To the editor:** Long-term persistence of vertically transferred maternal cells occurs ubiquitously  
16 in mammalian offspring (1). Presence of these exceptionally rare maternal microchimeric cells  
17 (MMc), with ensuing immunological tolerance to noninherited maternal antigen (NIMA), is  
18 associated with a variety of remarkable phenotypes including serological resistance to  
19 noninherited maternal HLA sensitization (2), improved long-term survival of NIMA-matched  
20 renal allografts (3), neonatal heart block (4), type 1 diabetes (5), and cross generational  
21 reproductive fitness with expanded accumulation of NIMA-specific regulatory T cells (6).  
22 Herein, we considered whether MMc may exert other physiological benefits beyond these  
23 immunological features linked with antigenicity.

24 A provocative consideration is whether phenotypically wildtype MMc can reduce disease  
25 severity in autosomal recessive disorders caused by defective or missing proteins. Given shared  
26 susceptibility to infection caused by complement deficiency in humans and mice (7), and  
27 enriched MMc in the liver where C3 and other complement components are produced (8), this  
28 hypothesis was investigated by evaluating complement levels and infection susceptibility of C3

29 NIMA (C3<sup>-/-</sup> mice born to C3<sup>+/-</sup> mothers) compared with genetically identical C3<sup>-/-</sup> mice born  
30 to complement deficient mothers, along with C3<sup>+/-</sup> littermate controls (**Figure 1A**;  
31 **Supplemental Figure 1**).

32 We found increased serum C3 levels in C3 NIMA compared with C3<sup>-/-</sup> mice born to complement  
33 deficient mothers, albeit at levels still considerably reduced compared with C3<sup>+/-</sup> controls  
34 (**Figure 1B**). C3<sup>-/-</sup> mice are highly susceptible to *E. coli*, and an intermediate dosage (15000  
35 CFUs) that accentuates this susceptibility was used for intravenous infection to further  
36 investigate functional consequences of complement producing MMc (**Supplemental Figure 2**).  
37 These experiments showed the normally high bacterial burden in tissues of infected C3<sup>-/-</sup> mice  
38 were reduced in C3 NIMA mice (**Figure 1C**), demonstrating that being born to complement  
39 sufficient mothers can dominantly impact infection susceptibility of otherwise genetically  
40 identical complement-deficient mice.

41 To verify importance of C3<sup>+/-</sup> MMc, we evaluated C3 levels and infection susceptibility after  
42 MMc depletion using antibody or pregnancy induced MMc displacement in male and female C3  
43 NIMA mice, respectively. For antibody MMc depletion, transgenic mice with constitutive cell  
44 surface expression of ovalbumin (OVA) (6, 9) were intercrossed with C3<sup>-/-</sup> mice to generate C3  
45 OVA NIMA offspring born to C3<sup>+/-</sup>-OVA<sup>+/-</sup> mothers (**Supplemental Figure 3**). Transforming  
46 OVA with C3 into NIMAs in this fashion allows MMc depletion using anti-OVA IgG, and  
47 verifying loss of MMc by quantifying OVA<sup>+</sup> genomic DNA in tissues such as heart, liver and  
48 uterus which consistently contain highest MMc levels (6, 9). These experiments showed C3<sup>+/-</sup>-  
49 OVA<sup>+/-</sup> MMc depletion reduces serum C3 to background levels (**Figure 1B**), and overturns  
50 infection resistance of C3 NIMA mice (**Figure 1C, 1D**).

51 Despite the ability to persist long-term, MMc are also susceptible to pregnancy induced  
52 displacement and replacement with fetal microchimeric cells (FMc) (9). To further investigate  
53 the necessity of C3<sup>+/-</sup> MMc in C3 NIMA females, we compared C3 levels and susceptibility  
54 after pregnancy sired by complement-deficient males with ensuing replacement by C3<sup>-/-</sup> FMc.  
55 C3 NIMA female mice postpartum after pregnancy sired by C3<sup>-/-</sup> males, with loss of C3<sup>+/-</sup>-  
56 MMc, contained only background serum C3 levels (**Figure 1B**), and infection susceptibility  
57 comparable to C3<sup>-/-</sup> controls (**Figure 1E, 1F**). Thus, complement producing MMc are

58 responsible for above background C3 levels and reduced infection susceptibility in complement  
59 deficient offspring.

60 Beyond complement deficiency, these results suggesting clinical phenotypes associated with  
61 missing or defective proteins in autosomal recessive disorders can be altered by functionally  
62 wildtype MMc opens up fundamental new ways for explaining why individuals with the same  
63 gene defect in many autosomal recessive disorders, including cystic fibrosis and sickle cell  
64 anemia, have widely varied disease severity (10, 11). In turn, these protective benefits associated  
65 with complement producing MMc highlight importance for further investigating how these cells  
66 work, including their cellular identity and phenotype heterogeneity, since expanding their  
67 accumulation beyond natural microchimeric levels represents an innovative approach for  
68 therapeutically reducing severity of common genetic disorders.

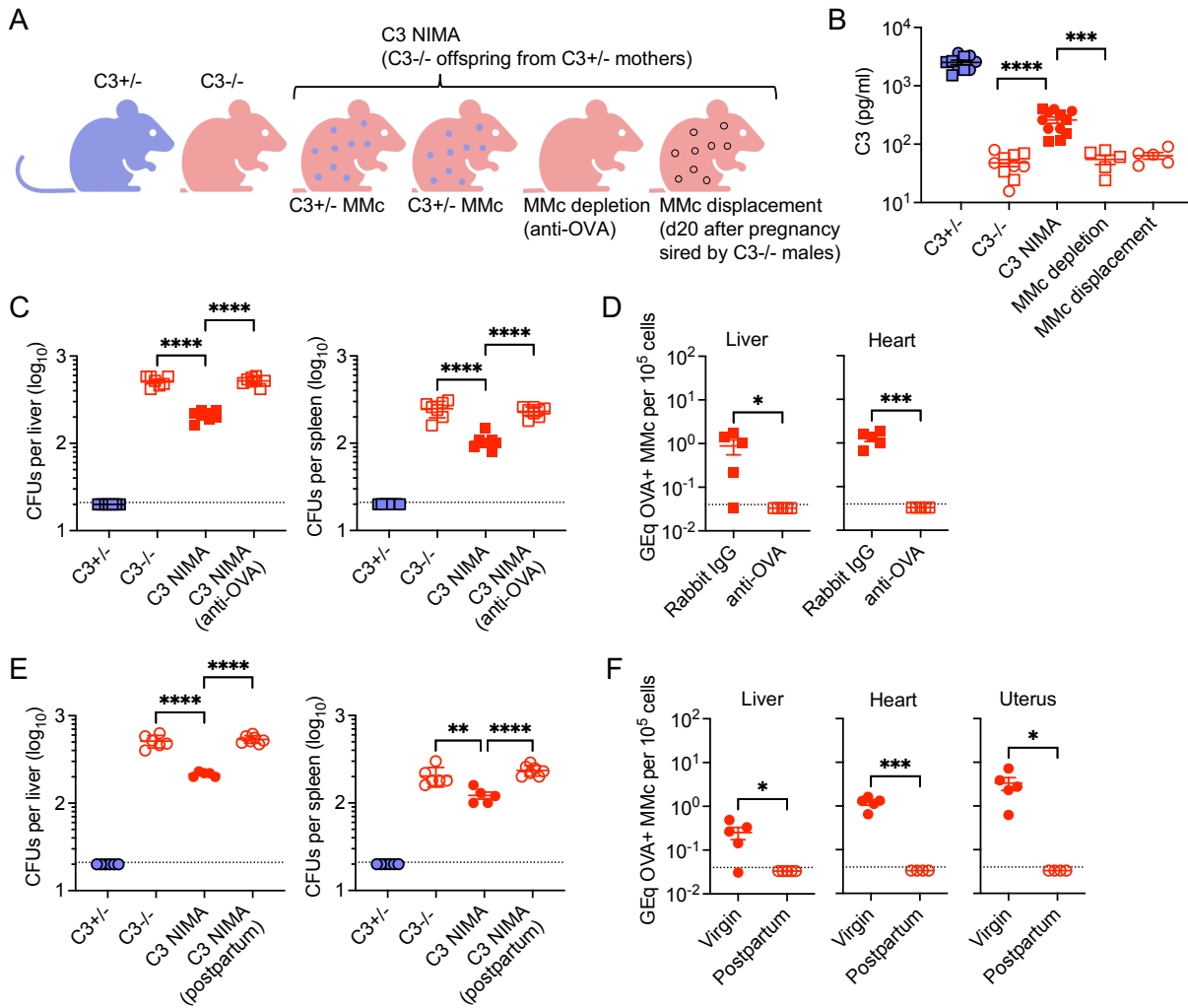
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## 70 REFERENCES

71

- 72 1. Nelson JL. The otherness of self: microchimerism in health and disease. *Trends Immunol.*  
73 2012;33(8):421-7.
- 74 2. Claas FH, Gijbels Y, van der Velden-de Munck J, and van Rood JJ. Induction of B cell  
75 unresponsiveness to noninherited maternal HLA antigens during fetal life. *Science.*  
76 1988;241(4874):1815-7.
- 77 3. Burlingham WJ, Grailer AP, Heisey DM, Claas FH, Norman D, Mohanakumar T, et al.  
78 The effect of tolerance to noninherited maternal HLA antigens on the survival of renal  
79 transplants from sibling donors. *N Engl J Med.* 1998;339(23):1657-64.
- 80 4. Stevens AM, Hermes HM, Rutledge JC, Buyon JP, and Nelson JL. Myocardial-tissue-  
81 specific phenotype of maternal microchimerism in neonatal lupus congenital heart block.  
82 *Lancet.* 2003;362(9396):1617-23.
- 83 5. Nelson JL, Gillespie KM, Lambert NC, Stevens AM, Loubiere LS, Rutledge JC, et al.  
84 Maternal microchimerism in peripheral blood in type 1 diabetes and pancreatic islet beta  
85 cell microchimerism. *Proc Natl Acad Sci U S A.* 2007;104(5):1637-42.
- 86 6. Kinder JM, Jiang TT, Ertelt JM, Xin L, Strong BS, Shaaban AF, et al. Cross-Generational  
87 Reproductive Fitness Enforced by Microchimeric Maternal Cells. *Cell.* 2015;162(3):505-  
88 15.
- 89 7. Wessels MR, Butko P, Ma M, Warren HB, Lage AL, and Carroll MC. Studies of group B  
90 streptococcal infection in mice deficient in complement component C3 or C4  
91 demonstrate an essential role for complement in both innate and acquired immunity. *Proc*  
92 *Natl Acad Sci U S A.* 1995;92(25):11490-4.
- 93 8. Alper CA, Johnson AM, Birtch AG, and Moore FD. Human C'3: evidence for the liver as  
94 the primary site of synthesis. *Science.* 1969;163(3864):286-8.

- 95 9. Shao TY, Kinder JM, Harper G, Pham G, Peng Y, Liu J, et al. Reproductive outcomes  
96 after pregnancy-induced displacement of preexisting microchimeric cells. *Science*.  
97 2023;381(6664):1324-30.
- 98 10. Tewari S, Brousse V, Piel FB, Menzel S, and Rees DC. Environmental determinants of  
99 severity in sickle cell disease. *Haematologica*. 2015;100(9):1108-16.
- 100 11. Drumm ML, Ziady AG, and Davis PB. Genetic variation and clinical heterogeneity in  
101 cystic fibrosis. *Annu Rev Pathol*. 2012;7:267-82.  
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105 **Figure 1. C3<sup>+/-</sup> MMc overrides complement deficiency in C3<sup>-/-</sup> mice.** (A) Schematic comparing  
 106 C3<sup>+/-</sup>, C3<sup>-/-</sup> and C3 NIMA mice, or C3 NIMA mice after MMc depletion or displacement. (B) Serum C3  
 107 levels in male (square) or female (circle) mice described in panel A. (C) *E. coli* CFUs after infection for  
 108 male C3<sup>+/-</sup>, C3<sup>-/-</sup>, C3 NIMA and C3 NIMA mice depleted of OVA+ MMc using anti-OVA IgG. (D)  
 109 Genome equivalents (GEq) OVA DNA specific to OVA+ MMc in male C3 OVA NIMA mice 14 days  
 110 after anti-OVA compared with isotype control IgG administration. (E) *E. coli* CFUs after infection for  
 111 female C3<sup>+/-</sup>, C3<sup>-/-</sup>, virgin C3 NIMA and C3 NIMA postpartum after pregnancy sired by C3<sup>-/-</sup> males. (F)  
 112 GEq OVA DNA specific to OVA+ MMc among female C3 OVA NIMA mice 20 days postpartum after  
 113 pregnancy sired by C3<sup>-/-</sup> males compared with age matched virgin control mice. Each point represents the  
 114 data from an individual mouse, combined from at least 2 independent experiments each with similar  
 115 results. Bar, mean ± standard error. \*P<0.05; \*\*P<0.01; \*\*\*P<0.005; \*\*\*\*P<0.001.