

Deletion of immune evasion genes provides an effective vaccine design for tumor-associated herpesviruses

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1 **Abstract**

2 Vaccines based on live attenuated viruses often induce broad, multifaceted immune
3 responses. However, they also usually sacrifice immunogenicity for attenuation. It is
4 particularly difficult to elicit an effective vaccine for herpesviruses due to an armament of
5 immune evasion genes and a latent phase. Here, to overcome the limitation of
6 attenuation, we developed a rational herpesvirus vaccine in which viral immune evasion
7 genes were deleted to enhance immunogenicity while also attaining safety. To test this
8 vaccine strategy, we utilized murine gammaherpesvirus-68 (MHV-68) as a proof-of-
9 concept model for the cancer-associated human γ -herpesviruses, Epstein-Barr virus and
10 Kaposi sarcoma-associated herpesvirus. We engineered a recombinant MHV-68 virus by
11 targeted inactivation of viral antagonists of type I interferon (IFN-I) pathway and deletion
12 of the latency locus responsible for persistent infection. This recombinant virus is highly
13 attenuated with no measurable capacity for replication, latency, or persistence in
14 immunocompetent hosts. It stimulates robust innate immunity, differentiates virus-specific
15 memory T cells, and elicits neutralizing antibodies. A single vaccination affords durable
16 protection that blocks the establishment of latency following challenge with the wild type
17 MHV-68 for at least six months post-vaccination. These results provide a novel approach
18 to effective vaccination against cancer-associated herpesviruses through the elimination
19 of latency and key immune evasion mechanisms from the pathogen.

20

21 **Keywords:** attenuation, γ -herpesviruses, immunogenicity, immune evasion, latency, T-
22 cell, tumorigenesis, type I interferon, vaccine

23 **Introduction**

24 Human γ -herpesviruses Epstein-Barr virus (EBV) and Kaposi sarcoma-associated
25 herpesvirus (KSHV) are associated with cancer, and with no effective vaccine remain a
26 global health challenge. Despite strong innate and adaptive immune responses, once
27 acquired, herpesviruses persist for the rest of the host's life. EBV is associated with
28 Burkitt's lymphoma, nasopharyngeal carcinoma (NPC), and Hodgkin's- and non-
29 Hodgkin's lymphomas¹⁻³ while KSHV is associated with Kaposi's sarcoma (KS), primary
30 effusion lymphoma (PEL), and multicentric Castleman's disease (MCD). These
31 malignancies frequently develop in AIDS patients⁴⁻⁶, but also in immunocompetent
32 people with more than 160,000 annual cancer cases associated with EBV and KSHV⁷.
33 Clearly, effective vaccines against human γ -herpesviruses would dramatically reduce the
34 incidence of malignancies associated with these viruses.

35

36 Herpesviruses establish persistent infections characterized by lytic replication and
37 latency. Lytic replication of α - and β -herpesviruses results in disease pathologies, such
38 as varicella and herpes zoster for Varicella-Zoster virus (VZV), cold sores and genital
39 lesions for herpes simplex virus (HSV), and congenital defects for cytomegalovirus
40 (CMV). In comparison, malignancies associated with γ -herpesvirus infection are linked to
41 viral latency. Viral genes expressed during latency promote the survival and proliferation
42 of infected cells with increased susceptibility to carcinogenic transformation. Therefore,
43 effective vaccine strategies against tumor-associated herpesviruses ideally should
44 prevent latent infections. The oncogenic potential of γ -herpesviruses has focused vaccine
45 research and development on protein subunit vaccines without the latency risk of live

46 viruses. Subunit anti-EBV vaccines have been based on the envelope protein gp350.
47 Antibodies against gp350 block EBV infection in B-cells⁸ which are long-term latency
48 reservoirs. Gp350-based vaccines protect against infectious mononucleosis (IM);
49 however, they do not influence the overall infection rate⁹ and thus are unlikely to prevent
50 EBV-associated cancers. Similarly, subunit vaccines against HSV-2 may reduce genital
51 lesions but do not prevent infection¹⁰. Therefore, a new strategy is required to establish
52 wide, durable immunity against herpesviruses.

53
54 Live viral vaccines simulate an infection presenting the entire viral antigen repertoire to
55 create stable, long lasting immune memory. Viruses can be attenuated by removing viral
56 genes essential for replication. However, replication-deficient viruses may undergo
57 recombination and regain replication capacity during propagation in complementing cells
58 expressing the missing genes. Furthermore, attenuation of replication competence may
59 compromise immunogenicity. An alternative approach is to selectively inactivate viral
60 genes involved in immune evasion in order to attenuate replication and enhance
61 immunogenicity. Viral antagonists of type I interferon (IFN-I) response are an important
62 class of immune evasion genes to consider. The IFN-I response is the first line of antiviral
63 defense in the host, and subverting the IFN-I response is critical for viruses to establish
64 infections in hosts. IFN-I response initiates a signaling cascade inducing the transcription
65 of over 300 genes that counteract viral infections¹¹⁻¹⁴ and also promotes adaptive immune
66 responses. Approximately 25% of genes encoded by γ -herpesviruses modulate host
67 immunity, including those that counteract the IFN-I response^{15,16}.

68

69 Here, we designed a viral vaccine that addresses both immunogenicity and safety. We
70 hypothesized that a recombinant herpesvirus lacking multiple IFN-I evasion genes and
71 deficient in latency can prime memory development in T and B cells despite attenuated
72 replication. However, human γ -herpesviruses are highly species-specific and cannot
73 infect small animals. To overcome this limitation, we utilized murine gammaherpesvirus
74 68 (MHV-68), closely related to EBV and KSHV¹⁷, to test the hypothesis. We show that
75 an MHV-68 virus engineered to be latency- and immune evasion-deficient is highly
76 attenuated in immunocompetent hosts yet a potent inducer of antiviral immunity.
77 Moreover, this recombinant virus elicits robust long-lasting protection against persistent
78 wild type viral infection.

79 **Results**

80 ***Construction of a virus deficient in immune evasion and persistence (DIP)***

81 In our previous genome-wide screen of MHV-68 open reading frames (ORFs), eight were
82 found to reduce IFN-I responses according to an IFN-stimulated response element (ISRE)
83 reporter assay¹⁸. We selected *ORF10*, *ORF36*, and *ORF54* for the insertion of
84 translational stop codons as these genes are dispensable for viral replication and are
85 conserved among MHV-68, KSHV, and EBV. We also inactivated K3, a viral inhibitor of
86 MHC class I antigen presentation pathway, by truncation to increase the immunogenicity
87 of the vaccine virus^{19,20}. We hypothesized that removal of these four immune evasion
88 genes would increase immunogenicity while attenuating replication of the vaccine virus
89 by inducing a robust IFN response and presenting all viral epitopes.

90

91 A critical safety component of our design is eliminating the latency of the vaccine virus.
92 In KSHV and MHV-68, the biphasic life cycle is regulated by RTA, the replication and
93 transcription activator, and by LANA, the latency associated nuclear antigen. The latter is
94 required for latency establishment^{21–23} while the former upregulates lytic genes^{24–26}. We
95 previously showed that abolishing LANA expression combined with constitutive RTA
96 expression results in a latency-deficient virus²⁷. Here, we replaced the latency locus
97 comprising ORF72, ORF73 (LANA), ORF74, and M11 with constitutively expressed RTA
98 driven by the phosphoglycerate kinase 1 (PGK) promoter in a two-tiered approach to
99 prevent persistent infection. Deletion of the latency locus, constitutive RTA expression,
100 and the removal of immune evasion genes created a live attenuated γ -herpesvirus
101 vaccine named DIP (deficient in immune evasion and persistence) (Fig. 1A).

102

103 ***DIP replication is attenuated in vitro***

104 Comparison of the *in vitro* growth kinetics of DIP in NIH3T3 fibroblasts with the wild type
105 (WT) virus showed that DIP replication was significantly attenuated. After infection at MOI
106 of 0.01, DIP yielded 300-fold and 40-fold less viral production than WT at 48 h and 72 h
107 post-infection, respectively (Fig. 1B). Pretreatment with IFN- β inhibited replication of WT
108 by 10-fold and DIP by 100-fold (Fig. 1C). This larger decrease in infectious DIP virion
109 production confirmed augmented susceptibility to the IFN-I response in the absence of
110 viral IFN evasion genes.

111

112 ***DIP produces no infectious virions in vivo***

113 We hypothesized that removal of the viral IFN-I evasion genes would generate a highly
114 attenuated vaccine *in vivo*. To test this, we infected C57Bl/6 mice intraperitoneally and
115 harvested their spleens 3 d after infection. While the WT virus produced 88 PFU/spleen,
116 no infectious virus was detected in the spleens of DIP-inoculated mice (Fig. 2A). We also
117 harvested spleens at later times post-infection and assessed spontaneously reactivating
118 virus by the infectious center assay and quantified latent viral genomes by qPCR. Viral
119 reactivation or latent virus was undetectable in the spleens of DIP-infected mice at 14 d
120 (Figs. 2B and 2C) and at 2 mo (Figs. 2E and 2F) after infection. Furthermore, no infectious
121 virion production in the lungs or latency establishment was observed in the spleens after
122 intranasal inoculation (Figs. S1A-D).

123

124 Latency establishment is associated with the expansion of V β 4-specific T-cells and
125 splenomegaly²⁸⁻³⁰. At 14 d post-infection, the spleens of WT-infected mice increased to
126 0.22 g on average while those of DIP-infected mice weighed 0.10 g (Fig. 2D), similar to
127 uninfected mice. At 2 mo post-infection, WT-infected mice still had significantly enlarged
128 spleens compared to those of DIP-inoculated mice (Fig. 2G).

129

130 To determine whether the IFN-I response contributed to the attenuation of the DIP virus,
131 we injected 10⁵ PFU of DIP intraperitoneally into the interferon- α/β receptor-deficient
132 (IFNAR $\alpha/\beta^{-/-}$) mice. DIP replication was rescued and 4 x 10⁵ infectious virions were
133 detected in the spleens at 3d post-infection (Fig. 2H). Infectious virions were also detected
134 in the livers and lungs but not the brains of IFNAR1^{-/-} mice. In contrast, no detectable
135 infectious virions were recovered from severe combined immune deficiency (SCID) mice.
136 Comprehensive analyses of the spleens, livers, brains, and lungs showed no evidence of
137 infectious virions in either C57BL/6 or SCID mice, both of which have intact IFN-I
138 responses (Fig. 2H).

139

140 ***DIP immunization prevents latent infection***

141 γ -herpesvirus associated malignancies are linked to latency^{31,32}. Therefore, the goal of
142 vaccination against γ -herpesviruses is to prevent latency establishment. We assessed
143 the level of protection conferred by DIP immunization against latent infection by WT
144 challenge. Mice were intraperitoneally injected with 1 x 10⁵ PFU DIP then intranasally
145 challenged 1 mo later with 5,000 PFU WT. Mock immunized mice presented an average
146 of 6 x 10² infectious centers per 2 x 10⁷ splenocytes whereas no viral reactivation was

147 detected in six of seven DIP-immunized mice 14 d after challenge (Fig. 3A). Analysis of
148 viral copy number confirmed that DIP immunization provided protection against splenic
149 latent infection (Fig. 3B). At 1 mo post-challenge, five of six (83.3%) DIP-vaccinated mice
150 were completely protected from latent infection (Fig. 3C). The remaining DIP-vaccinated
151 mouse had a 100-fold reduction in latently infected cells compared to mock immunized
152 mice. We also challenged the immunized mice 6 mo after a single vaccination and
153 measured viral latency at 28 d post-challenge. All six DIP-immunized mice were
154 completely protected against latent infection by a WT virus challenge (Fig. 3D).

155

156 ***DIP primes virus-specific T cells that limit WT infection***

157 We hypothesized that the DIP vaccine elicited a robust and functional T cell response
158 accounting for the long-lasting protection against WT challenge. We quantified virus-
159 specific CD8⁺ T-cells using tetramers for the MHV-68 epitopes, ORF6₄₈₇₋₄₉₅ and
160 ORF61₅₂₄₋₅₃₁. At 1 mo post-infection, WT and DIP induced similar frequencies of specific
161 T-cells to ORF6 and ORF61 (Figs. S2A & S2B). At 2 mo post-infection, the frequency of
162 ORF6-specific T cells increased two-fold in DIP compared to WT while the frequency of
163 ORF61-specific T-cells were similar (Figs. 4A & 4B). The effector/memory subtypes of
164 these virus-specific T cells were examined by the expression levels of IL7R α (CD127)
165 and killer cell lectin-like receptor (KLRG1) (Fig. S2D). The CD127^{high}KLRG1^{low} subset are
166 memory precursors effector cells (MPECs), which develop into long-lived memory cells,
167 whereas the CD127^{low}KLRG1^{high} subset, referred to as short-lived effector T cells
168 (SLECs), are terminally differentiated³³. We observed that DIP promoted the generation
169 of CD8⁺ MPECs. Significantly more ORF6-specific CD8⁺ T cells (54%) primed by DIP

170 differentiated into MPECs compared to those primed by WT (33%). However, no
171 difference was found between WT and DIP infection in terms of ORF61-specific T cells
172 (Figs. 4C and S2C).

173

174 We assessed the functions of these virus-specific T-cells by examining their abilities to
175 produce IFN- γ , TNF- α , and IL-2. Consistent with the tetramer-staining results, cells
176 producing IFN- γ , TNF- α , or IL-2 upon stimulation of the ORF6 peptide were twice as
177 frequent in DIP-infected mice as in WT-infected mice (Figs. 4D, 4F, & 4H). Cells producing
178 IFN- γ upon stimulation of the ORF61 peptide were at similar frequencies between WT
179 and DIP infection. However, a lower frequency of cells primed by DIP produced TNF- α in
180 response to the ORF61 peptide compared to those primed by WT (Figs. 4E & 4F). The
181 ORF61 peptide did not stimulate any cells from either WT or DIP-infected mice to produce
182 IL-2 (Fig. 4I). Therefore, despite its limited and transient antigen expression due to highly
183 attenuated replication, DIP still induces robust and functional T-cell responses.

184

185 To determine whether DIP-primed T-cells confer protection against WT challenge, we
186 harvested CD4⁺, CD8⁺ or total T-cells from mice infected 2 mo earlier with WT or DIP and
187 transferred 3×10^6 cells into naïve mice. These recipient mice were challenged with 5000
188 PFU WT 1 d after transfer. No significant difference was observed between WT- and DIP-
189 primed T cells in terms of donor cell expansion (Fig. S3). At 14d post-challenge, CD4⁺ T-
190 cell transfer had minimal impact on the number of latently infected cells (Fig. 5A) despite
191 evidence that CD4⁺ T cells are cytotoxic to herpesviruses^{34–36}. The transfer of WT-primed
192 CD8⁺ T-cells caused a five-fold reduction in reactivated latently infected cells. In contrast,

193 CD8+ T-cells primed by DIP failed to affect the latently infected cell pool (Fig. 5B).
194 However, naïve mice receiving DIP-primed total T-cells had a 30-fold reduction in the
195 number of reactivated latently infected cells, whereas transferring of WT-primed total T
196 cells caused a 20-fold reduction (Fig. 5C). The results indicate that virus-specific CD4+
197 and CD8+ T-cells act cooperatively to confer protection. Despite severe attenuation, DIP
198 vaccination elicited robust cellular immunity that inhibits the establishment of latency by
199 the challenge virus.

200

201 ***Optimal DIP-mediated protection requires both antibodies and T cells***

202 To determine whether DIP-elicited antibodies complemented the T-cell-mediated
203 protection, serum and total T-cell from DIP-infected mice were transferred to naïve mice.
204 This combination completely protected four of six mice against a 5000 PFU WT challenge
205 (Fig. 6A). The two unprotected mice had a significantly reduced number of reactivating
206 latently infected cells compared to the control. We examined the protective capacity of
207 antibodies by passively transferring DIP-immune serum to naïve mice. No significant
208 difference in protection was observed between those receiving DIP-immune serum and
209 those receiving serum from mock infected mice (Fig. 6B). Neutralizing activity in DIP-
210 immune serum was less than that in WT-immune serum at 2 mo post-infection (Fig. 6D)
211 despite relatively higher levels of virus-specific IgG in the serum DIP-infected mice (Fig.
212 6C). These results indicate that DIP-elicited humoral immunity collaborates with cellular
213 immunity to provide optimal protection.

214

215 ***DIP vaccine elicits robust inflammatory responses***

216 Despite its limited replication, DIP primed robust virus-specific immune responses and
217 conferred durable protection. Activation of innate immune response is essential for the
218 development of adaptive immunity⁵⁴⁻⁵⁶. WT MHV-68 avoids inducing inflammatory
219 cytokines in order to evade the immune system. *In vitro*, a high MOI (100 PFU/cell) was
220 required to elicit a measurable cytokine response in bone marrow-derived macrophages
221 (BMDM) and dendritic cells³⁷. We investigated whether DIP induces inflammatory
222 cytokines. BMDMs were infected with WT and DIP at MOI of 1 and cytokine RNA
223 expression was quantified at 24 h post-infection. IFN- β , TNF- α , IL-6, and IL-12p40 were
224 significantly upregulated in response to DIP infection compared to WT (Fig. 7A). In
225 addition, WT infection at MOI of 10 still did not induce cytokine expression (Fig. S4). IL-
226 12 is critical for Th1 polarization and cytotoxic cellular immune responses³⁸. We validated
227 the IL-12p40 RNA expression by measuring the protein with enzyme-linked
228 immunosorbent assay (ELISA). DIP induced 30-fold more IL-12p40 protein than WT
229 infection (Fig. 7B). The ability of DIP to stimulate the innate immune responses *in vivo*
230 was also determined. Two days after intraperitoneal injections of viruses, there were five
231 times as many peritoneal exudate cells (PECs) in DIP-infected as in mock-infected mice
232 and significantly more cells than in WT-infected mice (Fig. 7C). Flow cytometry analysis
233 of cellular compositions revealed that DIP significantly induced more plasmacytoid DCs
234 (pDCs) than WT (Fig. 7D). As pDCs produce IFN-I³⁹, we also detected the upregulation
235 of ISG54 and IFIT2 in the PECs of DIP-infected mice compared to those of WT-infected
236 mice (Fig. 7E). Taken together, the foregoing results indicate that DIP is highly efficacious
237 at inducing inflammatory responses.

238 **Discussion**

239 An effective γ -herpesvirus vaccine should protect against the establishment of latency
240 given the association between latent infection and tumorigenesis. Several vaccine
241 strategies targeting single viral antigens were previously tested in the MHV-68 mouse
242 infection model. These antigens reduced infectious mononucleosis-like symptoms of
243 lymphoproliferation but failed to limit establishment of latency⁴⁰⁻⁴⁵. This finding resembles
244 that reported for a clinical trial of EBV gp350-based vaccines⁹. The only vaccine strategy
245 proven to reduce long-term latent viral loads in the MHV-68 model was based on live
246 attenuated viruses designed to be latency-deficient^{27,46-50}. However, a major drawback of
247 latency-deficient viruses is the ability to undergo lytic replication⁵¹. In this study, we tested
248 a strategy to attenuate the *in vivo* replication of the vaccine virus by inactivating viral
249 antagonists of the IFN-I response. IFN-I is the first line in host antiviral defense and is
250 critical in the development of effective immune responses. IFNs bridge innate and
251 adaptive immunity by activating dendritic cells and inducing Th1 and potent antibody
252 responses⁵²⁻⁵⁵. IFN-I has been tested as an adjuvant for vaccines against several distinct
253 viruses including influenza, HIV, Ebola, CMV, and γ -herpesviruses⁵⁶⁻⁶⁰. Interestingly,
254 Aricò et al. used MHV-68 to demonstrate an increase in viral-specific antibody titers when
255 heat-inactivated virus was co-administered with IFN- α /- β ⁵⁶. We proposed that disarming
256 viral IFN-I evasion genes may facilitate the IFN-I response, providing the adjuvanticity
257 required for attenuated viral vaccines. In addition, we also inactivated viral inhibitor of
258 MHC class I presentation pathway and deleted the latency locus to increase the
259 immunogenicity and safety of the vaccine virus, DIP. The present study demonstrates
260 that DIP is highly attenuated yet maintains overall immunogenicity relatively similar to WT.

261 DIP cannot undergo productive infection or persist *in vivo*. Despite its attenuated
262 replication, DIP elicits robust innate immune responses (e.g. IL-12), memory T cells, and
263 virus-specific antibodies with neutralizing activity. Single DIP vaccination protected
264 against latency establishment following WT challenge. DIP-mediated protection was
265 durable as all immunized mice remained fully protected even 6 months after a single
266 vaccination.

267

268 Antibodies represent the first line of vaccine-mediated protection. Nevertheless, the ideal
269 prophylactic vaccine also induces protective cellular immunity. The adoptive transfer
270 experiments revealed that DIP-induced, virus-specific T cells and antibodies complement
271 each other to provide optimal protection (Fig. 6A). While CD4+ T-cells alone have little
272 protective capacity, they collaborate with CD8+ T-cells to provide protection (Fig. 5). This
273 CD4-CD8 collaboration occurred with either WT- or DIP-primed T-cells indicating the
274 capability of DIP to elicit effective adaptive immunity. It is recognized that CD4+ T-cells
275 optimize the development and maintenance of memory CD8+ T-cells^{61,62}. However, it is
276 unclear whether memory CD8+ T-cells absolutely require this help from CD4+ T-cells or
277 are simply enhanced by them⁶³. The expansion of donor CD8+ T-cells in mice receiving
278 total T cells did not surpass that in mice only receiving CD8+ T-cells (Fig. S3). Previous
279 work indicated that memory CD4+ T-cells enhance the functionality of memory CD8+ T-
280 cells⁶⁴⁻⁶⁶ but this enhancement was not examined in this study. Furthermore, the effect
281 of memory CD4+ T-cells observed here may not have been mediated by enhancing
282 memory CD8+ T-cells responses. Rather, CD4+ and CD8+ T-cells may target different
283 infected cells, complementing each other, to provide effective protection against latency

284 establishment in response to WT challenge. The mechanisms underlying the T-cell
285 collaboration identified herein merit further investigation. It is clear that a prophylactic
286 vaccine against γ -herpesvirus should prime both memory CD8+ and CD4+ T-cells.

287

288 A live viral vaccine induces a broad immune response against multiple viral targets
289 especially when the mechanisms required for protection against a pathogen are not
290 known. Furthermore, by mimicking an infection, a live vaccine stimulates multiple innate
291 immune responses, robustly induces inflammatory and immunomodulatory cytokines,
292 and provides adjuvanticity for long-lasting vaccine-mediated protective immunity.
293 Nevertheless, most viruses have evolved strategies to counteract the host innate immune
294 system. Data from BMDMs infected *in vitro* and PECs from infected mice indicates that
295 DIP induces a stronger inflammatory response than WT (Fig. 7), partially accounting for
296 the effective immunogenicity of DIP. DIP also recruits more pDCs than WT, which could
297 also explain the ability of DIP to elicit robust CD8+ T-cells and antibody responses in spite
298 of its highly attenuated replication⁶⁷⁻⁷⁰. Future experiments examining the role of specific
299 cytokines or pDCs via genetic knockout and antibody depletion approaches may reveal
300 how DIP-induced innate immune responses impact humoral and cellular immunity.
301 Increased inflammatory cytokine production favors SLEC generation whereas shortening
302 the duration of inflammation has been shown to accelerate MPEC development^{71,72}. DIP-
303 mediated inflammatory responses could be short-lived as DIP is highly attenuated *in vivo*.
304 The heightened but transient DIP-induced inflammation appears to prime a robust T-cell
305 response towards the MPEC phenotype.

306

307 The development of vaccines against human γ -herpesviruses has been hindered by their
308 restricted host range. Neither EBV or KSHV infects small animals. While the results from
309 mouse studies are not always directly translatable to humans, mouse models have been
310 instrumental in elucidating fundamental principles that cannot be directly tested in
311 humans. MHV-68 infection in mice provides a powerful, easily manipulated small animal
312 model for analyzing fundamental events associated with the infection and immune control
313 of γ -herpesviruses⁷³⁻⁷⁸. Moreover, the MHV-68 model serves to assess proof-of-concept
314 vaccine strategies⁷⁹. The results from the present study provides the guidance for a
315 rational design of effective live EBV and KSHV vaccines that are highly attenuated and
316 deficient in latency. Deletion of viral immune evasion genes may provide a strategy for
317 the construction of safe yet immunogenic live vaccines against other pathogens.
318

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332

333 **Author contributions**

334 G.B., N.F., R.S., and T.T.W. conceived and planned the experiments. G.B. and N.F.
335 carried out the experiment with the help of A.S., A.K.L., W.W.L., L.T., Y.H.K., and T.H.
336 G.B., N.F., R.S., and T.T.W. analyzed and interpreted the results. C.F.W., and M.A.B.
337 provided critical scientific advice to the research described in this manuscript. G.B., N.F.,
338 and T.T.W. wrote the manuscript with inputs from A.K.L., W.W.L., M.A.B., C.F.W., and
339 R.S.

340

341 **Author information**

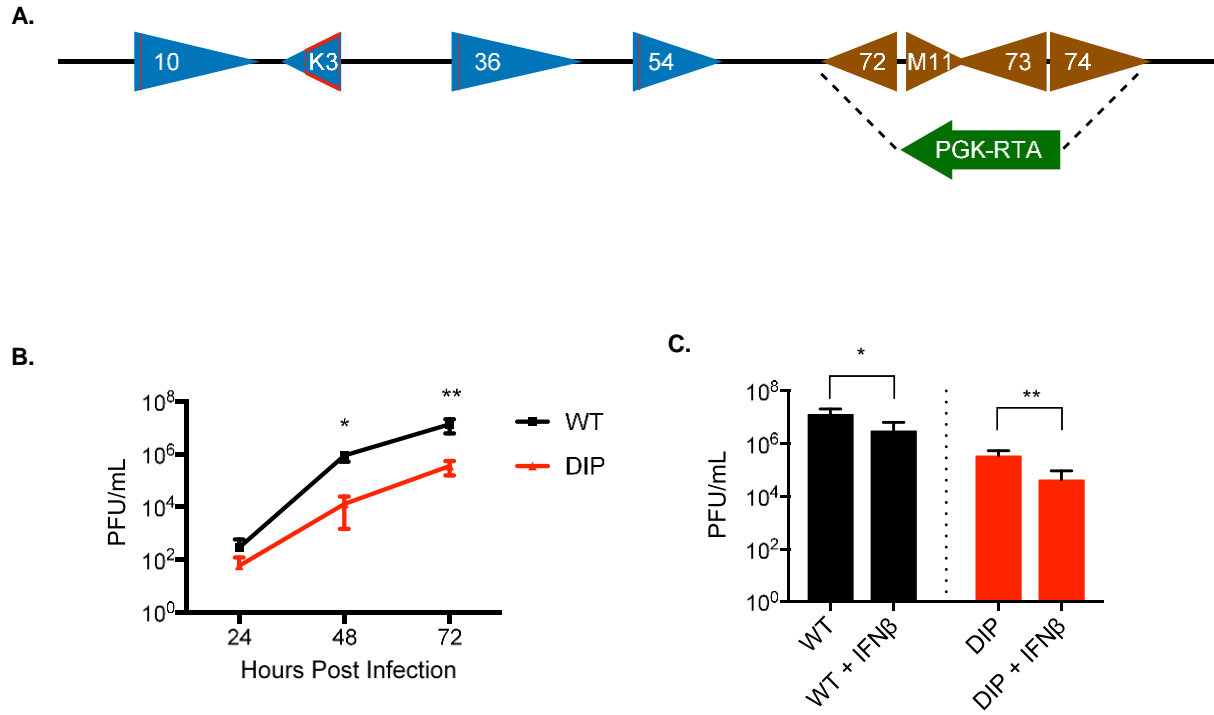
342 Correspondence and requests for materials should be addressed to T.T.W
343 (twu@mednet.ucla.edu).

344

345 **Competing interests**

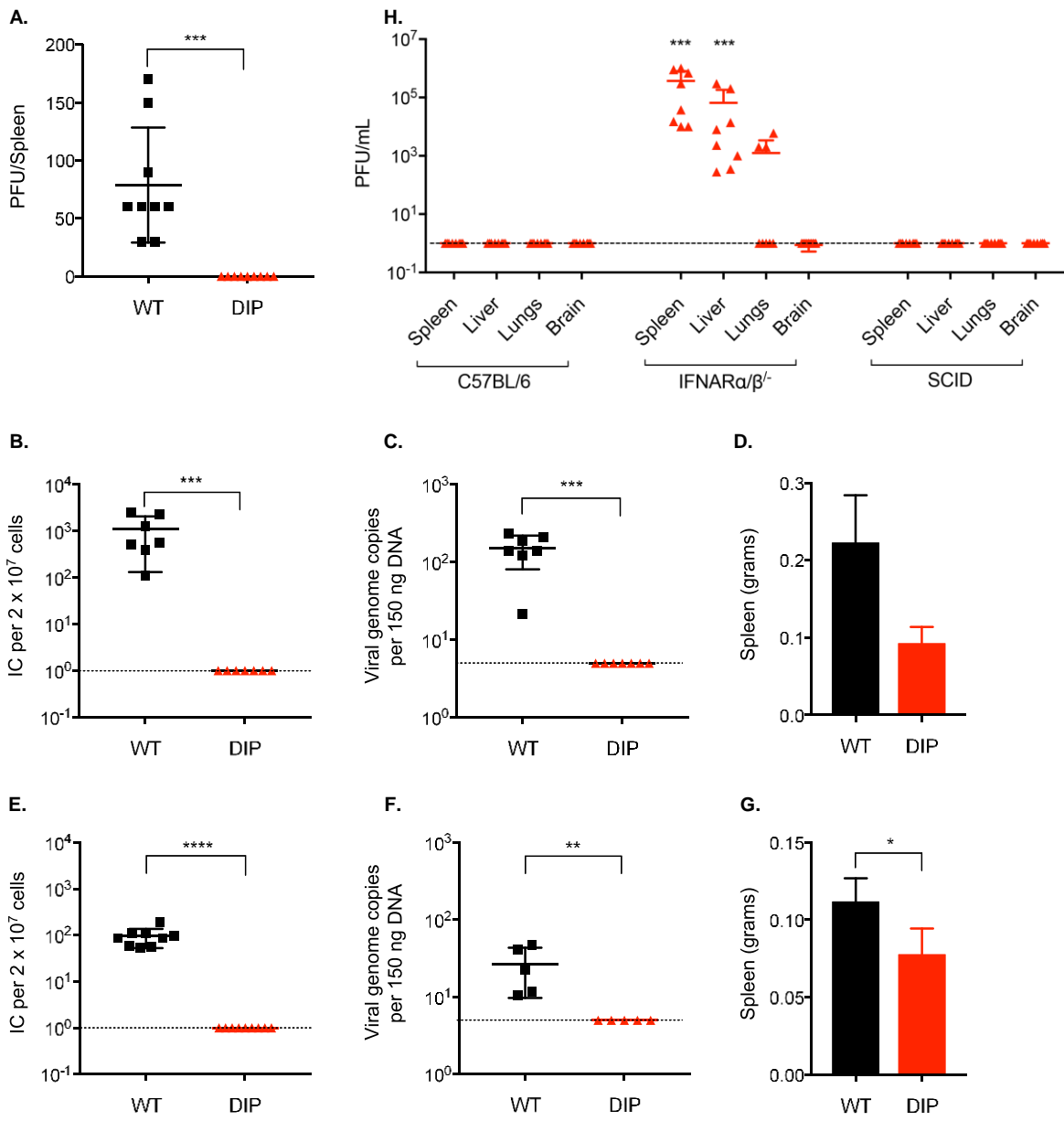
346 The authors declare no competing financial interests.

Figure 1



349 **Figure 1. Construction of DIP virus and its replication properties in vitro**
350 **(A)** Schematic representation of mutations introduced in the MHV-68 genome to generate
351 the DIP vaccine. Red lines indicate insertion of translation stop codons into ORF10,
352 ORF36, and ORF54. The open red tetragon indicates deletion of the coding sequence in
353 K3. The latency locus was replaced by the RTA cassette (arrowhead) constitutively driven
354 by the PGK promoter. **(B)** Growth curves of the WT and DIP viruses in 3T3 cells using
355 MOI = 0.01 and measured by plaque assay to quantify virion production. **(C)** NIH 3T3
356 cells were either mock treated or treated with 100 U mL⁻¹ IFN-β for 24 h then infected with
357 either WT or DIP virus at MOI = 0.01 for 72 h. Virion production was quantified with plaque
358 assays. All experiments were performed in triplicate and statistical significance was
359 analyzed by a two-tailed Student's *t*-test. Graphs represent means of triplicates with SD.
360

Figure 2

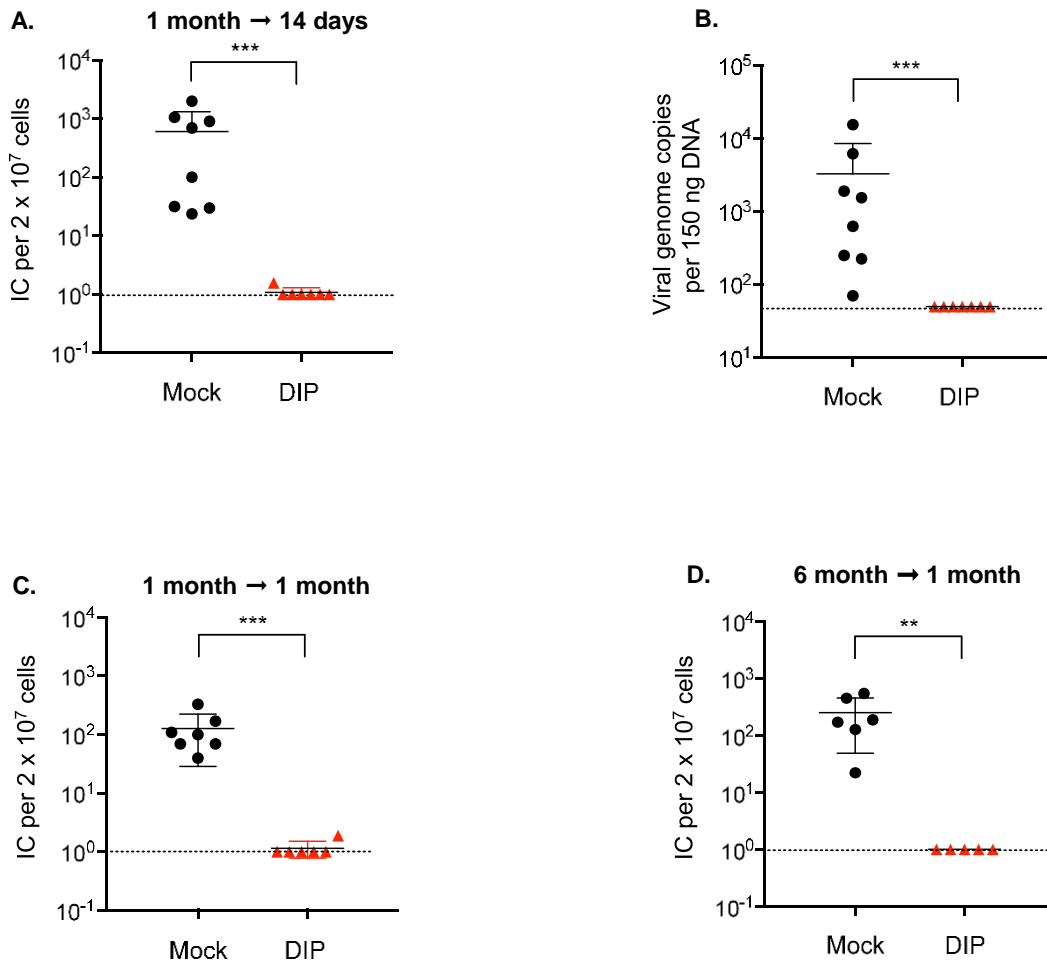


362 **Figure 2. DIP produces no infectious virions and is latency deficient in vivo**

363 All infections were performed intraperitoneally using 10^5 PFU WT or DIP. (A) Productive
364 infection in the spleens 72 h post-infection was assessed by plaque assay. (B) Latent
365 infection in the spleens at 14 d post-infection was evaluated by infectious center
366 assay and (C) qPCR analysis of viral DNA copy numbers. (D) Spleen weight at 14 d post-
367 infection was measured. No statistically significant difference was found between WT-
368 and DIP-infected mice. (E) Latent infection in the spleen at 2 mo post-infection was
369 measured by infectious center assay and (F) qPCR analysis of viral DNA copy numbers.
370 (G) Spleen weight at 2 mo post-infection was measured. (H) Spleens, livers, lungs, and
371 brains of DIP infected C57BL/6, IFNAR^{-/-}, and SCID mice were harvested at 3 d post-
372 infection. Infectious viruses were determined by plaque assay. The graphs except (A)
373 depicts the pooled data from 2 independent experiments using different numbers of mice
374 for each replicate. Symbols indicate individual mice and data are means and SD.
375 Statistical significance was determined by a two-tailed Student's *t*-test.

376

Figure 3

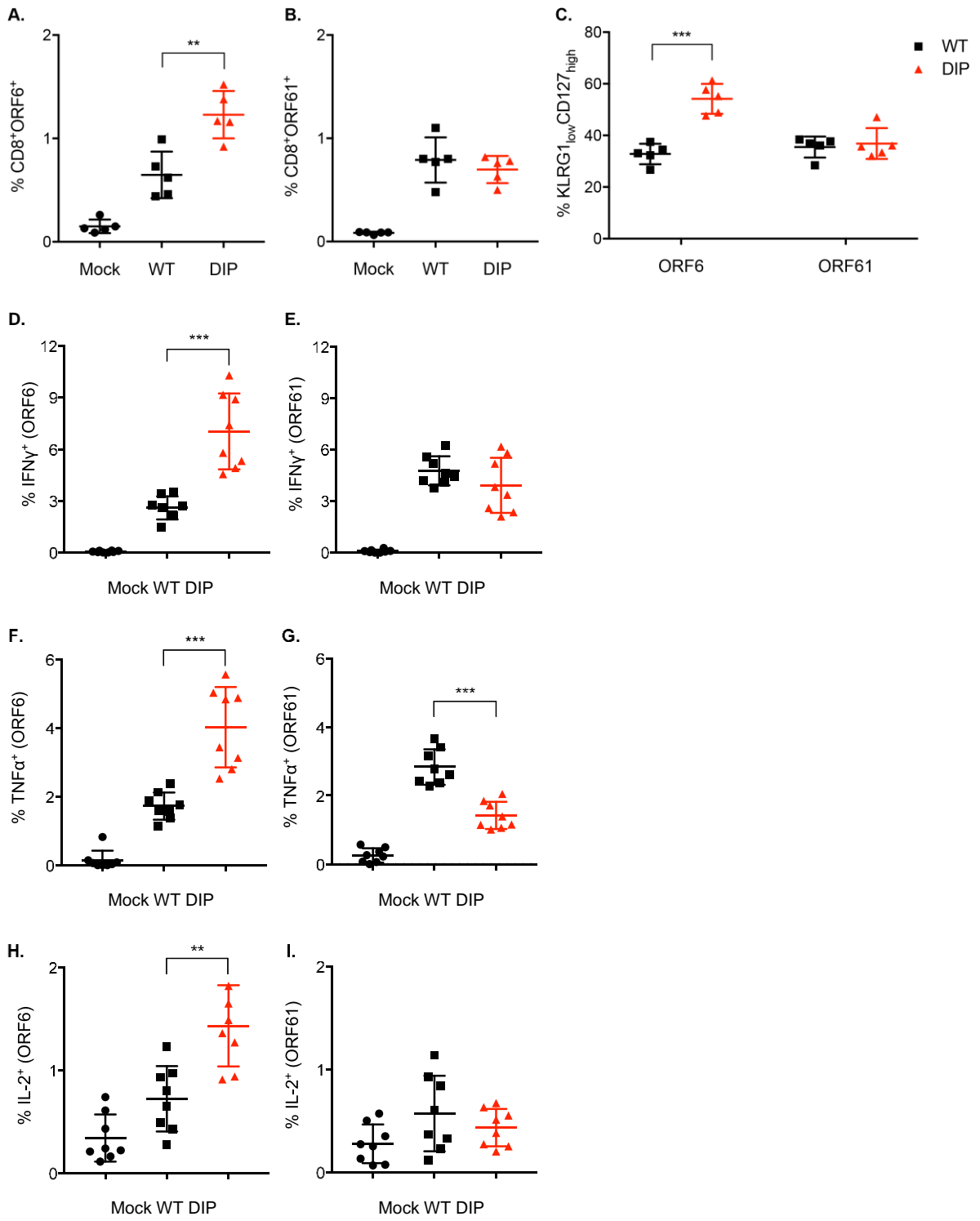


378 **Figure 3. DIP vaccination confers durable protection**

379 Mice were intraperitoneally vaccinated with 10^5 PFU DIP and challenged intranasally with
380 5×10^3 PFU WT virus at 1 (**A-C**) or 6 (**D**) mo post-vaccination. Latent infection in the
381 spleen was examined at 14 (**A, B**) or 28 (**C, D**) d after challenge. Viral loads were
382 determined by infectious center assay (**A, C, D**) and qPCR (**B**). Dotted line indicates
383 detection limit. The graph depicts the pooled data from 2 independent experiments using
384 different numbers of mice for each replicate. Data for individual mice, means, and SD
385 were plotted. Statistical significance was analyzed by a two-tailed Student's *t*-test.

386

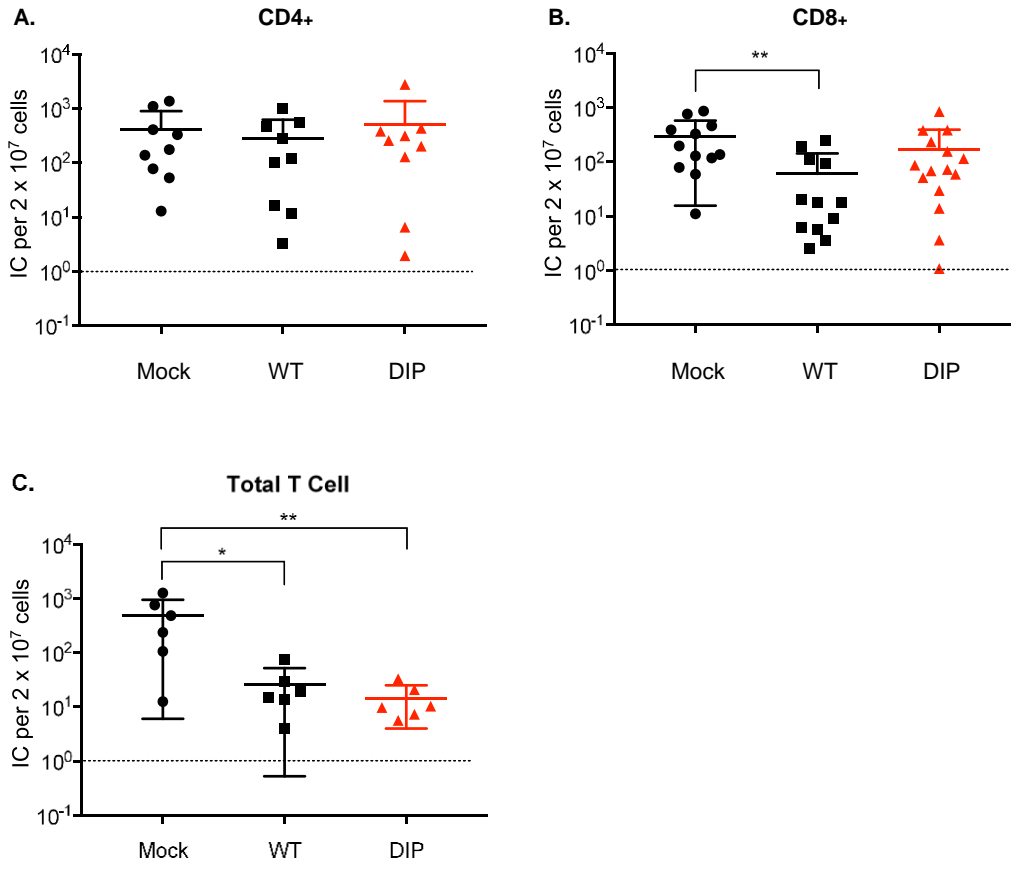
Figure 4



388 **Figure 4. DIP elicits robust virus-specific T cell immunity**

389 Mice were mock-infected or intraperitoneally injected with 10^5 PFU WT or DIP. **(A, B)** At
390 2 mo post-infection, splenocytes were harvested and examined for virus-specific CD8+ T
391 cells using the tetramers ORF6₄₈₇₋₄₉₅/Db and ORF61₅₂₄₋₅₃₁/Kb. **(C)** Tetramer-positive
392 CD8+ T cells were examined for KLRG1 and CD127 expression. **(D-I)** Splenocytes were
393 stimulated with ORF6₄₈₇₋₄₉₅ peptide **(D, F, H)** or ORF61₅₂₄₋₅₃₁ peptide **(E, G, I)** and stained
394 for intracellular IFN- γ **(D, E)**, TNF- α **(F, G)**, and IL-2 **(H, I)**.
395

Figure 5

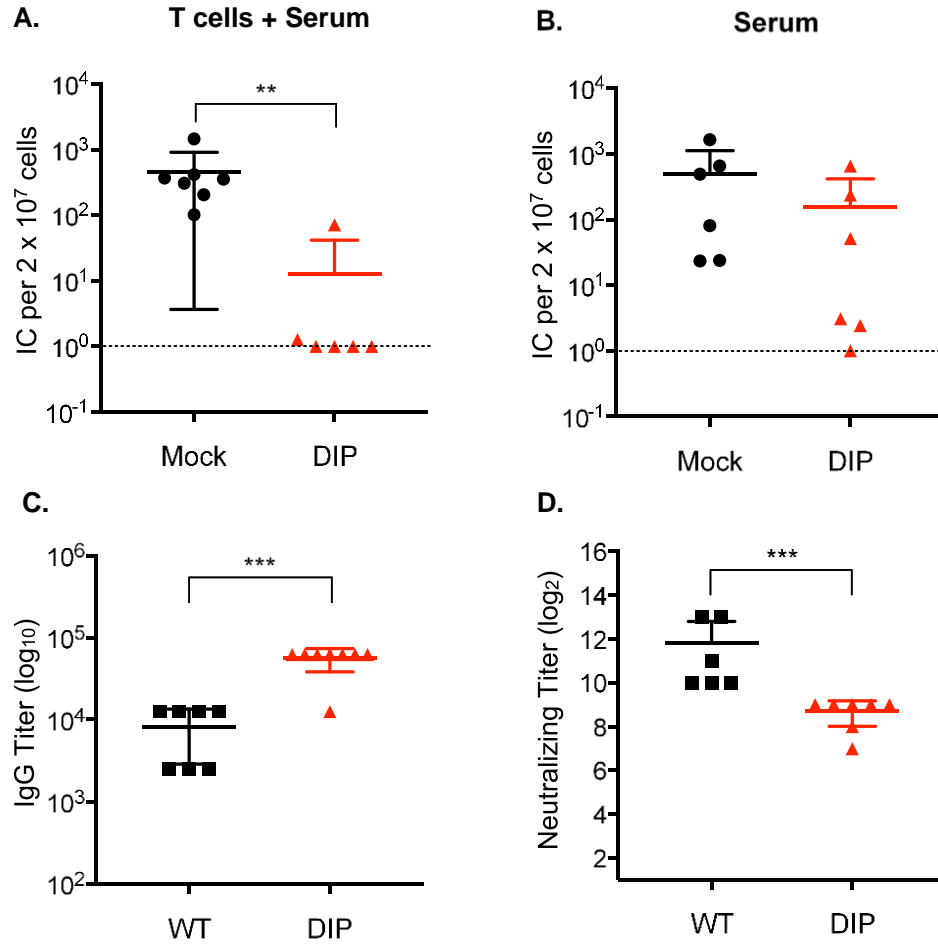


397 **Figure 5. CD4+ and CD8+ T-cells confer antiviral protection**

398 CD4+, CD8+, or total T cells were purified via negative selection from the spleens of
399 mock-infected mice or mice that were intraperitoneally infected with 10^5 PFU WT or DIP
400 2 mo previously. Three million CD4+ (**A**), CD8+ (**B**), or total T (**C**) cells were transferred
401 to a congenic mouse by tail vein injection. The recipient mice were intranasally challenged
402 with 5×10^3 PFU WT at 24 h post-transfer. Latent infection in the spleen at 14 d post
403 challenge was measured by infectious center assay. Pooled data from 2 independent
404 experiments using different numbers of mice for each replicate. Data for individual mice,
405 means, and SD were plotted. Statistical significance was analyzed by a two-tailed
406 Student's *t*-test.

407

Figure 6

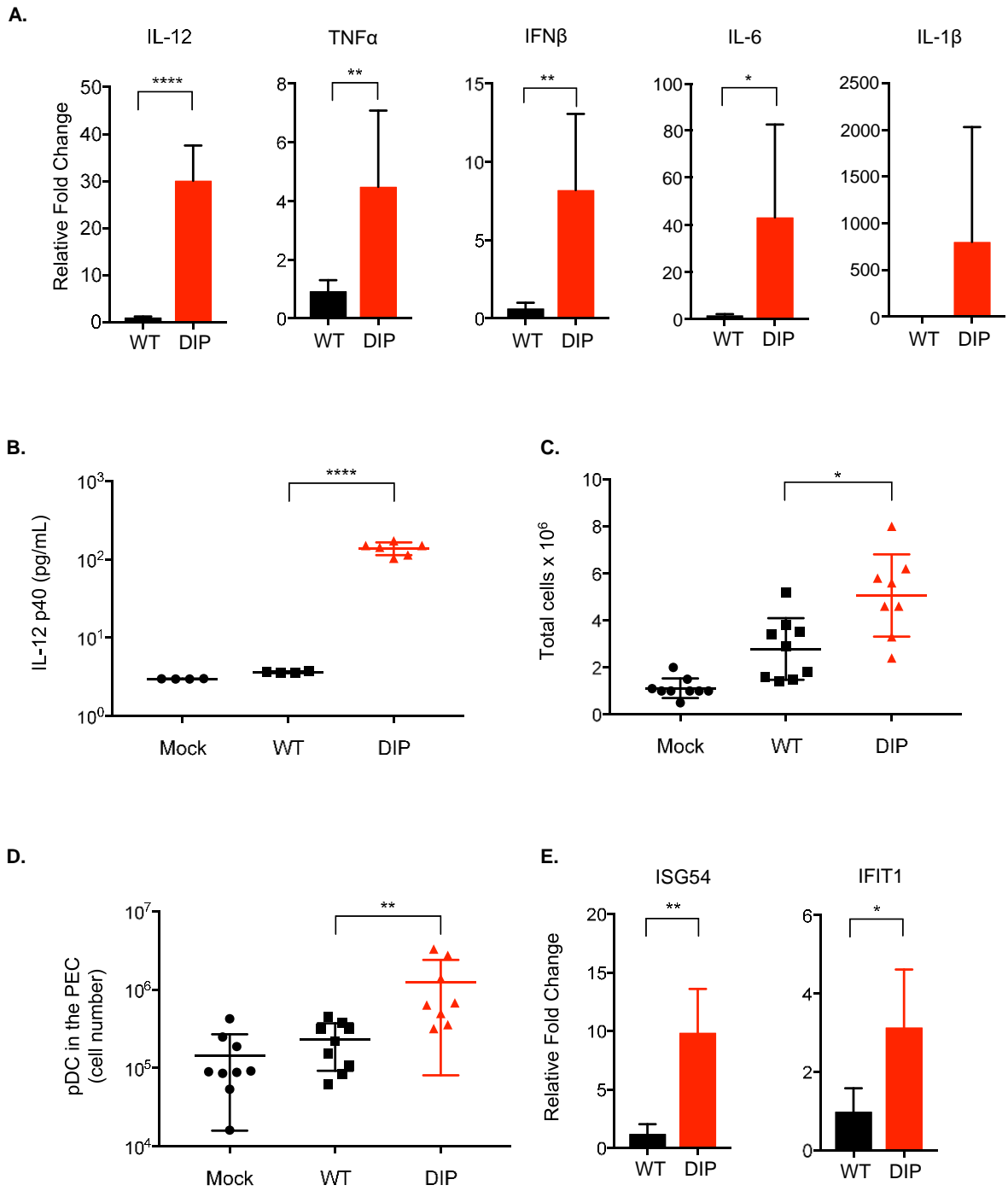


409 **Figure 6. DIP vaccination elicits protective antibodies**

410 Mice were intraperitoneally infected with 10^5 PFU WT or DIP 2 mo previously. **(A)** Total
411 T cells and sera isolated from mock- or DIP-infected mice were transferred to congenic
412 naïve mice by tail vein and intraperitoneal injections, respectively. Recipient mice were
413 intranasally challenged 24 h later with 5×10^3 PFU WT virus. Latent infection in the spleen
414 at 14 d post-challenge was assessed by infectious center assay. **(B)** Sera collected from
415 uninfected- or DIP-infected mice were transferred to naïve mice that were intranasally
416 challenged 24 h later with 5×10^3 PFU WT virus. Latent infection in the spleen at 14 d
417 post-challenge was evaluated by infectious center assay. **(C, D)** Sera collected from
418 infected mice were analyzed for virus-specific IgG by ELISA and for neutralizing activity.
419 Pooled data from 2 independent experiments using different numbers of mice for each
420 replicate. Means and SD were plotted. Statistical significance was analyzed by a two-
421 tailed Student's *t*-test.

422

Figure 7



424 **Figure 7. DIP elicits inflammatory and immunomodulatory cytokines**

425 Mouse BMDMs were infected with WT or DIP at MOI = 1 (triplicate). **(A)** Total RNA was
426 extracted 24 h post-infection for reverse transcription and qPCR to measure the
427 expression levels of IFN- β , IL-1 β , TNF- α , IL-6, IL-12, and β -actin. Cytokine RNA
428 expression was normalized against β -actin and the relative fold change was calculated
429 by comparison with mock-infected BMDM. **(B)** Supernatants were collected 24 h post-
430 infection to measure IL-12p40 production by ELISA. Mice were either mock-infected or
431 intraperitoneally injected with 10^5 PFU WT or DIP. PECs were collected at 48 h post-
432 infection. **(C)** Total cell numbers in the PECs were counted. **(D)** The pDCs were identified
433 by gating on the Lin⁻(CD3⁻CD19⁻NK1.1⁻)B220⁺CD11c^{Int}PDCA-1⁺ population. **(E)** Total
434 RNA was extracted from the PECs. RNA expression of ISGs was analyzed by quantitative
435 PCR. Means and SD were plotted. Statistical significance was analyzed by a two-tailed
436 Student's *t*-test.

437 **Materials & Methods**

438

439 ***Viruses and cells***

440 WT MHV-68 was obtained from the American Type Culture Collection (ATCC; Vr1465;
441 Manassas, VA, USA). WT and DIP viruses were propagated in 3T3 and Vero cells and
442 titered by plaque assay. Viruses were concentrated by high-speed centrifugation and
443 resuspended in serum-free Dulbecco's modified Eagle's medium (DMEM). Vero cells
444 were cultured in DMEM containing 10% (w/v) fetal bovine serum (FBS) supplemented
445 with penicillin and streptomycin. The 3T3 cells were cultured in DMEM containing 10%
446 (w/v) bovine calf serum (BCS) and 1% penicillin and streptomycin.

447

448 ***Plaque assay***

449 Each sample was serially diluted tenfold and incubated on Vero cells on 12-well plates in
450 duplicate. The inoculum was removed after 1 h of incubation and the cells were overlaid
451 with 1% (w/v) methylcellulose in DMEM containing 10% (w/v) FBS. Six days post-
452 infection, the cells were fixed with 2% (w/v) crystal violet in 20% (v/v) ethanol. Viral titers
453 were determined by counting plaque numbers. To determine viral titers in the mouse
454 tissues, 1-mL homogenates were prepared in a Dounce homogenizer (Thomas Scientific,
455 Swedesboro, NJ, USA) and used for the plaque assay. Plaques were counted and viral
456 titers in each tissue were expressed in PFU mL⁻¹.

457

458 ***In vitro growth curve***

459 The 3T3 cells were plated on media with or without IFN- β (100 U mL⁻¹) for 24 h. Cells
460 were infected at MOI = 0.01 with WT or DIP virus for 1 h at 37 °C. The inoculum was then
461 removed and the cells were washed twice with media before adding fresh media with or
462 without IFN- β (100 U mL⁻¹). Cells and supernatant were harvested 24 h, 48 h, and 72 h
463 post-infection for the plaque assay.

464

465 ***Construction of DIP vaccine***

466 The recA+ *Escherichia coli* GS500 harboring a BAC containing the WT MHV-68 genome
467 was used to construct recombinant MHV-68 by allelic exchange with conjugation-
468 competent *E. coli* GS111 containing the suicide shuttle plasmid pGS284⁷⁴⁻⁷⁶. For each
469 recombinant MHV-68, an overlap extension PCR was used to construct the unique shuttle
470 plasmid pGS284 harboring the desired mutation and a ~500-bp flanking region.
471 Sequences upstream of the desired mutation (A fragments) were amplified by AF and AR
472 primers. The downstream sequences (B fragments) were amplified by BF and BR primers
473 using wild type MHV-68 virion DNA as the template. The A and B fragments had > 20-bp
474 overlapping sequences. For the subsequent PCR reaction, the A and B fragments were
475 used as templates and amplified by AF and BR primers. The final PCR products were
476 digested with the appropriate enzymes and cloned into pGS284. To screen for the correct
477 mutation, restriction enzyme digestion was performed on the PCR products obtained
478 using the AF and BR primers on the BAC MHV-68 clones. Sequential allelic exchanges
479 were conducted to obtain the final recombinant clone containing all the designed
480 mutations (Fig. 1). After the desired recombinant clone was selected, the MHV-68 BAC
481 was purified and transiently transfected with Lipofectamine™ 2000 into 293T cells with

482 equal amounts of plasmid expressing Cre recombinase to remove the BAC sequence.
483 Three days post transfection, a single viral clone was isolated by limiting dilution. It was
484 then propagated for use in subsequent experiments. The viruses were quantified by
485 plaque assay and limiting dilution.

486 The ORF36 and ORF54 primers were used to construct the shuttle plasmids^{18,80}. Primers
487 used to construct the other shuttle plasmids are listed in Supplementary Table 1. Primers
488 1-8 were used to construct shuttle plasmids for the stop codon mutation. To construct
489 the shuttle plasmid to replace the latency locus with RTA expression driven by the PGK
490 promoter, four fragments were amplified with primers 9-16, A (ORF72), B (RTA coding
491 sequence and poly A tail), C (PGK promoter), and D (ORF74). The ABCD fused fragment
492 was then generated to be cloned into pGS284.

493

494 ***Mice***

495 The animal studies were approved by the Animal Research Committee at the University
496 of California, Los Angeles (UCLA), Los Angeles, CA, USA. Female C57BL/6J, SCID, and
497 B6.SJL-*Ptprc^a Pepc^b*/BoyJ mice were obtained from Jackson Laboratory, Bar Harbor,
498 ME, USA. IFNAR^{-/-} mice were donated by Genhong Cheng at UCLA. Mice aged 6-8 wks
499 were intraperitoneally infected with 10⁵ PFU virus in 200 μL. Intranasal vaccinations and
500 challenges were performed by anesthetizing the mice with isoflurane and administering
501 20 μL virus dropwise. At the endpoint, mice were euthanized and their tissues were
502 collected in 1 mL DMEM and homogenized with mesh filters and a Dounce homogenizer.
503 Tissue lysates were clarified by centrifugation and used in the plaque assays. Their DNA
504 was extracted with a DNeasy blood and tissue kit (Cat. No. 69504; Qiagen, Hilden,

505 Germany). For the infectious center assay and the flow cytometry study, single-cell
506 suspensions were obtained from the spleens and the red blood corpuscles were lysed in
507 ACK (ammonium-chloride-potassium) buffer.

508

509 ***Phenotyping virus-specific T cells***

510 Before staining, the splenocytes were incubated with FC block (No. 553142; BD
511 Bioscience, Franklin Lakes, NJ, USA). Tetramers were obtained from the NIH Tetramer
512 Core Facility, Atlanta, GA, USA. Allophycocyanin-conjugated MHCI tetramers specific for
513 the MHV68 epitopes D^b/ORF6487–495 (AGPHNDMEI), K^b/ORF61524–531 (TSINFVKI),
514 K^b/ORF75c940–947 (KSLTYYYKL), and K^b/ORF8₆₀₄₋₆₁₂ (KNYIFEEKL) were incubated
515 with splenocytes for 1 h at room temperature. Surface-staining with the following
516 antibodies was performed by incubation at 4 °C for 30 min: anti-KLRG1 (No. 46-5893;
517 eBioscience/Affymetrix, Santa Clara, CA, USA), anti-CD127 (No. 17-1273;
518 eBioscience/Affymetrix, Santa Clara, CA, USA), anti-CD8 (No. 48-0081;
519 eBioscience/Affymetrix, Santa Clara, CA, USA), anti-CD4 (No. 11-0042;
520 eBioscience/Affymetrix, Santa Clara, CA, USA), anti-CD3 (No. 25-0031;
521 eBioscience/Affymetrix, Santa Clara, CA, USA), anti-CD44 (No. 11-0441;
522 eBioscience/Affymetrix, Santa Clara, CA, USA), anti-CD62L (No. 83-062;
523 eBioscience/Affymetrix, Santa Clara, CA, USA), anti-CCR7 (No. 47-1971;
524 eBioscience/Affymetrix, Santa Clara, CA, USA), anti-CD45.1 (No. 47-0453;
525 eBioscience/Affymetrix, Santa Clara, CA, USA), and anti-CD45.2 (No. 12-0454;
526 eBioscience/Affymetrix, Santa Clara, CA, USA). For intracellular staining, BD Cytotfix and
527 Cytoperm (Cat. No. 554714; BD Bioscience, Franklin Lakes, NJ, USA) were used before

528 incubating splenocytes with anti-IFN- γ (No. 17-7311; eBioscience/Affymetrix, Santa
529 Clara, CA, USA), anti-TNF- α (No. 46-7321; eBioscience/Affymetrix, Santa Clara, CA,
530 USA), and anti-IL-2 (No. 25-7021; eBioscience/Affymetrix, Santa Clara, CA, USA)
531 antibodies at room temperature for 30 min. All samples were fixed in 1% (w/v)
532 paraformaldehyde (PFA). All experiments were analyzed on a SORP BD LSRII analytic
533 flow cytometer (BD Bioscience, Franklin Lakes, NJ, USA). Data were analyzed in FlowJo
534 (FlowJo LLC, Ashland, OR, USA).

535

536 ***Ex vivo T cell peptide stimulation***

537 B cells in splenocytes were depleted by incubation in flasks coated with AffiniPure goat
538 anti-mouse IgG (H+L) (Jackson ImmunoResearch Laboratories Inc., West Grove, PA,
539 USA) for 1 h at 37 °C. B-cell-depleted splenocytes from infected mice (CD45.2+) were
540 incubated with naïve splenocytes (CD45.1+) at a 1:1 ratio in culture media containing 10
541 U mL⁻¹ IL-12, 10 μ g mL⁻¹ brefeldin A, and 1 μ g mL⁻¹ peptide for 5 h at 37 °C. Splenocytes
542 were stained and processed for flow cytometry with the indicated tetramers and surface
543 marker antibodies.

544

545 ***Infectious center assay***

546 Serially diluted splenocytes were plated on a Vero cell monolayer and incubated overnight
547 at 37 °C. The splenocytes were aspirated then washed off by gentle agitation. The Vero
548 cells were overlaid with 1% (w/v) methylcellulose in DMEM containing 10% (w/v) FBS for
549 6 d before fixing with 2% (w/v) crystal violet in 20% (v/v) ethanol. Infectious centers
550 indicated by plaques were counted.

551

552 **Quantitative PCR (qPCR)**

553 The qPCR was performed on MJ Opticon 2 using PerfeCTA Fastmix (Quantabio, Beverly,
554 MA, USA). For the viral genome copy number analysis, 150 ng extracted DNA (~2 x 10⁴
555 cells) and the primers were annealed to the upstream of the ORF6 coding sequence
556 (ORF6: 5'-TGCAGACTCTGAAGTGCTGACT-3' and 5'-
557 ACGCGACTAGCATGAGGAGAAT-3') were used.

558

559 For the RNA expression analysis, cells were harvested in TRIzol (Thermo Fisher
560 Scientific, Waltham, MA, USA) for RNA extraction according to the recommended
561 protocol. Total RNA was treated with DNase and used for reverse transcription in a
562 qScript cDNA synthesis kit (Quantabio, Beverly, MA, USA) to generate cDNA for qPCR.

563

564 **Gene expression analysis by qPCR**

565 Cell lysates were stored in TRIzol at -80 °C. Isolated RNA was treated with DNase I then
566 used to generate cDNA in a qScript cDNA synthesis kit (Quantabio, Beverly, MA, USA)
567 followed by gene expression analysis with PerfeCTa Fastmix (Quantabio, Beverly, MA,
568 USA). The primers used in qPCR for IL-1 β , TNF- α , IL-6, IL-12, and β -actin are listed in
569 Supplementary Table 2.

570

571 **Infection of BMDM**

572 Cells were harvested from bone marrow and differentiated into macrophages (BMDM) by
573 incubation for 7 d in DMEM containing 20% (w/v) FBS, 5% (w/v) M-CSF, 1% (w/v)

574 penicillin and streptomycin, 1% (w/v) glutamine, and 0.5% (w/v) sodium pyruvate. The
575 BMDMs were infected with WT or DIP at MOI = 1. At 24 h post-infection, total RNA was
576 extracted with TRIzol. Supernatants were collected for analysis in an IL-12/IL-23 p40
577 (total) mouse uncoated ELISA kit (No. 88-7120-22; Thermo Fisher Scientific, Waltham,
578 MA, USA).

579

580 ***Neutralizing activity***

581 Twofold serially-diluted serum was incubated with 100 PFU WT virus for 1 h at 37 °C. The
582 mixture was plated on a Vero cell monolayer for 1 h at 37 °C then removed. The plate
583 was overlaid with 1% (w/v) methylcellulose in DMEM containing 10% (w/v) FBS for 6 d
584 before fixing with 2% (w/v) crystal violet in 20% (v/v) ethanol. The neutralizing titer was
585 taken as the highest dilution maintaining the ability of the diluted serum to reduce the
586 number of plaques by 50% relative to the virus mixture containing fourfold diluted mock
587 serum.

588

589 ***Virus-specific IgG ELISA***

590 A 5 µg mL⁻¹ WT virion antigen solution coated a 96-well plate which was then incubated
591 overnight at 4 °C. The plate was blocked overnight in PBS containing 1% (w/v) BSA and
592 0.05% (w/v) Tween-20. The plate was washed twice with PBS-T (PBS containing 0.5%
593 Tween-20). Mouse sera were diluted in ELISA buffer (PBS containing 0.1% BSA and
594 0.025% Tween-20) and incubated on the plate for 1 h at room temperature. The plate
595 was washed thrice with PBS-T and then once with PBS. A substrate solution consisting
596 of one tablet each of *o*-phenylenediamine and urea hydrogen peroxide (No. P9187;

597 Sigma-Aldrich Corp., St. Louis, MO, USA) in 10 mL ddH₂O, was added to the plate. The
598 plate was incubated for 30 min in the dark at 4 °C. The reaction was stopped by adding
599 4N H₂SO₄ and the plate was read at 490 nm and 620 nm. The virus-specific IgG titer was
600 taken as the highest dilution generating signals higher than those of the 1:50 diluted mock
601 serum.

602

603 ***Serum transfer***

604 Sera were obtained from mice at 2 mo post-infection. Then 200 µL pooled heat-
605 inactivated serum was intraperitoneally injected into naïve mice. After 24 h, the naïve
606 recipient mice were challenged intranasally with 5 x 10³ PFU WT. A second dose of 200
607 µL pooled heat-inactivated serum was intraperitoneally injected 7 d after the WT
608 challenge. Splenocytes were harvested 14 d after the challenge for the infectious center
609 assay.

610

611 ***Adoptive T cell transfer***

612 Splenocytes were isolated from mice at 2 mo post-infection and pooled from multiple
613 mice. Splenocytes were negatively selected for CD4, CD8, or total T cells using EasySep
614 isolation kits (Catalog Nos. 19765, 19853, and 19851; STEMCELL Technologies Inc.,
615 Vancouver, BC, Canada). Negative selection was confirmed by flow cytometry analysis
616 to > 90% purity. Three million cells in 100 µL were injected into the tail vein of each
617 B6.SJL-*Ptprca*^a *Pepcb*^b/BoyJ mouse. Twenty-four hours after T-cell transfer, the
618 recipient mice were intranasally challenged with 5 x 10³ PFU WT MHV-68. Spleens were
619 harvested 14 d post-challenge for the infectious center assay and flow cytometry to

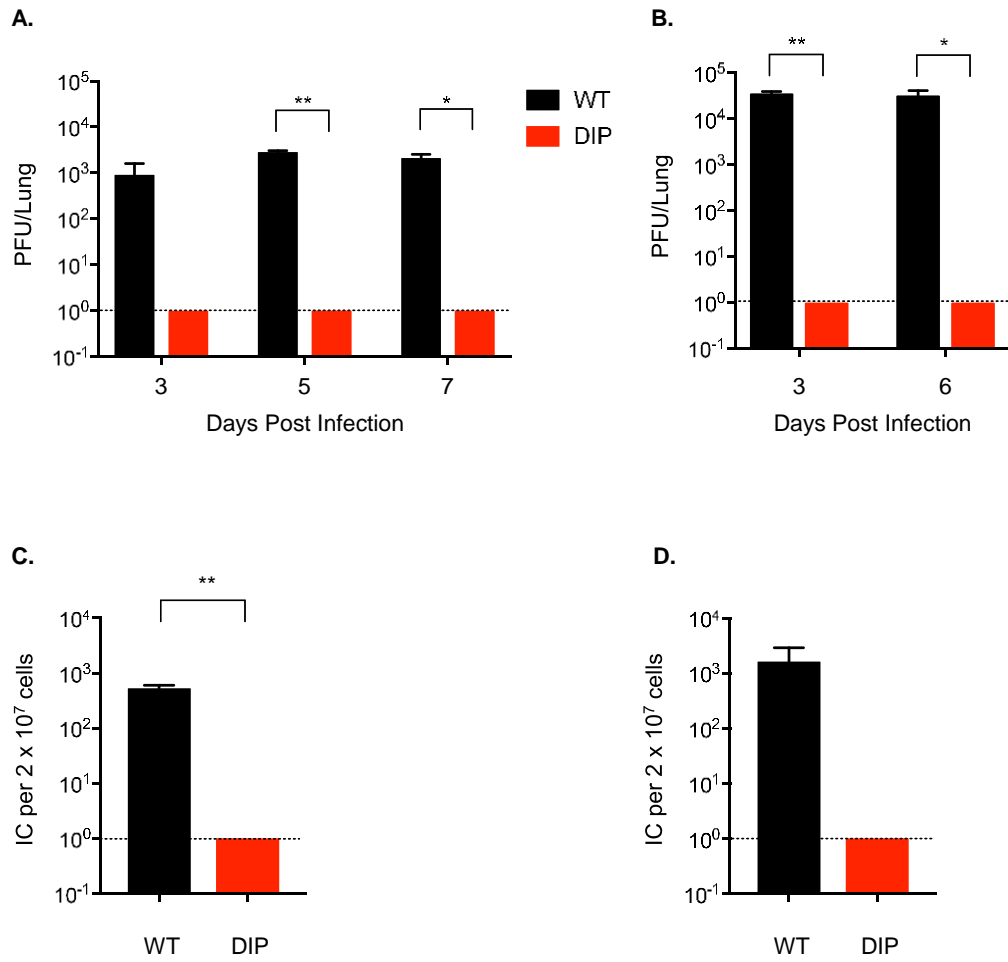
620 confirm the presence of donor T-cells in the recipient mice with anti-CD45.1 and anti-
621 CD45.2.

622

623 ***Statistical analysis***

624 Data are presented as means and their differences were analyzed by a two-tailed
625 unpaired Student's *t*-test unless otherwise indicated. $P < 0.05^*$, $P < 0.01^{**}$, $P < 0.001^{***}$,
626 and $P < 0.0001^{****}$.

Figure S1



629 **Figure S1. In vivo DIP virus is both replication- and latency deficient upon**
630 **intranasal inoculation**

631 Mice were intranasally inoculated with 5,000 (**A, C**) or 10^5 PFU (**B, D**) WT or DIP. (**A, B**)

632 Lungs (n = 3) were excised at the times indicated at the bottoms of the graphs for plaque

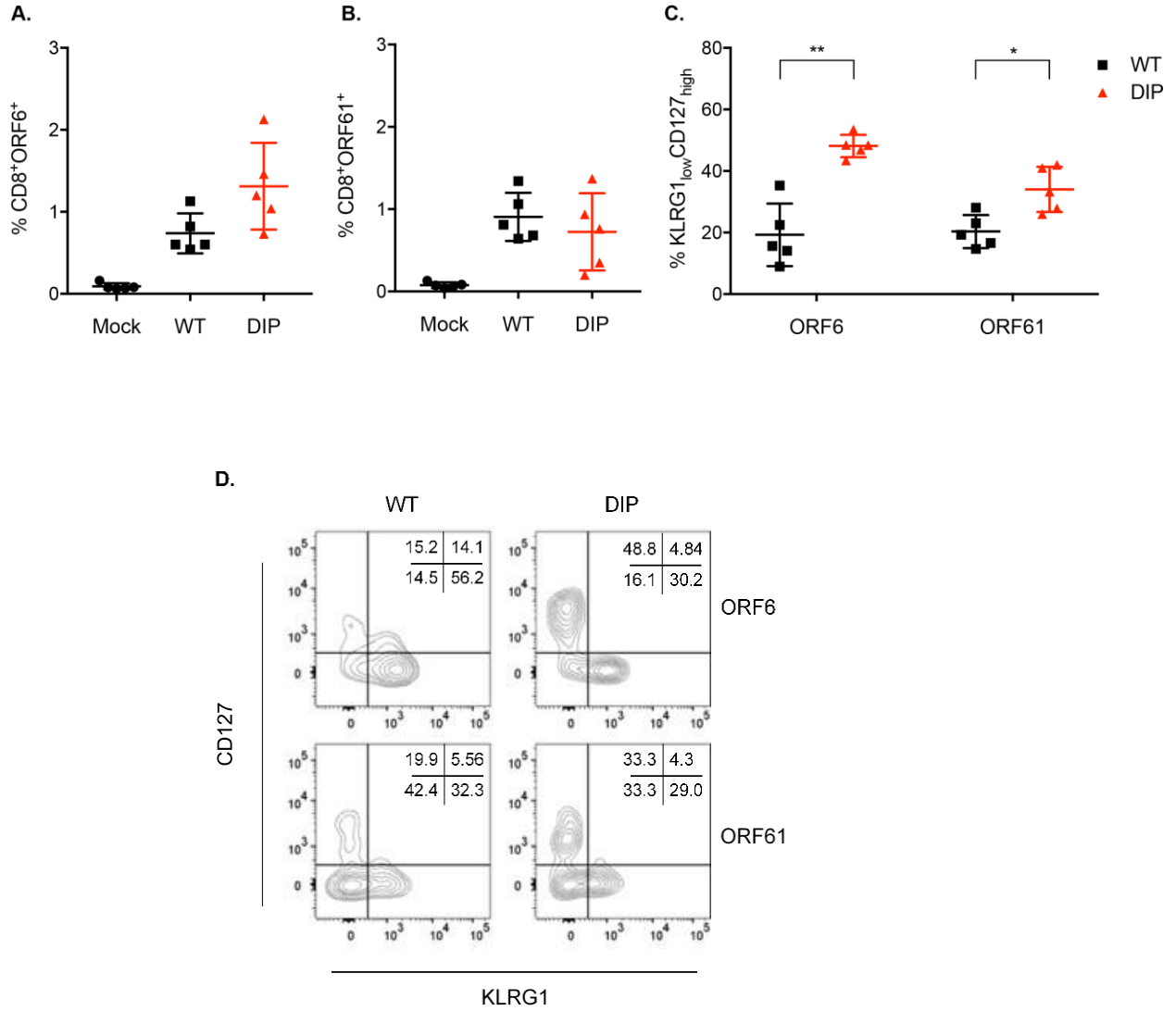
633 assay. (**C, D**) Spleens (n = 3) were excised 14 d post infection for infectious center assay.

634 Means and SD were plotted. Statistical significance was analyzed by a two-tailed

635 Student's *t*-test.

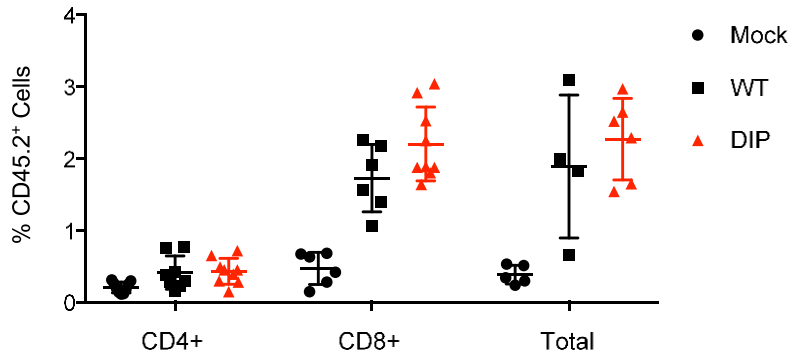
636

Figure S2



638 **Figure S2. DIP infection induces a robust, virus-specific T cell response**
639 Mice were either mock-infected or intraperitoneally infected with 10^5 PFU WT or DIP. (**A**,
640 **B**) At 1 mo post-infection, splenocytes were harvested and examined for virus-specific
641 CD8⁺ T cells using the tetramers ORF6₄₈₇₋₄₉₅/Db and ORF61₅₂₄₋₅₃₁/Kb. (**C**) Tetramer-
642 positive CD8⁺ T cells were examined for KLRG1 and CD127 expression. Data for
643 individual mice (n = 5), means, and SD were plotted. Statistical significance was analyzed
644 by a two-tailed Student's *t*-test. (**D**) Gating strategy to determine MPEC and SLEC
645 population frequencies using representative samples from WT- and DIP-inoculated mice.
646

Figure S3

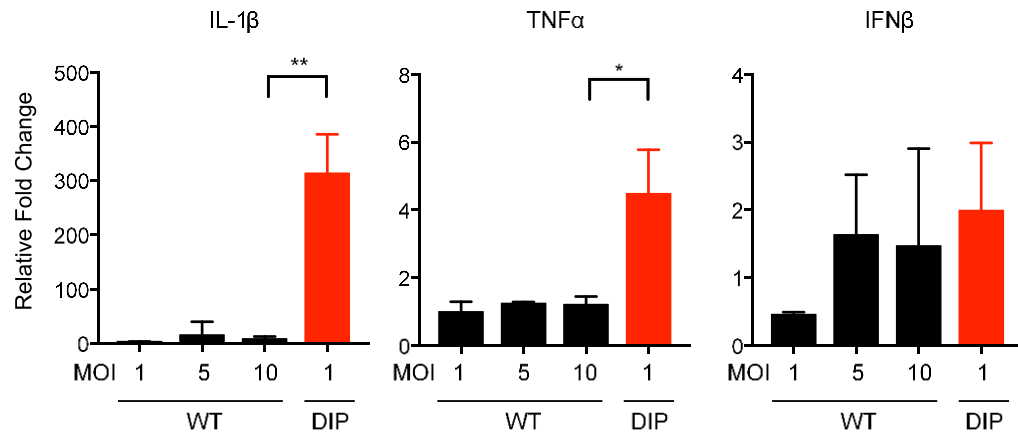


648 **Figure S3. WT- and DIP-primed T cells expand to the same extent in the recipient**
649 **mice**

650 CD4⁺, CD8⁺, or total T cells were purified via negative selection from the spleens of mock-
651 infected mice or mice intraperitoneally infected with 10⁵ PFU WT or DIP. Three million
652 purified cells were transferred to a congenic mouse by tail vein injection. At 14 d after
653 transfer, donor cells were analyzed by flow cytometry and the percentages are shown.
654 Data for individual mice, means, and standard deviations are plotted. Statistical
655 significance was analyzed by a two-tailed Student's *t*-test. No statistical significance was
656 determined for the percentages of WT- and DIP-primed donor cells.

657

Figure S4



659 **Figure S4. WT does not upregulate inflammatory cytokines to the same level as DIP**
660 Mouse BMDM were infected with WT or DIP at the MOIs indicated at the bottoms of the
661 graphs (triplicate). (A) Total RNAs were extracted 24 h post-infection to measure IFN β ,
662 IL-1 β , TNF- α , and β -actin expression. Cytokine RNA expression was normalized against
663 β -actin. Relative fold change was calculated by comparing to mock-infected BMDM.
664 Statistical significance was analyzed by a two-tailed Student's *t*-test.

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