Published in final edited form as:

Antiviral Res. 2012 April ; 94(1): 1–8. doi:10.1016/j.antiviral.2012.02.001.

A humanized IgG but not IgM antibody is effective in prophylaxis and therapy of yellow fever infection in an AG129/17D-204 peripheral challenge mouse model

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Abstract

Yellow fever virus (YFV), a member of the genus *Flavivirus*, is a mosquito-borne virus found in tropical regions of Africa and South America that causes severe hepatic disease and death in humans. Despite the availability of effective vaccines, YFV is responsible for an estimated 200,000 cases and 30,000 deaths annually. There are currently no prophylactic or therapeutic strategies approved for use in human YFV infections. Furthermore, implementation of YFV 17D-204 vaccination campaigns has become problematic due to an increase in reported post-vaccinal adverse events. We have created human/murine chimeric MAbs of a YFV-reactive murine monoclonal antibody (mMAb), 2C9, that was previously shown to protect mice from lethal YFV infection and to have therapeutic activity. The new chimeric (c)MAbs were constructed by fusion of the m2C9 IgG gene variable regions with the constant regions of human IgG and IgM and expressed in Sp2 murine myelomas. The 2C9 cMAbs (2C9-cIgG and 2C9-cIgM) reacted with 17D-204 vaccine strain in an enzyme-linked immunosorbent assay and neutralized virus *in vitro* similarly to the parent m2C9. Both m2C9 and 2C9-cIgG when administered prophylactically 24 hours prior to infection protected AG129 mice from peripheral 17D-204 challenge at antibody concentrations $\geq 1.27 \mu\text{g}/\text{mouse}$; however, the 2C9-cIgM did not protect even at a dose of $127 \mu\text{g}/\text{mouse}$. The 17D-204 infection of AG129 mice is otherwise uniformly lethal. While the m2C9 was shown previously to be therapeutically effective in YFV-infected BALB/c mice at day 4 post-infection, the m2C9 and 2C9-cIgG demonstrated therapeutic activity only when administered 1 day post-infection in 17D-204-infected AG129 mice.

Keywords

yellow fever; human monoclonal antibodies; treatment; prophylaxis

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1. INTRODUCTION

Yellow fever virus (YFV), a member of the genus *Flavivirus*, family *Flaviviridae*, is found in tropical regions of Africa and South America, and is transmitted to primates by mosquitoes. Despite the availability of effective vaccines, YF is still a matter of public health concern - responsible for an estimated 200,000 cases and 30,000 deaths annually - and is considered a re-emerging disease. This is due to the reinfestation of countries with the *Aedes aegypti* vector, lapses in implementation of preventative vaccination programs in endemic regions such as sub-Saharan Africa, and the lack of compliance by at-risk populations (Barnett, 2007; Monath, 2006). Given the increase in travel and commerce originating from once isolated, YFV-endemic regions, there is a growing concern regarding the potential of YFV to cause urban epidemics if introduced into geographical regions containing the *Ae. aegypti* mosquito vector and large concentrations of susceptible humans (e.g., the southern U.S.) (Gardner and Ryman, 2010; Gubler, 2001, 2002).

For those with access to vaccination, the use of the live-attenuated YF vaccine is contraindicated in a number of individuals. It is advised that infants under 6 months of age not be given the YF vaccine due to a risk of viral encephalitis developing in the child (Cetron et al., 2002; Staples et al., 2010). Also at risk are those individuals who suffer from hypersensitivity to eggs since the YF vaccine is prepared in embryonated eggs. The YF vaccine is not suitable for those who are immunocompromised due to AIDS or HIV infection, or whose immune system has been altered by either diseases such as leukemia and lymphoma or through drugs and radiation (Cetron et al., 2002; Staples et al., 2010). Studies have shown that persons aged ≥ 65 years are particularly susceptible to systemic adverse events following immunization with 17D-204 (Khromava et al., 2005; Martin et al., 2001; Massad et al., 2005; Monath et al., 2005). Additionally, recent reports suggest an increase in post-vaccinal severe adverse events during YF vaccination campaigns (Barrett et al., 2008; Barrett and Teuwen, 2009; Doblas et al., 2006; Engel et al., 2006; Ferguson et al., 2010).

Treatment options are limited for YFV-infected individuals (Monath, 2008). There is currently no specific treatment for YF, a disease that is estimated to have a case-fatality rate of approximately 20%. A small number of antivirals, such as ribavirin or recombinant interferon (IFN), have shown efficacy in reducing viremia and/or prolonging time to death in animal models of YF if administered early in the disease. However, the effectiveness of these therapies is dramatically reduced if given after the onset of clinical symptoms (Julander et al., 2007; Monath, 2008; Sbrana et al., 2004). The study by Sbrana et al., showed that early treatment with ribavirin appears to reduce liver damage associated with YFV infection (Sbrana et al., 2004); however, for the most part treatment of YFV-infected patients is limited to supportive therapy and is directed towards the organ systems involved.

Treatment of YFV infection with murine monoclonal antibodies (mMAbs) provides *in vivo* protection in a mouse model. Brandriss et al. showed that neutralizing mMAbs directed against the envelope (E) glycoprotein of the 17D-204 vaccine strain protected mice from lethal encephalitis if administered either 1 day prior to or 3 to 5 days after viral intracerebral (i.c.) challenge with 17D-204 vaccine (Brandriss et al., 1986; Schlesinger et al., 1983; Schlesinger et al., 1984). A similar study by Schlesinger et al. demonstrated that prophylactic administration of non-neutralising mMAbs directed against the 17D non-structural glycoprotein, NS1, also protected mice from lethal YF encephalitis (Schlesinger et al., 1986). Unfortunately, human treatment with mMAbs can be compromised by the human anti-mouse antibody response.

Advances in antibody engineering have now made it possible to produce human-murine chimeric (cMAbs) or fully human MAbs. Both cMAbs and hMAbs retain the specificity,

avidity and neutralizing activity of the mMABs they are derived from; however, cMABs and hMABs reduce the human anti-mouse antibody (HAMA) response in humans, and are more effective therapeutics than mMABs. In addition, MABs provide an infinite source of antibodies that are homogeneous in both specificity and affinity, thus making them an attractive substitute for human polyclonal sera for use in human therapeutics. A number of cMABs have been approved by the FDA for treatment or prophylaxis of various disorders including non-Hodgkin's lymphoma, renal allograft rejection, and rheumatoid arthritis (Gaffo et al., 2006; Leget and Czuczman, 1998; Lupo et al., 2008). Fully human and humanized MABs have been successfully used to protect against West Nile virus (WNV) and Venezuelan equine encephalomyelitis virus (VEEV) in both prophylactic and therapeutic animal models of infection (Hunt et al., 2011; Hunt et al., 2006; Hunt et al., 2010; Morrey et al., 2006; Morrey et al., 2007; Morrey et al., 2008; Nybakken et al., 2005; Oliphant et al., 2005). Clinical trials are currently being conducted to evaluate the safety and efficacy of humanized E16 antibody (MGAWN1) for use in WNV infections (Beigel et al., 2010). In this study, we report creation of human/murine chimeric cIgG and cIgM from a protective and therapeutic YFV-reactive mMAB, 2C9, that is specific for the envelope (E) protein for either wild-type or vaccine YF stains (Brandriss et al., 1986; Schlesinger et al., 1983). Sequencing of the E protein of 2C9 neutralization resistant 17D-204 virus identified amino acids 71 and 72 as being involved in 2C9 binding, therefore this MAB recognizes an E protein Domain II epitope (Lobigs et al., 1987). Both the mMAB and the cMAB 2C9-cIgG were shown to be effective prophylactically and therapeutically in a new YF mouse model of infection that utilizes 17D-204 vaccine peripherally-challenged, interferon receptor-deficient AG129 mice (Lee and Lobigs, 2008; Meier et al., 2009).

2. MATERIALS AND METHODS

2.1. Mice

The 129/Sv/Ev mice deficient for IFN- α/β and - γ receptors in combination (strain AG129) obtained from B & K Universal (Hull, United Kingdom) and housed in the Division of Vector-Borne Disease (DVBD) animal care facilities at CDC were used for all animal studies (Johnson and Roehrig, 1999). Mice were euthanized with isoflurane followed by cervical dislocation when signs of illness became obvious as indicated by reduced activity and increased huddling during normal activity hours, lack of appetite, and the development of neurologic signs such as hind leg paralysis or weakness. The use of animals for research purposes complied with all relevant federal guidelines and specific protocols were approved by the DVBD Institutional Animal Care and Use Committee.

2.2. Cells and virus

The Sp2/0-Ag14 (Sp2) murine myeloma and 2C9 murine hybridoma (Roehrig et al., 1980; Schlesinger et al., 1983) were propagated in hybridoma growth medium (HGM) unless noted otherwise. The YFV vaccine strain (17D-204) was obtained from the DVBD reference collection (Fort Collins, CO). A single pool containing 2.0×10^7 plaque-forming units per ml (PFU/ml) of 17D-204 grown in Vero cells with minimum essential medium (MEM; D-MEM supplemented with 10% FBS, 2 mM L-glutamine, 0.15% sodium bicarbonate, 100 U/ml penicillin G sodium, and 100 μ g/ml streptomycin sulfate) was used for all inoculations.

2.3. Assembly of human-murine chimeric (c)2C9 IgG and IgM plasmid constructs

The heavy and light variable regions (V_H and V_K) of the IgG gene were amplified from the m2C9 hybridoma and subsequently modified by PCR for insertion into the human IgG expression vector pdHL2 or the human IgM expression vector pJH2-24-95B1, referred to as pJH2 (Abbott Laboratories, Abbott Park, IL) as previously described (Hackett et al., 1998; Thibodeaux et al., 2010; Thibodeaux and Roehrig, 2009). Degenerate primers used for V_H

and V_K amplification and primers used to add partial 5' leader sequences, 3' splice donor junctions, and appropriate restriction sites to the variable regions are listed in Table 1. The V_H and V_K regions of m2C9 IgG were incorporated separately into pdHL2 or pJH2 by ligation as previously described (Hackett et al., 1998) generating plasmids pdHL2-2C9 and pJH2-2C9. Plasmids pdHL2-2C9 and pJH2-2C9 were used to transform *E. coli* DH5αE (Invitrogen) by electroporation following the manufacturer's protocol.

2.4. Sequencing

V regions were sequenced in triplicate to ensure sequence fidelity after initial amplification and again after PCR modification. Sequencing reactions were performed using the BigDye® Terminator v3.1 Cycle Sequencing Kit (Applied Biosystems, Foster City, CA) and sequence data analyzed using the ABI 3130xl Genetic Analyzer (Applied Biosystems).

2.5. Transfection of cells with clg plasmid constructs

Sp2 cells were prepared and transfected with plasmids pdHL2-2C9 or pJH2-2C9 as previously described (Thibodeaux et al., 2010). Transfected cells were expanded and screened for human IgG or IgM production by ELISA.

2.6. ELISA for detection of YFV-specific clg

To detect the presence of human YFV-specific cIgG antibody in samples of cell culture supernatants or titrate purified cIgG, a modification of the indirect IgG-ELISA protocol originally described by Johnson et al. was used (Johnson et al., 2000; Thibodeaux et al., 2010). The flavivirus cross-reactive 6GF4 cIgG and the alphavirus cross-reactive 1GD5 cIgG were used as positive and negative control antibodies in the ELISA (Thibodeaux and Roehrig, 2009). To detect the presence of cIgM, IgM antibody-capture (MAC) ELISA was used as previously described (Thibodeaux et al., 2011; Thibodeaux and Roehrig, 2009). The flavivirus cross-reactive 6ME2 cIgM and the alphavirus cross-reactive 1MD11 cIgM were used as positive and negative control antibodies in the ELISA (Thibodeaux and Roehrig, 2009). All ELISAs were performed in duplicate.

2.7. Antibody purification

Transfected Sp2 cells that tested positive for production of cIgG or cIgM were expanded and passaged three times in HGM with methotrexate followed by one additional passage in HGM without methotrexate. Supernatants from each passage were tested for human anti-YFV activity using the indirect IgG ELISA described above. Stably transformed cell lines that exhibited consistently high levels of human anti-YFV cMAb -- 2C9-H4 (cIgG) or 2C9-G9b (cIgM) -- were used for commercial ascites fluid production and antibody purification (QED Bioscience Inc., San Diego, CA). Ascitic fluid and purified IgG were also made from the parent m2C9 hybridoma by QED Biosciences, Inc. Antibodies that were purified from ascitic fluid and eluted in PBS were at final concentrations of 0.55 mg/ml (2C9-cIgG), 0.51 mg/ml (2C9 mMAb) and 0.92 mg/ml (2C9-cIgM).

2.8. YFV plaque reduction neutralization test (PRNT)

The methods for determining flaviviral neutralizing antibody titers have been published previously (Roehrig et al., 2008). Briefly antibody diluted in PBS was mixed with an equal volume of 17D-204 seed (1.0×10^3 PFU/ml) diluted in MEM. Antibody-virus mixtures were allowed to incubate at 37°C for 1 h prior to being dispensed onto 6-well Vero plates in aliquots of 200μl/well. Plates were overlaid with a mixture of agarose and growth medium and incubated at 37°C for 4 days, after which a second overlay containing 0.0053% neutral red (Sigma) was added. Plates were incubated at 37°C for an additional 24 h before plaque formation was recorded. The PRNT₅₀ of each antibody was defined as the titre of antibody

required to reduce total PFU/well by 50% compared to wells inoculated with 17D-204 alone. Starting antibody concentrations were standardized to 0.51 mg/ml. The humanized alphavirus cross-reactive 1GD5 cIgG was used as a negative control antibody (Thibodeaux et al., 2010).

2.9. Complementarity determining region (CDR) analysis

CDR determination was performed using the method described by Kabat (Johnson and Wu, 2000).

2.10. Nucleotide sequence accession numbers

The 2C9 V region sequences have been submitted to GenBank and assigned the following accession numbers: 2C9 V_H, **GU724339**; 2C9 V_K, **GU724340**.

2.11. Animal studies

All animal studies were performed under a DVBD/CDC IACUC-approved protocol. Mice used in all challenge experiments were between 6 and 8 weeks of age. All mice challenged with 17D-204 received a single intraperitoneal (i.p.) injection of 100 µl containing 2×10^5 PFU/mouse. Mice treated prophylactically or therapeutically with purified antibody received a single dose of antibody in a volume of 250 µl delivered by i.p. injection. Beginning at day 12 post-infection (p.i.) which coincided with the appearance of morbidity in untreated challenged mice, mice were euthanized upon exhibiting signs of morbidity; tissues were collected at time of sacrifice. Blood was collected via post-mortem cardiac puncture; serum was separated using microtainer tubes (Becton-Dickinson, Franklin Lakes, NJ). Serum samples from euthanized mice were analyzed for *de novo* production of murine anti-YFV IgM antibodies by MAC-ELISA as previously described to confirm viral infection (Thibodeaux and Roehrig, 2009).

Brain and liver samples were also collected and were manually chopped into small sections using a scalpel and rinsed in PBS prior to being placed into pre-weighed MagNA Lyser Green Bead tubes (Roche Diagnostics GmbH, Mannheim, Germany) containing 500 µl MEM (0% FBS) and stored at -70°C ; serum samples separated from blood were stored at -70°C . These tissue specimens were used in qRT-PCR to determine genome copy levels (see Viral Quantitation below).

2.11.1. Protective and therapeutic efficacy—AG129 mice challenged with 2×10^5 PFU/mouse 17D-204 via i.p. injection were administered a single i.p. injection of 127 µg m2C9 IgG, 2C9-cIgG, 2C9-cIgM, or PBS 1 day prior to infection, or one of the IgG MABs at 1, 3, or 5 days p.i. Individual mice were euthanized upon demonstrating signs of morbidity as described above. Brain samples were collected post-mortem and processed as described. Antibody-treated mice that failed to exhibit signs of morbidity were monitored for an additional 7 days following the sacrifice of the last untreated mouse before being euthanized for tissue collection (day 31). Additionally, prophylactic dose-response analysis was performed using diminishing concentrations of m2C9 and 2C9-cIgG.

2.11.2. Viral Quantitation

2.11.2.1. RNA Extraction from Tissues: For determination of viral genome content of mouse tissues, frozen samples of brain were thawed and homogenized for two 30 s cycles at 4,000 rpm using a Roche MagNA Lyser (Roche Diagnostics GmbH). Homogenized samples were centrifuged for 5 min at 10,000 rpm at room temperature (RT) and total RNA was purified from 50 µl of clarified homogenate using an RNeasy 96 Universal Kit (Qiagen, Valencia, CA) following the manufacturer's protocol. Viral RNA was extracted from 140 µl

serum samples using a QIAamp® Viral RNA Mini Kit (Qiagen) following the manufacturer's protocol. Viral RNA was purified from ten-fold serially-diluted 17D-204 in parallel with tissue samples using the RNeasy 96 Universal Kit for generation of a standard curve. The 17D-204 used for RNA standards was also titrated by plaque assay to correlate the viral genome equivalents with PFU/ml.

2.11.1.2. Quantitative Reverse-Transcription Polymerase Chain Reaction (qRT-PCR):

Primers and probe used for qRT-PCR were obtained from the CDC Biotechnology Core Facility (Atlanta, GA) and are detailed in Table 1. Each reaction mixture contained 5 µl of purified RNA; primers and probes were used at final concentrations of 1 µM for 8280F and 8354R and 0.2 µM for the 8308 probe. Amplifications were performed in an iQ™5 Real-Time PCR Detection System (Bio-Rad, Hercules, CA) using a QuantiTect® Probe RT-PCR Kit (Qiagen) under the following conditions: 50°C for 30 min, 95°C for 12 min 30 s, followed by 45 cycles of 94°C for 15 s, 55°C for 1 min with continuous fluorescence data collection. Each RNA sample was tested in duplicate (growth curve) or triplicate (protection study) and virus genome equivalents were determined by extrapolation from the standard curves generated within each experiment.

3. RESULTS

3.1. Development of cIgG and cIgM for YFV

We have shown previously that m2C9 was capable of protecting mice from lethal i.c. challenge with 17D-204 when administered prophylactically or therapeutically to BALB/c or CD-1 mice (Brandriss et al., 1986). Using the pdHL2 expression vector, which contains genomic clones of both the kappa (C_K) and IgG1 constant ($C_{\gamma 1}$) region genes of human immunoglobulin, we constructed a 2C9 cIgG antibody that maintained the specificity and protective capacities of the m2C9 parent. Additionally, to determine the effect that a YFV-neutralizing IgM containing the same antigen binding site as the YFV 2C9-cIgG would have on infection, we also constructed a 2C9-cIgM antibody. Multiple clones of each V gene amplicon were sequenced to ensure against possible DNA polymerase-induced errors, Fig. 1.

Cell-free supernatants of Sp2 cells transfected with pdHL2-2C9 or pJH2-2C9 were analyzed by ELISA for the presence of human-murine chimeric Ig approximately two weeks after transfection. A total of three wells, H4, F6, and G9, for cIgG and three wells, G-9, G-9b, and H-11, for cIgM of transfected Sp2 cells tested positive for YFV-specific antibody by ELISA. Single stably transformed cell lines, 2C9-H4 (cIgG) and 2C9-G9 (cIgM) that consistently secreted high levels of anti-YFV antibody over multiple passages were sent to a commercial company for production and purification of the antibodies used in this study. The reactivities of the purified 2C9-cIgG and m2C9 were similar by ELISA against YFV 17D-204 virus (data not shown). More interestingly, the reactivities of the 2C9-cIgG and 2C9-cIgM with YFV-17D were verified by ELISA and also shown to be identical to one another, Fig. 2. The similar binding curves of purified and standardized 2C9-cIgG and 2C9-cIgM suggests similar binding avidities with 17D-204. This result was a bit unexpected given the difference in size between IgG and IgM.

3.2. 2C9-H4 and 2C9-G9 Neutralizing Activity in vitro

The YFV-neutralizing titers of purified 2C9-cIgG and 2C9-cIgM were compared to that of the parent m2C9 by PRNT in Vero cell cultures, Fig. 2, inset. The PRNT₅₀ end-point reciprocal titre for the 2C9-cIgG (128) was two-fold less than the parent m2C9 MAb (256) and two-fold more than the 2C9-cIgM (68). The alphavirus cIgG 1GD5 was used as a negative control antibody and did not neutralize YFV.

3.3. Prophylaxis and therapy of YFV infections using 2C9-clgG and 2C9-clgM

The 2C9-cIgG, 2C9-cIgM, and their m2C9 parent were assayed for their efficacy in prophylaxis and therapy of 17D-204 infection of AG129 mice. Prior to testing, 127 µg/mouse of 2C9-cIgG was administered via i.p. injection to small groups of AG129 mice to ensure that no adverse reactions resulted and no adverse reactions were observed (data not shown).

Using the AG129/17D-204 peripheral challenge model, the 2C9-cIgG and m2C9 parent MAbs were first tested for their ability to limit YF disease when administered 24 h prior to virus challenge. Protection was defined as increased average survival time (AST) and decreased morbidity of treated mice compared to non-treated mice. The aggregate results of 2–4 independent experiments are summarized in Fig. 3a and Table 2. Mice receiving either 2C9-cIgG or m2C9 24 h prior to infection with 17D-204 showed a significant increase in AST (m2C9 AST=30.4±2.5 days; 2C9-cIgG AST=28.3±4.5 days) compared to infected mice receiving PBS alone (AST=17.3±4.3 days). Additionally, 95% of animals receiving m2C9 and 72% of animals receiving 2C9-cIgG survived viral challenge. Interestingly, the 2C9-cIgM failed to protect significant numbers of mice (2/10) from viral challenge and did not significantly increase ASTs (20.6±5.9 days vs. 17.3±4.3 days for PBS control animals) even though it contained the same antigen-combining site as 2C9-cIgG and had the same virus-neutralizing activity.

Mice that were treated with m2C9 one day after viral challenge demonstrated a significantly increased AST (25.8±8.5 days) compared to untreated mice (AST=17.3±4.3 days) Table 2 and Fig. 3b. Mice receiving 2C9-cIgG at one day p.i. demonstrated a slightly increased AST (19.4±7.0). Seventy percent of animals receiving m2C9 and 20% of animals receiving 2C9-cIgG 1 day after infection survived viral challenge. Survival curves are shown in Fig. 3b. Infected mice receiving either m2C9 or 2C9-cIgG at days 3 or 5 p.i. (not shown) did not show an increase in AST compared to PBS treated mice. ASTs were slightly shorter for mice administered MAb at 3 days p.i. compared to PBS control animals (15.8 and 15.3 days vs. 17.3 days, Table 2), however these differences were not statistically significant (t-tailed t-test, $p \leq 0.05$). Because of its low level of prophylactic capacity, the therapeutic capacity of the 2C9-cIgM was not tested.

Dose-response titrations of the protective capacity of m2C9 and 2C9-cIgG indicated that m2C9 was more effective in protecting mice than was 2C9-cIgG. At a dose of 1.27 µg/mouse, m2C9 protected 60% of challenged mice while 2C9-cIgG protected 20% of mice, Table 3. Both antibodies protected efficiently at the higher doses of 127 or 12.7 µg/mouse.

There were no significant differences in viral load in brain tissue between mice showing morbidity in either antibody-treated or untreated groups, both demonstrating 6.6 to 6.9 log₁₀ genomic equivalents/g of brain tissue, Table 3. Variation within groups was high, however, with 17D-204 vRNA below detectable levels in some mice. Mice surviving viral challenge demonstrated significantly reduced viral load compared to non-surviving mice given the same treatment. In fact, most surviving mice, regardless of IgG treatment, were essentially virus free at the end of the experiment. Interestingly, the two surviving mice treated prophylactically with 2C9-cIgM were not virus free, but they did have significantly reduced viral loads in the brain when compared to the mice that did not survive challenge (6.0 vs. 7.0 log₁₀ genomic equivalents/g of brain tissue, Table 3). Serum samples taken from all mice were positive for murine anti-YFV IgM as determined by MAC-ELISA, indicating that all mice were infected and mounted an antibody response (data not shown).

4. DISCUSSION

In this study, use of the AG129/17D-204 model permitted us to analyze the protective capacity of our MAbs in a neurotropic model of infection that resembles the original i.c. BALB/c 17D-204 infection model used previously (Brandriss et al., 1986). In this case, however, we were able to use a peripheral route of viral challenge. These earlier studies determined that m2C9 could protect 100% of mice (40 µg m2C9/mouse) or 60% of mice (4 µg m2C9/mouse) subsequently challenged i.c. with 17D-204. Additionally, m2C9 cured 90% of 17D-204-infected mice when 40 µg 2C9/mouse were administered by day 4 p.i. The use of specific antibody for prophylaxis or therapy of viral infection has been explored for a number of other flaviviruses with varying degrees of success. Numerous studies have illustrated that treatment of flavivirus infections with immune serum or mMAbs (both neutralizing and non-neutralizing) can result in protection in rodents. However, evidence also suggests that successful treatment depends on the dosage and timing of antibody administration. In the case of encephalitic flaviviruses, passive mMAb therapy is usually only effective if administered before viral replication in the brain (Mathews and Roehrig, 1984; Roehrig et al., 2001). Other factors to consider in the use of passive antibody therapy for flaviviruses include the possibility of antibody dependent enhancement (ADE) of infection arising from the presence of sub-neutralizing concentrations of antibody as is seen during secondary DENV infection.

Although ADE has not been described for YFV, a similar phenomenon, referred to as antibody-dependent early death (ADEED), has been described in models of YFV infection involving passive antibody therapy of encephalitic disease in rodents. We did not observe ADEED when using lower, partially-protective doses of antibody (1.27 µg/mouse), as compared to PBS control animals (Table 3). This syndrome results from specific reactions between virus and antibody in the infected brain and is not mediated through Fc- and complement receptor-bearing macrophages (Barrett and Gould, 1986; Gould et al., 1987). ADEED has been reported only with selected virus/antibody combinations, usually with antibodies demonstrating little or no YFV-neutralizing activity (Barrett and Gould, 1986). A study by Gould et al. suggested that neutralizing antibodies normally capable of protecting mice from virus challenge enhanced neurovirulence only if they were given at a time after infection when infection in the mouse brain was established (Gould and Buckley, 1989).

For this study, the m2C9 MAb chosen for humanization is one of only a handful of YFV type-specific mMAbs currently in existence and is 1 of only 3 mMAbs capable of neutralizing both the wt YF-Asibi and 17D-204 vaccine strains *in vitro* and demonstrating protective activity against both strains *in vivo*. The growth curve of 17D-204 in AG129 brain tissue indicates that viral dissemination to and replication in the brain occurs as early as day 3 p.i. (Lee and Lobigs, 2008; Meier et al., 2009). Administration of 2C9 MAbs at either days 3 or 5 p.i. did not provide significant protection in this model, Table 2 and Fig. 3b.

The observation that m2C9 (and to a lesser extent 2C9-cIgG) only provided protection to AG129 mice when administered 1 day prior to or 1 day after infection is in contrast to our previous studies (Brandriss et al., 1986). Although m2C9 and 2C9-cIgG shared similar 17D-204 neutralizing capacity *in vitro*, it is conceivable that the humanization of 2C9 may have reduced the protective capacity of this antibody in murine models of YF disease. A study by Schlesinger and Chapman reported that the loss or absence of the murine IgG Fc region resulted in a loss in the protective capacity of 2E10 mMAb in mice (Schlesinger and Chapman, 1995). The inability of 2C9-cIgM to protect animals from viral challenge even though its antigen recombining site was identical to that of 2C9-cIgG supports the involvement of the IgG Fc portion in protection. Considering that mouse IgG2a and human IgG1 are structurally similar and analogous in that both are able to fix complement and bind

to human Fc γ -RIA (CD64) with very high affinity, it is likely that human IgG1 binds tightly to mouse CD64. An alternative explanation for the observed decrease in protective activity could also be attributable to the different mouse strains used in the protection assays.

Alternatively, the AG129 mice used in this protection study have defects in their type I and II IFN responses, unlike the BALB/c or CD-1 mice used in the initial characterization of m2C9. It is likely that the early protection afforded mice by the mouse IFN response also increases the protective capacity of the 2C9 MAbs. To address this question we will analyze the protective and therapeutic activities of these MAbs in the hamster model of YFV infection. This model uses the wt YFV-Asibi strain as the challenge virus (Julander et al., 2007; Sbrana et al., 2006; Tesh et al., 2001; Xiao et al., 2001)

If the reduced protective activity of 2C9-cIgG in AG129 mice is due to incompatibility between the cMAb and murine immune system or is a result of the immune deficiencies in the mouse strain used, then it is likely that the 2C9-cIgG will be more effective in a human host as is intended. In any event, we are currently investigating these possibilities and are also humanizing additional YFV-protective mMABs for combination therapy with 2C9-cIgG in order to reduce the risk of selection for YFV neutralization-escape variants in response to treatment with a single MAb.

The involvement of flaviviral IgM in protection and recovery from viral infection has not been well studied. In a previous study with YFV it was noted that a 400 μ g of a highly neutralizing YFV mMAb, 8A3, was not able to passively protect mice from 17D-204 challenge (Brandriss et al., 1986). Additionally, the IgM fraction of immune sera cannot facilitate immune-enhancement through Fc-receptor binding to cells (Schlesinger and Brandriss, 1981). A subsequent study investigated the role of IgM in protecting WNV-challenged mice defective in their immune response (Diamond et al., 2003). This study suggested that IgM was critical in limiting early viral replication and could protect animals from virus challenge. Not unexpectedly, passive transfer of a virus-reactive, but non-neutralizing IgM MAb had no effect on viral replication (Diamond et al., 2003). Because we had the ability to re-engineer this mMABs, we created an IgM with the same Fab as the paired IgG. This cIgM retained the virus-neutralizing and virus binding activity of 2C9-cIgG, Fig. 2.

While treatment with this 2C9-cIgM reduced viral loads in surviving mice, it was at least 100-fold less protective than 2C9-cIgG. It is well known that both IgG and IgM enter the blood stream following i.p. inoculation (Maitta et al., 2004; Faria-Neto et al., 2006; Dadchova et al., 2007; Nakouzi et al., 2008; and Han, 2010) and have pharmacological activity. Therefore, the lack of *in vivo* protective capacity of the 2C9-cIgM could be due to either its inability to cross the blood-brain barrier (unlike IgG), or its lack of a free Fc-region, while maintaining its *in vitro* virus-neutralizing capacity. In this regard, we have shown previously that monoclonal IgG antibody with high *in vitro* virus-neutralizing activity, loses its capacity to protect mice from a challenge with Venezuelan equine encephalitis – a neurotropic alphavirus – if the Fc portion of the antibody is removed prior to i.p. inoculation (Mathews et al., 1985).

In this study we have demonstrated that the AG129/17D-204 model of YF is an effective model for evaluating the protective activity of a MAbs. The 2C9-cIgG shows promise as a possible prophylactic or therapeutic agent for YFV infection and may be useful in individuals for whom vaccination with 17D-204 is contraindicated or for unvaccinated individuals who contract YFV infections. Furthermore, both the 2C9-cIgG and 2C9-cIgM may have utility as YFV-specific human immunoglobulins for use in serodiagnosis of YFV infection.

Acknowledgments

This research was supported by NIH/NIAID grant U54AI-065357 to the Rocky Mountain Regional Center of Excellence for Biodefense and Emerging Infectious Disease Research (<http://www.rmrc.colostate.edu/>), the U.S. Centers for Disease Control and Prevention.

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There are no specific treatments for human yellow fever (YF) infection.

We have created human/murine chimeric IgG (cIgG) and cIgMantibodies for YF virus.

The cIgG prophylactically protects AG129 mice from peripheral 17D-204 challenge.

The cIgGs therapeutically active at 1 day post-infection with 17D-204.

The cIgM has no antiviral activity *in vivo*.

Fig. 1.
Nucleotide and deduced amino acid sequences of m2C9 variable regions. (Panel a), Variable region of the κ -chain. (Panel b), Variable region of the γ -chain. +1 denotes start of mature protein; solid line box, CDR 1; dotted box, CDR 2; dashed box, CDR 3; the splice points of J κ 2 to C κ (A) and of J H_2 to the C region gene (B) are indicated by the arrows.

Fig. 2. End-point titration of purified cMAbs. Symbols: 2C9-cIgG (●); 2C9-cIgM (◆); cIgG 1GD5 (alphavirus cross-reactive) (■); cIgM 1MD11 (alphavirus cross-reactive) (▲). IgM cMAbs were titrated in MAC-ELISA and the absorbance was measured at 450nm; IgG cMAbs were titrated in indirect ELISA and absorbance was measured at 405nm and titration results were superimposed. Insert: YFV neutralization end-point titres of MAbs m2C9, 2C9-cIgG, 2C9-cIgM, and cIgG 1GD5, expressed as PRNT₅₀ reciprocal end-points.

Fig. 3.

Survival curves for antibody-treated mice. Panel a, survival of AG129 mice that were passively transferred intraperitoneally 127 μ g of m2C9 (■), 2C9-cIgG (●), 2C9-cIgM (◆), or PBS (▲), 24 hours prior to i.p. challenge with 17D-204. Clinical signs began after 11 days p.i. so graphs show surviving animals from days 12 to 31, after which experiments were terminated. Panel b, survival of AG129 mice challenged i.p. with 17D-204 and administered i.p. 127 μ g of m2C9 (■) or 2C9-cIgG (●) at 24 h or m2C9 (□) or 2C9-cIgG (○) 72 h p.i. A survival curve for PBS control mice is included for comparison (▲). Mice were monitored as in Panel a.

Table 1

Oligonucleotide primers for amplification and modification of antibody V-regions

Primer Name	Sequence ^a
First Round Primers	
M-IgG2a	5' ATTCGGAT <u>AGATCT</u> AGTGGATAGACCGATGG 3'
MK-REV	5' ATTCGGAT <u>AGATCT</u> TGGATGGTGGGAAGATG 3'
MHV-5	5' ACTAGTCGACATGGACTCCAGGCTCAAT TTAGTTTTCCTT 3'
MKV-4	5' ACTAGTCGACATGAGGGCCCTGCTCACTTTTTTGGCTTCTTG 3'
Second Round Primers	
VK-5'	5' TCACGAAGTCTAGACTGGACATGAGGGCCCTGCTCACTTTTTTGG 3'
VK-3'	5' GAATCTATGGATCCTGACACACTTACGCTTTATTTCCAACCTTGTCCTCCGA 3'
VH-5'	5' <u>ACACTATA</u> CTCGAGCTCACGATGAACCTTGGGCTCAGCTTGATTT 3'
VH-3'	5' <u>TTCAGATCA</u> AGCTTGACACACTTACCTGAGGAGACGGTGACTGAGGTTC 3'

^aRestriction endonuclease sites are underlined

Table 2

Prophylactic and therapeutic activity of m2C9, 2C9-H4 cIgG, and 2C9-H4 cIgM in 17D-204-challenged AG129 mice.

Treatment ^a	Survivors/Total Mice (% Survivors)	Brain Viral Load ^b		
		Survivors	Dead	AST ^c
PBS Controls	1/27 (4)	n.a.	6.9 (0.6)	17.3 (4.3)
Prophylaxis Day -1				
m2C9 IgG	18/19 (95) ^d	1.2 (0.8) ^e	7.0	30.4 (2.5) ^d
2C9-H4 IgG	18/25 (72) ^d	0.7 (0.4) ^e	6.7	28.3 (4.5) ^d
2C9-H4 IgM	2/10 (20)	6.0 (0.1) ^f	7.7 (0.3)	20.6 (5.9)
Treatment Day 1				
m2C9 IgG	7/10 (70) ^d	2.0 (2.0) ^e	7.4 (3.0)	25.8 (8.8) ^d
2C9- H4 IgG	3/15 (20)	1.7 ^d	6.7 (0.5)	19.4 (7.0)
Treatment Day 3				
m2C9 IgG	0/10 (0)	n.a.	6.6 (0.3)	15.8 (2.8)
2C9-H4 IgG	0/15 (0)	n.a.	6.7 (4)	15.3 (2.0)

^a Mice were inoculated i.p. with 127 µg purified MAb 24 hours pre-infection or 24 or 72 hours post-infection by an i.p. viral challenge.

^b Expressed as mean (s.d.) genomic copies (log₁₀)/g tissue.

^c Experiments were terminated at 31 days post-infection. Average survival time (AST) in days (s.d.).

^d $\rho < 0.05$ by two-tailed t-test statistic vs. PBS control mice.

^e $\rho < 0.05$ by two-tailed t-test statistic vs. paired PBS control mice or paired dead mice.

^f $\rho < 0.05$ by two-tailed t-test statistic vs. paired dead mice only.

Table 3

Prophylactic dose-response comparison of m2C9 and 2C9-H4 IgG in 17D-204 challenged AG129 mice.

Group	μ g Antibody ^a	Survivors/Total Mice (% Survivors)	Brain Viral Load ^b		AST ^c
			Survivors	Dead	
PBS Controls	n/a	0/5 (0)	n.a.	6.9 (0.6)	18.8 (5.9)
2C9-H4 IgG	127	4/5 (80) ^d	2.2 (2.5) ^{d,e}	6.74	28.8 (4.9) ^d
	12.7	10/10 (100) ^d	1.0 (0.4) ^d	n.a.	31 (0) ^d
	1.27	2/10 (20)	5.1 (0.7) ^{d,e}	6.7 (0.3)	19.2 (6.9)
m2C9 IgG	127	5/5 (100) ^d	0.9 (0.3) ^d	n.a.	31 (0) ^d
	12.7	8/10 (80) ^d	1.5 (1.3) ^{d,e}	6.5 (0.1)	27.0 (6.9) ^d
	1.27	6/10 (60) ^d	3.9 (2.4) ^{d,e}	6.7 (0.5)	26.1 (6.8) ^d

^a Mice were inoculated i.p. with varying doses of purified MAb 24 hours prior to i.p. virus challenge. Experiments were terminated at 31 days post-infection.

^b Expressed as mean (s.d.) of the genomic copies (log10)/g tissue.

^c Average survival time (AST) in days (s.d.).

^d $p < 0.05$ by two-tailed t-test statistic vs. paired PBS control mice.

^e $p < 0.05$ by two-tailed t-test statistic vs. paired dead mice.