



A Murine Monoclonal Antibody With Potent Neutralization Ability Against Human Adenovirus 7

Rong Wang[†], Jiansheng Lu[†], Quan Zhou, Lei Chen, Ying Huang, Yunzhou Yu and Zhixin Yang^{*}

Laboratory of Protein Engineering, Beijing Institute of Biotechnology, Beijing, China

OPEN ACCESS

Edited by:

Milan Surjit,
Translational Health Science and
Technology Institute (THSTI), India

Reviewed by:

Santosh Dhakal,
Johns Hopkins University,
United States
Yucel Aydin,
Tulane University, United States

*Correspondence:

Zhixin Yang
yy_xiao@126.com

[†]These authors have contributed
equally to this work

Specialty section:

This article was submitted to
Virus and Host,
a section of the journal
Frontiers in Cellular and Infection
Microbiology

Received: 17 August 2019

Accepted: 21 November 2019

Published: 04 December 2019

Citation:

Wang R, Lu J, Zhou Q, Chen L,
Huang Y, Yu Y and Yang Z (2019) A
Murine Monoclonal Antibody With
Potent Neutralization Ability Against
Human Adenovirus 7.
Front. Cell. Infect. Microbiol. 9:417.
doi: 10.3389/fcimb.2019.00417

B1-type human adenoviruses (HAdVs) HAdV-3, HAdV-7, and HAdV-55 have caused epidemics in North America, Asia, and Europe. However, to date, no adenovirus vaccines or antiviral drugs have been approved for general use. In the present work, a scFv-phage immune library was constructed and mouse monoclonal antibody (MMAb) 10G12 was obtained through selection. 10G12 is specific for HAdV-7 and binds the hexon loop1 and loop2 (LP12), resulting in strong neutralization activity against HAdV-7. Additionally, it is stable in serum and buffer at various pH values. The findings provide insight into adenovirus and antibody responses and may facilitate the design and development of adenovirus vaccines and antiviral drugs.

Keywords: human adenovirus type 7 (HAdV-7), mouse monoclonal antibody (MMAb), immune library, neutralizing antibody, antiviral drugs

INTRODUCTION

Human adenoviruses (HAdVs), non-enveloped, icosahedral, double-stranded DNA viruses spanning > 85 genotypes, are classified into seven species (A–G) (Yoshitomi et al., 2017). HAdV infection is characterized by a broad spectrum of disease symptoms in humans, including sore throat, pneumonia, fever, and acute otitis media, with most cases involving gastrointestinal symptoms that vary with infection genotype (Arnold et al., 2010; Kunz and Ottolini, 2010). Symptoms are generally mild and self-limiting in immune-competent adults, but outbreaks of acute respiratory diseases (ARDs), such as community-acquired pneumonia (CAP), can occur in newborns, school students, and military recruits (Tan et al., 2016). B1 type adenoviruses HAdV-3, HAdV-7, and HAdV-55 are responsible for most epidemics in North America, Asia, and Europe (Choi et al., 2005; Zhang et al., 2006; James et al., 2007; Selvaraju et al., 2011; Tang et al., 2011; Gopalkrishna et al., 2016). To date, no vaccines for the general population available for HAdVs, and only vaccines against HAdV types 4 and 7 have been developed for the USA military (Russell et al., 2006; Kajon et al., 2015). Additionally, no antiviral drugs or efficient antiviral therapies have been approved for treating HAdVs (Echavarría, 2008).

Immune reconstitution plays a critical role in controlling AdV infection, and serotype-specific neutralizing monoclonal antibodies (MAbs) correlate with clearance of AdV (Heemskerck et al., 2005; Echavarría, 2008). The adenovirus capsid is formed from three major proteins (hexon, penton base, and fiber) and four minor proteins (IIIa, VI, VIII, and IX). Hexon, the most abundant capsid protein, recruits cytoplasmic dynein, a crucial component for transporting viral capsids along microtubules (Scherer and Vallee, 2015). Hexon is also an important antigen for neutralizing antibodies against HAdV-3, -5, -7, -14, and -55 (Sumida et al., 2005; Tian et al., 2011; Yu et al., 2013; Su et al., 2016). Type-specific neutralization epitopes on hexon proteins of many adenoviruses

are primarily located in seven hypervariable regions (Rux et al., 2003; Pichla-Gollon et al., 2007; Yuan et al., 2009; Bradley et al., 2012; Qiu et al., 2012; Tian et al., 2018b). Hexon stimulates type-specific neutralizing Abs (NAbs), whereas fiber induces Abs with cross-neutralizing activity against HAdV-14 and HAdV-55 (Feng et al., 2018). However, to date, only a few neutralization epitopes on the HAdV fiber knob region have been identified.

In the present study, scFv 10G12 was screened from an scFv-phage antibody immune library and subcloned to generate an MAb. We identified MAb 10G12 as a potent antibody that effectively targets HAdV-7 *in vitro* at low concentrations by binding to hexon loop1 and loop2 (LP12). MAb 10G12 displayed good stability in serum and phosphate buffer (PB) at different pH values.

MATERIALS AND METHODS

Cell Lines and Viruses

HEK293F and A549 cells (ATCC, USA) were cultured in Dulbecco's modified Eagle's medium (DMEM) containing 10% fetal bovine serum (FBS) (Excell, China). FreeStyle™ 293-F cells (Invitrogen, USA) were cultured in FreeStyle™ 293 Expression Medium (12338; Gibco, USA). Cells were incubated at 37°C in a 5% CO₂ atmosphere. The HAdV-7 GZ6965 strain (human/CHN/GZ6965/2001) used herein was obtained as described previously (Qiu et al., 2012) and maintained in our laboratory. The HAdV-55 strain was isolated from a patient and kindly provided by Prof. Hongbin Song (Center for Disease Control and Prevention of Chinese PLA, Beijing, China). HAdV-7 and HAdV-55 were propagated in HEK293-F cells grown in DMEM containing 2% FBS. When 75–95% of cells exhibited typical cytopathic effects (CPEs) consistent with HAdV infection, the cell suspension was frozen at –80°C and thawed three times, centrifuged at 4,000 g for 5 min, and the supernatant was inactivated and purified using standard CsCl gradient centrifugation (Wu et al., 2002). The obtained virus particles were aliquoted and stored at –80°C.

Construction and Selection of scFv-phage Antibody Immune Libraries

Preparation and characterization of the scFv-phage display library was subsequently performed. Female BALB/c mice at 6–8 weeks old were immunized with inactivated HAdV-7. Pre-immune sera were collected from mouse tails and used as negative controls. A 100 µg sample of inactivated HAdV-7 emulsified in Freund's complete adjuvant (Sigma, USA) was intraperitoneally injected, followed by four boosters of the same dose at 2-week intervals. Spleens were harvested 3 days after the final booster, and total RNA was isolated from spleen cells and was reverse transcribed into cDNA (K1621, Thermo Scientific, USA). Primers for reverse-transcription were PmCGR (TGCATTTGAACTCCTTGCC) and PmCKR (CCATCAATCTTCCACTTGAC). Full-length variable light (V_L) and variable heavy (V_H) chain genes were amplified by overlap-extended PCR and the scFv fragment was cloned into phage display vector pADSCFV-S. Competent *Escherichia coli* HST08 Blue cells were transformed with the ligation mixture by

electroporation. Transformed cells were titrated on agar plates to determine the library size, and colony PCR was performed on a selection of colonies to determine the presence of DNA inserts in the vector. Harvested cells samples harboring the final scFv antibody gene library were combined, aliquoted, and stored at –80°C.

Purified HAdV-7 (300 ng, 100 µl) in PBS was incubated in a microtiter plate well overnight at 4°C, then blocked with 3% BSA in TBS (50 mM Tris-HCl pH 7.5, 150 mM NaCl) for 2 h at 37°C. A 100 µl sample of phage library at 1.9×10^7 plaque-forming units (pfu) per ml was added and incubated for an additional 2 h at 37°C after a washing step. After washing, wells with TBST (TBS containing 0.05% Tween-20), bound phage was eluted with 120 µl 0.1 M glycine-HCl (pH 2.2) and neutralized with 15 µl 1 M Tris-HCl (pH 9.0). After eluting, phage was amplified by infecting *E. coli* XL1-Blue cells, and four rounds of panning were carried out. Positively selected phages were amplified and resulting scFv was subjected to DNA sequence.

MAb Generation of scFv

PCR was performed to amplify the full-length variable light (V_L) and variable heavy (V_H) chain genes of positively selected phages. PCR products were digested with restriction endonucleases *Sal* I and *Age* I, then cloned separately into pMABG1 and pMABKa vectors containing a mouse immunoglobulin constant gene. Recombinant antibodies were obtained as IgG1 molecules, regardless of their original isotype. FreeStyle 293-F cells were transfected with equal quantities of plasmids encoding heavy and light chains using a FectoPRO transfection kit (116-001, Polyplus-Transfection, French) for antibody expression. At 4 days after transfection, antibody-containing supernatants were harvested, and antibodies were purified using HiTrap MabSelect Xtra (28-4082-60, GE Healthcare, USA).

Expression and Purification of Loop1 and Loop2 (LP12) and Fiber

Viral DNA was extracted from A549 cells infected with HAdV-7 or HAdV-55 using DNAzol (Vigorous, Beijing, China) following the manufacturer's instructions. Genes encoding the hexon LP12 fragment and fiber were amplified by PCR and inserted into the pTIG-TRX vector. Primers used for PCR are listed in **Table 1** (7LP12, LP12 of HAdV-7; 55LP12, LP12 of HAdV-55; 7Fiber, Fiber of HAdV-7; 55Fiber, Fiber of HAdV-55). The pTIG-TRX-LP12 plasmid was transformed into *E. coli* BL21 (DE3) cells (TransGen, Beijing, China) for expression of His-tagged fusion protein. Transformed cells were cultured in Luria-Bertani medium containing ampicillin at 37°C, and recombinant expression was induced with 0.6 mM isopropyl b-D-thiogalactoside (IPTG) when the absorbance at 600 nm (OD₆₀₀) reached 0.4–0.6. After reducing the temperature from 37°C to 16°C, cells were cultured for a further 16 h. Bacteria were lysed by ultrasonic treatment, and recombinant protein was purified by Ni-agarose resin.

ELISA Assay

Wells of ELISA assay plates (9018, Costar, USA) were coated with 200 ng antigen and incubated overnight at 4°C. Wells were

TABLE 1 | Sequences of primers used for PCR.

Primer name	Primer sequences (5'-3')
7LP12-F	AGGAATCTAATGGGATCCACCACCATCATCATCATCATCTCAGTGGATAGTTACA
7LP12-R	GTGCTCGAGCTCGAGCTATTAATTGTCCATTGGGTCAAG
55LP12-F	AGGAATCTAATGGGATCCACCACCATCATCATCATAGTTTCAAACCTATTCTGGTAC
55LP12-R	GTGCTCGAGCTCGAGCTATTACCGCCGTTTCATGTAGTCGTA
7Fiber-F	AGGAATCTAATGGGATCCACCACCATCATCATCATACCAAGAGAGTCCGGCTCA
7Fiber-R	GGTGTCTCGAGTCATTAGTCGTCTTCTCTGTATGTA
55Fiber-F	AGGAATCTAATGGGATCCACCACCATCATCATCATACCAAAGAGTCCGGCTCAGT
55Fiber-R	GGTGTCTCGAGTCATTAGTCGTCTTCTCTGTATGTA

then blocked with 200 μ l of 5% (w/v) skimmed milk-PBS for 2 h at 37°C, and 200 μ l of antibody was added and incubated for 2 h at 37°C. Plates were washed three times with PBS-Tween (0.1% v/v), and goat anti-mouse horseradish peroxidase (HRP)-conjugated IgG antibody (1:5,000, v/v) was added and incubated for 1 h at 37°C. Finally, three rounds of washing with PBS-Tween were carried out, and detection at 492 and 630 nm was performed using OPD chromogen substrate.

To examine whether MAb 10G12 was specific for HAdV-7, inactivated HAdV-7 and HAdV-55 were used as antigens, and inactivated influenza virus H3N2 (A/swine/Colorado/1/77) (Karasin et al., 2000) and a synthetic polypeptide antigen of foot-and-mouth disease virus (FMDV) (ETQVQRRQHTDVSFILDRFVKVTPKDQINALDLMQTPAHTEPGSRVTNVRGDLQVLAQKAARTLPPGSRHKQKIVAPVKQLL) served as negative controls. The MAb antibody 10G12 was used at a concentration of 7.5 μ g/mL. To examine the affinity of MAb 10G12 for HAdV-7, inactivated HAdV-7 was used as antigen, and the MAb 10G12 antibody was 2-fold serially diluted from an initial concentration of 80 μ g/mL to 19.07 pg/mL. To identify which epitope MAb 10G12 binds, fiber, and the hexon LP12 from HAdV-7 and HAdV-55 were used as antigens, and MAb 10G12 antibody employed at 7.5 μ g/mL.

Virus Neutralization Test

For *in vitro* adenovirus neutralization experiments, 100 μ l of A549 cells (3×10^5 cells/ml) were seeded in each well of 96-well plates incubated overnight at 37°C in a 5% CO₂ atmosphere. Purified 10G12 was serially diluted 2-fold from 25 to 0.1 μ g/ml in DMEM, and 50 μ l aliquots of each dilution were mixed with 50 μ l HAdV-7 or HAdV-55 with 100TCID₅₀. Anti-DENV1 (Lu et al., 2018) and anti-EGFR (CN102993305B) antibodies served as negative controls. Antibody-virus mixtures were incubated at 37°C for 1 h, transferred to 96-well plates containing 85–95% confluent A549 cell monolayers, and cultured in DMEM without Phenol Red or serum for 72 h. Infected cells were observed under a microscope and the number of holes in cells with lesions was counted.

To test the ability of MAb to rescue HAdVs infection, 100 μ l of A549 cells (3×10^5 cells/ml) were seeded in each well of 96-well plates and incubated overnight at 37°C in a 5% CO₂ atmosphere. Next, 100 μ l samples of HAdV-7 or HAdV-55 with 100TCID₅₀ were added to 96-well plates containing 85–95% confluent A549 cell monolayers and incubated at 37°C for 1 h. Purified 10G12 was serially diluted 2-fold from 25 to 0.1 μ g/ml in DMEM without Phenol Red and serum, and 100 μ l aliquots of each dilution were added to 96-well plates and incubated for 1 week at 37°C. Anti-DENV1 and anti-EGFR antibodies served as negative controls. Infected cells were observed under a microscope and the number of holes in cells with lesions was counted.

MAb Stability Analysis

To test the stability of MAb in serum, purified 10G12 was diluted in fetal bovine serum (FBS, Excell) to 25 μ g/ml and incubated for 3, 7, or 10 days at 37°C. ELISA assays were then performed to detect whether samples still efficiently recognized HAdV-7.

To test the MAb stability in PB at different pH values, purified 10G12 was diluted in PB at pH 6.0, 6.5, 7.0, 7.5, and 8.0, and incubated at 37°C for 5 or 8 days. ELISA assays were then performed to detect whether samples still efficiently recognized HAdV-7.

Western Blotting Assay

Purified antigens were quantified using a NanoDrop One^c instrument (Thermo, USA). A 7.5 μ g sample of reduced protein was separated by SDS-PAGE and subsequently transferred to a polyvinylidene fluoride (PVDF) membrane. Primary antibody 10G12 (1 mg/mL) was diluted 1:1,000, and secondary antibody HRP-goat anti-mouse immunoglobulin G (IgG) (ZSGB-BIO) was diluted 1:5,000. Signals were detected using Western HRP Substrate Peroxide solution (Millipore).

Statistical Analysis

All experiments were repeated at least three times, except for the stability assay. Data are presented as means \pm standard deviation (SD). Statistical significance was determined using GraphPad Prism 5.0 software. An affinity graph was plotted, and EC₅₀ values were determined using GraphPad Prism 5.0 software, too. The significance of differences in protective effects compared with controls was evaluated using two-tailed Student's *t*-tests, and *p*-values < 0.05 were considered statistically significant.

RESULTS

Construction and Selection of scFv-phage Antibody Immune Libraries

Female BALB/c mice at 6–8 weeks old were immunized with inactivated HAdV-7, and spleens were harvested for RNA extraction after four booster injections. Genes encoding V_L and V_H chains were amplified by PCR, and DNA fragments of the expected size (350 bp) were obtained. Overlay-extended PCR was performed to generate scFv DNA fragments of ~750 bp, which were then cloned into the phage vector pADSCFV-S. The final

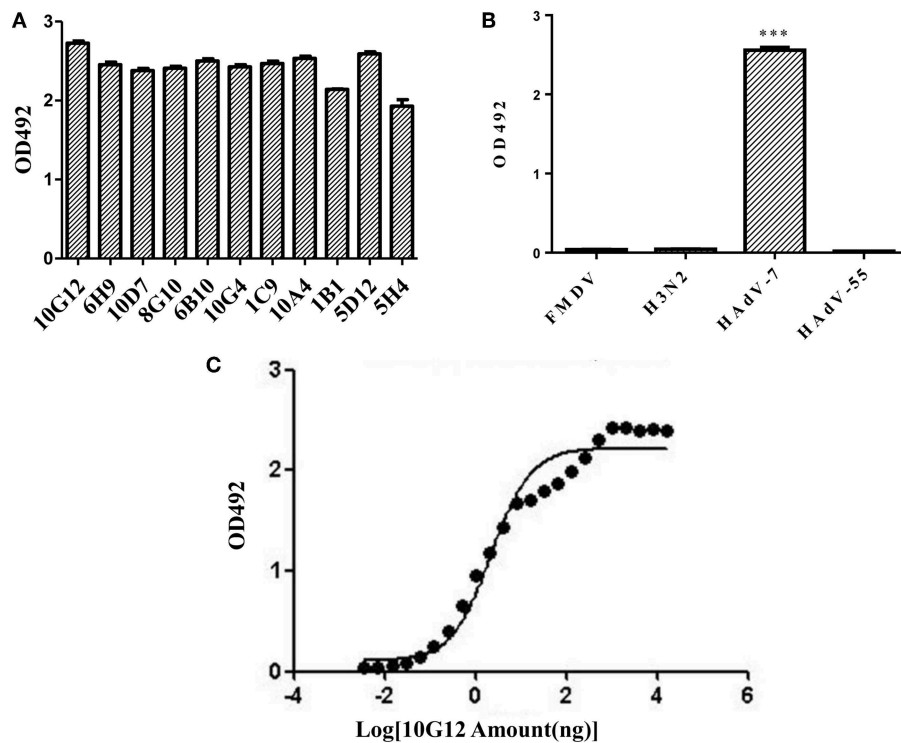


FIGURE 1 | Identification of mouse monoclonal antibodies (MMABs) against HAdV-7. **(A)** Screening of scFv-displaying phage by ELISA. After three round of panning, 11 positive clones were identified that recognized HAdV-7. **(B)** ELISA analysis of binding between MAb 10G12 and various antigens. The synthetic polypeptide antigen of foot-and-mouth disease virus (FMDV) and influenza A virus H3N2 (*A/swine/Colorado/1/77*) served as negative controls. HAdV-55 was tested for potential cross-reactivity. Results are presented as means \pm SD from three independent experiments (** $p < 0.001$ vs. negative controls calculated by *t*-tests). **(C)** Affinity curve of the ELISA results for binding between HAdV-7 and serially diluted MAb 10G12. ELISA assays were performed with 200 ng of inactivated HAdV-7 per well. MAb 10G12 was serially diluted from an initial concentration of 80 μ g/mL.

scFv antibody gene library consisted of 1.9×10^7 independent clones, with 80% correctness.

In total, 11 positive clones were identified from samples after the third round of panning against HAdV-7, and their ability to interact with HAdV-7 is shown in **Figure 1A**. These phagemids were extracted and each insert was sequenced. The results revealed five unique full-size scFv sequences among the 11 clones (10G12, 6H9, 10D7, 8G10, 6B10, 10G4, and 1C9 shared the same sequence; 5H4 had the wrong sequence). V_H and V_L of 10G12, 10A4, 1B1, and 5D12 were recloned into pMABG1 or pMABKa to generate murine IgG1 molecules. Although these four antibodies were specific to HAdV-7, 10A4, 1B1, and 5D12 did not exhibit neutralizing activity (data not shown). Thus, subsequent experiments only characterized 10G12.

MMAB 10G12 Is Specific for HAdV-7

To examine whether MAb 10G12 is specific for HAdV-7, an ELISA assay was performed. Since positive clones bound to various unrelated viruses in the previous selection of the scFv-phage antibody library, two unrelated antigens (influenza virus H3N2 and the synthetic polypeptide antigen of FMDV) were included as negative controls. Additionally, since there are more than 85 HAdV genotypes, HAdV-55 was tested for potential cross-reactivity with MAb 10G12. As shown in

Figure 1B, MAb 10G12 bound to inactivated HAdV-7, but not to inactivated influenza virus H3N2, the synthetic polypeptide antigen of FMDV, or HAdV-55. Furthermore, ELISA assay was performed to test the affinity of MAb 10G12 for HAdV-7. Based on the absorbance at 492 nm, and affinity graph was plotted using GraphPad Prism 5.0 software (**Figure 1C**). The resulting EC_{50} values indicated that the affinity between MAb and HAdV-7 was 0.14 nM.

In vitro Neutralizing Activity and Therapeutic Effects of MAb 10G12

To examine the neutralization potential of MAb 10G12, *in vitro* adenovirus neutralization experiments were performed using A549 cells. Purified 10G12 was serially diluted 2-fold from 25 to 0.1 μ g/ml in DMEM, and 50 μ l aliquots of each dilution were mixed with 50 μ l HAdV-7 or HAdV-55 with 100TCID₅₀. The antibody-virus mixtures were transferred to A549 cells, and every dilution included eight replicates. After 72 h, infected cells were observed under the microscope, and wells containing surviving cells were counted. Two antibodies (anti-DENV and anti-EGFR) served as negative controls. As shown in **Table 2**, 50 μ l aliquots of 10G12 with 0.4 μ g/ml could neutralize 100% of 50 μ l HAdV-7 with 100TCID₅₀, and all cells at this dilution

TABLE 2 | *In vitro* neutralization activity of MAb 10G12 (surviving cell holes/total).

Virus	Antibody	Antibody concentration ($\mu\text{g/ml}$)								
		25	12.5	6.25	3.2	1.6	0.8	0.4	0.2	0.1
No	10G12	8/8	8/8	8/8	8/8	8/8	8/8	8/8	8/8	8/8
HAdV-7	Anti-DENV1	2/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8
HAdV-7	Anti-EGFR	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8
HAdV-7	No	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8
HAdV-7	10G12	8/8	8/8	8/8	8/8	8/8	8/8	8/8	4/8	0/8
HAdV-55	10G12	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8

TABLE 3 | *In vitro* therapeutic effects of MAb 10G12 (surviving cell holes/total).

Virus	Antibody	Antibody concentration ($\mu\text{g/ml}$)								
		25	12.5	6.25	3.2	1.6	0.8	0.4	0.2	0.1
No	10G12	8/8	8/8	8/8	8/8	8/8	8/8	8/8	8/8	8/8
HAdV-7	Anti-DENV1	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8
HAdV-7	Anti-EGFR	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8
HAdV-7	No	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8
HAdV-7	10G12	8/8	8/8	8/8	8/8	6/8	2/8	0/8	0/8	0/8
HAdV-55	10G12	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8	0/8

had no lesions. Even 50 μl aliquots of 10G12 with 0.2 $\mu\text{g/ml}$ could neutralize 50% of 50 μl HAdV-7 with 100TCID₅₀, but cells at this dilution had partial lesions. Furthermore, even 10G12 at 25 $\mu\text{g/ml}$ was unable to neutralize HAdV-55 with 100TCID₅₀. These findings indicate that MAb 10G12 exhibited strong neutralization activity against HAdV-7 and poor cross-reactivity with HAdV-55.

Next, the ability of MAb 10G12 to rescue HAdVs infection was investigated to explore potential therapeutic effects. HAdV-7 or HAdV-55 with 100TCID₅₀ were added to 96-well plates containing 85–95% confluent monolayers of A549 cells and incubated at 37°C for 1 h. Purified 10G12 was then serially diluted 2-fold from 25 to 0.1 $\mu\text{g/ml}$, added to infected A549 cells and incubated for 1 week at 37°C. Anti-DENV1 and anti-EGFR antibodies served as negative controls. Infected cells were observed under a microscope and the number of holes in cells with lesions was counted. As shown in **Table 3**, even at 1 h after infection, 3.2 $\mu\text{g/ml}$ 10G12 could rescue 100% of cells infected with 100TCID₅₀ HAdV-7, and none of the cells at this dilution displayed lesions. Even 0.8 $\mu\text{g/ml}$ 10G12 could protect 25% of cells infected with 100TCID₅₀ HAdV-7. Furthermore, even 25 $\mu\text{g/ml}$ 10G12 was unable to rescue A549 cells infected with 100TCID₅₀ HAdV-55. These findings indicate that MAb 10G12 exhibited potent therapeutic effects against HAdV-7.

MAb 10G12 Is Stable in Serum and PB at Different pH Values

To test MAb stability in serum, purified 10G12 was diluted in FBS to 25 $\mu\text{g/ml}$ and incubated in 37°C for 3, 7, and 10 days. ELISA assays were then performed to detect whether samples still efficiently recognized HAdV-7. As shown in **Figure 2A**,

compared with day 0, the binding activity of samples diluted in FBS after 3, 7, and 10 days decreased by a statistically significant amount. However, even after 10 days of incubation at 37°C, the binding activity was only decreased 20%. Additionally, from day 3 to 10, the binding activity did not decrease with increasing incubation duration. Differences between fetal bovine serum and the mouse monoclonal antibody may explain this decrease in binding activity. The results indicate that MAb 10G12 was relatively stable in serum.

To test the MAb stability in PB at different pH values, purified 10G12 was diluted in PB at pH 6.0, 6.5, 7.0, 7.5, and 8.0, and incubated at 37°C for 5 or 8 days. ELISA assays were then performed to detect whether samples still efficiently recognized HAdV-7. As shown in **Figure 2B**, samples diluted in PB at pH 6.0, 6.5, 7.0, 7.5, and 8.0 after 8 days of incubation still had the same binding activity as those before incubation. These results indicate that MAb 10G12 is stable in serum and PB at different pH values.

MAb 10G12 Binds Hexon Loop1 and Loop2

Hexon is an important antigen of neutralizing antibodies, and fiber also has a few neutralization epitopes. To determine to which epitope MAb 10G12 binds, fragments comprising loop1 and loop2 (LP12) of hexon and fiber from HAdV-7 and HAdV-55 were amplified by PCR and subcloned into pTIG-TRX. Recombinant proteins were expressed and purified (**Figure 3A**), and the results of western blotting showed that MAb 10G12 bound LP12 of HAdV-7 but not HAdV-55 or fiber (**Figure 3B**). When using reduced protein samples for SDS-PAGE, 10G12 should bind the linear epitopes of 7LP12. Thus, ELISA assay plates were coated with LP12 or fiber from HAdV-7 and HAdV-55 at 200 ng per well, and the affinity for MAb 10G12 was measured. As shown in **Figure 3C**, MAb 10G12 bound LP12 of HAdV-7 but not fiber, consistent with the results of western blotting in **Figure 3B**. However, the ELISA result showed that 10G12 bound LP12 of HAdV-55 with weaker affinity than HAdV-7. Combined with the western blotting results in **Figure 3B**, this indicates that 10G12 might bind some spatial epitopes of 55LP12. However, 10G12 did not neutralize HAdV-55 (**Table 2**), suggesting that these might be non-neutralizing spatial epitopes. In summary, MAb 10G12 clearly bound the hexon LP12 region.

DISCUSSION

In this study, we identified antibody MAb 10G12 that binds specifically to HAdV-7 through the hexon LP12 region. MAb 10G12 exhibited strong neutralization activity against HAdV-7 and was stable in serum and PB at different pH values.

Despite more than 85 genotypes have been identified for HAdVs, few neutralizing antibodies have been reported. Tian et al. (2018a) reported that the recombinant trimeric HAdV-11 fiber knob region is responsible for cross-neutralizing antibody responses against HAdV-11, -7, -14p1, and -55 in mice. Three neutralizing MAbs, 6A7, 3F11, and 3D8, were

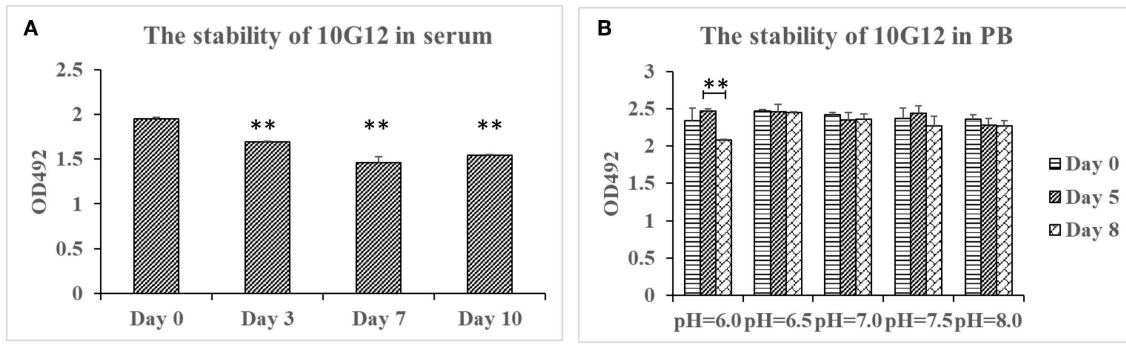


FIGURE 2 | Stability of 10G12 in serum and PB at different pH values. **(A)** ELISA analysis of the binding between HAdV-7 and MAb 10G12 in serum. Purified 10G12 was diluted in fetal bovine serum to 25 μ g/ml and incubated at 37°C for 3, 7, and 10 days. ELISA assays were performed with 200 ng of inactivated HAdV-7 per well. **(B)** ELISA analysis of binding between HAdV-7 and MAb 10G12 in PB at different pH values. Purified 10G12 was diluted in PB at pH 6.0, 6.5, 7.0, 7.5, and 8.0 and incubated at 37°C for day 5 and 8. ELISA assays were performed with 200 ng inactivated HAdV-7 per well. Data were obtained from two separate experiments, and results are presented as means \pm SD (** p < 0.01 vs. day 0 values calculated by *t*-test).

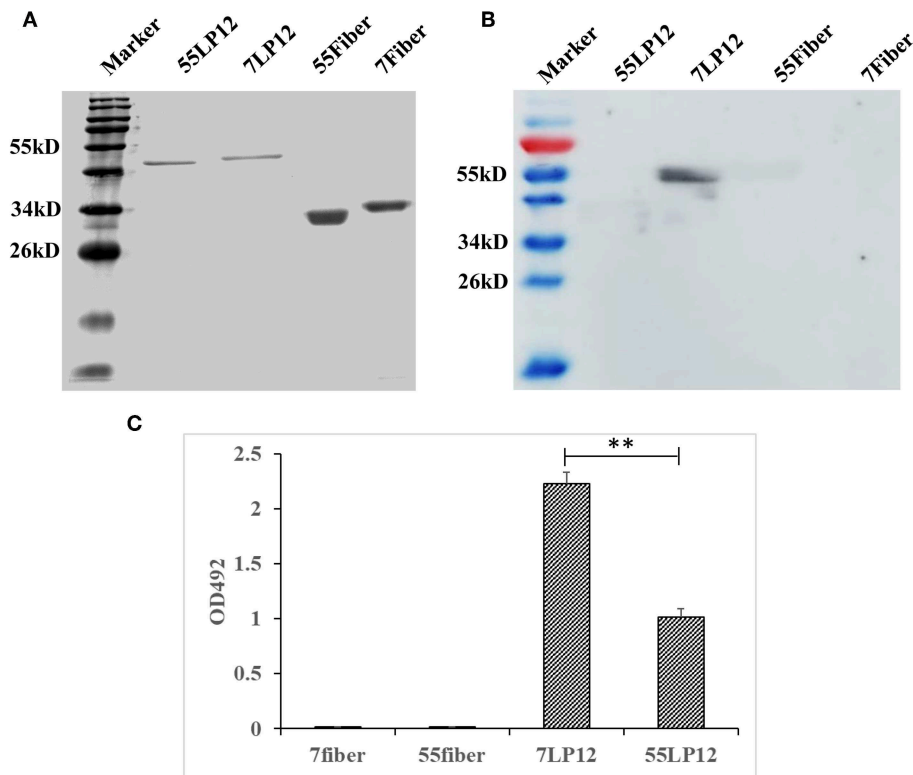


FIGURE 3 | Binding of MAb 10G12 to loops 1 and 2 (LP12) of hexon. **(A)** Reducing polyacrylamide gel electrophoresis analysis of purified antigens. A 5 μ g sample of reduced Fiber and LP12 of HAdV-7 and HAdV-55 was separated by SDS-PAGE and subsequently stained with Coomassie Brilliant Blue. **(B)** Western blotting analysis of binding between 10G12 and various antigens. A 7.5 μ g sample of reduced Fiber and LP12 of HAdV-7 and HAdV-55 was separated by SDS-PAGE and subsequently transferred to a PVDF membrane. Primary antibody 10G12 (1 mg/mL) was diluted 1:1,000, and secondary antibody HRP-goat anti-mouse IgG was diluted 1:5,000. Signals were detected using Western HPR Substrate Peroxide solution. **(C)** ELISA analysis of binding between 10G12 and various antigens. Data were obtained from three separate experiments, and results are presented as means \pm SD (** p < 0.05 between 7LP12 and 55LP12, calculated by *t*-tests).

obtained via mouse hybridoma fusion, of which 3F11 and 3D8 cross-neutralized HAdV-11,-7, and -55, but not HAdV-14p1 (Tian et al., 2018a). Feng et al. (2018) previously demonstrated that a fiber-specific antibody in sera contributed

to cross-neutralizing activity against HAdV-14 and HAdV-55. However, HAdV-55 is a recombinant chimera of HAdV-11 and -14 that contains the fiber gene from HAdV-14. Our current study shows that 10G12 does not neutralize HAdV-55, and

has poor cross-reactivity, probably through interaction with non-neutralizing epitopes.

MMAb 10G12 is a murine neutralizing antibody like 6A7, 3F11, and 3D8 (Tian et al., 2018a). Monoclonal antibodies of mouse origin can induce human anti-mouse antibody (HAMA) responses, which restricts the use of MAb in humans (Hertel et al., 1990). The first MAb, Muromonab, has been on the market since 1992, and humanized antibodies based on murine MAb are increasingly being developed (Makulska-Nowak, 1993). In future work, MAb 10G12 may be further humanized for therapeutic use.

In this study, we expressed and purified the recombinant LP12 hexon region. ELISA assay plates were coated with 200 ng of inactivated HAdV-7 or LP12 per well, and the affinity for MAb remained the same (Figures 1B, 3C). This indicates that LP12 is present in the main neutralization epitopes of HAdV-7. Previous studies confirmed that hexon protein, the most abundant capsid protein, is the predominant target of neutralizing antibodies (NAbs) recognizing HAdV-3, -5, -7, -14, and -55 (Sumida et al., 2005; Tian et al., 2011; Bradley et al., 2012; Yu et al., 2013; Su et al., 2016; Feng et al., 2018). Additionally, the affinity results indicate that LP12 could substitute for inactivated HAdV-7 in initial screening and testing.

In summary, MAb 10G12 was specific for HAdV-7 and displayed good stability. In future work, we will explore whether MAb 10G12 can provide protection against HAdV-7 *in vivo*, and if so, MAb may be further humanized for use as a therapeutic agent.

REFERENCES

- Arnold, J., Jánoska, M., Kajon, A. E., Metzgar, D., Hudson, N. R., Torres, S., et al. (2010). Genomic characterization of human adenovirus 36, a putative obesity agent. *Virus Res.* 149, 152–161. doi: 10.1016/j.virusres.2010.01.011
- Bradley, R. R., Maxfield, L. F., Lynch, D. M., Iampietro, M. J., Borducchi, E. N., and Barouch, D. H. (2012). Adenovirus serotype 5-specific neutralizing antibodies target multiple hexon hypervariable regions. *J. Virol.* 86, 1267–1272. doi: 10.1128/JVI.06165-11
- Choi, E. H., Kim, H. S., Eun, B. W., Kim, B. I., Choi, J. Y., Lee, H. J., et al. (2005). Adenovirus type 7 peptide diversity during outbreak, Korea, 1995–2000. *Emerg. Infect. Dis.* 11, 649–654. doi: 10.3201/eid1105.041211
- Echavarría, M. (2008). Adenoviruses in immunocompromised hosts. *Clin. Microbiol. Rev.* 21, 704–715. doi: 10.1128/CMR.00052-07
- Feng, Y., Sun, X., Ye, X., Feng, Y., Wang, J., Zheng, X., et al. (2018). Hexon and fiber of adenovirus type 14 and 55 are major targets of neutralizing antibody but only fiber-specific antibody contributes to cross-neutralizing activity. *Virology* 518, 272–283. doi: 10.1016/j.virol.2018.03.002
- Gopalkrishna, V., Ganorkar, N. N., and Patil, P. R. (2016). Identification and molecular characterization of adenovirus types (HAdV-8, HAdV-37, HAdV-4, HAdV-3) in an epidemic of keratoconjunctivitis occurred in Pune, Maharashtra, Western India. *J. Med. Virol.* 88, 2100–2105. doi: 10.1002/jmv.24565
- Heemskerk, B., Lankester, A. C., van Vreeswijk, T., Beersma, M. F., Claas, E. C., Veltrop-Duits, L. A., et al. (2005). Immune reconstitution and clearance of human adenovirus viremia in pediatric stem-cell recipients. *J. Infect. Dis.* 191, 520–530. doi: 10.1086/427513
- Hertel, A., Baum, R. P., Auerbach, B., Herrmann, A., and Hör, G. (1990). The clinical relevance of human anti-mouse-antibody (HAMA) in immunoscintigraphy. *Nuklearmedizin* 29, 221–227. doi: 10.1055/s-0038-1629535

DATA AVAILABILITY STATEMENT

The data used to support the findings of this study are available from the corresponding author upon request.

ETHICS STATEMENT

The animal study was reviewed and approved by Academy of Military Medical Sciences (AMMS; ID: SYXK2012–05).

AUTHOR CONTRIBUTIONS

JL and ZY conceived this study. YH, YY, and ZY carried out experiments. RW, YH, LC, and QZ performed data analysis. RW and ZY drafted, wrote, edited, and reviewed the manuscript. ZY acquired funding. JL and QZ provided resources. RW, YH, YY and ZY supervised the work.

FUNDING

This work was supported by 2015ZX09J15105-003-003.

ACKNOWLEDGMENTS

The authors thank Prof. Hongbin Song (Center for Disease Control and Prevention of Chinese PLA, Beijing, China) for kindly providing the HAdV-55.

- James, L., Vernon, M. O., Jones, R. C., Stewart, A., Lu, X., Zollar, L. M., et al. (2007). Outbreak of human adenovirus type 3 infection in a pediatric long-term care facility—Illinois, 2005. *Clin. Infect. Dis.* 45, 416–420. doi: 10.1086/519938
- Kajon, A. E., Hang, J., Hawksworth, A., Metzgar, D., Hage, E., Hansen, C. J., et al. (2015). Molecular epidemiology of adenovirus type 21 respiratory strains isolated from US military trainees (1996–2014). *J. Infect. Dis.* 212, 871–880. doi: 10.1093/infdis/jiv141
- Karasin, A. I., Schutten, M. M., Cooper, L. A., Smith, C. B., Subbarao, K., Anderson, G. A., et al. (2000). Genetic characterization of H3N2 influenza viruses isolated from pigs in North America, 1977–1999: evidence for wholly human and reassortant virus genotypes. *Virus Res.* 68, 71–85. doi: 10.1016/S0168-1702, 00154-4
- Kunz, A. N., and Ottolini, M. (2010). The role of adenovirus in respiratory tract infections. *Curr. Infect. Dis. Rep.* 12, 81–87. doi: 10.1007/s11908-010-0084-5
- Lu, J., Wang, R., Xia, B., Yu, Y., Zhou, X., Yang, Z., et al. (2018). Potent neutralization ability of a human monoclonal antibody against serotype 1 Dengue virus. *Front. Microbiol.* 9:1214. doi: 10.3389/fmicb.2018.01214
- Makulska-Nowak, E. (1993). Muromonab CD3—the first monoclonal antibody used in humans. *Pol. Tyg. Lek.* 48, 857–859.
- Pichla-Gollon, S. L., Drinker, M., Zhou, X., Xue, F., Rux, J. J., Gao, G. P., et al. (2007). Structure-based identification of a major neutralizing site in an adenovirus hexon. *J. Virol.* 81, 1680–1689. doi: 10.1128/JVI.02023-06
- Qiu, H., Li, X., Tian, X., Zhou, Z., Xing, K., Li, H., et al. (2012). Serotype-specific neutralizing antibody epitopes of human adenovirus type 3 (HAdV-3) and HAdV-7 reside in multiple hexon hypervariable regions. *J. Virol.* 86, 7964–7975. doi: 10.1128/JVI.07076-11
- Russell, K. L., Hawksworth, A. W., Ryan, M. A., Strickler, J., Irvine, M., Hansen, C. J., et al. (2006). Vaccine-preventable adenoviral respiratory illness in US military recruits, 1999–2004. *Vaccine* 24, 2835–2842. doi: 10.1016/j.vaccine.2005.12.062

- Rux, J. J., Kuser, P. R., and Burnett, R. M. (2003). Structural and phylogenetic analysis of adenovirus hexons by use of high-resolution x-ray crystallographic, molecular modeling, and sequence-based methods. *J. Virol.* 77, 9553–9566. doi: 10.1128/JVI.77.17.9553-9566.2003
- Scherer, J., and Vallee, R. B. (2015). Conformational changes in the adenovirus hexon subunit responsible for regulating cytoplasmic dynein recruitment. *J. Virol.* 89, 1013–1023. doi: 10.1128/JVI.02889-14
- Selvaraju, S. B., Kovac, M., Dickson, L. M., Kajon, A. E., and Selvarangan, R. (2011). Molecular epidemiology and clinical presentation of human adenovirus infections in Kansas City children. *J. Clin. Virol.* 51, 126–131. doi: 10.1016/j.jcv.2011.02.014
- Su, X., Tian, X., Jiang, Z., Ma, Q., Liu, Q., Lu, X., et al. (2016). Human adenovirus serotype 3 vector packaged by a rare serotype 14 hexon. *PLoS ONE* 11:e0156984. doi: 10.1371/journal.pone.0156984
- Sumida, S. M., Truitt, D. M., Lemckert, A. A., Vogels, R., Custers, J. H., Addo, M. M., et al. (2005). Neutralizing antibodies to adenovirus serotype 5 vaccine vectors are directed primarily against the adenovirus hexon protein. *J. Immunol.* 174, 7179–7185. doi: 10.4049/jimmunol.174.11.7179
- Tan, D., Zhu, H., Fu, Y., Tong, F., Yao, D., Walline, J., et al. (2016). Severe community-acquired pneumonia caused by human adenovirus in immunocompetent adults: a multicenter case series. *PLoS ONE* 11:e0151199. doi: 10.1371/journal.pone.0151199
- Tang, L., Wang, L., Tan, X., and Xu, W. (2011). Adenovirus serotype 7 associated with a severe lower respiratory tract disease outbreak in infants in shaanxi province, China. *Virol. J.* 8:23. doi: 10.1186/1743-422X-8-23
- Tian, X., Fan, Y., Liu, Z., Zhang, L., Liao, J., Zhou, Z., et al. (2018a). Broadly neutralizing monoclonal antibodies against human adenovirus types 55, 14p, 7, and 11 generated with recombinant type 11 fiber knob. *Emerg. Microbes Infect.* 7:206. doi: 10.1038/s41426-018-0197-8
- Tian, X., Qiu, H., Zhou, Z., Wang, S., Fan, Y., Li, X., et al. (2018b). Identification of a critical and conformational neutralizing epitope in human adenovirus Type 4 hexon. *J. Virol.* 92:e01643-17. doi: 10.1128/JVI.01643-17
- Tian, X., Su, X., Li, H., Li, X., Zhou, Z., Liu, W., et al. (2011). Construction and characterization of human adenovirus serotype 3 packaged by serotype 7 hexon. *Virus Res.* 160, 214–220. doi: 10.1016/j.virusres.2011.06.017
- Wu, H., Dmitriev, I., Kashentseva, E., Seki, T., Wang, M., and Curiel, D. T. (2002). Construction and characterization of adenovirus serotype 5 packaged by serotype 3 hexon. *J. Virol.* 76, 12775–12782. doi: 10.1128/JVI.76.24.12775-12782.2002
- Yoshitomi, H., Sera, N., Gonzalez, G., Hanaoka, N., and Fujimoto, T. (2017). First isolation of a new type of human adenovirus (genotype 79), species human mastadenovirus B (B2) from sewage water in Japan. *J. Med. Virol.* 89, 1192–1200. doi: 10.1002/jmv.24749
- Yu, B., Dong, J., Wang, C., Zhan, Y., Zhang, H., Wu, J., et al. (2013). Characteristics of neutralizing antibodies to adenovirus capsid proteins in human and animal sera. *Virology* 437, 118–123. doi: 10.1016/j.virol.2012.12.014
- Yuan, X., Qu, Z., Wu, X., Wang, Y., Liu, L., Wei, F., et al. (2009). Molecular modeling and epitopes mapping of human adenovirus type 3 hexon protein. *Vaccine* 27, 5103–5110. doi: 10.1016/j.vaccine.2009.06.041
- Zhang, Q., Su, X., Gong, S., Zeng, Q., Zhu, B., Wu, Z., et al. (2006). Comparative genomic analysis of two strains of human adenovirus type 3 isolated from children with acute respiratory infection in southern China. *J. Gen. Virol.* 87, 1531–1541. doi: 10.1099/vir.0.81515-0

Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Copyright © 2019 Wang, Lu, Zhou, Chen, Huang, Yu and Yang. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.