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Sudan ebolavirus long recovered survivors produce GP-specific Abs that are of the IgG1 subclass and preferentially bind Fc γ RI

Olga Radinsky¹, Avishay Edri¹, Michael Brusilovsky¹, Shlomit Fedida-Metula¹, Ariel Sobarzo¹, Orly Gershoni-Yahalom¹, Julius Lutwama⁴, John Dye³, Leslie Lobel^{1,4} & Angel Porgador^{1,2}

Ebolavirus is a highly lethal pathogen, causing a severe hemorrhagic disease with a high fatality rate. To better understand immune correlates of protection by virus specific IgG, we investigated the evolution of the Fc γ receptors (Fc γ Rs)-activating capabilities of antiviral IgG in serum samples of long recovered survivors. To this end, longitudinal serum samples from survivors of Sudan ebolavirus (SUDV) infection, studied over years, were examined for the presence of Ebola-GP specific IgG subclasses, and for their binding to Fc γ Rs. We developed a cell-based reporter system to quantitate pathogen-specific antibody binding to Fc γ RIIIA, Fc γ RIIA, Fc γ RIIB and Fc γ RI. With this system, we demonstrate that anti-GP-specific stimulation of the Fc γ RI reporter by survivors' sera was substantially high one year after acute infection, with a slight reduction in activity over a decade post infection. We further demonstrate that GP-specific IgG1 is by far the seroprevalent subclass that retained and even enhanced its presence in the sera, over ten years post infection; the prevalence of other GP-specific IgG subclasses was considerably reduced over time. In accordance, GP-specific Fc γ RI reporter response and GP-specific total IgG1 subclass correlated in the studied group of Ebola survivors. These observations are important for further informing Ebola vaccine and therapeutic development.

Ebolavirus hemorrhagic fever (EHF) is a severe disease, caused by a members of the filoviridae family, with an as yet undefined reservoir and a high case fatality rate¹. Recent outbreaks in West Africa have demonstrated the significant human and societal burden of outbreaks of this virus^{2,3}. Defining a comprehensive profile of the native humoral and cellular immune responses, which correlate with protective immune responses, is key for effective countermeasure development. Studies that examined the pathogenesis of ebolavirus infection in humans indicate that recovery is largely dependent upon, and associated with, the development of both cell-mediated and humoral immune responses⁴⁻⁶. Previous studies that examined survivors and asymptomatic cases demonstrated the presence of significant levels of virus-specific IgM and IgG associated with a temporary, early and strong inflammatory response⁷⁻⁹. In addition, recent evidence from long recovered SUDV survivors has demonstrated several distinctive profiles of immunity, which included persistent and strong IgG neutralizing humoral immunity more than a decade post infection in some survivors^{10,11}. However, other studies have also documented a significant number of convalesced patients with no residual humoral or cell mediated memory immune responses¹². As such, it is clear that a comprehensive picture of immunity to ebolavirus is lacking, as well as an understanding of the interplay between components of the human immune system. To shed greater light on immune factors that correlate with survival, we describe herein a novel study of immune responses in Sudan ebolavirus survivors, which suggest a coordinated response between the humoral recognition and activation components of immunity in ebolavirus hemorrhagic fever (EHF).

¹The Shraga Segal Department of Microbiology, Immunology and Genetics, Faculty of Health Sciences, Ben-Gurion University of the Negev, Beer Sheva, Israel. ²National Institute for Biotechnology in the Negev, Ben-Gurion University of the Negev, Beer Sheva, Israel. ³Virology Division – U.S. Army Medical Research Institute of Infectious Diseases 1425 Porter St., Fort Detrick, Frederick, Maryland, 21701, USA. ⁴Department of Arbovirology, Emerging and Re-emerging Infection, Uganda Virus Research Institute, Entebbe, Uganda. Olga Radinsky and Avishay Edri contributed equally to this work. Correspondence and requests for materials should be addressed to L.L. (email: leslie.lobel@gmail.com) or A.P. (email: angel@bgu.ac.il)

Human Fc γ receptors (Fc γ Rs) are a family of proteins that bind specifically to the Fc region of IgGs eliciting various immunological responses¹³. Measuring the Fc γ R-activating capabilities of antiviral IgG augments definition of immune correlates of protection against infections and/or infection-induced disease progression. Three different types of Fc γ receptors are displayed on the cell surface of human leukocytes: Fc γ RI (CD64), Fc γ RII (types A, B, and C, collectively known as CD32), and Fc γ RIII (types A and B, collectively known as CD16)¹⁴. Binding affinity of human IgG Fc to a corresponding Fc γ R is dictated by both the IgG-Fc subclass (IgG1, IgG2, IgG3 and IgG4) and changes in a single N-linked glycan located in the CH2 domain of the IgG Fc^{15–18}. For example, IgG1 is considered as the subclass with the highest affinity to Fc γ Rs^{19–21}; yet, fucose, galactose and sialic acid modifications decrease or increase its affinity to Fc γ RIII and Fc γ RII²².

Destruction of IgG-coated targets by cell-mediated pathways begins with an interaction between the IgG Fc region and Fc γ Rs on the surface of leukocytes. As such, several studies analyzed binding of pathogen-specific antibodies to Fc γ Rs^{23–26}. Mahan *et al.* demonstrated that dramatic differences exist in bulk IgG glycosylation among individuals in distinct geographical locations, however, HIV-specific immunization is able to overcome these differences and elicit antigen-specific antibodies with similar antibody glycosylation²⁶. Their data strongly suggest that the immune system naturally drives antibody glycosylation in an antigen-specific manner. We aimed to further study Fc γ Rs' binding of pathogen-specific Abs by developing a cell-based reporter system to quantitate antibody binding to Fc γ RIII, Fc γ RII and Fc γ RI. We then investigated sera from SUDV survivors for the SUDV glycoprotein-specific Ab response. Long recovered survivors of SUDV infection, with no additional clinical reported exposures, enables assessment of long-term B-cell memory to an isolated single infection. We observed that IgG1 is the dominant GP-specific IgG subclass that significantly remains detectable more than a decade post infection. Interestingly, it correlates with prominent binding to Fc γ RI, as compared to binding to Fc γ RIIIA, Fc γ RIIA and Fc γ RIIB.

Results

Development of four Fc γ Rs reporter system. Murine BW5147 thymoma cells, transfected with chimeras composed of CD3 ζ chain fused to the extracellular part of a certain surface receptor, can be stimulated to secrete endogenous murine IL-2 following activation of the fused surface receptor²⁷. Several studies employed these cells to study various chimeric receptors, particularly human cell surface receptors, taking advantage of the lack of endogenous TCR and ζ as well as the studied receptor in BW5147 cells^{28,29}. We developed four BW5147-based human Fc γ R reporters based on transfection of chimeric CD3 ζ -Fc γ RIIIA (CD16A), CD3 ζ -Fc γ RIIA (CD32A), CD3 ζ -Fc γ RIIB (CD32B) and CD3 ζ -Fc γ RI (CD64). Since BW5147 lacks expression of endogenous murine Fc γ receptors, we did not expect cross-reactivity derived from the function of endogenous murine Fc γ Rs³⁰. Figure 1a shows the homogenous high level expression of the transfected chimeric human Fc γ Rs in the selected BW5147 clones. To verify that binding to the transfected receptors activates the different Fc γ R reporters to secrete murine IL-2, we incubated the cells in wells coated with commercial mAbs recognizing the specific receptors (clone 3G8 anti-human Fc γ RIIIA, clone FUN-2 anti hFc γ RIIA and anti hFc γ RIIB, & clone 10.1 anti hFc γ RI). Substantial levels of murine IL-2 were secreted by the activated reporters (Fig. 1b). We then aimed to assess the response of the four reporters to the Fc of IgGs in human sera employing the protocol depicted in Fig. 1c. 96-well plates were pre-coated with commercial goat(Fab)₂-anti-human IgG(Fab)₂, then blocked and coated with titrated amounts of human sera, followed by addition of reporter cells. IL-2 secreted into the supernatant represents binding of the transfected Fc γ receptors by the upward-pointing Fc of the sera IgGs interacting with the plastic-coated anti human IgG(Fab)₂. IgG1 constitutes more than 70% of the IgGs in normal human sera, and IgG1-Fc affinity to hFc γ RI is the highest, followed by its affinity to hFc γ RIIIA^{19,21,31}. In accordance, when tested on a control human serum, Fc γ RI reporter manifested the highest levels of secreted IL-2 followed by the Fc γ RIIIA reporter (Fig. 1d).

Fc γ Rs' binding potency of long recovered SUDV survivors is similar to non-infected controls. Following testing of the Fc γ R reporters, we further investigated binding to the reporters of sera from human survivors of SUDV hemorrhagic fever. For controls, sera was collected from unexposed individuals who were from the same districts where the outbreak occurred but were never diagnosed with Ebola virus infection and/or were negative by PCR for SUDV¹². We compared sera from 14 non-infected and 14 ebolavirus decade long survivors (Fig. 2). We chose to compare sera from long recovered survivors to non-infected individuals to avoid the short-term systemic effect of the Ebola disease on circulating IgG. Sera was tested in two dilutions (1:5 K and 1:25 K) except for the Fc γ RIIB reporter (1:1 K and 1:5 K). Overall, substantial IL-2 secretion was recorded for Fc γ RIIIA and Fc γ RI reporters in both non-infected and survivor groups. IL-2 secretion recorded for the Fc γ RIIA reporter was nearly 3-fold lower in both groups, whereas IL-2 secretion recorded for the Fc γ RIIB was significantly lower as compared to the other 3 reporters (Fig. 2). To summarize, reporter-binding profiles of circulating IgG from survivors and non-infected groups did not manifest differences that could point to a specific advantage/characteristic of the survivor group.

Fc γ RI response is consistently prominent in GP-specific Ab response in long recovered SUDV survivors. We therefore further assessed the binding to Fc γ R reporters of Ebola GP-Gulu specific antibodies found in survivors' sera. We employed the protocol depicted in Fig. 3a. 96-well plates were pre-coated with recombinant GP-Gulu antigen, then blocked and coated with titrated amounts of sera, followed by addition of reporter cells. IL-2 secreted into the supernatant represented the binding of the transfected Fc γ receptors by the upward-pointing Fc of the IgGs interacting with the plastic-coated GP-Gulu antigen. We again compared sera from non-infected and ebolavirus survivors; sera were tested at two dilutions (1:200 and 1:400). Specificity of the method is demonstrated in Fig. 3b, which depicts representative serum from a 1-year survivor and from a non-infected control. As expected, serum from the non-infected control induced GP-specific zero to null IL-2

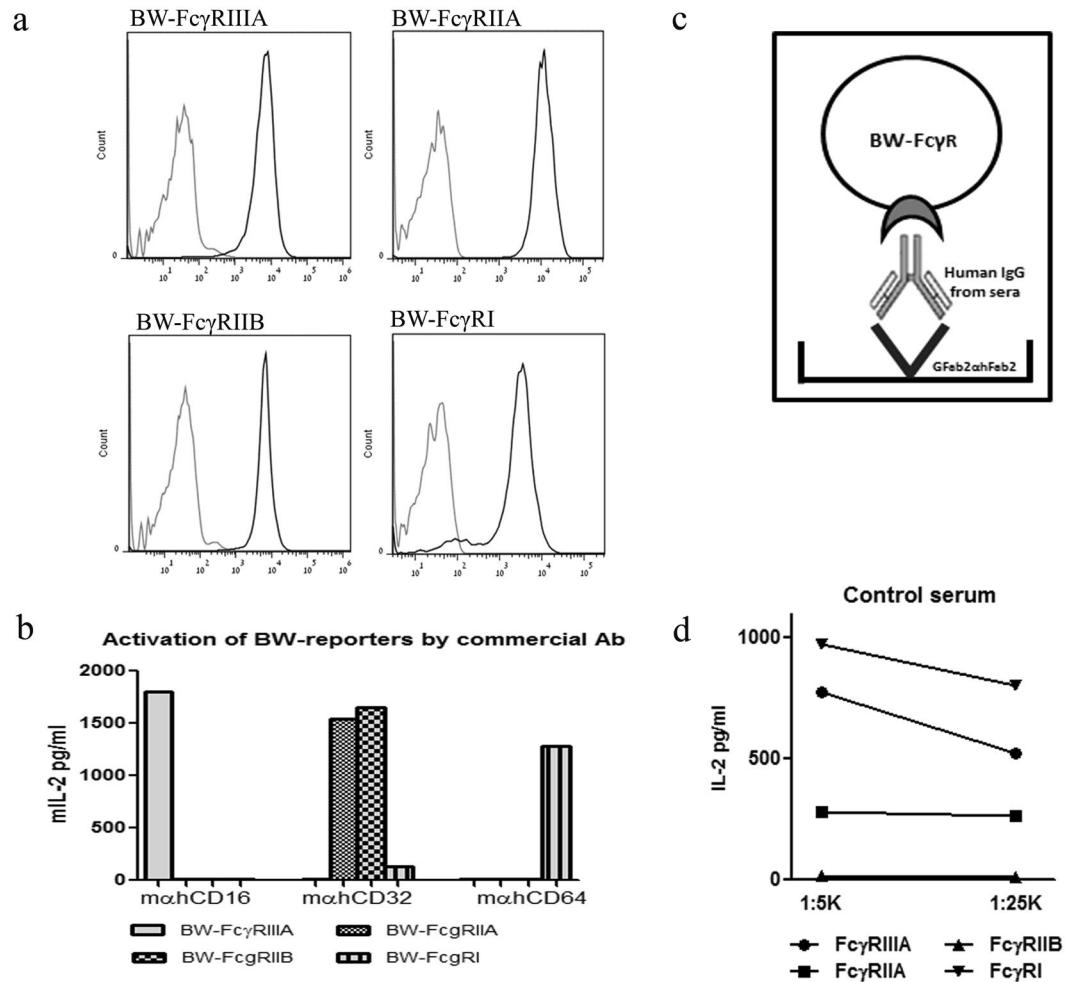


Figure 1. Production and activation of the BW-Fc γ R reporter cells by IgG. **(a)** Detection of stable transfected Fc γ Receptors was screened by flow cytometry using combined staining with anti-human CD-16/ 32/ 64 -black line, secondary APC- grey line. **(b)** Activation of BW reporter cells by commercial mAb. 96-well culture plates were pre-coated with Goat anti-mouse IgG (2 μ g/ml), blocked, washed and coated with mAbs specific for each Fc γ R 250 ng/ml. Following additional washing, BW-Fc γ R cell lines were added, 50×10^3 cell/well. mIL-2 secretion was determined by ELISA kit after 16 h incubation. **(c)** IgG mediated activation process diagram of BW-Fc γ R reporter cells. Coating of plates with goat Fab2 portion anti human Fab2 of IgG, adding human sera containing IgG followed by seeding with reporter cells 50×10^3 cells/well. mIL-2 secretion of activated cells determined by ELISA after 16 h incubation. **(d)** Activation of BW reporter cells by control sera (1:5K and 1:25K dilution). mIL-2 secretion was determined by ELISA after 16 h incubation.

secretion from all four reporters. Serum from 1-year survivors induced a very marginal IL-2 GP-specific response from the Fc γ RIIA and Fc γ RIIB reporters; GP-specific IL-2 secretion from the Fc γ RIIIA reporter reached a level of 40 pg/ml. In contrast, GP-specific stimulation of the Fc γ RI by sera from the 1-year survivor reached a level of 1400 pgs/ml. Note that differences could not be attributed to a prominent difference in the overall sensitivity of the reporters since (i) mAb-based activation of all four reporters induced similar IL-2 levels (Fig. 1b); and (ii) non-antigen specific activation by sera IgG of Fc γ RIIIA and Fc γ RI reporters induced similar IL-2 intensities (Fig. 2).

The prominent differences in GP-specific stimulation of the 4 reporters shown for the 1-year survivor, motivated us to follow specific survivors over a period of 10 years from the acute ebolavirus infection. Figure 4 summarizes the GP-specific reporter responses to sera obtained from seven Ebola survivors one, ten and thirteen years post infection (2001, 2010 and 2013, respectively). The lack of GP-specific stimulation of the Fc γ RIIA and Fc γ RIIB reporters was observed for sera from all 7 survivors and at all time points tested (Fig. 4). GP-specific stimulation of Fc γ RIIIA reporter was low but consistent over the years. GP-specific stimulation of the Fc γ RI reporter by survivors' sera was substantially higher 1 year post infection, and although with a slight reduction remained very high even 13 years post infection.

GP-specific IgG1 is the seroprevalent subclass retained in long recovered SUDV survivors, correlating with Fc γ RI response. To complement the reporter-derived GP-specific responses, we analyzed

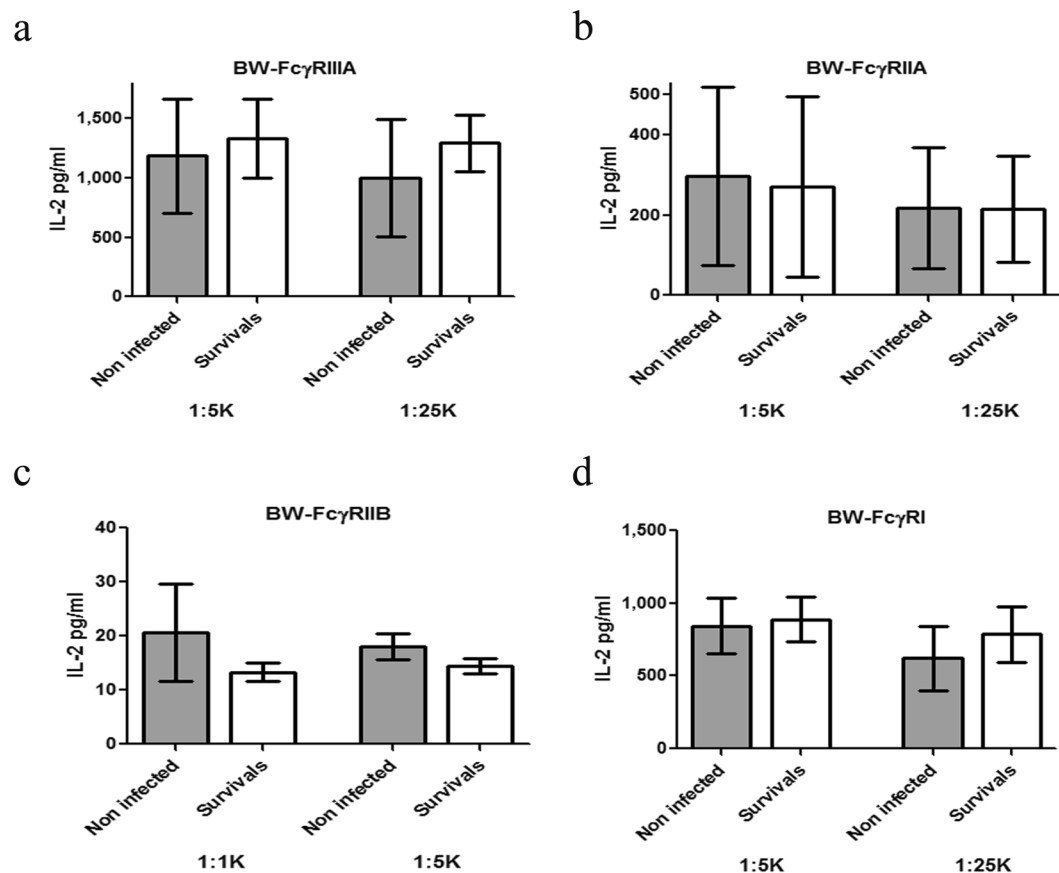


Figure 2. Binding and activation of the reporters by sera from survivors of SUDV outbreak. Total IgG dependent activation through Fc γ Rs with goat-anti human IgG. After pre-coating with goat-anti human IgG (2 μ g/ml) the activation assay was performed (as described in the legend of Fig. 1) using a total of 14 non-infected and 14 survivors' serum samples (including 5 serum samples from a longitudinal study for the GP-Gulu specificity). 1:1 K and 1:5 K sera dilution was used for BW-Fc γ RIIB response, and 1:5 K and 1:25 K for BW-Fc γ RIIA/IIA/I response. mIL-2 secretion was determined by ELISA kit after 16 h incubation.

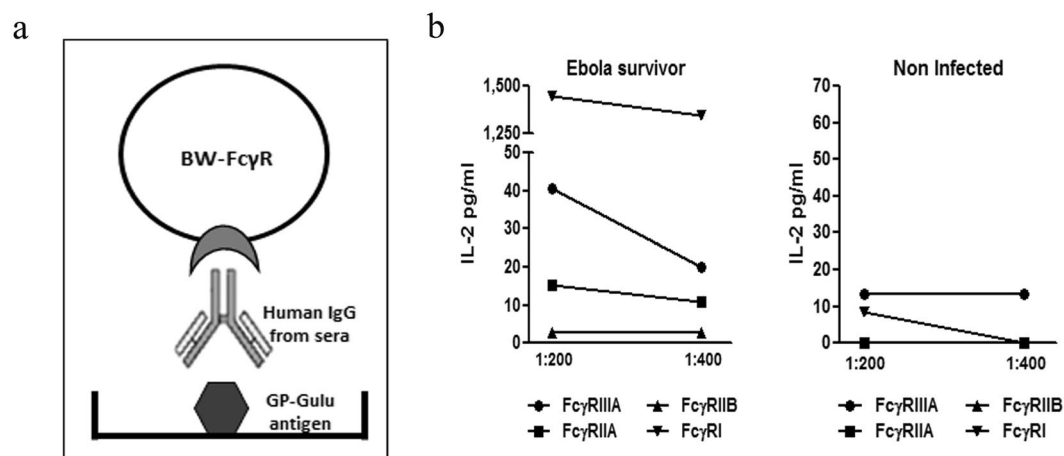


Figure 3. Binding of Ebola GP-specific antibodies found in survivors sera to Fc γ R reporters. Serum samples from non-infected controls was compared to SUDV survivors by its BW-Fc γ R activation capacity. (a) Diagram of antigen specific IgG mediated activation of BW-Fc γ R cells. Coating of plates with GP-Gulu antigen (5 μ g/ml), addition of human sera following seeding with reporter cells at 50×10^3 cells/well. mIL-2 secretion of activated cell determined by ELISA after 16 h incubation. (b) Serum samples from SUDV survivor and non-infected patient diluted 1:200 and 1:400 were added to pre-coated plates with GP-Gulu antigen (5 μ g/ml). After activation assay, mIL-2 secretion was determined by ELISA after 16 h.

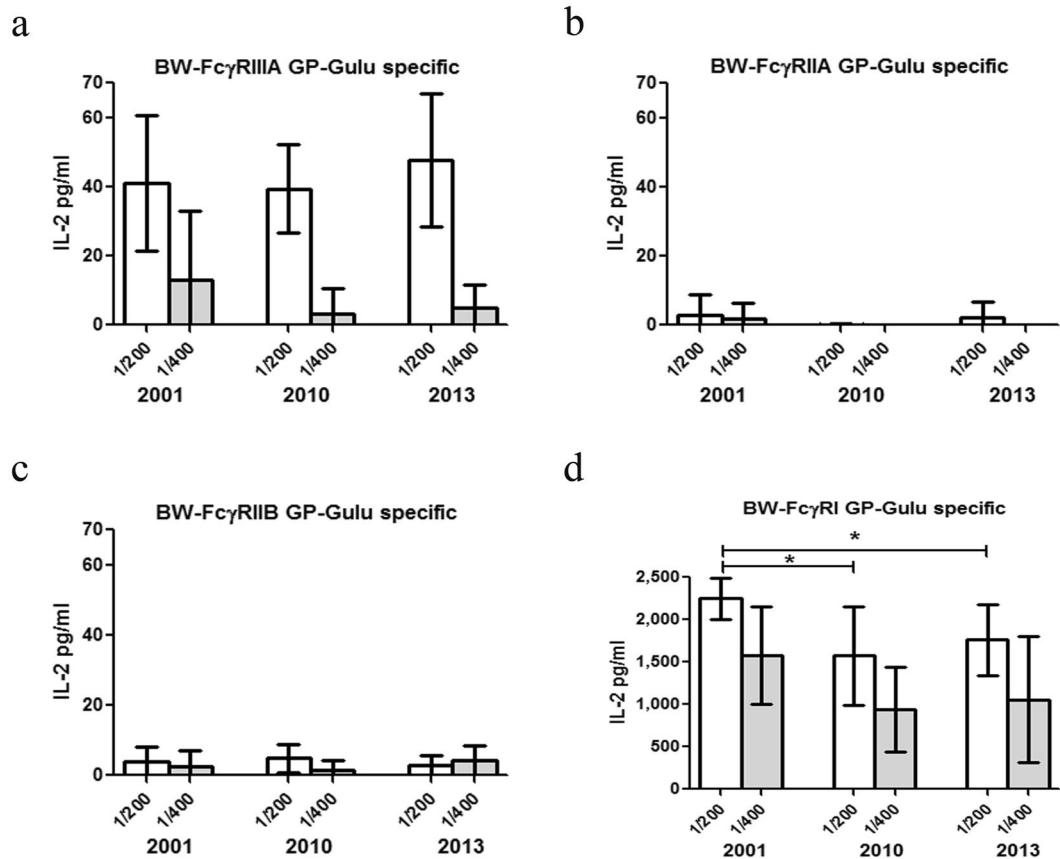
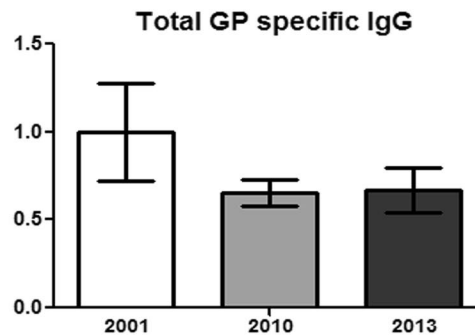


Figure 4. GP-specific reporters' responses to SUDV survivors' sera from longitudinal studies. 96 well culture plates were pre-coated with GP-Gulu antigen (5 μ g/ml) and the activation assay was performed using sera samples from SUDV survivors collected during the period of 2001–2013. The sera samples were diluted 1:200 and 1:400 for all BW-Fc γ R cells. Supernatant was collected after 16 h and analyzed for mIL-2 by ELISA. n(2001/2013) = 7, n(2010) = 6. Two-tailed Mann Whitney test, *P value < 0.05; P value (2001–2010) = 0.0221, P value (2001–2013) = 0.0111.

by ELISA the titer of GP-specific total IgG and IgG subclasses at these 3 time points following disease. The slight reduction between 2001 to 2010 & 2013 observed for the response of the Fc γ RI reporter (~25%) was also observed for total IgG binding to GP-coated wells (Fig. 5a). However, when we tested the GP-specific binding of different IgG subclasses, we observed different binding phenotypes. GP-specific IgG4 subclass binding was none to negligible in 2001 and remained none in 2010 and 2013 (Fig. 5b). Three of 7 survivors demonstrated low to moderate levels of GP-specific IgG3 subclass binding in their 2001 sera, which was reduced to low to background levels in sera from 2010 and 2013. Five of the 7 survivors' moderate to high levels of GP-specific IgG2 subclass binding in their 2001 sera was reduced (except one) to background levels in 2010 and 2013. In contrast, GP-specific response of the IgG1 subclass demonstrated the opposite observation. Three of the 7 survivors demonstrated substantial enhancement of their GP-specific IgG1 binding over the years. One exceptionally high responder (survivor 33) demonstrated high responses for all subclasses in its 2001 sera, and these responses were reduced but still the highest in 2013 for the IgG1 and the IgG2 subclasses (Fig. 5b).

Finally, we next determined whether the different parameters that we measured for the 7 survivors correlate. We employed the Spearman Correlation assay to analyze the associations of GP-specific responses of 3 reporters (Fc γ RIIIA, Fc γ RIIB & Fc γ RI) and the GP-specific levels of total IgG and 4 IgG subclasses (IgG1, 2, 3 & 4). We did not analyze associations for the Fc γ RIIIA reporter since most GP-specific results of this reporter had the zero value. Associations were studied for the 7 SUDV survivors, of which we had sera from 2001, 2010 and 2013. Of 15 cross-tabulations in the Spearman rank-order correlation coefficients table (Fig. 6a for sera from 2013), only two were significantly associated at the level of $p \leq 0.05$. These two significant associations were between (i) GP-specific Fc γ RI reporter and GP-specific total IgG (spearman r value = 0.8214, Fig. 6b), and (ii) GP-specific Fc γ RI reporter and GP-specific total IgG1 subclass (spearman r value = 0.8571, Fig. 6c). Similar significance and association results were obtained when we tested the sera from 2010 ($r = 0.9429$ and $r = 0.9429$, respectively, Fig. 6). Interestingly, no significant associations were observed for the 2001 serum samples that were taken within the first year post acute infection.

a



b

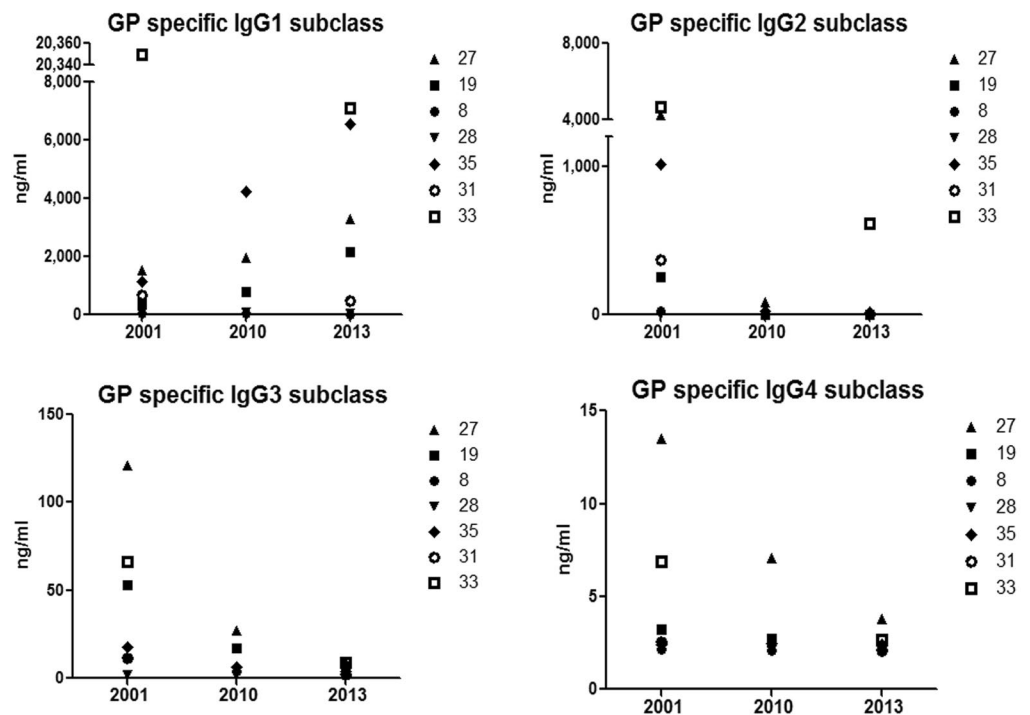


Figure 5. Analysis of total GP-specific IgG, and IgG subclasses in sera from SUDV survivors. To test total levels of GP-specific IgG (a) and IgG subclasses (b) in sera samples of SUDV survivors from the longitudinal study (2001-2010-2013), plates were pre-coated with GP-Gulu antigen (5 μ g/ml), washed, blocked and seeded with sera samples (triplicates) diluted 1:100 for subclass analysis and 1:400 for total IgG analysis. Total levels of GP-specific IgG from 2010 and 2013 normalized to the level of GP-specific IgG from 2001.

Discussion

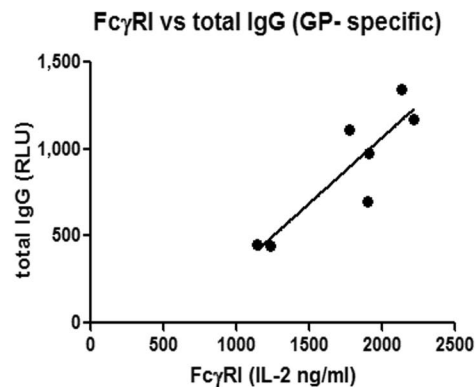
In this study, we completed a longitudinal analysis of sera from long-recovered survivors of SUDV infection for the presence of Ebola-GP specific IgG subclasses, and for binding to Fc γ receptors. Our analysis indicated that IgG1 is the dominant GP-specific IgG subclass that remains significantly noticeable more than a decade post infection. We further demonstrated that GP-specific Abs within sera manifest prominent binding to Fc γ RI, as compared to binding to Fc γ RIIIA, Fc γ RIIA and Fc γ RIIB. To investigate the binding of IgG (Ag-specific or non-specific) to Fc γ receptors, we developed a reporter system based on the expression of CD3zeta-fused Fc γ Rs in BW cells, which upon activation of the receptor secrete murine IL-2. The use of BW cells expressing zeta-fused Fc γ receptor for studying receptor functions was previously described by our laboratory and others (29), (31), (36). Recently, Corrales-Aguilar *et al.* published a similar BW-based system and Promega introduced a Jurkat-based Fc γ R reporter system^{25, 32, 33}. These published reporters include the Fc γ RIIIA, Fc γ RIIA and Fc γ RI and we developed similar reporters with the addition of the inhibitory Fc γ R, Fc γ RIIB. Likewise, we demonstrated that these reporters facilitate reproducible and less variable quantitative measurements of Fc γ R function.

Binding to Fc γ R, which elicits its function, represents IgG-effector functions. Two Ab-Fc modifications mediate its binding to Fc γ R and thus derived functionality: changes in antibody subclass and changes in a single N-linked glycan located in the CH2 domain of the IgG Fc³⁴. In this regard, we found that GP-specific IgG1 is by

a

	IgG GP specific	IgG1 subtype	IgG2 subtype	IgG3 subtype	IgG4 subtype
BW-FcγRIIIA	r=-0.03571 p=0.9635	r=-0.2500 p=0.5948	r=-0.07412 p=0.9063	r=0.2500 p=0.5948	r=-0.1071 p=0.8397
BW-FcγRIIB	r=-0.2143 p=0.6615	r=-0.3214 p=0.4976	r=-0.2224 p=0.6615	r=-0.2500 p=0.5948	r=-0.5357 p=0.2357
BW-FcγRI	r=0.8214 p=0.0341	r=0.8571 p=0.0238	r=0.6301 p=0.1389	r=0.4643 p=0.3024	r=0.6071 p=0.1667

b



c

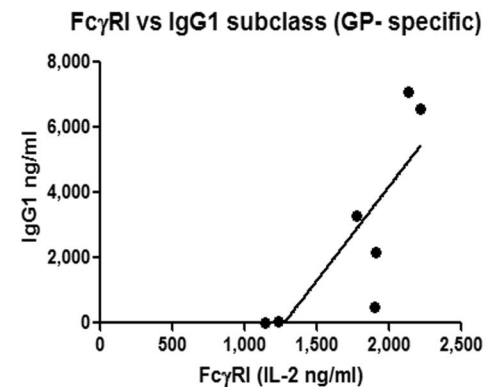


Figure 6. Correlation of reporters' response and IgG levels. (a) Spearman rank-order correlation coefficients table for associations between reporters' response and IgG subclasses is depicted in the 15 cells representing the tabulations. Each cell contains the corresponding spearman r-value and the probability of the association. The two grey-shaded cells have statistically significant associations, whereas the 13 clear cells have non-significant associations. (b,c) Spearman regression analysis for FcγRI and total IgG (b)/IgG1 subclass (c).

far the seroprevalent subclass that retained and even enhanced its presence in the sera, over ten years post infection; the prevalence of other GP-specific IgG subclasses was considerably reduced over time. The unanimity of IgG1 seroprevalence results within all tested survivors is in contrast to results reported for HCMV, in which both HCMV-specific IgG1 and IgG3 retained sera activity over the years and with significant variability among individuals³⁵. This variability was attributed to recurrence rate of HCMV, in which re-infections of HCMV resulted in elevation of HCMV-specific IgG3; on the other hand, persistence of IgG1 in sera was ascribed to no-recurrence of the HCMV. Therefore, GP-specific IgG1 seroprevalence in long-time survivors of SUDV could be a function of non-recurrence of the virus as reported for these survivors¹². The recent outbreak of the Makona-type EBOV, is followed by reports of recurrence of the virus in survivors and prolonged periods of virus secretion from immune-privileged sites³⁶. Hence, studying the GP-specific IgG1 and IgG3 in long-recovered Makona survivors could reveal a different phenotype as compared to the SUDV survivors.

For both HCMV and HIV-1, it was reported that pathogen-specific IgG3 Abs have greater neutralizing potential than IgG1^{37,38}. In accordance, IgG3 was secreted from a significant number of GP-specific hybridomas, generated from memory B cells of SUDV survivors secreting neutralizing mAbs (Leslie Lobel, unpublished results). We recently published that SUDV survivors, with no clinically different history, can be divided into groups with SUDV-specific neutralizing Abs, SUDV-specific non-neutralizing Abs, and with no SUDV-specific Abs¹⁰. Hence, the seroprevalence of GP-specific IgG1 that possibly lacks GP-mediated neutralizing activity is not associated with different clinical manifestations within the survivors.

In summary, we investigated GP-specific IgG responses in SUDV infected survivors, and demonstrated that over a decade post-acute disease, GP-specific IgG responses are significant primarily for the IgG1 subclass, and that FcγRI is the prominent FcγR associated with binding of the GP-specific IgGs in these survivors, from acute-infection to more than a decade post infection. Based on these results, Ebolavirus vaccine development should be informed by, and further investigated, with respect to the induced GP-specific IgG1 subclass and binding potency to FcγRI.

Materials and Methods

Cell lines. Mouse BW5147 thymoma cells (ATCC TIB-47™) and FcγR transfectants thereof were maintained in RPMI 1640 medium containing 10% (v/v) FCS, penicillin, streptomycin, glutamine, and sodium pyruvate (1 mM). BW-derived reporter cell lines were maintained in RPMI medium containing selection antibiotics.

Human sera samples. Serum samples from infected and non-infected patients were obtained from a banked collection, as well as from surviving patients (in 2000, 2003, and 2010), in collaboration with the Uganda Virus Research Institute (Entebbe, Uganda), and shipped to Ben-Gurion University, Israel. Sera from negative controls were collected from individuals who were from the districts where the outbreak occurred but were never diagnosed with ebolavirus infection and/or were negative by polymerase chain reaction (PCR) for SUDV¹². All serum samples used in our study were gamma irradiated and obtained under the appropriate informed consent.

Ethics statement. The study was approved by the Helsinki committees of the Uganda Virus Research Institute in Entebbe, Uganda (reference number GC/127/13/01/15), Soroka Hospital, Beer-Sheva, Israel (protocol number 0263-13-SOR) and the Ugandan National Council for Science and Technology (UNCST) (registration number HS1332). A written informed consent was translated for and signed by each subject; a personal health questionnaire was completed for each subject at the time of blood collection. We confirm that all experiments were performed in accordance with relevant guidelines and regulations.

Sample collection. Whole blood samples were obtained following antecubital venipuncture. Samples were directly aspirated into sterile glass vacuum tubes containing freeze-dried sodium heparin (final heparin concentration 14.3 units/ml, Becton Dickinson, Franklin Lakes, NJ, USA) and kept at 4 °C until further analysis.

Production and selection of functional BW-reporter cell lines. Cloned sequences encoding the human extracellular portion of Fc γ R_s fused to murine CD3 ζ chain were ordered from (Invitrogen, CA USA). Fc γ R_s included Fc γ R_{IIIA} (higher affinity variant with V158³⁹), Fc γ R_{IIA}, Fc γ R_{IIIB} and Fc γ R_I. Extracellular portions of Fc γ R_{IIA} cDNA (NM_001136219.1; (34Q-217G)), Fc γ R_{IIIB} cDNA (NM_004001.4; (43Q-223 G)) and Fc γ R_I cDNA (NM_000566.3; (16Q-92H)) were fused to mouse CD3 ζ chain cDNA (NM_001113391.2; (31L-164R)) making Fc γ R_{IIA}- ζ , Fc γ R_{IIIB}- ζ and Fc γ R_I- ζ respectively. The Fc γ R- ζ sequences were cloned into pHAGE2⁴⁰. HEK-293T cells were transiently transfected with plasmids of interest and lentiviral packaging plasmids, and retrovirus (RV)-containing supernatants were harvested, aliquoted and employed for transduction of BW5147 thymoma cells. BW-Fc γ R_{IIIA} transfectants have been previously described⁴¹. Following selection with Puromycin (Invivogen, CA USA) /G418 (Sigma-Aldrich, Israel), stable transfectants were screened by flow cytometry using clone 3G8 mouse anti-human CD16, clone FUN-2 mouse anti-human CD32 and clone 10.1 mouse anti-human CD64 (Biolegend, CA USA) using secondary goat anti-mouse IgG APC (Jackson immunoresearch, PA USA). BW5147 expressing receptor fused to murine CD3 ζ secrete IL-2 following activation of the receptor⁴¹. To further test function of transfectants, cells were incubated with purified human IgGs, followed by testing with a murine IL-2 commercial sandwich ELISA kit (Biolegend, CA USA).

Ebolavirus Sudan (Gulu) GP antigen and activation of the BW-Fc γ R reporter cells by IgG. Evaluation of IgG dependent activation of the BW-Fc γ R cells was prepared by incubating the cell lines with human sera and analyzing the supernatant from activated cells for mIL-2 by ELISA assay. 96 well cell culture plates were pre-coated with goat-Fab₂ anti human IgG Fab₂ 2.5 μ g/ml (Jackson immunoresearch, PA USA) or Ebola GP-Gulu antigen 2.5 μ g/ml⁴² diluted in 1xPBS and incubated for 2 h, blocked with 1% BSA (Sigma-Aldrich, USA) for 1 h. Human diluted samples sera (1:1 K, 1:5 K and 1:25 K; 1:200 and 1:400 dilutions respectively) were added and incubated for 1 h. BW-Fc γ R cells were supplemented, 50*10³ cells/well and incubated in RPMI 10% (v/v) FCS medium for 16–18 h. Each step was followed by a primary wash step with 1xPBS at 200 μ l/well. All incubations were at 37 °C in an atmosphere of 5% CO₂.

Ebolavirus Sudan (Gulu) GP antigen (Ebola GP-Gulu) consists of residues 1-649 of the glycoprotein, which comprises the complete ectodomain. It was expressed in HEK293T cells as a secreted protein with a HIS tag and purified over nickel chromatography by the protein core laboratory of the Nation Cancer Institute, NIH.

Evaluation of BW-Fc γ R activation by mIL-2 measurement. Supernatant of activated cells (as described above) was collected after 16–18 h incubation and analyzed for mIL-2 by ELISA assay. 96 well ELISA plates were pre-coated with purified anti-mouse IL-2 (Biolegend, CA USA) using coating buffer (0.1 M, Na₂HPO₄; pH9.0), blocked with 10% FBS in PBST (0.05% Tween-20), coated with collected supernatant followed by addition of biotinylated anti-mouse IL-2 (Biolegend, CA USA), and then SA-HRP (Jackson immunoresearch, PA USA) and TMB (Dako, Denmark) for detection of mIL-2.

Total antigen specific IgG and antigen specific IgG subclass. To estimate Ebola antigen specific IgG total and subclass dependent activation profile: 96 well U shape plates were pre-coated with Ebola GP-Gulu antigen, blocked with 10% skim milk and human sera added (1:400 and 1:100 sera dilution respectively) followed by anti-human IgG 1–4 (Southern Biotech, AL USA) for IgG subclass, and then anti-human HRP (Jackson immunoresearch, PA USA) and peroxidase solution with luminol enhancer solution, respectively (SuperSignal West Pico, USA). Relative light units (RLU) acquired by Infinite F200 pro Elisa reader (Tecan, Austria).

References

1. Jens Kuhn, H. & Calister, C. H. *FILOVIRUSES: A Compendium of 40 Years of Epidemiological, Clinical, and Laboratory Studies*. *Archives of virology* Supplement (2008).
2. Dixon, M. G. & Schafer, I. J. Ebola viral disease outbreak—West Africa, 2014. *MMWR. Morbidity and mortality weekly report* **63**, 548–51 (2014).
3. Kuhn, J. H. *et al.* Evaluation of Perceived Threat Differences Posed by Filovirus Variants. *Biosecurity Bioterrorism Biodefense Strateg. Pract. Sci.* **9**, 361–371 (2011).
4. Mohamadzadeh, M., Chen, L. & Schmaljohn, A. L. How Ebola and Marburg viruses battle the immune system. *Nat. Rev. Immunol.* **7**, 556–567 (2007).

5. Wong, G., Kobinger, G. P. & Qiu, X. Characterization of host immune responses in Ebola virus infections. *Expert Rev. Clin. Immunol.* **10**, 1–10 (2014).
6. Mohamadzadeh, M., Chen, L., Olinger, G. G., Pratt, W. D. & Schmaljohn, A. L. Filoviruses and the balance of innate, adaptive, and inflammatory responses. *Viral Immunol.* **19**, 602–612 (2006).
7. McElroy, A. K. *et al.* Human Ebola virus infection results in substantial immune activation. *Proc. Natl. Acad. Sci. USA* **112**, 4719–24 (2015).
8. Bellan, S. E., Pulliam, J. R. C., Dushoff, J. & Meyers, L. A. Ebola control: Effect of asymptomatic infection and acquired immunity. *The Lancet* **384**, 1499–1500 (2014).
9. Leroy, E. *et al.* Human asymptomatic Ebola infection and strong inflammatory response. *Lancet* **355**, 2210–2215 (2000).
10. Sobarzo, A. *et al.* Persistent Immune Responses after Ebola Virus Infection. *N. Engl. J. Med.* **369**, 492–493 (2013).
11. Sobarzo, A. *et al.* Immune Memory to Sudan Virus: Comparison between Two Separate Disease Outbreaks. *Viruses* **7**, 37–51 (2015).
12. Sobarzo, A. *et al.* Profile and persistence of the virus-specific neutralizing humoral immune response in human survivors of Sudan ebolavirus (Gulu). *J. Infect. Dis.* **208**, 299–309 (2013).
13. Berken, A. & Benacerraf, B. Properties of antibodies cytophilic for macrophages. *J. Exp. Med.* **123**, 119–44 (1966).
14. Caaveiro, J. M. M., Kiyoshi, M. & Tsumoto, K. Structural analysis of Fc/Fc γ R complexes: A blueprint for antibody design. *Immunological Reviews* **268**, 201–221 (2015).
15. Shibata-Koyama, M. *et al.* The N-linked oligosaccharide at Fc γ RIIIa Asn-45: An inhibitory element for high Fc γ RIIIa binding affinity to IgG glycoforms lacking core fucosylation. *Glycobiology* **19**, 126–134 (2009).
16. Lu, J. & Sun, P. D. Structural mechanism of high affinity Fc γ RI recognition of immunoglobulin G. *Immunological Reviews* **268**, 192–200 (2015).
17. Hayes, J. M. *et al.* Fc gamma receptor glycosylation modulates the binding of IgG glycoforms: A requirement for stable antibody interactions. *J. Proteome Res.* **13**, 5471–5485 (2014).
18. Anthony, R. M., Wermeling, F. & Ravetch, J. V. Novel roles for the IgG Fc glycan. *Ann. N. Y. Acad. Sci.* **1253**, 170–180 (2012).
19. Smith, K. G. C. & Clatworthy, M. R. Fc γ RIIB in autoimmunity and infection: evolutionary and therapeutic implications. *Nat. Rev. Immunol.* **10**, 328–343 (2010).
20. Hogarth, P. M. & Pietersz, G. A. Fc receptor-targeted therapies for the treatment of inflammation, cancer and beyond. *Nat. Rev. Drug Discov.* **11**, 311–331 (2012).
21. Shields, R. L. *et al.* High Resolution Mapping of the Binding Site on Human IgG1 for Fc γ RI, Fc γ RII, Fc γ RIII, and FcRn and Design of IgG1 Variants with Improved Binding to the Fc γ R. *J. Biol. Chem.* **276**, 6591–6604 (2001).
22. Subedi, G. & Barb, A. The immunoglobulin G1 N-glycan composition affects binding to each low affinity Fc γ receptor. *MAbs* **0**, (2016).
23. Parcina, M. *et al.* Staphylococcus aureus-Induced Plasmacytoid Dendritic Cell Activation Is Based on an IgG-Mediated Memory Response. *J. Immunol.* **181**, 3823–3833 (2008).
24. DiLillo, D. J., Palese, P., Wilson, P. C. & Ravetch, J. V. Broadly neutralizing anti-influenza antibodies require Fc receptor engagement for *in vivo* protection. *Journal of Clinical Investigation* **126**, 605–610 (2016).
25. Corrales-Aguilar, E. *et al.* Highly individual patterns of virus-immune IgG effector responses in humans. *Med. Microbiol. Immunol.*, doi:10.1007/s00430-016-0457-y (2016).
26. Mahan, A. E. *et al.* Antigen-Specific Antibody Glycosylation Is Regulated via Vaccination. *PLoS Pathog.* **12** (2016).
27. Wegener, A. M. K. *et al.* The T cell receptor/CD3 complex is composed of at least two autonomous transduction modules. *Cell* **68**, 83–95 (1992).
28. Yagüe, J. *et al.* The T cell receptor: the α and β chains define idotype, and antigen and MHC specificity. *Cell* **42**, 81–87 (1985).
29. Letourneur, F. & Malissen, B. Derivation of a T cell hybridoma variant deprived of functional T cell receptor β and γ chain transcripts reveals a nonfunctional β -mRNA of BW5147 origin. *Eur. J. Immunol.* **19**, 2269–2274 (1989).
30. Daeon, M., Yodoi, J., Neaupoort-Sautes, C., Moncuit, J. & Fridman, W. H. Receptors for immunoglobulin isotypes (FcR) on murine T cells. I. Multiple FcR expression on T lymphocytes and hybridoma T cell clones. *Eur. J. Immunol.* **15**, 662–667 (1985).
31. Vidarsson, G., Dekkers, G. & Rispens, T. IgG subclasses and allotypes: From structure to effector functions. *Front. Immunol.* **5** (2014).
32. Tada, M., Ishii-Watabe, A., Suzuki, T. & Kawasaki, N. Development of a cell-based assay measuring the activation of Fc γ RIIa for the characterization of therapeutic monoclonal antibodies. *PLoS One* **9** (2014).
33. Cheng, Z. J. Development of a bioluminescent cell-based bioassay to measure Fc receptor functionality in antibody-dependent cellular cytotoxicity. In *American Association of Cancer Research (AACR) Annual Meeting* (2012).
34. Ackerman, M. E. *et al.* Natural variation in Fc glycosylation of HIV-specific antibodies impacts antiviral activity. *J. Clin. Invest.* **123**, 2183–2192 (2013).
35. Simon, B., Kundi, M. & Puchhammer-Stöckl, E. Association of HCMV specific IgG subclass antibody levels with gender and age. *Exp. Gerontol.* **48**, 472–475 (2013).
36. Nelson, E. A. *et al.* Clomiphene and its isomers block Ebola virus particle entry and infection with similar potency: Potential therapeutic implications. *Viruses* **8** (2016).
37. Scharf, O. *et al.* Immunoglobulin G3 from polyclonal human immunodeficiency virus (HIV) immune globulin is more potent than other subclasses in neutralizing HIV type 1. *J. Virol.* **75**, 6558–65 (2001).
38. Gupta, C. K., Leszczynski, J., Gupta, R. K. & Siber, G. R. IgG subclass antibodies to human cytomegalovirus (CMV) in normal human plasma samples and immune globulins and their neutralizing activities. *Biologicals* **24**, 117–24 (1996).
39. Bowles, J. A. & Weiner, G. J. CD16 polymorphisms and NK activation induced by monoclonal antibody-coated target cells. *J. Immunol. Methods* **304**, 88–99 (2005).
40. Mostoslavsky, G. *et al.* Efficiency of transduction of highly purified murine hematopoietic stem cells by lentiviral and oncoretroviral vectors under conditions of minimal *in vitro* manipulation. *Mol. Ther.* **11**, 932–940 (2005).
41. Mandelboim, O. *et al.* Recognition of haemagglutinins on virus-infected cells by NKp46 activates lysis by human NK cells. *Nature* **409**, 1055–60 (2001).
42. Sobarzo, A. *et al.* Profiling the native specific human humoral immune response to Sudan Ebola virus strain Gulu by chemiluminescence enzyme-linked immunosorbent assay. *Clin. Vaccine Immunol.* **19**, 1844–1852 (2012).

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Author Contributions

O.R. and A.P. designed the experiments. L.L., A.E. and S.F.M. helped with designing the IgG subclass experiments. O.R. and A.E. performed the experiments. O.R. and M.B. constructed the Fc reporter system. O.G.Y. helped with

performing experiments. A.S., S.F.M., A.E., J.L. and J.D. collected and sorted the samples. O.R. and A.P. wrote the primary manuscript and L.L. and A.P. further edited primary manuscript.

Additional Information

Competing Interests: The authors declare that they have no competing interests.

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