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Viral Control in Chronic HIV-1 Subtype C Infection is associated with Enrichment of p24 IgG1 with Fc Effector Activity

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Abstract

Objective—Post-infection HIV viral control and immune correlates analysis of the RV144 vaccine trial indicate a potentially critical role for Fc receptor-mediated antibody functions. However, the influence of functional antibodies in clade C infection is largely unknown.

Design—Plasma samples from 361 chronic subtype C-infected, antiretroviral therapy-naïve participants were tested for their HIV-specific isotype and subclass distributions, along with their Fc receptor-mediated functional potential.

Method—Total IgG, IgG subclasses and IgA binding to p24 clade B/C and gp120 consensus C proteins were assayed by multiplex. Antibody-dependent uptake of antigen-coated beads and Fc receptor-mediated NK cell degranulation were evaluated as surrogates for antibody-dependent cellular phagocytosis (ADCP) and antibody-dependent cellular cytotoxicity (ADCC) respectively.

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Author contributions:

AWC, JMM, PG, BW, TN, GA conceived the study, AWC, JMM, BN, AL, HR, YR conducted experiments. AWC, JMM, MG, TR conducted the analysis. AWC, JMM, TN, GA wrote the manuscript.

Conflict of interest

The authors declare no conflict of interest.

Results—p24 IgG1 was the only subclass associated with viral control ($p=0.01$), with higher p24-specific ADCP and ADCC responses detected in individuals with high p24 IgG1. Although p24 IgG1 levels were enriched in subjects with elevated Gag-specific T cell responses, these levels remained an independent predictor of low viral loads ($p=0.04$) and high CD4 counts ($p=0.004$) after adjusting for Gag-specific T cell responses and for protective HLA class I alleles.

Conclusion—p24 IgG1 levels independently predict viral control in HIV-1 clade C infection. Whether these responses contribute to direct antiviral control via the recruited killing of infected cells via the innate immune system or simply mark a qualitatively superior immune response to HIV, is uncertain, but highlights the role of p24-specific antibodies in control of clade C HIV-1 infection.

Keywords

HIV-1 subtype C; chronic; ADCP; NK cell degranulation; antibody; p24; Gag

Background

In 2016 alone, there were an estimated 1.8 million new HIV-1 infections globally; a figure that highlights the relentless spread of this virus and the urgent need for an efficacious vaccine^[1]. In 2009, the RV144 trial, a phase III HIV-1 human vaccine trial, was the first to show modest efficacy^[2]. Immune correlates analysis identified binding antibodies to the viral envelope variable loops 1 and 2 as the primary correlates of reduced risk from infection^[3]. However, in secondary analyses, Fc-mediated antibody-dependent cellular cytotoxicity (ADCC) preferentially mediated by IgG3 was also associated with reduced risk of infection in the absence of IgA^[4, 5], suggesting a potentially critical role for non-neutralizing antibody Fc effector activities in protection from HIV-1 infection.

In addition to ADCC, antibodies mediate additional extra-neutralizing functions relevant in HIV-1 infection, such as antibody-mediated cellular phagocytosis (ADCP), through their capacity to interact with Fc receptors expressed on innate immune cells^[6, 7]. Importantly, both the magnitude and breadth of the ADCC response has been linked to slow disease progression^[8–12], and higher ADCC activity has been observed in rare individuals capable of controlling viral replication in the absence of antiretroviral therapy (ART)^[13]. Furthermore, HIV-1 transmission from mothers to their breast-feeding infants is significantly reduced in women with high ADCC activity in breast milk^[14]. In animal models, vaccine-induced ADCC and antibody-dependent cell-mediated viral inhibition responses have been associated with protection and reduced viral set point^[15–20]. Moreover, prevention of simian-human immunodeficiency virus infection via vaccination has also been linked to enhanced ADCP activity^[19, 20], which was also enriched in RV144 vaccinees^[5, 21]. Together, these observations have renewed interest in dissecting the importance of Fc effector functions in natural infection, particularly against clades that contribute to the greatest HIV-1 burden globally.

The antibody response to HIV-1 proteins is largely dominated by IgG1 antibodies, with variable levels of other antibody subclasses (IgG2, IgG3 and IgG4)^[22]. In natural infection, however, levels of gp120-specific antibodies do not correlate with markers of disease

progression despite their potential antiviral activity^[23–31]. In contrast, anti-p24 antibody levels have been shown to correlate with viral control in several studies, largely focused on subtype B cohorts (summarized by French et al.^[32]). However, because p24 is not expressed on the cell surface, the direct contribution of p24-specific antibodies to antiviral control is unclear, but has also been proposed as a surrogate marker of an intact CD4 T cell response^[33].

Because the humoral immune response may differ between clade B and C infected individuals^[34], here we aimed to define the Fc receptor-mediated functional profiles of gp120- and p24-specific antibody responses in the context of HIV-1 subtype C infection. Similar to previous studies^[32, 35], we found that levels of p24-specific but not gp120-specific IgG1 were associated with lower viral loads and higher CD4 counts. Furthermore, individuals with high p24 IgG1 levels had higher Fc-mediated responses that showed a trend towards viral control and higher CD4 counts. The association between levels of p24 IgG1 and viral loads and CD4 counts was maintained even after adjusting for the impact of Gag-specific CD4 and CD8 T cell responses and protective HLA class I alleles (HLA-I). These data indicate that p24-specific IgG1 may independently contribute to viral control and further studies are warranted to ascertain their direct or indirect roles in antiviral immunity.

Materials and Methods

Study Subjects

361 archived plasma samples from the Sinikithemba cohort^[36], a study of HIV-1 subtype C–infected, ART-naïve individuals, established in Durban, South Africa were included in this study. All study participants provided written informed consent, and the study protocol was approved by the Biomedical Research Ethics Committee of the University of KwaZulu-Natal.

Customized IgG Subclass Binding Assay

To determine the relative levels of each HIV-1 specific isotype/subclass in plasma, we adopted a customized subclass binding assay described previously^[37]. We used carboxylated microspheres (luminex) coupled to HIV-1 recombinant proteins of interest: p24 subtype B/C and gp120 consensus C (Immune Technology Corp).

Antibody-Dependent Cellular Phagocytosis (ADCP)

The ability of plasma antibodies to mediate ADCP was determined as previously described using gp120 consensus C and p24 subtype B/C protein (Immune Technology Corp)^[38]. Antigen-bound fluorescent neutravidin beads (Invitrogen) were incubated with 1:100 dilution of plasma antibodies in the presence of THP-1 cells. HIV negative plasma and media alone were used as negative controls and pooled HIV positive plasma was used as a positive control. The cells were analyzed for bead uptake by flow cytometry and the phagocytic score representing iMFI values (integrated Mean Fluorescence Intensity=frequency x MFI)^[39] was calculated. Flow cytometry gating strategy is shown in Supp Figure 1A.

While previous studies^[38] have confirmed that gp120-specific ADCP activity is mediated by Fc γ R mechanisms, this has not been confirmed when using p24 protein as an antigen. Therefore as an additional control, 6 randomly selected HIV positive plasma samples were purified for IgG using Melon Gel (Thermo Fisher Scientific). THP-1 cells were pre-incubated with Fc γ RII (Abcam) or Fc γ RIII (Sigma) blocking antibodies for 30 min prior to assaying for p24-specific ADCP responses as described above (Supp Figure 1B).

Natural Killer (NK) Cell degranulation as a surrogate for ADCC

We used a modified ELISA-based protocol for detection of CD107a as a surrogate marker of NK cell-mediated cytotoxicity^[40]. This assay has previously been shown to be associated with Fc γ RIIIa binding capacity of antigen-specific antibodies^[41, 42]. A 96-well plate was coated overnight at 4°C with 150ng of recombinant protein per well. 2% BSA blocked plates were used as antigen controls. The next day the plates were washed 6 times with PBS, 50 μ l of plasma (diluted 1:100) was added to each well, and incubated at 37°C for 2 hours. HIV negative plasma samples or media alone were used as negative controls, while HIVIG (pooled HIV Immunoglobulin G, NIH AIDS Reagents Program) was used as a positive control. The plates were washed and 5 \times 10⁴ NK cells enriched via negative selection from healthy blood donors (RosetteSep, Stemcell Technologies,) were added to each well in the presence of Brefeldin A (Biolegend), Golgi stop, and anti-CD107a-PE-Cy5 (BD Biosciences). The plate was incubated for 5 hours at 37°C and 5% CO₂. Following incubation, cells were stained with anti-CD3-AF700, anti-CD56-PE-Cy7, anti-CD16-APC (BD), fixed with Perm A, permeabilized using Perm B (Invitrogen), and stained with anti-IFN γ -APC and anti-MIP1 β -PE (BD). The cells were then fixed with 2% paraformaldehyde and analyzed by flow cytometry (gating strategy shown in Supp Figure 2).

Gag-Specific CD4+ and CD8+ T cell responses

Gag-specific IFN- γ CD4+ T cell responses were detected by intracellular cytokine staining (ICS) using freshly isolated PBMCs as previously described^[43] staining for anti-human CD4 APC and anti-human IFN γ FITC. Gag-specific IFN γ CD8+ T cell were detected by IFN γ enzyme-linked immunosorbent spot (ELISpot) using overlapping peptides spanning Gag subtype C as previously described^[44].

Statistical Analysis

Non-parametric Spearman's rank tests were used to test for correlations, and a 2-tailed Mann Whitney test was used to evaluate unmatched groups. p values less than 0.05 were considered significant. To assess the effect of p24 IgG1 on viral load and CD4 count, multiple linear regression was used. IgG1p24 values were z-score normalized (standardized) to have a mean of zero and a standard deviation of one. We considered a variable to be a potential confounder of the effect of p24 IgG1 if it was shown to have a significant relationship with the outcome in univariate analysis or if its inclusion in the model resulted in a 10% or greater change in the estimated coefficients. A square root transformation was applied to CD4 count and a log transformation was applied to viral loads, which resulted in approximately normally distributed values. Analysis was performed using Graphpad Prism version 6 (Graphpad Software) and Stata Statistical Software: Release 13 (StataCorp LP).

Results

p24 IgG1 levels predict viral control in chronic HIV-1 subtype C infection

Previous studies, largely in clade B infected cohorts, have shown that p24- but not gp120-specific antibody levels are a prognostic marker of disease progression^[25, 28–32]. Thus, to first define whether similar relationships were observable in clade C infection, a cross-sectional study including a cohort of 361 HIV-infected subjects (median viral load: 4.8 log₁₀ HIV-RNA copies/ml, interquartile range (IQR) 3.9–5.3; median CD4 count: 339 cells/mm³, IQR 226–492) were profiled for gp120- or p24-specific antibodies. HIV-specific IgG to both proteins were detected in all individuals, however, neither gp120- nor p24-specific antibody levels correlated with either viral loads or CD4 counts (Fig 1A and C-left).

Next we determined whether relative differences in gp120- or p24-specific Ig subclass specific responses correlated with measures of viral control. Similar to reports from other subtypes^[22], the antibody responses directed at both gp120 and p24 were dominated by IgG1 responses. IgG3 directed to p24 was also observed in all individuals, and gp120-IgG3 responses were detected in most individuals (99%). Conversely, levels of other subclasses were lower and less frequently detected: p24- and gp120-specific IgA were detected in 94% and 99% of the cohort respectively. Conversely, levels of other subclasses were lower and less frequently detected: gp120- and p24-specific IgA were detected in 94% and 99% of the cohort respectively, while gp120- and p24-specific IgG2 were detected in 80% and 94% of subjects respectively ($p=0.0032$), and gp120- and p24-specific IgG4 were detected in 99% and 90% respectively ($p=0.0052$) (Fig. 1).

To further probe the relationship between antigen-subclass specific responses and viral control, the relationships between individual antigen-specific subclasses and markers of disease progression were assessed. Higher gp120-specific IgG2 and IgG4 responses were associated with higher viral loads ($p=0.0002$, $r=0.206$ and $p=0.035$, $r=0.115$ respectively) (Fig. 1A) and were negatively associated with CD4 counts ($p=0.014$, $r=-0.133$ and $p=0.034$, $r=-0.115$ respectively) (Fig. 1B). Similarly, increasing levels of IgG3 responses targeting p24 were also associated with higher viral loads ($p=0.0002$, $r=0.205$) (Fig. 1C) and lower CD4 counts ($p=0.006$, $r=-0.151$) (Fig. 1D). In contrast, Gag p24 IgG1 levels were negatively correlated with viral loads ($p=0.01$, $r=-0.139$) (Fig. 1C) and positively correlated with CD4 counts ($p=0.025$, $r=0.121$) (Fig. 1D). These data suggest that higher levels of p24-IgG3 subclass selection profiles in chronic subtype C infection are associated with poor viral control, while p24-specific IgG1 responses in chronic subtype C infection may be a marker of viral control.

Elevated p24 IgG1 levels track with enhanced Fc effector activity

While antibodies directed against internal structural proteins of the virion have a limited capacity to mediate neutralization, these antibodies may contribute to viral control via non-neutralizing antiviral functions^[45, 46]. Thus, we next profiled the ADCP and ADCC activity of these antibodies and found that p24-specific phagocytic activity was detectable in all individuals while only low-level p24-specific ADCC activity was observed, in a minority of subjects. Detectable p24-specific NK cell degranulation (CD107a) was observed in 34% of

the tested subjects, and IFN γ and MIP1 β secretion was detected in 47% and 16% of the subjects, respectively, with strong correlations observed between all three functions ($p=0.004$ to $p<0.0001$) (Fig 2A). We then determined the relationship between antibody levels and functional activity and found that p24 IgG1 was positively correlated with both ADCP phagosome and CD107a expression ($p<0.0001$ and $p=0.033$ respectively, Fig 2B). Interestingly, there was a clear distribution of high and low IgG1 levels when considering ADCP phagosome. Using the median value as a cutoff for p24 IgG1 levels, we found that individuals with high p24 IgG1 levels were able to mediate significantly higher ADCP and ADCC (CD107a) responses ($p<0.0001$ Fig. 2C-left and $p=0.027$ respectively; Fig. 2C-right). These data suggest that p24 antibodies may be effective at mediating Fc effector functions.

Higher p24-specific Fc effector activity trends with markers of disease progression

Given that higher p24 IgG1 levels were associated with higher Fc effector activity, we next investigated the relationship between p24-specific Fc effector activity and markers of disease progression. While we did not observe a significant relationship between the expression of CD107a and viral load ($p=0.562$, $r=-0.055$) (Fig. 3B), a trend towards higher CD4 counts ($p=0.065$, $r=0.172$) was observed (Fig. 3D). In addition, the magnitude of p24-specific ADCP responses trended towards a negative correlation with viral load ($p=0.086$, $r=-0.157$; Fig. 3A) and a positive correlation with CD4 count ($p=0.082$, $r=0.158$; Fig. 3C), suggesting that p24-specific antibody functions are associated, albeit weakly, to viral control.

p24 IgG1 levels are not a simple surrogate of protective HLA class I antiviral activity

To next determine whether p24-specific antibodies may contribute to antiviral control, we examined the relationship between p24-specific IgG1 responses and traditional markers of HIV-1 control. Specifically, particular HLA class I alleles have been linked to slower disease progression, through the induction of superior CD8+ T cell activity in this cohort^[36, 44, 47]. Of the 361 individuals, those with protective HLA class I alleles (HLA-B*57:01, HLA-B*58:01, HLA-B*81:00 and HLA-B*81:01) had significantly lower viral loads ($p=0.002$) and higher CD4 counts ($p=0.003$). Thus to define whether p24 IgG1 was independently associated with antiviral control individuals with protective HLA-I alleles were removed from the analysis ($n=24$). Interestingly, individuals with high p24 IgG1 levels maintained higher ADCP ($p<0.0001$) and ADCC ($p=0.035$), (Fig. 4A, B) and continued to exhibit lower viral loads ($p=0.004$) and higher CD4 counts ($p=0.019$) (Fig. 4C, D), indicating that the relationship between p24 specific IgG1 responses and enhanced viral control was not solely influenced by protective HLA-I alleles and therefore may represent an independent predictor of enhanced viral control.

p24 IgG1 levels independently mark protective HIV-1 control

Both CD4+ and CD8+ T cell responses have also been linked to protective immunity in this cohort^[36, 43, 48]. Moreover, because enhanced helper CD4 T cell responses may promote more effective antibody responses in addition to viral control it is possible that elevated p24 antibodies may simply represent a marker of preserved T cell immunity rather than a direct antiviral mechanism of control. Thus, to parse out the relationship between T cell mediated antiviral control and p24-specific antibody responses, we next examined the inter-relationship between p24 IgG1 responses and both arms of the T cell response in these

individuals. The frequency of Gag-specific CD4 T cells (by ICS) and magnitude of CD8 T cells (by ELISpot) was associated with lower viral loads, ($p < 0.0001$ for both) and higher CD4 counts ($p = 0.0002$ and 0.138) respectively in these individuals, as previously observed^[36]. Moreover, when individuals were split into those with high or low p24 IgG1 levels, subjects with high p24 IgG1 levels had significantly higher Gag-specific CD4 ($p = 0.037$) and CD8 ($p = 0.049$) responses (Fig. 4E, F) suggesting that p24-specific humoral immune responses are induced in parallel to protective Gag T cell immunity.

However, to gain insights into whether p24-specific IgG1 immunity was simply a surrogate of a more effective T cell immune response or if it represented an independent marker of enhanced viral control, factors that were associated with viral control and elevated CD4 counts in univariate analyses including IgG1p24 levels, CD4/CD8-T cell Gag responses (dichotomized as having any or no response) and the presence of protective HLA-I alleles, were considered in a multivariate analysis. The CD4 T cell Gag response remained a significant predictor of viral load and CD4+ T cell counts ($p = 0.002$ and $p < 0.001$, respectively) (Table 1). Yet, most interestingly, even after adjusting for CD4 or CD8 Gag responsiveness and carriage of protective HLA-I alleles, IgG1 p24 antibody responses remained significantly associated with lower viral loads ($p = 0.030$) and higher CD4 counts ($p = 0.005$, Table 1). Hence, p24 IgG1 antibodies independently predict viral control and preservation of CD4 cells and likely complement T cell-mediated antiviral immunity in the control of viral replication.

Discussion

Despite the growing appreciation for the role of non-neutralizing antibody functions in both antiviral control^[9–13] and protection from HIV-1 infection^[4, 5, 14], little is known about the role of these functions in the context of HIV-1 subtype C viral control, which predominates the current global epidemic. Thus, here, we aimed to define the role of Fc-mediated antibody activities in chronic subtype C infection in a cross-sectional cohort of HIV-1-infected subjects at the heart of the South African epidemic, in Durban. Similar to previous studies of other viral subtypes^[23–31], the levels of gp120-specific total IgG and other IgG subclasses did not correlate with viral control. Instead, as previously reported^[32] p24-specific IgG1 responses were associated with antiviral control and slower progression to disease. Moreover, these antibodies have the capacity to mediate both cytotoxicity and clear infected cells via phagocytosis, providing two potential antiviral mechanisms by which these antibodies may contribute to antiviral control. Finally, while elevated CD4 and CD8 T cell responses were induced among individuals with the highest p24-specific IgG1 responses, multivariate modeling demonstrated that the p24-specific IgG1 levels independently predicted HIV-1 control, suggesting for the first time that p24-specific functional IgG1 antibodies may contribute directly to antiviral immunity in a manner complementary to known mechanisms of cellular immune control of HIV-1.

The selection of the poorly functional gp120 IgG2 and IgG4 was associated with poor viral control, likely by dampening antibody functionality due to their lower affinity for Fc-receptors, as has been previously reported^[5]. However, interestingly, while IgG3, the most functional antibody subclass, has been linked to reduced risk of HIV-1 infection in RV144

vaccinees^[4, 5], here, elevated gp120- or p24-specific IgG3 levels were not observed in subjects with markers of slower disease progression. Rather, elevated p24-specific IgG3 responses were associated with higher viral loads and lower CD4 T cell counts. As IgG3 is the first subclass in the immunoglobulin heavy locus, the selection of IgG3 antibodies is a signature of recently selected B cells, that have not rearranged to IgG1, IgG2, or IgG4^[49]. Thus, these data suggest that p24 IgG3 antibody responses in chronic infection may represent a marker of a less mature B cell response, which lack provision of B cell cross switch recombination to other subclasses, while, in contrast, enhanced p24 IgG1 responses may contribute most effectively to post-infection antiviral immunity in the setting of Clade C HIV-1 infection.

While the association between p24-specific antibody levels and markers of disease control has been reported in several studies^[32], the mechanism by which p24-specific IgG1 antibodies contribute to antiviral immunity remains unclear. One possibility includes a scenario where higher virus loads drive the formation of p24-containing immune complexes; although, this hypothesis has been refuted by a study done by Fenouillet et al. who showed that acid dissociation of immune complexes did not increase p24 levels^[50]. Alternatively, p24-specific IgG1 antibodies may only be a marker of an intact immune response, where preserved p24-specific CD4+ helper T cell responses sustain elevated levels of p24-specific B cell responses p24-specific IgG1 but do not individually contribute to viral control^[27, 51]. Our data, however, challenges this last hypothesis. Although we show that although p24-specific IgG1 levels are associated with higher CD4 T cell numbers (Fig. 1D) as well as elevated CD4+ T cell help (Fig. 4E), the ability to predict lower viral loads is sustained after controlling for CD4 T cell numbers and Gag-specific CD4 responses (Table 1). Thus, our data indicates that p24-specific IgG1 responses, beyond being only a marker of a more effective overall immune response, may also contribute independently to viral control.

Whether Gag is exposed on the virion surface remains a subject of debate. Although rare studies^[52-54] have detected cell surface p24 on infected cells, these studies have only been conducted on cell lines, and surface expression was only observed at high, non-physiological rates of infection. However, it is possible that Gag may be exposed during delivery of HIV-1 Gag to the plasma membrane for virion assembly^[55], particularly in the setting of early apoptosis when the inner-cell lipid layer flips out to expose phosphatidyl-serine, potentially exposing cytosolic anchored membrane proteins as well^[56]. It is also plausible that in addition to native protein expression on the cell surface, processed Gag peptides may be recognized by antibodies in the groove of MHC molecules on the surface of infected cells to induce ADCP or ADCC, though this has yet to be demonstrated. Importantly, Fc effector responses to other non-envelope protein derived-peptides have been reported^[12, 45, 46, 57-59]. Indeed Tjiam et. al. recently showed p24 IgG1 and IgG2 enhanced dendritic cell-mediated opsonophagocytic responses in individuals who controlled viremia^[46] and these same opsonophagocytic responses were associated with early viral control, interestingly specifically mediated by p24 IgG1^[45]. In our study, only p24-specific IgG1 levels were associated with higher ADCP and ADCC responses that trended towards viral control. Unlike Tjiam's study, p24-specific IgG2 levels did not correlated with Fc function, potentially due to the different assays used, which measure different Fc functions and utilize different effector cells. Although in our study, these p24-specific Fc effector functions did

not achieve significance with markers of disease progression (Fig 4, p -value range=0.062–0.086), antibodies are capable of mediating multiple other Fc-driven effector functions beyond ADCP and ADCC, including complement fixation, neutrophil activation, DC-activation as well as recruiting multiple other Fc γ R-expressing effector cells. Furthermore, all our assays were tested utilizing total plasma, which includes multiple different immunoglobulin Isotypes including IgA, IgM, IgE along with other serum proteins including complement which were not assessed. Future studies testing for alternative Fc effector functions using purified IgG and isolated p24-specific IgG1 could provide greater insights.

Additionally, antibodies have recently been shown to not only drive extra-cellular antiviral activity, but also to target pathogens within cells. In fact, the intracellular host restriction factor TRIM21 is an intracellular cytoplasmic Fc receptor that can drive rapid and profound autophagy. In this context, interactions between TRIM21 and cytosolic antibody-coated antigens triggers an innate immune intracellular signaling cascade that ultimately results in the induction of autophagy^[60]. Thus, it is plausible that p24-specific antibodies able to gain access to the cytoplasm may drive the rapid elimination of infected cells. Taken together, these observations suggest that p24-specific IgG1 antibodies may contribute to antiviral control via a spectrum of diverse mechanisms, thereby driving viral control by unconventional Fc-driven means.

This study used a large cohort that has been well characterized for cellular responses associated with disease outcome, enabling a holistic understanding of the relationship between HIV-specific antibody responses and function with markers of slower disease progression. Importantly, the study demonstrated the independent predictive value of p24-specific IgG1 responses on markers of disease progression, highlighting a potential role for Gag-specific antibodies in antiviral control. However because this study used cross-sectional samples from a cohort of chronically infected individuals with unknown dates of infection, it remains unclear whether p24-specific antibodies actively contribute to reduced disease progression. Thus future longitudinal studies on acutely infected patients may shed light on the impact of p24-specific IgG1 antibodies on rates of disease progression.

In summary, these data confirm the importance of antibody isotype selection in HIV-1 clade C infection and show that p24-specific IgG1 and not gp120-specific antibodies are associated with markers of slower HIV-1 disease progression. Moreover, p24-specific antibodies have the capacity to mediate Fc effector functions and predict viral control independent of Gag-specific CD4 and CD8 T cell responses or the presence of protective HLA-I alleles. Determining both the mechanisms of antiviral activity of Gag-specific antibodies will provide us with a greater understanding of its potential as a target for an antibody-based vaccine, particularly in light of its more conserved nature.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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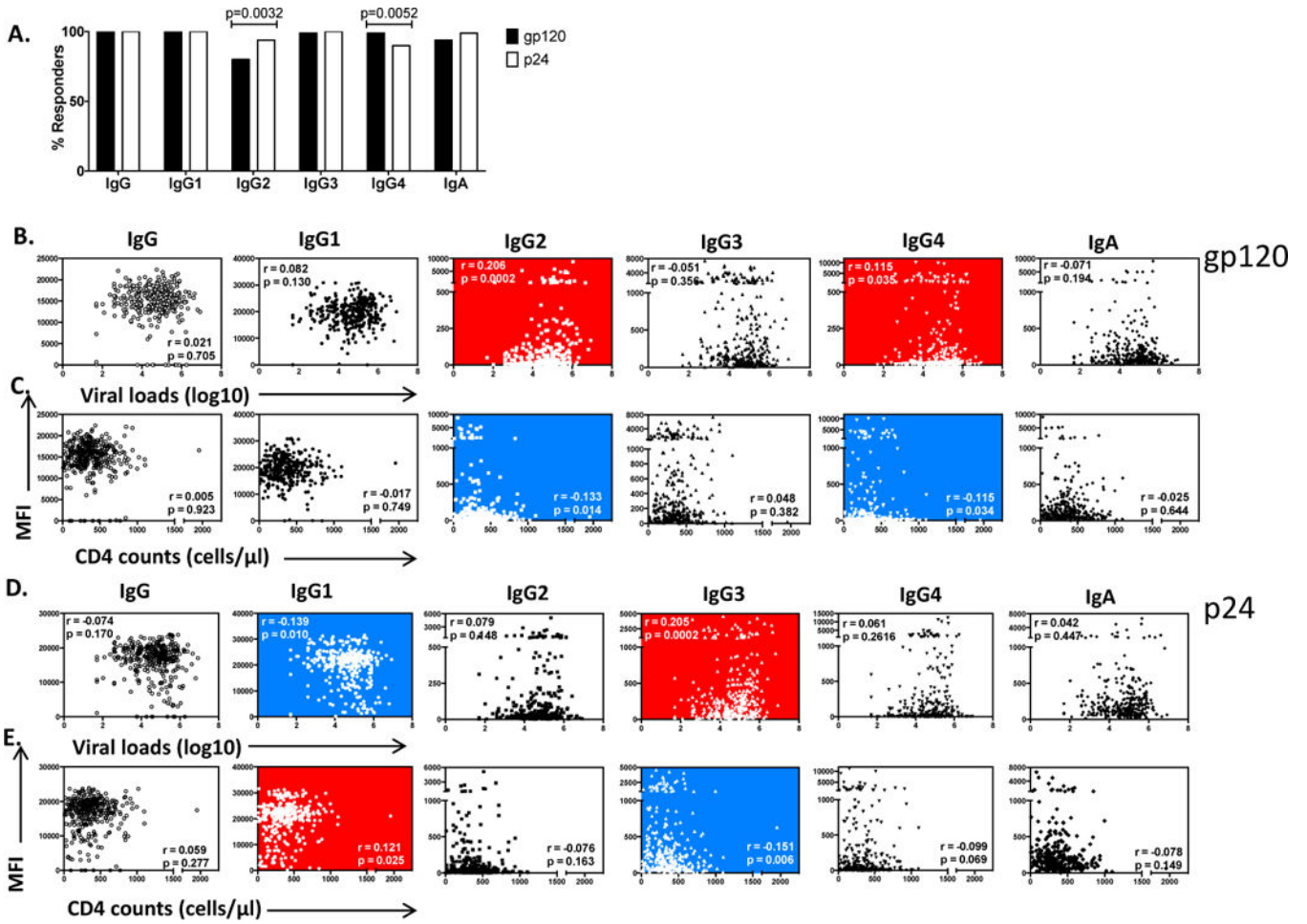


Figure 1. p24 IgG1 levels predict viral control in chronic HIV-1 subtype C infection
 Using recombinant protein coupled to fluorescent beads and diluted plasma, a customized luminex assay was performed to determine the levels of HIV-1 gp120- and p24 specific antibodies. A) describes % of positive responders. A positive threshold was determined as 3× mean +2SD of HIV negatives. The relationship between the mean fluorescence intensity (MFI) of gp120 total IgG and isotype specific binding antibodies and viral loads or CD4 counts is depicted in panels B) and C) while p24 is shown in panel D) and E). Plots with red and blue background represent significant positive and negative correlations respectively while a white background denotes non-significant relationships. p-values were calculated by Spearman’s correlation.

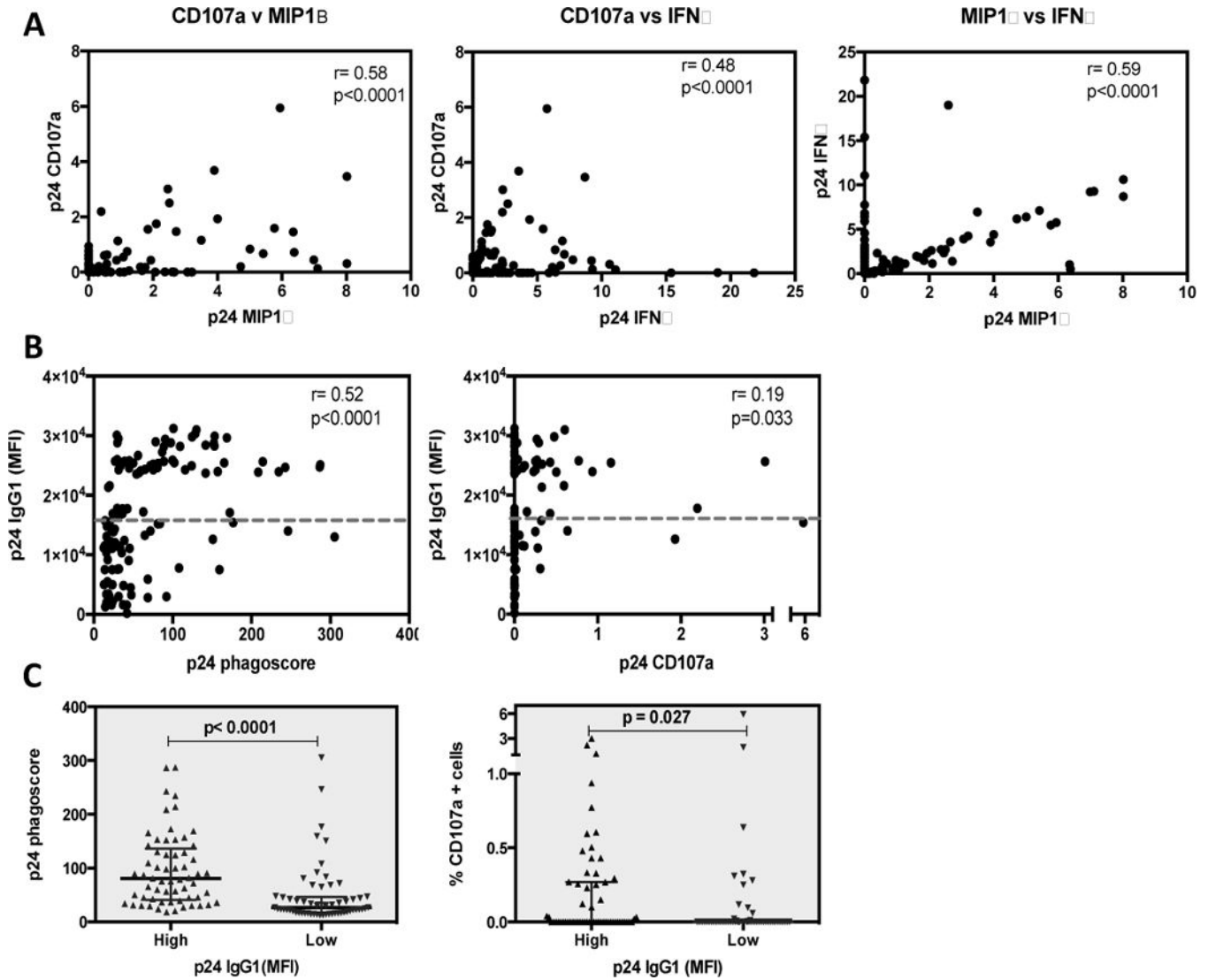


Figure 2. p24 IgG1 levels correlate with both ADCP phagosome and CD107a expression levels (A) Antibody mediated NK cell activation measured by expression of CD107a degranulation as a surrogate for ADCC, IFN γ cytokine secretion and MIP1 β were assayed in the presence of p24 protein as a surrogate for ADCC activity. All measurements of NK activation were high correlated with each other. (B) Antibody dependent cellular phagocytosis (ADCP) of fluorescently labelled bead conjugated to p24 protein by monocyte cell line THP-1 was assessed. The relationship between the mean fluorescence intensity (MFI) of p24 specific IgG1 with ADCP phagosome (left) and CD107a levels (right) are depicted. The grey dotted line on each graph represents the median p24 specific IgG1 levels. (C) Using the median, individuals were dichotomized into 2 groups with high and low p24 IgG1. The differences in magnitude of p24 ADCP (left) and ADCC (right) responses in the groups are shown respectively. Correlations between two variables were calculated by Spearman's test. Difference between two groups were calculated by 2-tailed Mann-Whitney test.

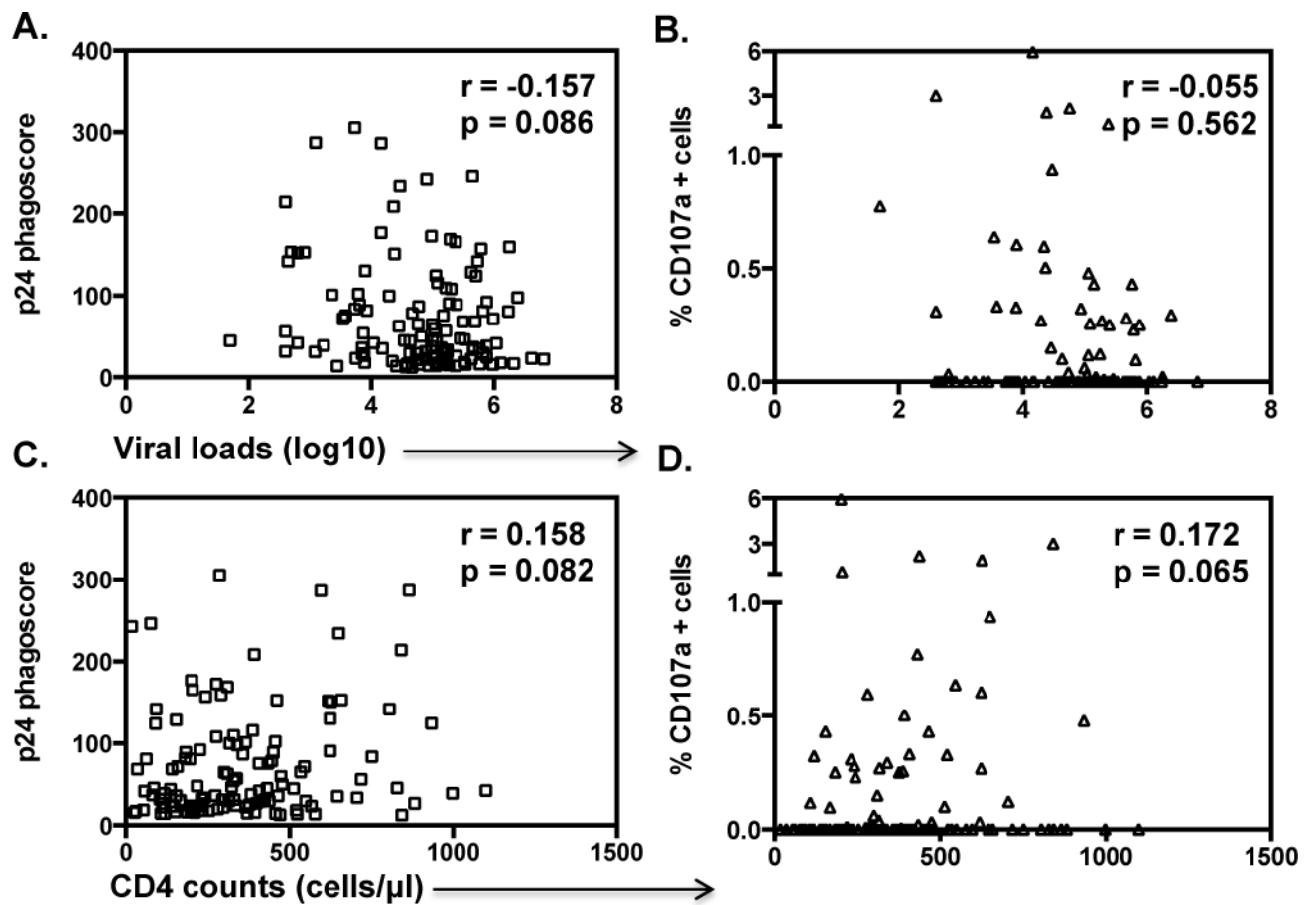


Fig. 3. Higher p24-specific Fc effector activity trends with markers of disease progression
 Diluted plasma was used to cross link and initiate phagocytosis of FITC fluorescent beads conjugated to p24 protein (ADCP) by THP-1 cells. Antibody mediated NK activation (detection of CD107a used as a surrogate for ADCC) was measured in the presence of p24 protein. The relationship between the magnitude of ADCP and ADCC responses and viral loads shown in panels A and B while panels C and D show the relationship with CD4 counts. p-values were calculated by Spearman's correlation.

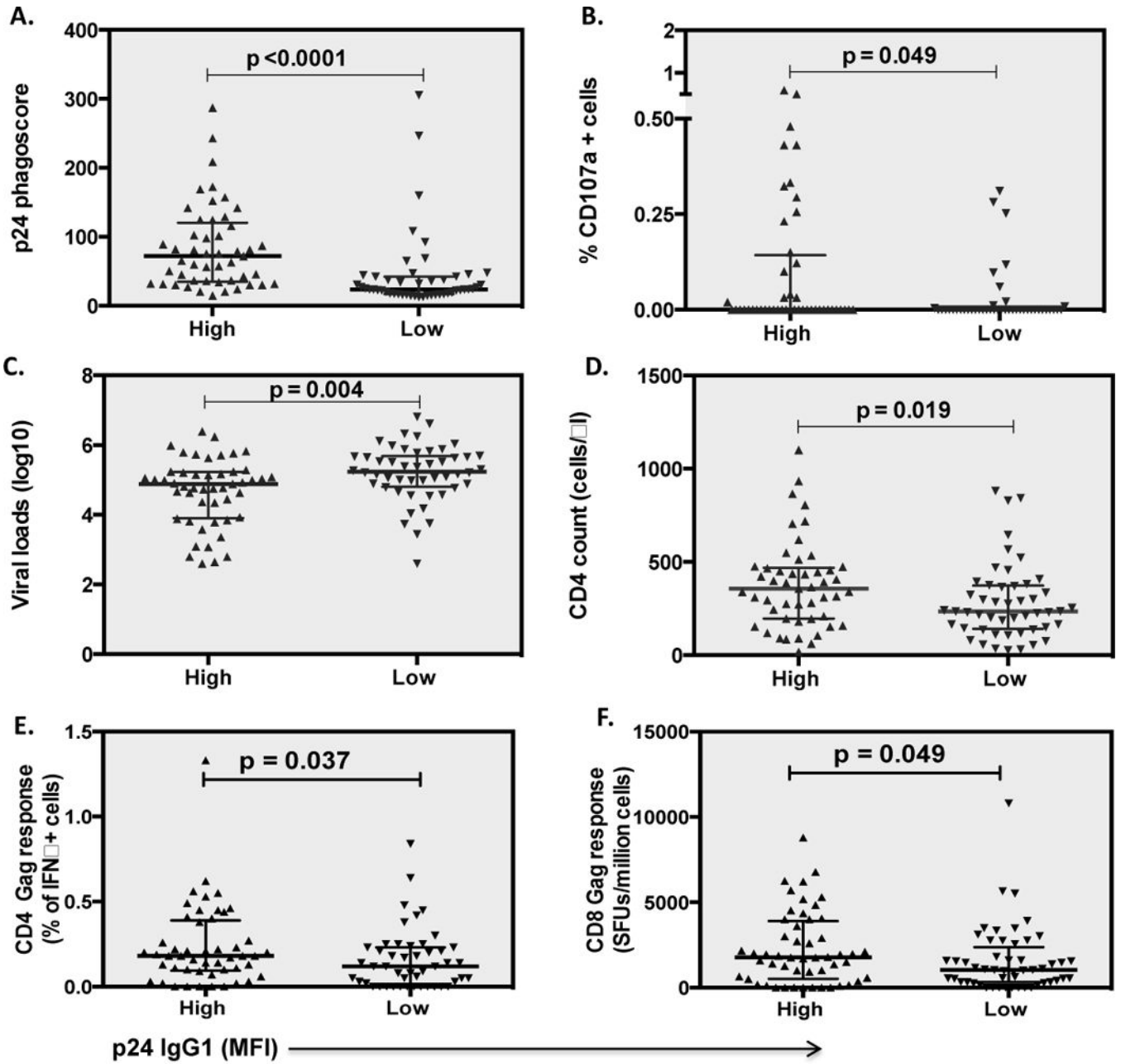


Figure 4. Relationship between HLA alleles, CD4 and CD8 Gag responses and Gag p24 IgG1 levels
 Patients with protective HLA alleles (HLA-B*57:01, HLA-B*58:01 and HLA-B* 81:00/01) were removed from this analysis and individuals with no protective alleles were dichotomized into groups of high and low p24 IgG1 levels. The panels show the difference in the magnitude of ADCP (A) and ADCC (B) responses, viral loads (C) and CD4 counts (D). Gag CD4 and CD8 activity was measured by ICS and ELISpot respectively and reported as % of cells positive for IFN γ or as number of spot forming units (SFUs/million PBMCs) respectively. Differences in CD4 and CD8 Gag responses in the dichotomized

groups of individuals with high and low p24 IgG1 respectively are shown in panels A and B. p-values were calculated by 2-tailed Mann-Whitney test

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Table 1

Univariate and multivariate analysis of relationships with Viral Loads or CD4 counts

Variable	Unadjusted		Adjusted	
	Coefficient (SE)	p value	Coefficient (SE)	p value
Viral load ^a				
p24 IgG1 ^b	-0.14 (0.05)	0.009	-0.13 (0.06)	0.030
CD4 Gag	-0.46 (0.12)	<0.001	-0.40 (0.13)	0.002
CD8 Gag	-0.42 (0.16)	0.008	-0.31 (0.16)	0.052
Protective HLA	-0.56 (0.13)	<0.001	-0.37 (0.14)	0.012
CD4 count ^c				
p24 IgG1 ^b	0.87 (0.32)	0.007	0.96 (0.34)	0.005
CD4 Gag	3.71 (0.70)	<0.001	3.55 (0.72)	<0.001
CD8 Gag	1.95 (0.94)	0.038	1.61 (0.89)	0.072
Protective HLA	1.50 (0.81)	0.066	1.29 (0.81)	0.115

^aLog transformed^bz-score normalized^csquare root transformed