

1 **The association of antibody immunity with cryptococcal antigenemia and mortality in a**
2 **South African cohort with advanced HIV disease**

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19 **Running title:** Antibody immunity and CrAg in PLWH.

1 **Abstract**

2 **Background:**

3 Asymptomatic cryptococcal antigenemia (positive blood cryptococcal antigen [CrAg]) is
4 associated with increased mortality in HIV-positive individuals even after adjusting for CD4
5 count and despite receiving antifungal treatment. The association of antibody immunity with
6 mortality in HIV-positive adults with cryptococcal antigenemia is unknown.

7 **Methods:**

8 *Cryptococcus* spp. capsular glucuronoxylomannan (GXM)- and naturally occurring β -glucans
9 (laminarin, curdlan)-binding antibodies were measured in blood samples of 197 South Africans
10 living with HIV who underwent CrAg screening and were prospectively followed up to 6
11 months. Associations between antibody titers, CrAg status and all-cause mortality were sought
12 using logistic and Cox proportional hazards regression, respectively.

13 **Results:**

14 Compared to CrAg-negative (n=130), CrAg-positive individuals (n=67) had significantly higher
15 IgG1 (median, 6,672 [interquartile range, IQR, 4,696-10,414] vs. 5,343 [3,808-7,722] μ g/ml;
16 $P=0.007$), IgG2 (1,467 [813-2,607] vs. 1,036 [519-2,012] μ g/ml; $P=0.01$), GXM-IgG (1:170 [61-
17 412] vs. 1:117 [47-176]; $P=0.0009$) and lower curdlan-IgG (1:47 [11-133] vs. 1:93 [40-206];
18 $P=0.01$) titers. GXM-IgG associated directly with cryptococcal antigenemia adjusted for CD4
19 count and antiretroviral therapy use (odds ratio, 1.64 for each log increase in titer; 95%
20 confidence interval [CI], 1.21-2.22). Among CrAg-positive individuals, GXM-IgG inversely
21 associated with mortality at 6 months adjusted for CD4 count and active or prevalent
22 tuberculosis (hazard ratio, 0.50; a two-fold reduction per log increase in titer; 95% CI, 0.33-
23 0.77).

1 **Conclusions:**

2 The finding of an inverse association of GXM-IgG with mortality in CrAg-positive individuals
3 suggests that GXM-IgG titer may have prognostic value in such individuals. Prospective
4 longitudinal studies of antibody levels to investigate this hypothesis and identify mechanisms by
5 which antibody may protect against mortality are warranted.

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8 **Keywords:** cryptococcal antigenemia, HIV/AIDS, serology, GXM-IgG, mortality
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1 **Introduction**

2 Despite an ongoing international effort to end cryptococcal-associated deaths [1], cryptococcosis
3 remains a major cause of morbidity and mortality among people living with human
4 immunodeficiency virus (HIV). Most cases occur in sub-Saharan Africa, where substantial
5 research efforts have advanced treatment and prevention of cryptococcosis [2, 3]. The
6 implementation of cryptococcal antigen (CrAg) screening in individuals with CD4 counts <200
7 cells/mm³ makes it possible to identify HIV-positive individuals at risk for developing
8 cryptococcal meningitis (CM) before the onset of symptoms, thereby facilitating earlier
9 diagnostic testing and treatment [2].

10 While the incidence of CM has declined with CrAg screening and preemptive treatment
11 programs, mortality remains higher in HIV-positive individuals with asymptomatic cryptococcal
12 antigenemia than those without cryptococcal antigenemia, even after adjustment for CD4 count
13 and despite preemptive antifungal therapy [3, 4]. Non-CD4 T-cell associated factors are likely to
14 affect the pathogenesis of cryptococcosis since cryptococcosis occurs in HIV-positive
15 individuals with CD4 T-cell counts >200 cells/mm³ [5] and in HIV-negative individuals [6].
16 Previous studies show that *Cryptococcus neoformans* (CN) capsular glucuronoxylomannan
17 (GXM) and β -glucan (laminarin, a branched β -[1-3]-glucan and curdlan, a linear β -[1-3]-glucan)-
18 binding antibody levels associate with CrAg or CM status in HIV-infected and HIV-uninfected
19 individuals [6-9]. In this study, we investigated associations between antibody immunity,
20 cryptococcal antigenemia, and mortality in a South African cohort with advanced HIV disease
21 with and without asymptomatic cryptococcal antigenemia.

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23

1 **Methods**

2 *Study design.* This is a retrospective study of serum samples obtained at enrollment of a
3 prospective cohort study of asymptomatic CrAg-positive and CrAg-negative HIV-positive adults
4 followed over 6 months after CrAg screening at two large urban hospitals in Johannesburg,
5 South Africa [4]. Participants were screened for asymptomatic/subclinical CM (positive
6 cerebrospinal fluid [CSF] using culture, India Ink microscopy, or CrAg test) when logistically
7 possible. All participants were started on antifungal therapy, and then antiretroviral therapy
8 (ART) as per local management guidelines (National Department of Health, South Africa, 2015).
9 Baseline demographic, clinical, and mortality data for up to 6 months were collected.

10

11 *Study population.* Eligible individuals were HIV-positive adults ≥ 18 years with CD4 count < 100
12 cells/mm³. Consecutive CrAg-positive and 1:2 ratio of CrAg-negative individuals attending
13 study sites following CrAg screening were invited to participate. Exclusion criteria were a
14 previous diagnosis of CM, or symptoms or signs of cryptococcal meningitis.

15

16 *Sample collection and processing.* Serum samples collected at baseline were frozen and stored at
17 -20°C . Before analysis, they were heat-inactivated at 56°C for 30 minutes to inactivate HIV
18 and/or other heat-sensitive viruses as well as complement and stored at 4°C .

19

20 *Antibody measurements.* Serum immunoglobulin (Ig), cryptococcal capsular GXM
21 polysaccharide-, and β -glucan (laminarin, curdlan)-binding antibodies were analyzed. The latter
22 bind conserved microbial determinants that are found on the cryptococcal cell wall and that of
23 other fungi [10]. IgM, IgG1, and IgG2 concentrations were measured using a Luminex platform

1 (Austin, TX) and quantified in units of $\mu\text{g/ml}$ as previously described [7, 11]. GXM-
2 IgM/IgG/IgA and β -glucan-binding IgM/IgG were detected by antigen-capture enzyme-linked
3 immunosorbent assays (ELISA) using plates coated, respectively, with GXM (from CN strain
4 24067) and laminarin or curdlan and reported as inverse titers as previously described [7, 11].
5 These assays have been used extensively for the detection of human GXM- and β -glucan-binding
6 antibodies [11-15].

7
8 *Ethics.* Samples were obtained during a cohort study [4], approved by the Ethics Committees at
9 the University of the Witwatersrand and the London School of Hygiene and Tropical Medicine.
10 All patients provided written or witnessed verbal informed consent, and de-identified serum
11 samples were studied under an Albert Einstein College of Medicine Institutional Review Board-
12 approved protocol (1989-228). Samples were transported outside of South Africa with the
13 permission of the local research ethics committee.

15 *Statistical analysis.*

16 *Bivariate analysis.* Baseline demographic, clinical characteristics, and antibody levels stratified
17 by plasma CrAg status were compared using the Wilcoxon rank-sum test for continuous and
18 Fisher's exact or χ^2 test for categorical variables. All tests were 2-sided ($\alpha=0.05$). Correlations
19 between measured antibody variables and between antibody variables and plasma CrAg titer
20 were assessed using Spearman's correlation coefficient.

21
22 *Model building.* Association between antibody levels and cryptococcal antigenemia was
23 estimated using multivariable logistic regression and association between antibody levels and

1 time to mortality was estimated using Cox proportional hazards regression. *A priori* decision was
2 made to include CD4 count in the model. Other variables were included in the full model if *P*-
3 value <0.20 in the univariable model. Variables assessed include ART use prior to enrollment,
4 tuberculosis diagnosed within 6 months prior to, or on the day of enrollment, and log-
5 transformed antibody levels. Total IgG1 or IgG2 were not included in the full model as they may
6 biologically be in the same causal pathway as the antigen-specific antibodies. Variables were
7 selected by backward elimination, checked for confounding and interaction with the main
8 independent variable of interest, GXM-IgG. In the survival analysis, GXM-IgG was evaluated as
9 both a continuous and a categorical variable in tertiles to demonstrate dose-response as Kaplan-
10 Meier curves. Hosmer-Lemeshow goodness of fit and Schoenfeld residuals were assessed in the
11 logistic regression and Cox model, respectively.

12
13 *Mediation analysis.* We fitted a linear regression model for the mediator (log of GXM-IgG) with
14 the exposure status (CrAg positive vs. negative) as the independent variable. The Cox
15 proportional hazards model for the outcome (mortality by 6 months) was built with the exposure
16 status and the mediator as independent variables and potential confounders as covariables. The
17 model also included the exposure-mediator interaction term to assess for potential interactions.
18 Regmedint package in R was used for the analysis [16, 17].

19
20 *Principal component analysis (PCA).* Baseline antibody markers were analyzed using PCA to
21 reduce complex correlated datasets into a series of linear, non-correlated principal components
22 (PCs) and avoid multiple comparisons as described [6]. Log-transformed antibody markers were

1 normalized to the mean. Differences in top PC scores by CrAg status (positive vs. negative) and
2 mortality were examined using a linear regression model.

3

4 All statistical analyses were performed in R and Stata/IC 16.1 software (College Station, TX:
5 StataCorp LLC).

6

7 **Results:**

8 *Study cohort.* Demographics and clinical characteristics of the 130 CrAg-negative and 67 CrAg-
9 positive cases are shown in Table 1. The median age was 39 years, 51% were male, 97% self-
10 identified as Black race, with no difference between the CrAg status groups. Compared to CrAg-
11 negative individuals, CrAg-positive individuals had lower CD4 counts (median 27 vs. 40.5
12 cells/mm³; $P=0.003$) and a greater proportion were on ART prior to enrollment (23.9% vs.
13 10.0%; $P=0.01$). The median plasma CrAg titer was 1:40 (interquartile range [IQR], 5-320). The
14 prevalence of tuberculosis within the past 6 months did not differ between the groups. There
15 were significantly more deaths in the CrAg-positive than CrAg-negative group (25.4% vs. 8.5%;
16 hazard ratio [HR], 3.43; 95% confidence interval [CI], 1.61-7.33) (Table 1, Supplementary Table
17 1).

18

19 *Antibody marker levels.* In the bivariate analysis, CrAg-positive individuals had higher inverse
20 titers of GXM-IgG (median, 169.5 [IQR, 61.1-411.9] vs. 117.3 [47.0-176.3]; $P=0.0009$) and
21 concentrations of IgG1 (6,672.2 [4,695.5-10,414] vs. 5,343.4 [3,807.7-7,721.9] $\mu\text{g/ml}$; $P=0.007$)
22 and IgG2 (1,467.4 [812.8-2,607] vs. 1,036.0 [519.2-2,012.4] $\mu\text{g/ml}$; $P=0.01$), and lower curdlan-
23 IgG (46.6 [11.1-133.0] vs. 93.2 [39.6-205.9]; $P=0.01$) than CrAg-negative individuals (Figure 1).

1 There were no between-group differences in laminarin-binding antibodies. There was
2 collinearity among GXM-IgG, GXM-IgM, and GXM-IgA (Supplementary Figure 1).

3
4 *Associations of antibody markers with CrAg status.* Higher GXM-IgG and lower curdlan-IgG
5 were each associated with CrAg-positive status adjusted for CD4 count and ART use prior to
6 enrollment in the multivariable logistic regression model (adjusted odds ratio [aOR], 1.64; 64%
7 increase in odds per log increase in GXM-IgG titer; 95% CI, 1.21-2.22 and aOR, 0.71; 29%
8 reduction in odds per log increase in curdlan-IgG titer; 95% CI, 0.58-0.87, respectively) (Figure
9 2).

10
11 *Correlation between plasma CrAg titer and antibody levels.* In the CrAg-positive group (n=67),
12 there were inverse correlations between CrAg titer and IgG1 ($\rho=-0.33$; $P=0.007$), IgG2 ($\rho=-0.29$;
13 $P=0.02$), laminarin-IgG ($\rho=-0.24$; $P=0.05$), and while not significant, GXM-IgG ($\rho=-0.19$;
14 $P=0.13$) (Supplementary Table 2).

15
16 *Associations of antibody markers with mortality.* Variables significantly associated with death
17 within 6 months in the univariable analysis included: positive plasma CrAg (HR, 3.43; 95% CI,
18 1.61-7.33), CrAg titer (HR, 1.19; 19% increase in hazard per log increase in titer; 95% CI, 1.05-
19 1.36), tuberculosis diagnosis within the prior 6 months (HR, 2.66; 95% CI, 1.26-5.59), and
20 GXM-IgM (HR, 0.67; 33% decrease in hazard per log increase in titer; 95% CI, 0.45-1.00)
21 (Supplementary Table 1). In the multivariable Cox proportional hazards analysis, there was an
22 interaction between CrAg status and GXM-IgG (interaction term, $P=0.04$), so the model was
23 stratified by CrAg status. In CrAg-positive individuals, GXM-IgG was inversely associated with

1 mortality in both unadjusted (HR, 0.55; a 1.8-fold reduction per log increase in titer; 95% CI,
2 0.36-0.84) and adjusted (HR, 0.50; a two-fold reduction per log increase in titer; 95% CI, 0.33-
3 0.77) model for CD4 count and tuberculosis (Table 2, Figure 3). GXM-IgG was not associated
4 with mortality in the CrAg-negative cohort. GXM-IgM was not associated with mortality in the
5 multivariable model.

6
7 *Mediation analysis.* Mediation analysis [16, 18] was performed to examine whether the influence
8 of the exposure (CrAg status) on the potential mediator (GXM-IgG) explains some or all of the
9 significance of the association between an exposure and an outcome (mortality) statistically. A
10 significant interaction between the exposure and mediator variable was detected and CrAg status
11 was significantly associated with GXM-IgG and mortality. The direct effect of CrAg status on
12 mortality was 1.71 (95% CI, 0.88, 2.54), the indirect effect mediated by GXM-IgG was -0.35
13 (95% CI, -0.67, -0.03), and -57% of the total effect on mortality was mediated by GXM-IgG
14 (95% CI, -1.21, 0.08) (Supplementary Figure 2).

15
16 *Principal component analysis.* Most of the variance in the dataset was explained by PC1 (35%)
17 and PC2 (19%), which were used to explore associations between antibody markers and CrAg
18 status and mortality. PC1 was composed primarily of GXM-IgA, GXM-IgG, and GXM-IgM.
19 PC2 was composed of curdlan-IgG, laminarin-IgG and a negative loading of laminarin-IgM
20 (Supplementary Figure 3). Among CrAg-positive individuals, PC1 score had a significant
21 inverse association with 6-month mortality; mean PC1 score was 0.32 among the CrAg-positive
22 individuals who survived, and -0.66 among the CrAg-positive individuals who died, a difference
23 of 0.98; 95% CI, 0.04, 1.92; $P=0.04$, indicating that higher levels of GXM-IgA, GXM-IgG, and

1 GXM-IgM were associated with higher survival rates in this population (Supplementary Table 3,
2 Supplementary Figure 4). There was no association between PC score and mortality in CrAg-
3 negative group. No association was found between PC score and CrAg status in the overall
4 cohort.

6 **Discussion**

7 In this retrospective serological study of HIV-positive CrAg-positive and negative
8 participants with CD4 <100 cells/mm³, higher enrollment GXM-IgG and lower curdlan-IgG
9 titers were associated with CrAg-positive status, but not CrAg titer. In addition, for CrAg-
10 positive participants, enrollment GXM-IgG, prior to antifungal therapy, was inversely associated
11 with mortality and a partial mediator on the causal pathway between CrAg and mortality by 6
12 months. This suggests that lower levels of GXM-IgG may in part contribute to excess mortality
13 in CrAg-positive HIV-positive individuals [3, 4] and echoes a pre-AIDS-era study in which an
14 absence of cryptococcal antibodies was a poor prognostic factor in patients with cryptococcal
15 meningitis [19].

16 Our data show that HIV-positive CrAg-positive participants with CD4 counts <100
17 cells/mm³ had higher IgG1 and IgG2 levels than CrAg-negative participants. However, IgG2
18 levels in both CrAg-positive and CrAg-negative participants were lower than normal HIV-
19 negative adult values. HIV is associated with hypergammaglobulinemia, generally manifested as
20 increased IgG1, but IgG2 is decreased [20]. IgG2 is the predominant IgG subclass of human
21 antibodies to capsular polysaccharides, including GXM [13, 21]. Thus, as in a prior study [7],
22 higher IgG2 levels in CrAg-positive than CrAg-negative participants may reflect a cryptococcal
23 response. We did not determine GXM-IgG subclasses in this study.

1 Higher GXM-IgG in CrAg-positive participants parallels multiple previous studies that
2 found higher GXM-IgG titers in HIV-positive and HIV-negative individuals with, than without,
3 cryptococcosis [7-9, 15]. Notably, in participants with cryptococcal antigenemia, lower GXM-
4 IgG was associated with mortality, while in an unadjusted analysis of the entire cohort, lower
5 GXM-IgM was associated with mortality. These associations were supported by the PCA, in
6 which the top principal component, primarily GXM-binding antibodies, was inversely associated
7 with CrAg titer and mortality in CrAg-positive individuals, although the GXM-IgG specific
8 signal in predicting CrAg-positive status was reduced by adding noise from GXM-IgM and
9 GXM-IgA. While association is not causation, these data suggest that GXM-binding antibodies
10 may have a protective role in those with cryptococcal antigenemia. This is consistent with a large
11 body of data showing that GXM monoclonal antibodies (mAbs) mediate beneficial effects,
12 including anticryptococcal activity *in vitro* [22], GXM clearance, including in a clinical trial in
13 patients with cryptococcal meningitis [23], and protection against lethal cryptococcal infection in
14 experimental models [24].

15 As previously reported [7], we found CrAg-negative participants had detectable GXM-
16 binding antibodies. Most likely, this reflects cryptococcal infection early in life or latency [25]
17 and/or ongoing environmental exposure, though we cannot rule out cross-reactivity with an
18 unknown determinant [21]. GXM-binding antibodies have been detected in HIV-negative and
19 HIV-positive individuals with and without a history of cryptococcosis as well as young children
20 [8, 14]. Thus, higher GXM-IgG titers in CrAg-positive compared to CrAg-negative individuals
21 most likely reflect an immune response to fungal growth. HIV infection impairs responses to
22 new, but not previously encountered antigens [20]. Although fungal growth is not always
23 detected (or quantified if present) in CrAg-positive persons, it is logical to posit that serum

1 cryptococcal antigenemia emerges when host defense is unable to contain latent or newly
2 acquired cryptococcal infection. Thus, an increase in GXM-IgG might signal cryptococcosis risk
3 in CrAg-negative individuals. Consistent with this idea, anti-mannan antibodies, which can
4 protect mice against experimental candidiasis [26], were detected in patients with candidiasis
5 prior to the development of positive blood cultures [27].

6 β -glucan-binding antibodies are part of the natural serum antibody repertoire, which
7 binds conserved microbial determinants found on the cryptococcal cell wall and other fungi and
8 is the first line of defense against pathogens [10, 28]. Although we did not find associations
9 between laminarin- or curdlan-binding antibodies and mortality, lower curdlan-IgG was
10 associated with CrAg status and laminarin-IgG inversely correlated with CrAg titer, a correlate
11 of risk for meningitis [29] and death [30]. Previous reports show associations between lower
12 laminarin-antibody titers and CrAg-positive status [7] as well as HIV-associated cryptococcus-
13 associated immune reconstitution inflammatory syndrome [11]. Evidence that naturally
14 occurring β -glucan-binding antibodies could contribute to host defense against cryptococcosis
15 includes that laminarin-binding antibodies have direct anti-cryptococcal effects *in vitro* [31],
16 indirect effects on the inflammatory response [11], and protect mice *in vivo* [32, 33]. Similar to
17 our findings in CrAg-positive participants, candidemic patients had higher β -glucan-associated-
18 protein MP65 titers than non-candidemic patients and survivors had higher titers than non-
19 survivors [34].

20 We found increased mortality in CrAg-positive participants with tuberculosis and lower
21 curdlan-IgM in participants with than without tuberculosis (Supplementary Table 4). Notably,
22 mycobacterial cell walls contain multiple glucans [35] and decreased levels of serum
23 mycobacterial cell wall-binding IgG were associated with disseminated tuberculosis [36]. Given

1 that tuberculosis is an independent risk factor for cryptococcosis [37, 38], more work is needed
2 to identify possible shared susceptibility factors.

3
4 *Strength and limitations*

5 We used well-established antigen-capture ELISAs to detect GXM-, laminarin-, and
6 curdlan-binding antibodies [9, 12, 15]. While previous studies examined the relationship between
7 antibody markers and cryptococcal disease status, we sought associations between antibody
8 markers and all-cause mortality. We used mediation analysis to establish that GXM-IgG partially
9 mediated a causal association between CrAg and mortality and PCA to eliminate
10 multicollinearity between antibody variables. Use of these platforms led to a novel result that
11 implicates GXM-binding antibody levels in survival of HIV-positive CrAg-positive individuals.

12 Our study also has limitations. Serum CrAg may have affected GXM-binding antibody
13 measurements, we did not assay for immune complexes. Since our assays were performed on a
14 single sample from one point in time, our data does not address longitudinal or temporal
15 variations in antibody levels. We did not perform functional assays and did not investigate the
16 relationship between B cells and cryptococcal disease status [39] or antibody levels as we did not
17 have access to peripheral blood mononuclear cells. Misclassification bias is possible. Though
18 rare, cryptococcal meningitis can occur in CrAg-negative individuals [40] and this may have
19 affected the difference in antibody levels by CrAg status. ART use prior to enrollment associated
20 with cryptococcal antigenemia, most likely reflecting prolonged immune suppression in ART-
21 experienced, non-adherent individuals. While we adjusted for ART use, our understanding of
22 possible interactions between ART and antibody levels is limited. There may have been other

1 unmeasured or latent confounders that could shed more light on the association between
2 antibody markers, ART, and mortality.

3

4 *Conclusion:*

5 Levels of IgG1, IgG2, GXM-IgG and β -glucan binding antibodies differed in CrAg-
6 positive and CrAg-negative individuals, GXM-IgG associated with CrAg status as did lower
7 curdlan-IgG, and lower GXM-IgG associated with an increased risk of all-cause mortality in
8 CrAg-positive individuals. These findings support the hypothesis that GXM- and β -glucan-
9 binding antibodies may enhance cryptococcal host defense, highlighting the need for mechanistic
10 and larger-scale prospective longitudinal clinical studies. Such studies may provide a better
11 understanding of the antifungal activities of GXM- and β -glucan-binding antibodies, identify
12 much needed biomarkers of cryptococcal pathogenesis and mortality, and inform opportunities
13 for earlier risk stratification and therapeutic intervention.

14

15

16

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5
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22

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1 Table 1. Baseline and clinical characteristics of participants by CrAg status

Covariables	Total (n=197)	CrAg-negative (n=130)	CrAg-positive (n=67)	P-value
Age, year	39.0 (32.4-47.1)	39.1 (33.0-47.8)	38.7 (31.7-47.1)	0.67
Sex				
Male	100 (50.8)	70 (53.9)	30 (44.8)	0.23
Female	97 (49.2)	60 (46.2)	37 (55.2)	
Race				
Black	191 (97.0)	125 (96.2)	66 (98.5)	0.67
Others	6 (3.1)	5 (3.9)	1 (1.5)	
CD4 count (cells/mm ³)	33 (12-61)	40.5 (16-64)	27 (7-40)	0.003
Plasma CrAg titers	1:40 (5-320)	n/a	1:40 (5-320)	
Baseline blood fungal growth	5 (2.5)	0	5 (7.5)	0.002
Tuberculosis in the past 6 mo. ¹	51 (25.9)	33 (25.4)	18 (26.9)	0.82
Tuberculosis treatment	29 (14.7)	18 (13.9)	11 (16.4)	0.63
ART prior to enrollment	29 (14.7)	13 (10.0)	16 (23.9)	0.01
Vital status at 6 mo.				
Alive	160 (81.2)	113 (86.9)	47 (70.2)	0.01
Dead	28 (14.2)	11 (8.5)	17 (25.4)	

Unknown	9 (4.6)	6 (4.6)	3 (4.5)
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1

2 Note: Categorical variables are summarized with counts, percentages and *P*-values based on
3 Fisher's exact or χ^2 test. Continuous variables are summarized with medians, interquartile range
4 and *P*-values based on Wilcoxon rank-sum test. *P*-values < 0.05 bolded.

5

6 ¹Includes pulmonary, disseminated, or extrapulmonary *Mycobacterium tuberculosis* infection
7 diagnosed within 6 months prior to, or on the day of enrollment.

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9 Abbreviations: ART, antiretroviral therapy; CrAg, cryptococcal antigen; HIV, human
10 immunodeficiency virus; mo, months; n/a, not applicable.

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1 Table 2. Stratified Cox proportional hazards regression model estimating GXM-IgG as a
2 predictor of mortality by 6 months following enrollment.

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	CrAg-negative (n=130)		CrAg-positive (n=67)	
	HR (95% CI)	P-value	HR (95% CI)	P-value
Death, no. (%)	11 (8.5)		17 (25.4)	
Model 1 – Unadjusted	1.03 (0.58-1.80)	0.93	0.55 (0.36-0.84)	0.006
Model 2 – Adjusted CD4 count	1.04 (0.59-1.82)	0.90	0.56 (0.36-0.86)	0.009
Model 3 – Adjusted CD4 count and tuberculosis	1.03 (0.60-1.76)	0.92	0.50 (0.33-0.77)	0.002

4

5 Note: GXM-IgG was log-transformed and used as a continuous variable. *P*-values < 0.05 bolded.

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7 Abbreviations: CI, confidence interval; CrAg, cryptococcal antigen; GXM,

8 glucuronoxylomannan; HR, hazard ratio.

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1 **Figure legends**

2 Figure 1. Total and antigen-specific antibody levels in 130 CrAg-negative and 67 CrAg-positive
3 individuals at baseline. Total Ig levels were determined by Luminex and GXM, laminarin, and
4 curdlan antibody titers were determined by enzyme-linked immunosorbent assay. Ig
5 concentrations in units of $\mu\text{g/ml}$ (A) and the inverse of GXM (B), laminarin (C), and curdlan (D)
6 titers, depicted as medians and interquartile ranges, are shown on the y-axis for each group
7 shown on the x-axis. CrAg status is represented by (+) or (-) to indicate positive or negative
8 CrAg status, respectively. * $P < 0.05$; ** $P \leq 0.001$. Wilcoxon rank-sum test. Abbreviation: CrAg,
9 cryptococcal antigen; GXM, glucuronoxylomannan; Ig, immunoglobulin.

10

11 Figure 2. Forest plots of the univariable (A) and multivariable (B) logistic regression analysis
12 estimating the association between antibody markers and plasma CrAg status in a cohort of 130
13 CrAg-negative and 67 CrAg-positive individuals. Antibody variables were log-transformed.
14 Covariables with $P < 0.20$ in the univariable analysis were included in the multivariable analysis
15 (CD4 count and ART use). Tuberculosis includes pulmonary, disseminated, or extrapulmonary
16 infection diagnosed within 6 months prior to enrollment. ART use includes treatment history
17 prior to enrollment. P -values < 0.05 bolded. Abbreviations: aOR, adjusted odds ratio; ART,
18 antiretroviral therapy; CI, confidence interval; CrAg, cryptococcal antigen; GXM,
19 glucuronoxylomannan; OR, odds ratio.

20

21 Figure 3. Kaplan-Meier Curves for survival by baseline GXM-IgG categories. The range of
22 inverse titers of GXM-IgG was categorized into tertiles; low (1.9-62.2), medium (62.8-172.0),
23 and high GXM-IgG (176.3-3,453.3). Curves are shown for the 67 CrAg-positive individuals

1 (Panel A) and 130 CrAg-negative individuals (Panel B). GXM-IgG was log-transformed in the
2 survival analysis. Abbreviations: GXM, glucuronoxylomannan.

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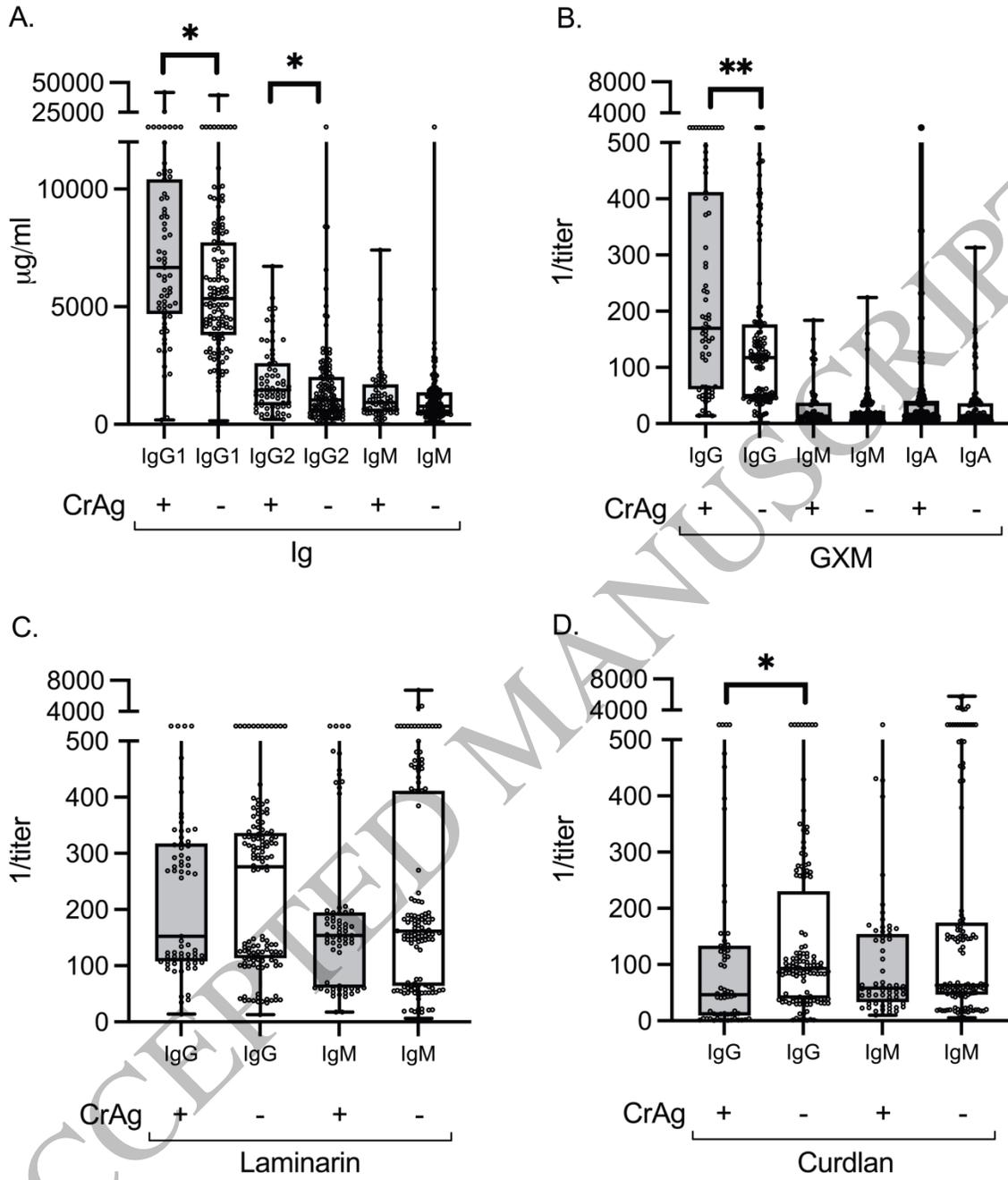
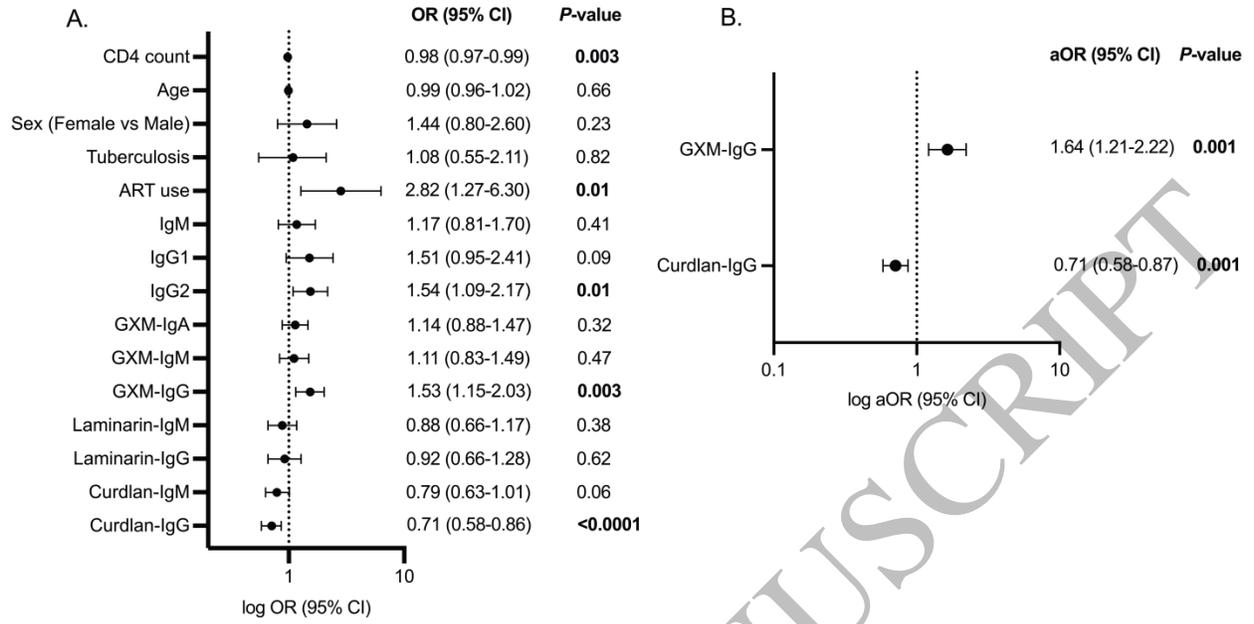


Figure 1
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Figure 2
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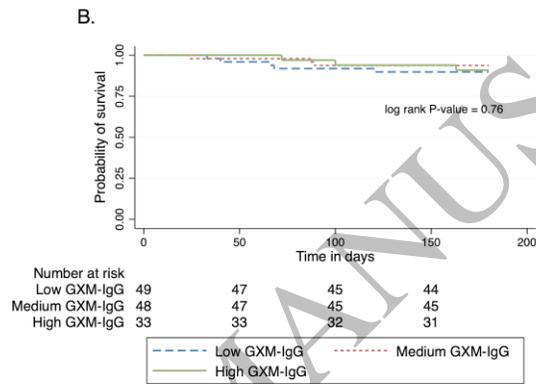
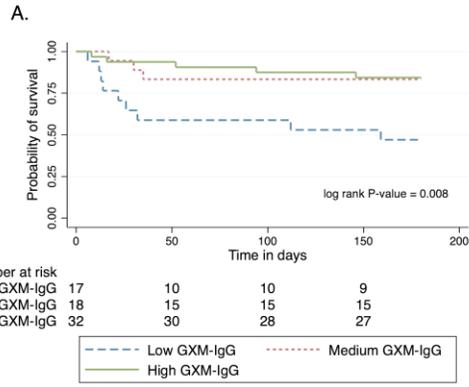


Figure 3
76x110 mm (x DPI)

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