

1 **Dynamics of *Mycobacterium tuberculosis* Ag85B revealed by sensitive ELISA**

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20 Running title: Ag85B secretion and trafficking

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28 Word count for the abstract: 233

29 Word count for the text: 3,812

30

31 **Abstract**

32 Secretion of specific proteins contributes to pathogenesis and immune responses in tuberculosis and  
33 other bacterial infections, yet the kinetics of protein secretion and fate of secreted proteins in vivo are  
34 poorly understood. We generated new monoclonal antibodies that recognize the *M. tuberculosis*  
35 secreted protein, Ag85B, and used them to establish and characterize a sensitive ELISA to quantitate  
36 Ag85B in samples generated in vitro and in vivo. We found that nutritional or culture conditions had  
37 little impact on secretion of Ag85B, and that there is considerable variation in Ag85B secretion by  
38 distinct strains in the *M. tuberculosis* complex: compared with the commonly-used H37Rv strain  
39 (Lineage 4), *M. africanum* (Lineage 6) secretes less, and two strains from Lineage 2 secrete more  
40 Ag85B. We also used the ELISA to determine that the rate of secretion of Ag85B is 10- to 100-fold  
41 lower than that of proteins secreted by gram-negative and gram-positive bacteria, respectively. ELISA  
42 quantitation of Ag85B in lung homogenates of *M. tuberculosis* H37Rv-infected mice revealed that  
43 although Ag85B accumulates in the lungs as the bacterial population expands, the amount of Ag85B  
44 per bacterium decreases nearly 10,000-fold at later stages of infection, coincident with development of  
45 T cell responses and arrest of bacterial population growth. These results indicate that bacterial protein  
46 secretion in vivo is dynamic and regulated, and quantitation of secreted bacterial proteins can  
47 contribute to understanding pathogenesis and immunity in tuberculosis and other infections.

48 **Importance**

49 Bacterial protein secretion contributes to host-pathogen interactions, yet the process and  
50 consequences of bacterial protein secretion during infection are poorly understood. We developed a  
51 sensitive ELISA to quantitate a protein (termed Ag85B) secreted by *M. tuberculosis* and used it to find  
52 that Ag85B secretion occurs with slower kinetics than for proteins secreted by gram positive and gram  
53 negative bacteria, and that accumulation of Ag85B in the lungs is markedly regulated as a function of  
54 the bacterial population density. Our results demonstrate that quantitation of bacterial proteins during  
55 infection can reveal novel insights into host-pathogen interactions.

56

57 **Introduction**

58 *Mycobacterium tuberculosis* employs secretion of specific proteins (estimated to include up to  
59 ~25% of the bacterial proteome (1)) to survive, interact with host targets during infection (2, 3),  
60 manipulate its intracellular niche (2-7), and induce protective and pathogenic immune  
61 responses (8). Among the proteins that are most abundant in *M. tuberculosis* culture  
62 supernatants are members of a family of three closely related proteins, the antigen 85 (Ag85)  
63 complex consisting of Ag85A, Ag85B, and Ag85C (9). All three of these proteins exhibit  
64 enzymatic activity as mycolyl transferases, in which they catalyze transesterification reactions  
65 to synthesize trehalose monomycolate (TMM), trehalose dimycolate (TDM), and mycolated  
66 arabinogalactan (10, 11). Because of these enzymatic activities and their importance in  
67 constructing the mycobacterial envelope, Ag85A, Ag85B, and Ag85C have been considered  
68 potential drug targets for treatment of tuberculosis (10).

69 Due to their ability to induce adaptive CD4 and CD8 T lymphocyte responses in a broad range  
70 of vertebrate hosts, Ag85A and Ag85B have been investigated as antigens for tuberculosis  
71 vaccines, and are prominent components of at least seven candidate vaccines in various  
72 stages of development (<http://www.aeras.org>). Quantitative assays of mRNA have revealed  
73 that the genes encoding Ag85A and Ag85B (*fbpA* and *fbpB*, respectively) are expressed at  
74 high levels by bacteria in the lungs early after aerosol infection of mice, but their mRNA  
75 expression decreases markedly after the recruitment of antigen-specific effector T cells to the  
76 lungs (12-14). Consistent with the results of bacterial RNA quantitation, CD4 T cells specific for  
77 Ag85B are activated in the lungs between two and three weeks after infection of mice, but their  
78 activation markedly decreases concurrent with decreased bacterial expression of the *fbpB*  
79 gene (12, 15).

80 Despite considerable knowledge of the properties of the *fbpA* and *fbpB* genes and the  
81 antigenicity of their products, there is less information on the secretion, *in vivo* expression, and

82 trafficking of the Ag85A or Ag85B proteins. Because of interest in Ag85B as a vaccine and/or  
83 diagnostic antigen, we generated new monoclonal antibodies to Ag85B and used them to  
84 establish a highly sensitive and specific ELISA. We then employed the ELISA in studies of  
85 secretion and trafficking of the Ag85B protein in vitro and in vivo.

86

87 **Results**

88 *Generation and characterization of monoclonal antibodies to Ag85B*

89 Monoclonal antibodies were generated using standard methods (16), from mice immunized  
90 with purified recombinant *M. tuberculosis* Ag85B (rAg85B), expressed in *E. coli*. Three  
91 monoclonal antibodies (mAbs), termed 710, 711, and 712, were selected for characterization.  
92 When examined by direct ELISA using wells coated with purified rAg85A, rAg85B, or rAg85C,  
93 and varying concentrations of antibody, all three mAbs recognized Ag85B, and yielded  
94 equivalent signals (Figure 1A). The three mAbs also recognized Ag85A in the direct ELISA,  
95 although recognition of Ag85A required higher antibody concentrations and reached lower  
96 maximum signal intensity than with Ag85B. mAb 710 generated a higher signal intensity, and  
97 at lower antibody concentrations compared with mAbs 711 and 712; the latter two mAbs also  
98 exhibited detectable binding to Ag85A when used at concentrations  $\geq 1 \mu\text{g/ml}$  (Figure 1B). In  
99 contrast, mAbs 711 and 712 did not bind to Ag85C in direct ELISA at any antibody  
100 concentration examined, while mAb 710 bound Ag85C at antibody concentrations as low as  
101  $0.01 \mu\text{g/ml}$  (Figure 1C). Testing of the mAbs at a fixed concentration of  $1 \mu\text{g/ml}$  on a dilution  
102 series of recombinant protein in direct ELISA, revealed that mAb 710 bound to Ag85A, Ag85B,  
103 and Ag85C at lower antigen concentrations than did mAb 711 or 712; mAbs 711 and 712 were  
104 indistinguishable in this assay. All three mAbs bound to Ag85B at lower antigen  
105 concentrations than required for binding to Ag85A or Ag85C (Figure 1D-E). Together, these  
106 results indicate that mAbs 710, 711, and 712 preferentially recognize Ag85B, although each of  
107 the mAbs also binds Ag85A and Ab85C when these antigens are present at high  
108 concentrations.

109 *Sandwich ELISA for quantitation of Ag85B*

110 When the three mAbs were used in pairwise combinations, using one mAb for capture, and  
111 another mAb conjugated to horseradish peroxidase (HRP) for detection of bound antigen, the

112 highest sensitivity for detection of Ag85B was obtained when mAb 710 was used as the  
113 capture antibody (Figure 2A). Although the differences were slight, sensitivity appeared to be  
114 greater when mAb 711 rather than mAb 712 was used as the detecting antibody. Compared  
115 with Ag85B capture by mAb 710, mAb 711 capture allowed detection of bound Ag85B by mAb  
116 710 but not by mAb 712. When mAb 712 was used for capture, mAb 710 was able to bind  
117 Ag85B, but mAb 711 was not. Together, these data indicate that the epitopes recognized by  
118 mAbs 711 and 712 overlap or may even be identical, while the epitope recognized by mAb 710  
119 is distinct from those of mAbs 711 and 712. Despite detectable binding of all three mAbs to  
120 Ag85A in the direct ELISA (Figure 1B), none of the combinations of capture or detecting  
121 antibodies yielded a signal when Ag85A was used as the antigen in the sandwich ELISA  
122 (Figure 2B). Likewise, despite binding of mAb 710 to Ag85C by direct ELISA, none of the  
123 capture or detecting antibody combinations resulted in a detectable signal in the sandwich  
124 ELISA when Ag85C was used as the antigen (Figure 2C). Together, these results suggest that  
125 the epitope recognized by mAb 710 is at least partially shared by Ag85A, Ag85B, and Ag85C,  
126 while the epitope(s) recognized by mAbs 711 and 712 is more specific to Ag85B.

127 To confirm specificity of the ELISA with the combination of mAbs 710 and 711 using native *M.*  
128 *tuberculosis* proteins, we examined culture filtrates of wild-type H37Rv and of our previously-  
129 characterized Ag85B-deficient (*fbpB* null) mutant strain of H37Rv (12). This yielded no  
130 detectable signal in the ELISA when culture filtrates of the Ag85B-deficient bacteria were  
131 examined, despite the detection of an abundant signal when wild-type culture filtrates were  
132 examined (Figure 2D). Since the Ag85B-null bacteria retain the capacity to synthesize and  
133 secrete Ag85A and Ag85C (17), these results provide further evidence that the sandwich  
134 ELISA with mAbs 710 and 711 is highly specific for Ag85B.

135 The assay in this form, with mAb 710 as the capture antibody and HRP-coupled mAb 711 as  
136 the detection antibody, has been run in our laboratory >30 times to determine Ag85B content

137 in various samples. Each assay comprised at least one standard curve with rAg85B per ELISA  
138 plate, starting at 10 ng/ml or above and including 2-fold dilutions to loss of signal (generally,  
139  $\leq 20$  pg/ml). Three representative assay standard curves are shown in Figure 2E. Ag85B in  
140 samples was quantitated based on either a dose-response sigmoidal regression curve or a  
141 linear regression in the linear proportion of the curve (0 ng/ml – 2.5 ng/ml). With both methods  
142 regression curves had  $R^2$  values  $> 0.99$ . When necessary, samples were diluted for ELISA to  
143 be in linear range of the standard curve and either at least two technical replicates or biological  
144 replicates (when available) were run.

145 *Carbon source effects, strain-dependent variation, and kinetics of Ag85B secretion*

146 We used the ELISA to examine whether Ag85B secretion is governed by bacterial growth  
147 conditions. When grown in rich broth (Middlebrook 7H9 medium with 10% albumin-dextrose-  
148 catalase), a 10-fold difference in Tween 80 concentration (0.5% vs 0.05%) did not affect the  
149 amount of Ag85B secreted by *M. tuberculosis* H37Rv in over a 24 h period (Figure 3A). When  
150 H37Rv was grown in minimal defined media (Sauton's medium, with 0.05% Tween 80, 0.5%  
151 BSA with 0.05% tyloxapol), or 7H9 broth with the addition of 0.2% acetate, dextrose or glycerol  
152 as carbon sources, the addition of acetate or glycerol did not alter Ag85B secretion, whereas  
153 0.2% dextrose increased Ag85B secretion approximately 2-fold (Figure 3A). These results  
154 indicate that bacterial metabolism as dictated by alternative carbon sources can affect the rate  
155 and/or amount of Ag85B secretion, but that the concentration of Tween 80 had little  
156 measurable effect.

157 We previously reported that *M. africanum* expresses lower levels of Ag85B than *M.*  
158 *tuberculosis* H37Rv, as detected by immunoblotting and by the magnitude of in vivo antigen-  
159 specific CD4 T cell responses (18). To verify this result and determine whether other strain-  
160 dependent variation in Ag85B secretion exists, we used the Ag85B-specific ELISA to assay  
161 culture filtrates from phylogenetically distinct bacterial isolates. This confirmed that *M.*

162 *africanum* secretes significantly lower quantities of Ag85B compared with *M. tuberculosis*  
163 H37Rv (Figure 3B). We also found that two distinct isolates of *M. tuberculosis* from Lineage 2  
164 (includes the Beijing family) secrete significantly higher quantities of Ag85B than H37Rv  
165 (Figure 3B). These results indicate substantial variation in secretion of Ag85B, according to the  
166 bacterial strain, possibly in a lineage-dependent manner.

167 To further characterize the properties of Ag85B, we determined the kinetics of its secretion by  
168 *M. tuberculosis* H37Rv. Washed mid-log phase bacteria were suspended in fresh 7H9 media  
169 at  $10^8$  CFU/ml, and culture filtrates were harvested at multiple intervals and assayed by ELISA.  
170 The rate of accumulation of Ag85B in culture filtrates was linear ( $r^2 = 0.93$ ) during the 24-hour  
171 period of sampling, at  $1.8 \pm 0.2$  ng/ml/hr (Figure 3C). Since the mature Ag85B protein has a  
172 molecular weight of 34,580, this is equivalent to 52 fmol per  $10^8$  CFU/hr, or approximately 300  
173 molecules secreted per bacterial cell/hr.

174 *Cell-free Ag85B during infection in vivo and in vitro*

175 Despite their biological activity and roles in pathogenesis and immune responses, little is  
176 known of the in vivo fate or distribution of secreted *M. tuberculosis* proteins, especially during  
177 infection. Therefore, we examined supernatants of lung homogenates obtained at various  
178 intervals after infecting C57BL/6 mice with *M. tuberculosis* H37Rv by aerosol. We found that  
179 Ag85B was detectable in lung homogenate supernatants in some, but not all, mice as early as  
180 4-8 days post-infection, followed by a progressive increase between 14 and 21 days post-  
181 infection (Figure 4A). Since the samples for assay were taken during the progressive growth  
182 phase of the bacteria in the lungs, the quantity of Ag85B detected is a function of the amount  
183 secreted by each bacterium and by the number of bacteria, which increases progressively until  
184 approximately 21 days post-infection (19). Therefore, we normalized the concentration of  
185 Ag85B in lung homogenates by the number of bacteria present in the lungs at each time point  
186 sampled. This revealed a progressive decrease in the amount of Ag85B relative to the

187 bacterial population size, commencing between 8 and 11 days post infection (Figure 4B). This  
188 result is consistent with results of Ag85B RNA quantitation on bacterial populations in the lungs  
189 of immunocompetent mice (12-14), and with results of studies indicating that activation of  
190 Ag85B-specific CD4 T cells in vivo decreases markedly after 2-3 weeks of infection, due to  
191 limited availability of antigen (12, 15).

192 There are several potential sources of free Ag85B in the lungs. First is that some of the  
193 bacteria may be extracellular and may secrete Ag85B directly to the intercellular spaces in  
194 lung tissues. Second is that, as we have recently reported, Ag85B and other secreted  
195 mycobacterial proteins can be exported from infected cells by a vesicular transport pathway  
196 (20, 21). A third potential mechanism is release of Ag85B associated with bacterial membrane  
197 vesicles (22) and/or exosomes shed by infected cells (23, 24). A fourth possible mechanism is  
198 that Ag85B synthesized by intracellular bacteria may be released from dying infected cells. We  
199 investigated the latter possibility, using bone marrow-derived dendritic cells infected with *M.*  
200 *tuberculosis* H37Rv. Under conditions of the multiplicities of infection, bacterial strain, and time  
201 points used, we observed a range of loss of cell viability as reflected by luminescence assay of  
202 ATP in cell lysates, after harvesting conditioned medium for assay of Ag85B by ELISA. This  
203 revealed a correlation between loss of cell viability and the quantity of Ag85B in conditioned  
204 media ( $r^2 = 0.5777$ ;  $p < 0.0001$ ) (Figure 5), indicating that Ag85B can be released from dead  
205 and/or dying infected cells in a form that remains detectable by ELISA. This suggests that  
206 death of infected cells can be a source of the Ag85B detected in cell-free homogenates of the  
207 lungs of infected mice, as in Figure 4.

208

209 **Discussion**

210 In this work, we developed three new murine monoclonal antibodies by immunizing mice with  
211 recombinant *M. tuberculosis* Ag85B, and used them to develop a sensitive and specific ELISA  
212 for quantitation of Ag85B in various biological samples. We found that, compared with the  
213 H37Rv strain from the *M. tuberculosis* complex lineage 4, two isolates from lineage 2 secreted  
214 greater quantities, while a lineage 6 (*M. africanum*) isolate secreted lesser quantities of Ag85B.  
215 Since Ag85B plays a role in synthesis of trehalose dimycolate, and trehalose dimycolate is a  
216 potent proinflammatory mediator during *M. tuberculosis* infection (25-28), the results suggest  
217 that differences in Ag85B secretion may contribute to strain-dependent differences in the  
218 proinflammatory properties of distinct strains and lineages in the *M. tuberculosis* complex (29).  
219 Strain variability in secretion may also influence the frequency and magnitude of T cell  
220 responses to Ag85B, which may provide an explanation for the finding that fewer human  
221 subjects exhibit detectable responses to Ag85B compared with other antigens such as ESAT-6  
222 and/or CFP-10 (30, 31). Likewise, bacterial strain-dependent variation in secretion of Ag85B  
223 may influence the protective efficacy of TB subunit vaccines that include Ag85B: even if a  
224 vaccine generates immune responses to Ag85B, vaccination may provide little protection  
225 against infection with mycobacterial strains that express and secrete lesser quantities of  
226 Ag85B. In studies of subunit vaccines that include Ag85B, it may be informative to  
227 characterize Ag85B expression and secretion in the isolates of *M. tuberculosis* from subjects  
228 that develop breakthrough infections despite apparently-appropriate immune responses to the  
229 vaccine antigen.

230 We also used the ELISA to determine the rate of secretion by *M. tuberculosis* H37Rv in broth  
231 culture as approximately 300 molecules of Ag85B per bacterial cell per hour. This is 10 to  
232 100-fold lower than the rate of translocation of *E. coli* proOmpA (4.5 mol/min) (32), or the  
233 secretion of Staphylococcal toxic shock toxin ( $1.1 \times 10^4$  molecules/CFU/hr) (33). To fully

234 understand the basis for this difference will require determination of the rates of protein  
235 synthesis in *M. tuberculosis* compared with that of other bacteria, and characterization of the  
236 factors and mechanisms that determine the rate of protein secretion by distinct secretion  
237 systems in bacteria. Given the important roles of secreted protein virulence factors in *M.*  
238 *tuberculosis* and other bacterial pathogens, better understanding of protein secretion may  
239 reveal new targets for therapeutic modulation and reduction of disease and pathogen  
240 transmission.

241 A third major finding in these studies is that *M. tuberculosis* Ag85B can be found in  
242 homogenized lung tissue supernatants from infected mice. This finding has several potential  
243 implications. If data from studies in human subjects provide supportive data, then detecting  
244 Ag85B in respiratory secretions may provide a rapid and economical approach to diagnosis of  
245 pulmonary tuberculosis and may also be useful in monitoring responses to treatment. Another  
246 is that the presence of extracellular Ag85B in lung tissue may make Ag85B available for  
247 uptake and processing and presentation by uninfected dendritic cells and macrophages in the  
248 lungs. Since we have reported that direct recognition of infected cells is required for optimal  
249 CD4 T cell control of intracellular *M. tuberculosis* (34), acquisition and presentation of Ag85B  
250 by uninfected cells in the lungs may provide antigen-loaded decoys for Ag85B-specific CD4 T  
251 cells that reduce the frequency of recognition of infected cells by those T cells. This is  
252 consistent with our recent finding in mouse lungs (35) and the finding in lungs of *M.*  
253 *tuberculosis*-infected rhesus macaques (36) that only a small fraction of the T cells in the lungs  
254 are in close contact with *M. tuberculosis*-infected cells. Our associated finding in this work that  
255 death of *M. tuberculosis*-infected host cells can be associated with release of Ag85B to the  
256 extracellular space indicates that the presence of Ag85B (and other mycobacterial proteins) in  
257 extracellular tissue compartments in the lungs may be the consequence of secretion by  
258 extracellular bacteria, vesicular transport from infected cells (21), carriage by exosomes (37),

259 38), and release by dying or dead host cells. Since Ag85B (39, 40), ESAT-6 (41) and other  
260 mycobacterial proteins have been reported to act on host cells to modulate inflammation, our  
261 findings provide additional evidence for the plausibility that extracellular mycobacterial proteins  
262 contribute to the pathogenesis of TB, and may be susceptible to therapeutic modulation.

263 **Materials and Methods**

264 *Ethics statement*

265 All animal experiments were done in accordance with procedures approved by the New York  
266 University School of Medicine Institutional Animal Care and Use Committee (Laboratory  
267 Animal Care Protocol 150502-01), which conformed to the guidelines provided by the Guide  
268 for the care and Use of Laboratory Animals of the National Institutes of Health.

269 *Bacterial strains*

270 The stocks of *M. tuberculosis* H37Rv, H37Rv: $\Delta fbpB$  (Ag85B-null), 4334, and *M. africanum*  
271 1182 used in our laboratory and for these studies have been previously described (19, 42-44).  
272 The *M. tuberculosis* HN878 strain was obtained from BEI Resources.

273 *Recombinant Ag85B*

274 RV1886c-ss.pET23b was transformed into *E. coli* BL21(DE3) pLysS (Invitrogen) and induced  
275 with 0.8 mM IPTG for 4 h at 37° C. Cultures were lysed with 50 mM Tris pH 7.0, 150 mM NaCl,  
276 lysozyme, benzonase, and 1 mM PMSF for 30 minutes at 22°C on an orbital shaker. The  
277 lysate was spun 10,000 x g for 30 minutes, sterile filtered, and loaded onto an AKTA FPLC  
278 His-Trap column (GE). The column was washed with 50 mM imidazole, and recombinant  
279 protein was eluted with 250 mM imidazole. Purity was assessed by SDS-PAGE.

280 *mAb generation and selection*

281 Balb/C mice were immunized with purified recombinant Ag85B (100 µg/mouse x 2 injections  
282 followed by 50 µg/mouse for 2 additional injections) subcutaneously in TiterMax Gold  
283 (TiterMax, Norcross, GA, USA), followed by 50 µg/mouse given intravenously 3 days before

284 harvesting and using spleen cells in fusions with P3X63Ag8 myeloma cells. Hybridoma  
285 supernatants were screened for recognition of rAg85B-coated wells by ELISA.

286 *Direct ELISA for characterization of individual mAbs*

287 Ag85A (BEI Resources NR-14871), Ag85B (our purified recombinant), or Ag85C (BEI  
288 Resources NR-14858) were used to coat wells at 0.5  $\mu$ g/well in PBS and incubated overnight  
289 at 4°C. The plates were washed 3 times with PBS pH 7.4 with 0.05 % Tween 20, and blocked  
290 with PBS containing 1.0 % BSA for 1 hour. A starting concentration of 5  $\mu$ g/ml of mAb 710,  
291 711, or 712 was serially diluted and 200  $\mu$ l/well incubated at room temperature for 2 hours.  
292 Plates were washed 5 times and incubated with goat anti-mouse IgG HRP (MP Biomedicals)  
293 for 1 hour at room temperature. Plates were washed 7 times and developed with TMB  
294 substrate according to the manufacturer's instructions (BD). The reaction was stopped with 2M  
295 sulfuric acid and absorbance read at 450 nm with a Synergy H1 micro plate reader (BioTek).

296 *Mycobacterial culture filtrates*

297 *M. tuberculosis* strains H37RV, 4334, and HN878, or *M. africanum* strain 1182, were  
298 inoculated from a 1 ml frozen stock of approximately  $3 \times 10^8$  CFU/ml into 10 ml of Middlebrook  
299 7H9 media supplemented with 0.5 % albumin, 0.2 % dextrose and 0.3 mg/100 ml catalase (in  
300 the text, termed 7H9) and grown to late log phase. The cultures were then passaged and  
301 grown once to mid log phase. Cultures were then collected and spun at 150 x g. The collected  
302 supernatant was then spun at 3,750 x g for 5 minutes and washed with PBS. The pellets were  
303 then resuspended in fresh 7H9 media and grown for 24 h. The cultures were pelleted and the  
304 media was sterile filtered.

305 *Sandwich ELISA development and optimization*

306 Individual mAbs were used at 2.5  $\mu$ g/ml in 50  $\mu$ l 0.05 M carbonate-bicarbonate buffer to coat  
307 wells overnight at 4° C. The plates were washed 3 times with PBS (pH 7.4) with 0.5% Tween  
308 20 and blocked with PBS containing 1.0% BSA for 1 hour. Antibodies for detection were

309 conjugated to HRP according to manufacturer's recommendations (Abcam). Briefly, 1  $\mu$ l of  
310 Modifier was mixed with 10  $\mu$ l of antibody and then added to a vial of HRP Mix. The vial was  
311 incubated overnight at room temperature in the dark. After incubation, 1  $\mu$ l of Quencher was  
312 added and mixed. The conjugated antibodies were diluted to 0.5  $\mu$ g/ml in PBS with 1% BSA  
313 and stored in aliquots at -20 °C until first use. Upon thawing, conjugated antibodies were  
314 stored at 4 °C.

315 For ELISA, samples were added to plates coated with the designated antibody and incubated  
316 at room temperature for 2 hours, then plates were washed 5 times. The labeled, detecting  
317 antibody was added and the plates were incubated at room temperature for 1 hour. Plates  
318 were washed 7 times and developed with substrate according to the manufacturers'  
319 instructions (BD). The reaction was stopped with 2M sulfuric acid and absorbance read at 450  
320 nm.

321 *Sandwich ELISA application*

322 The routine sandwich ELISA used the procedures as described above, with mAb 710 as the  
323 capture antibody used to coat wells, and HRP-conjugated mAb 711 used to detect bound  
324 antigen. Signal generation for the HRP reaction used TMB substrate (Thermo Scientific).

325 *Effects of medium and carbon source*

326 After washing in PBS, bacteria were resuspended in 6 ml of distilled water and split among 6  
327 bottles. 10 ml of media was added to the bottles. The media were Middlebrook 7H9 with 10%  
328 ADC and either 0.05% or 0.5% Tween 80; or Sauton's broth with 0.05% Tween 80, 0.5% BSA,  
329 and 0.05% tyloxapol, with 0.2% (w/v) acetate, glycerol, or dextrose (adapted from (45)), and  
330 bacteria were incubated for 24 h prior to preparation of culture filtrates.

331 *Preparation of lung homogenate supernatants from *M. tuberculosis*-infected mice*

332 C57BL/6 mice were infected by the aerosol route with ~60 CFU/mouse of *M. tuberculosis*  
333 H37Rv; lungs were harvested and single-cell suspensions were prepared as previously

334 described (46). After sampling the cell suspensions for determination of bacterial CFU, an  
335 aliquot of each sample was sterile-filtered, and Ag85B concentration determined by ELISA.

336 *Culture, infection, and analysis of bone marrow-derived dendritic cell death and release of*  
337 *Ag85B.*

338 Bone marrow-derived dendritic cells (BMDC), generated as previously described (20), were  
339 seeded in 96-well tissue culture treated plates (Corning) at  $2 \times 10^6$ /well and rested for 2 hours,  
340 then infected overnight with different multiplicities of infection (1, 2, 4 and 8) with *M.*  
341 *tuberculosis* H37Rv, treated with Amikacin (200  $\mu$ g/ml, 40min) in BMDC medium (RPMI1640  
342 supplemented with 10% heat-inactivated FBS, 2 mM L-glutamine, 1 mM sodium pyruvate, 1x  
343  $\beta$ -mercaptoethanol, 10 mM HEPES and 12 ng/ml recombinant mouse GM-CSF), washed three  
344 times in PBS and further cultured in fresh BMDC medium. CM was harvested 16, 24, 34 and  
345 48 hours later, sterile filtered and Ag85B in CM quantified by sandwich ELISA. At each harvest  
346 time point infected BMDC were assayed for cell death by CellTiter-Glo (Promega) according to  
347 the manufacturer's instructions and signal read as Luminescence with a Synergy H1 micro  
348 plate reader (BioTek). For each harvest time point signal from uninfected cells was considered  
349 as 100 % viability for determination of loss of viability of infected cells.

350 *Statistical analyses*

351 All statistical analyses were performed using Prism 7 (Graphpad). The specific tests used for  
352 data analysis are specified in the individual figure legends.

353 **Funding Information**

354 Supported by funds from the National Institutes of Health (R01 AI051242 and R01 AI124471;  
355 J.D.E.) and the Swiss National Science Foundation (P2BSP3-165349; M.B.).

356 **Acknowledgements**

357 We thank Smita Srivastava, Ph.D., for her contributions to the initial stages of this project.

358

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499

500 **Figure Legends**

501 **Figure 1. Recognition of Ag85A, Ag85B, and Ag85C by monoclonal antibodies 710, 711,  
502 and 712. Panels A-C:** Wells of 96-well plates were coated with Ag85B (panel A), Ag85A  
503 (panel B), or Ag85C (panel C), each at 0.5 µg/ml. Monoclonal Abs 710, 711, or 712 were  
504 added at the indicated concentrations, and binding was detected after washing, using  
505 horseradish peroxidase (HRP)-conjugated goat anti-mouse IgG and TMB substrate. Data  
506 shown in Panels A-C are representative of three independent experiments with one technical  
507 replicate per experiment per condition. **Panels D-F:** Individual wells were coated with a dilution  
508 series of Ag85B (panel D), Ag85A (panel E), or Ag85C (panel F), starting at 20 µg/ml.  
509 Monoclonal Abs 710, 711, or 712 were added at a concentration of 1 µg/ml, and binding was  
510 detected after washing, using horseradish peroxidase (HRP)-conjugated goat anti-mouse IgG  
511 and TMB substrate. Data in panels D-F are representative of two independent experiments  
512 with one technical replicate per experiment per condition.

513 **Figure 2. Characterization of sandwich ELISA for Ag85B.** Each mAb was tested as the  
514 capture antibody, with either of the remaining two mAbs (conjugated to HRP) used for  
515 detection. A) Detection of rAg85B added to mAb-coated plates at the concentrations indicated  
516 on the X axis. B) Detection of rAg85A under the same conditions as in panel A. The inset  
517 shows the same data on a contracted scale, to reveal small differences. C) Detection of  
518 rAg85C under the same conditions as in panel A. The inset shows the same data on a  
519 contracted scale, to reveal small differences. D) Specificity of ELISA using mAb 710 for  
520 capture and HRP-conjugated mAb 711 for detection of antigen in culture filtrates of wild-type  
521 *M. tuberculosis* H37Rv or *M. tuberculosis* H37Rv with a targeted deletion of the gene encoding  
522 Ag85B. The H37Rv culture filtrate was diluted to give a signal in the linear range of the ELISA  
523 standard curve, while the H37RvΔAg85B culture filtrate was assayed undiluted. The data  
524 shown are mean (bar) and standard deviation (error bar) of biological triplicate values. E)

525 Standard curves of rAg85B in Ag85B ELISA. Shown are curves of three independent  
526 experiments with two independent standard curves each. Shown are mean (dot) and standard  
527 deviation (error bar) when large enough to depict.  
528 Data shown in Panels A-C are representative of three independent experiments with one  
529 technical replicate per experiment per condition.

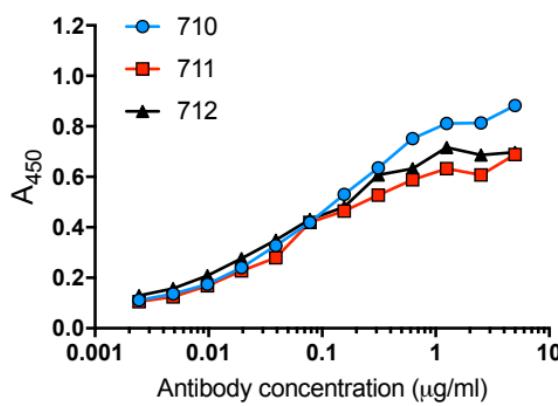
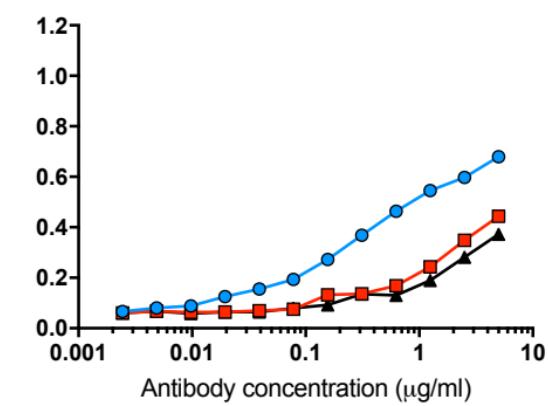
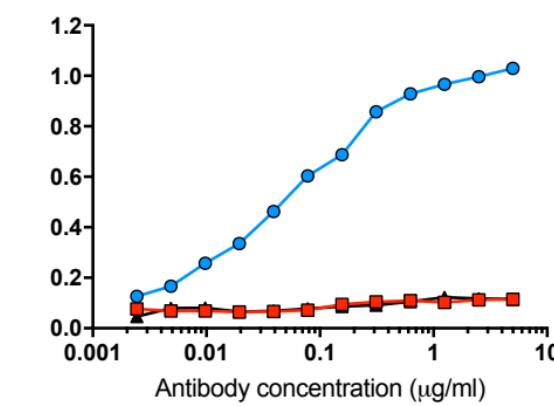
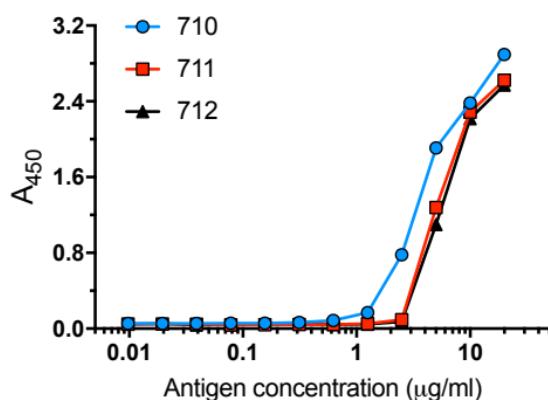
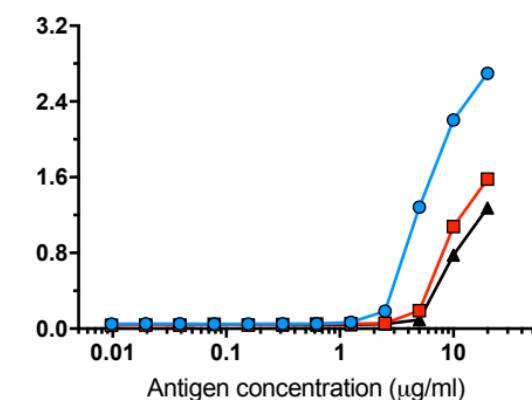
530 **Figure 3. Effects of carbon source and *M. tuberculosis* strain, and kinetics of Ag85B  
531 secretion.** A) Effects of media composition and carbon source on Ag85B secretion. Mid-log  
532 phase bacteria ( $A_{600} = 1.0$ ) grown in 7H9 medium were washed 5 times, then diluted 10-fold  
533 into the indicated media. After 24 h, the  $A_{600}$  was determined for each culture, and sterile  
534 filtrates were prepared for analysis by mAb 710-711 ELISA. Results of Ag85B quantitation by  
535 ELISA were normalized by the  $A_{600}$  of the individual culture. 7H9 0.05% T80: Middlebrook 7H9  
536 with 10% ADC and 0.05% (v/v) Tween 80; 7H9 0.5% T80: same as preceding, but with 0.5%  
537 Tween 80; Sau: Sauton's medium with 0.05% Tween 80; Ace: 7H9 media with acetate; Dex:  
538 7H9 media with dextrose; Gly: 7H9 media with glycerol. Show are mean (bar) and standard  
539 deviation (error bar) of biological triplicate values. Statistical comparison was done using one-  
540 way ANOVA; the adjusted p value after Dunnett's post-test for multiple comparisons was  
541 applied is shown for the effect of dextrose. Other effects were not significant after adjusting for  
542 multiple comparisons. B) Mycobacterial strain-dependent variation of Ag85B secretion. Mid-log  
543 phase cultures were collected and the bacteria from each strain were washed and  
544 resuspended in fresh 7H9 broth. After 24h, the bacteria were pelleted by centrifugation and  
545 Ag85B in culture filtrates was quantitated by ELISA. Show are mean (bar) and standard  
546 deviation (error bar) of biological triplicate values. The adjusted p values shown are for  
547 comparison of each strain with H37Rv, and were determined by one-way ANOVA with  
548 Dunnett's post-test. C) Kinetics of Ag85B secretion by *M. tuberculosis* H37Rv. Mid-log phase  
549 growing H37Rv was washed, added to fresh 7H9 medium at  $1 \times 10^8$  CFU/ml, and incubated for

550 24 h. Individual cultures were sampled at the designated time points and culture filtrates were  
551 assayed by ELISA. Statistical analysis by linear regression was determined using Prism 7.  
552 Dashed lines indicate 95% confidence interval. Figure A shows one representative of three  
553 independent experiments. Figures B and C show data of experiments each done once.

554 **Figure 4. Quantitation of Ag85B in supernatants of lung homogenates from mice**  
555 **infected with *M. tuberculosis* H37Rv.** A) *M. tuberculosis* colony-forming units (CFU) in the  
556 lung samples used for assays in panel B. B) Ag85B concentrations quantitated by ELISA in  
557 lung homogenate supernatants; C) Ag85B concentrations as in Panel B, normalized by the  
558 number of bacteria (CFU) in the same lung homogenates. Shown are individual data points,  
559 mean  $\pm$  standard deviation. This experiment was done once.

560 **Figure 5. Death of *M. tuberculosis*-infected primary dendritic cells releases Ag85B.** Bone  
561 marrow-derived dendritic cells were infected with *M. tuberculosis* H37Rv at different  
562 multiplicities of infection (1, 2, 4 and 8). At designated time points (12 h grey dots, 16 h  
563 medium blue dots, 24 h red dots, 36 h light blue dots, 48h dark blue dots), medium was  
564 removed for quantitation by ELISA, and dendritic cell viability was assessed by luminescence  
565 assay of ATP of cell lysates. Results of the two assays on a given sample were plotted as a  
566 data point, and Pearson correlation was determined for the whole dataset. Shown are  
567 representative results of two similar experiments.

568

**A****Ag85B****B****Ag85A****C****Ag85C****D****Ag85B****E****Ag85A****F****Ag85C**