

Opsonic Monoclonal Antibodies Directed Against MTB Enhance Blood Clearance in a Quantitative qPCR Mouse Model

Clara J. Sei^{1*}, Luke T. Daum², Richard F. Schuman³, Afia Mesadieu¹, Nimisha Rikhi¹, Gerald W. Fischer¹

¹Longhorn Vaccines and Diagnostics, Gaithersburg, Maryland, USA ²Longhorn Vaccines and Diagnostics, San Antonio, Texas, USA ³Antibody and Immunoassay Consultants, Rockville, Maryland, USA

BACKGROUND

In sub-Saharan Africa, Mycobacterium tuberculosis (MTB) is an important cause of bacteremia and sepsis. Patients with HIV and MTB bacteremia have a high mortality with many deaths occurring within 18 days of presentation. Therefore, rapid diagnosis of MTB bacteremia is critical and can be achieved using real-time quantitative PCR (qPCR). Treatment of MTB sepsis with immunotherapeutic agents may enhance clearance of MTB bacteremia and augment therapy for MDR and XDR TB.

METHODS

Monocional Antibody (MAB) Identity and Characterization: Hybridoma supernatant and purified MABs Go9, AB3 and IG7 were screened by EUSA for their binding activity to several MTB strains (Erdman, HN878, CDC1551) and *M. smegmatis*. The latter nonpathogenic strain was used as a surrogate *Mycobacteria* strain in the opsonophagocytic assay.

Opsonophagocytic Killing Activity (OPKA): The functional activity of MABs GG9, AB9 and JG7 (@25--0.06ug/mL) was evaluated using Human Leukemia Promyelocytic cell line (HL50) and Human Histiccytic Lymphoma cell line (U-937), against Mycobacterium smegmatis. MABs were reacted with differentiated effector cells in the presence (for HL60s) or absence (for U-937s) of complement component C1q and M. smegmatis. OPKA was defined as the percentage of the average CFU counts in test sample wells (with MAB) divided by average control CFU counts (without MAB). When bacterial CFU was reduced by greater than 50%, OPKA was considered antibody enhanced².

MTB Challenge/MAB Treatment: Female ICR mice were given opsonic anti-MTB MAB GGG intraperitoneally at docs levels 10mg/kg, 5mg/kg, 1mg/kg, and placebo (P85) twenty-four hours prior to MTB HN878 challenge. Challenge doses were at 10^5 and 10°8 CFU/mL, administered intravenously. Whole blood specimens were collected at 0, 4, and 24 hours post-challenge, placed in K2-EDTA and then 0.1 mL added into 1 mL PrimeStore MTM* prior to shipping at ambient temperature from Gaithersburg, Maryland to San Antonio, Texas for quantitative qeCR testing.

DNA Extraction/qPCR: Total nucleic acid (DNA) was extracted from 0.2mL murine blood specimen using PrimeXtract[™]. Real-time qPCR was performed by adding 2.5 µL of specimen into 7.5uL of PrimeMix[®] MTB Multiplex (IS6110/151081) on an ABI 7500.

MTB Blood Clearance: The level of MTB in murine blood was monitored by qPCR and total clearance was defined as cycle threshold (Cr) = 40 for both IS-6110 and IS-1081 targets. Group average C_T values, standard deviation and percentage of mice with total clearance was recorded.

RESULTS

MABs GG9, AB9 and JG7 bind to killed *Mycobacterial tuberculosis* strains (Erdman, HN878, and CDC1551; Figure 1) and live *M. smegmatis* (Figure 2) as screened using ELISA.

REFERENCES

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² Fleck RA, Romero-Steiner S and Nahm MH. (2005) Use of HL-60 Cell Line to Measure Opsonic Capacity of Pneumococcal Antibodies. CLIN DIAGN LAB IMMUN, p. 19-27.





MABs GG9, AB9 and JG7 have shown OPKA > 50% (across high and low concentrations) using both granulocytes and macrophages. Peak OPKA for GG9 was 67% (lug/mL); AB9, 75% (25ug/mL); and JG7, 81% (0.06ug/mL) using differentiated HLG0 cells (Figure 3). With U-937s, MABs GG9 and AB9 had peak OPKA of 55% at 1.5ug/mL (data not shown).



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Female ICR mice given MAB GG9 at doses 10mg/kg and 5mg/kg had enhanced blood clearance of killed HM878 bacilli by twenty-four hours post-challenge with both a low (10^5 CFU/mL) and high (10^8 CFU/mL) MTB dose compared to the placebo (PBS) groups (Figure 4). Treatment with GG9 at 5mg/kg showed total clearance ($c_1 = 40$ for (S5110) in 7/9 mice.



At twenty-four hours post MTB challenge, a steady decline in HN878 bacilli with decreased C_{τ} values was noted in MAB treated mice compared to PBS control (Figure 5). Duplicate averages with standard error are indicated.



CONCLUSIONS

- ✓ Anti-MTB MABs GG9, AB9 and JG7 were identified that bind to MTB strains and promote mycobacterial OPKA.
- ✓ Using qPCR to detect and monitor killed MTB in murine blood is an efficient methodology to identify MABs that enhance clearance of MTB.
- Anti-MTB MABs, like GG9, may enhance clearance of MTB bacteremia and might provide useful adjunctive therapy for MTB sepsis.
- Anti-MTB MABs may also be useful for therapy in patients with MDR and XDR TB.
- Current studies are being conducted using live MTB strains to further identify potential therapeutic MAB candidates.