OHet72 and Current Tuberculosis Treatments: Identifying Synergy Margaret Bourlon and Lucila Garcia-Contreras Department of Pharmaceutical Sciences, University of Oklahoma Health Sciences Center

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BACKGROUND

- •Tuberculosis (TB) is the second leading cause of death from a single infectious agent, after COVID-19.¹
- •Drug susceptible TB is treated with oral administration of 4 drugs for 6-9 months, whereas multi-drug resistant TB requires 5-10 drugs administered for 18-24 months (figure 1). Both regimens cause severe adverse effects.



•OHet72 is a novel anti-cancer compound³ that also has potent anti-TB activity in vitro with no apparent side effects

METHODS

- •The Microtiter alamar blue assay (MABA)⁴ was used to determine the minimum inhibitory concentration (MIC) of OHet72, first line TB drugs (isoniazid, rifampicin, pyrazinamide), and second line TB drugs (levofloxacin, moxifloxacin).
- •Each drug was diluted, and bacteria was added. MABA reagent was added after 1 week. The %reduction of reagent was calculated, and the MIC was considered the lowest concentration of drug that resulted in no bacterial growth.



Figure 2: Microtiter alamar blue assay (MABA) method

•Drug interactions were investigated using a MABA checkerboard assay⁵. Two drugs are diluted in different orientations and drug interactions are quantified by calculating the fractional inhibitory concentration index (FICI)⁶



MIC B alone

Figure 3: Checkerboard MABA assay template (left), representation of results (right), and fractional inhibitory concentration index (FICI) equation (bottom)

MIC A alone

Aim

Identify the type of drug interaction (synergy, addition, or antagonism) that may exist between OHet72 and five drugs currently used against Mycobacterium tuberculosis (MTB).

RESULTS

Table 1: Anti-TB drugs, their experimental MIC determined by the MABA assay against H37Ra MTB, and the reported literature values⁷.

Drug	Experimental MIC (µg/mL)	Literature MIC (µg/mL)
Isoniazid	0.1149 ± 0.046	0.1 - 0.2
Rifampicin	0.002 ± 0.00	0.004 – 0.032
Pyrazinamide	240.0 ± 83.56	16 – 512
Levofloxacin	0.3125 ± 0.00	0.125 – 0.5
Moxifloxacin	0.078 ± 0.00	0.032 – 8.0
OHet72	0.929 ± 0.5223	



Tukey's multiple comparisons test

- [Drug B] when administered with drug interactions levofloxacin, & moxifloxacin • OHet72 may target MTB cell wall

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Discussion

•Synergy describes the phenomenon where two or more drugs in combination have a therapeutic effect greater than the sum of the effects of the individual drugs⁸

•The actual drug concentrations of OHet72 and rifampicin in combination that are required to elicit a given therapeutic effect are lower than the expected concentrations based on the effects when the drugs are administered alone



Figure 5: (A) Graphical representation of synergistic (green lines) and antagonistic (orange/red lines) drug pairings compared to and additive reference (blue line); and (B) an isobologram of OHet72 and rifampicin pairing showing drug combinations required to reach specified therapeutic effects in an additive model (solid lines) and the actual concentrations of the drug required to elicit the same effects (dashed lines)

CONCLUSION

• OHet72 would be most effective rifampicin due to the synergistic • OHet72 has additive interactions with isoniazid, pyrazinamide,



Figure 6: Hypothesis of OHet72 mycobacterial target

REFERENCES

S., Weerasekare, G. M., Gale, J. B., Patterson, M. K., Jr, Wang, B., Wang, Rowland, T. C., DiSivestro, P., Lindamood, C., 3rd, Hill, D. L., & Berlin, K. D. Biologically active heteroarotinoids exhibiting anticancer activity and decreased

1. Global tuberculosis report 2022. Geneva: World Health Organization; 2022. Licence: CC 5. Hsieh, M. H., Yu, C. M., Yu, V. L., & Chow, J. W. (1993). Synergy assessed by checkerboard. A critical analysis. Diagnostic microbiology and infectious disease, 16(4), 6. Gómara, M., & Ramón-García, S. (2019). The FICI paradigm: Correcting flaws in antimicrobial in vitro synergy screens at their inception. Biochemical pharmacology, 163, Heinrichs, M. T., May, R. J., Heider, F., Reimers, T., B Sy, S. K., Peloguin, C. A., 8 Derendorf, H. (2018). Mycobacterium tuberculosis Strains H37ra and H37ry have equivalent minimum inhibitory concentrations to most antituberculosis drugs. International journal of mycobacteriology, 7(2), 156-161. . Dartois, V. A., & Rubin, E. J. (2022). Anti-tuberculosis treatment strategies and drug

development: challenges and priorities. Nature reviews. Microbiology, 20(11), 685–701

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