



Oxazolidinone Antibiotics Reversibly Inhibit Mitochondrial Metabolism in Human Cell Lines

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Introduction	Results		
• Oxazolidinone antibiotics inhibit bacterial protein synthesis but also mitochondrial DNA-encoded protein synthesis (MEPS) in eukaryotic cells [1], which has been associated with the development of severe adverse effects (myelosuppression [2], lactic acidosis [3] and neuropathies [4]) upon prolonged treatments.	1. Inhibition of CYTOX I expression (A) and cytochrome <i>c</i> -oxidase activity (B)	3. Recovery of CYTOX I expression (A), cytochrome <i>c</i> -oxidase activity (B), and of mitochondrial respiration (C) in HL-60 cells	
	HL-60 THP-1	mitochondrial respiration (C) in FIL-60 cens	
	A. CYTOX I expression $\begin{array}{c} \downarrow \\ 125 \\ \hline \\ $	A. CYTOX I expression	

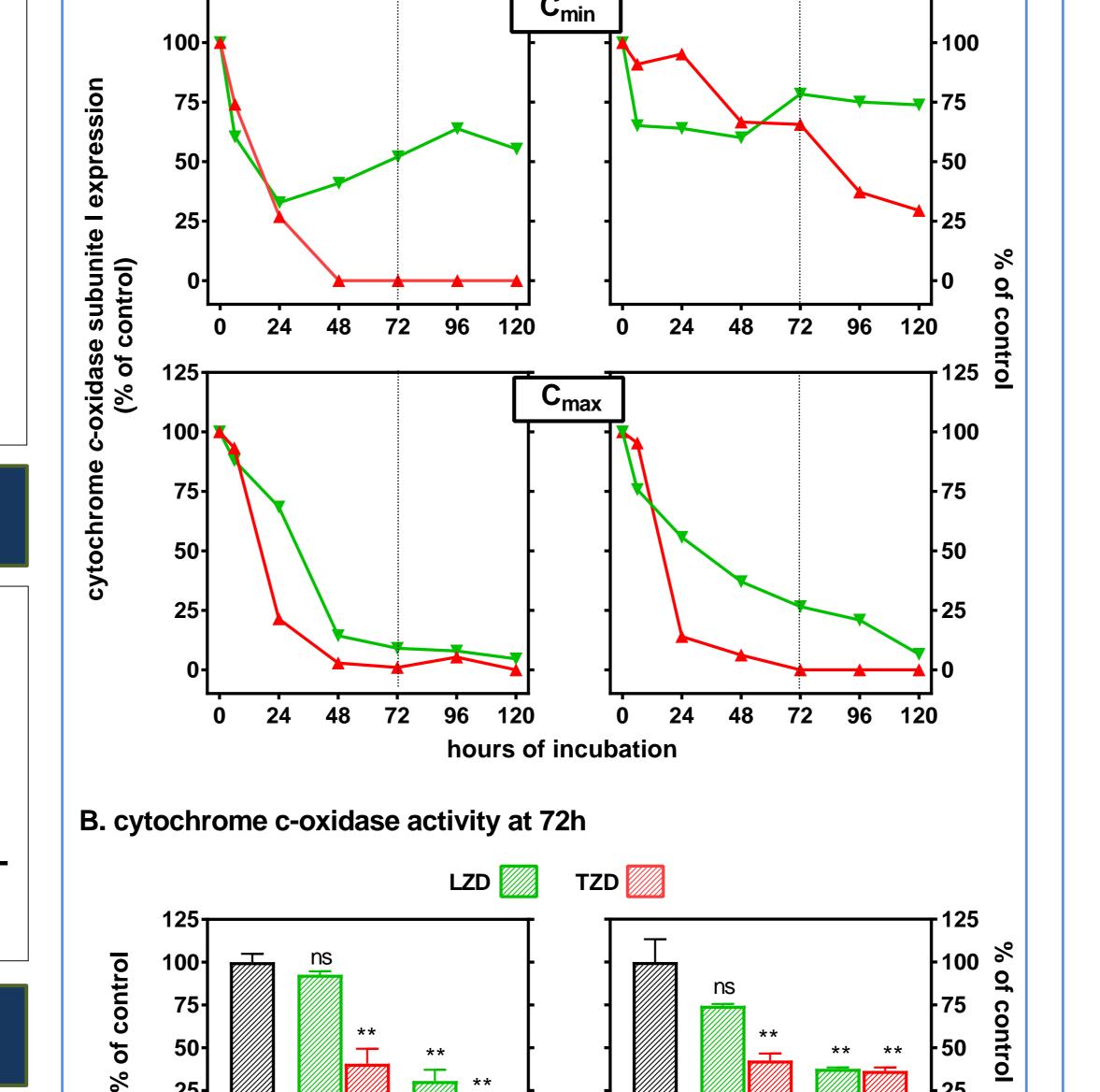
- Linezolid (LZD; approved in 2000) and tedizolid (TZD; approved by FDA in 2014 and EMA in 2015) are the two oxazolidinones currently on the market.
- TZD shows lowers minimal inhibitory concentrations (MICs) towards susceptible bacteria than LZD due to increasing binding to bacterial ribosomes [5] and is also a more potent inhibitor of MEPS [6].

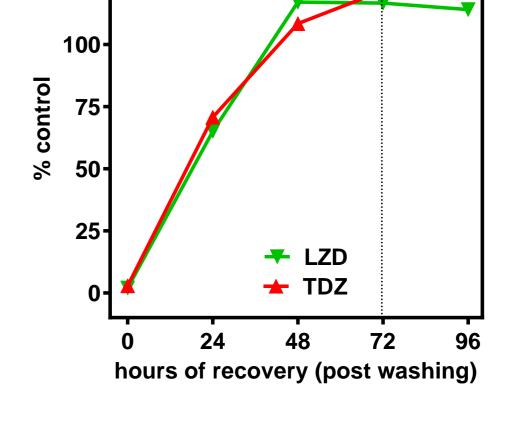
Aims of the study

- To compare the inhibitory potential of TZD and LZD towards the expression of a key protein encoded by the mitochondrial genome
- To assess its impact on mitochondrial respiration and metabolism in cultured human cells exposed to clinicallyrelevant concentrations of these drugs.

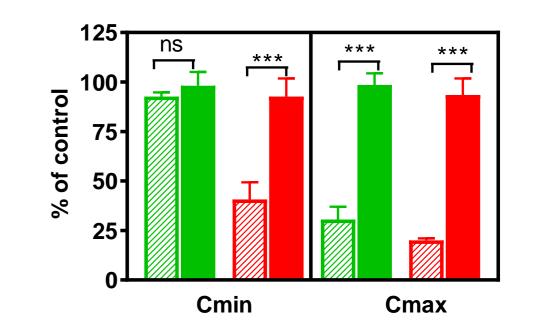
Methods

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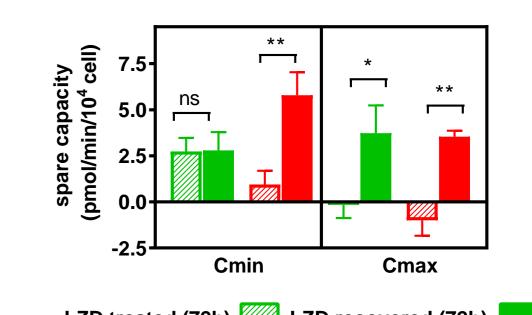




B. cytochrome *c*-oxidase activity



C. mitochondrial respiration



- Cells: Human promyelocytes (HL-60) and monocytes (THP-1)
- *Mitochondrial protein expression*: western blot of cytochrome *c*oxidase subunit I (CYTOX I) and succinate dehydrogenase (SDH), both encoded by mitochondrial and nuclear genome, with normalization using Tom 20 (outer membrane protein).
- Basal mitochondrial oxygen consumption rate (OCR) and reverse *capacity*: Seahorse XF96 bioanalyzer.
- Cytochrome c-oxidase activity: decrease of OD₅₅₀ of cytochrome c.
- Autophagy: increase in level of lipidated protein LC3II (Western blot, normalized to actin) in the presence of leupeptin (cathepsin B inhibitor).

Treatments:

• Incubation with 2.5 or 15 mg/L for LZD and 0.5 or 3 mg/L for TZD (total concentration, corresponding to C_{min} and C_{max} in patients receiving approved daily dosages (LZD, 600 mg BID; TZD, 200 mg qD)

References

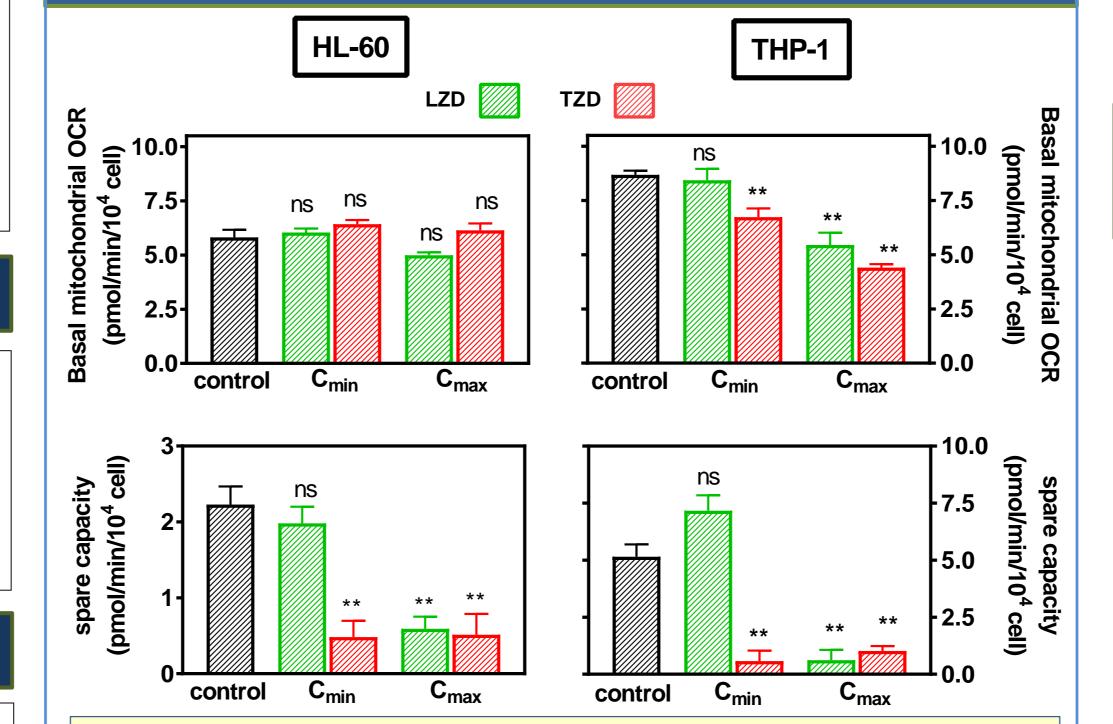
(1) Nagiec et al, Antimicrob Agents Chemother 49, 3896-902 (2005) – PMID 16127068 (2) Hachem *et al*, Clin Infect Dis 37, e8-11 (2003) – PMID 12830431 (3) Apodaca et al, N Engl J Med 348, 86-87 (2003) - PMID 12510056 (4) Bressler et al, Lancet Infect Dis 4, 528-531 (2004) - PMID 15288827 (5) Im et al, Eur J Med Chem 46, 1027-1039 (2011) – PMID 21292356 (6) Flanagan et al, Antimicrob Agents Chemother 59, 178-185 (2015) – PMID 25331703

Acknowledgments and Conflicts of Interest



- TZD is a more potent inhibitor of the expression of CYTOX I than LZD at both C_{min} and C_{max} (the expression of SDH was not impaired at either concentrations.
- Cytochrome c-oxidase enzyme activity (at 72h) was significantly decreased in both cell lines with TZD being globally more inhibitory than LZD.

Changes in mitochondrial metabolism (basal respiration and reserve capacity)





• CYTOX I expression, cytochrome *c*-oxidase activity, and mitochondrial respiration were all three recovered upon the drug withdrawal.

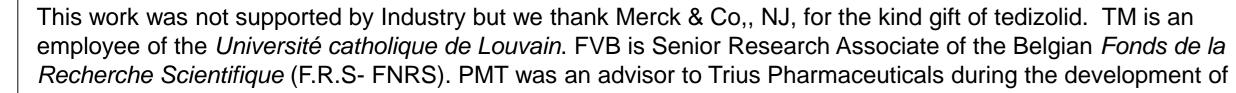
4. Autophagy (not illustrated)

• No increase in LC3II over leupeptin control was observed after 48 h of incubation with either TZD or LZD at C_{max} .

Conclusion

Oxazolidinones cause mitochondrial metabolic dysfunction probably due to the impairment of the expression of proteins encoded by mitochondrial genome, which may explain the development of the severe side effects associated with their use.

In a biological context, oxazolidinones may stand as a useful tool to explore the metabolic and functional consequences of impaired



tedizolid and is a current speaker for Bayer in connection with tedizolid.

