

Large Extracellular Cord Formation in a Zebrafish Model of Mycobacterium kansasii Infection

Johansen Matt, Kremer Laurent

► To cite this version:

Johansen Matt, Kremer Laurent. Large Extracellular Cord Formation in a Zebrafish Model of Mycobacterium kansasii Infection. Journal of Infectious Diseases, 2020, pp.jiaa187. 10.1093/infdis/jiaa187. inserm-02868836

HAL Id: inserm-02868836 https://inserm.hal.science/inserm-02868836

Submitted on 15 Jun2020

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

1	A zebrafish model of Mycobacterium kansasii infection reveals large
2	extracellular cord formation
3	
4	Matt D. Johansen ¹ and Laurent Kremer ^{1, 2 #}
5	
6	¹ Institut de Recherche en Infectiologie de Montpellier, Centre National de la Recherche
7	Scientifique UMR 9004, Université de Montpellier, Montpellier, France
8	² INSERM, Institut de Recherche en Infectiologie de Montpellier, Montpellier, France
9	
10	[#] To whom correspondence should be addressed:
11	Tel: (+33) 4 34 35 94 47; E-mail: laurent.kremer@irim.cnrs.fr
12	
13	
14	
15	
16	Running title: Mycobacterium kansasii cords in vivo

17 Abstract (100 words)

Mycobacterium kansasii is a slow-growing non-tuberculous mycobacteria responsible for co-infections particularly in patients with human immunodeficiency virus. To date, our knowledge of *M. kansasii* infection has been hampered due to the lack of an effective animal model to study pathogenesis. Here, we show the zebrafish embryo is permissive to M. kansasii infection, causing a chronic infection and forming granulomas. Upon macrophage depletion, we identified *M. kansasii* forms extracellular cords, resulting in acute infection and rapid larval death. These findings highlight the feasibility of zebrafish to study *M. kansasii* pathogenesis, and for the first time identify extracellular cords in this species. Keywords: Mycobacterium kansasii, granuloma, infection, cords, zebrafish, macrophage

39 Background (2000 words)

Mycobacterium kansasii is a slow-growing photochromogenic non-tuberculous mycobacteria 40 (NTM). This opportunistic pathogen is a common source of severe pulmonary infections in 41 patients with human immunodeficiency virus (HIV), chronic obstructive pulmonary disorder 42 and generalised immunosuppression [1-3]. It is regarded as one of the phylogenetically closest 43 44 species to *Mycobacterium tuberculosis* and is clinically indistinguishable based on clinical symptoms and radiographic features [2]. Previous research has reported >50% mortality rates 45 in patients with HIV and *M. kansasii* co-infections [1], representing a significant public health 46 47 burden in at-risk populations.

Despite the significance of *M. kansasii* infections in specific patient subpopulations, 48 there has been very little research exploring pathogenicity of this species. Previous reports 49 50 demonstrated that type I M. kansasii ATCC 12478 is able to rupture the phagosomal membrane similarly to *M. tuberculosis* [4], yet it is only able to cause a persistent non-replicating infection 51 in a murine model [2]. As such, there is an urgent need to explore alternative animal models to 52 study *M. kansasii* pathogenesis. In recent years, the zebrafish has emerged as an important 53 infection model to investigate mycobacterial infections, recapitulating important aspects of the 54 55 human mycobacterial infection, such as granuloma formation [5-7]. Advancements in genetics have allowed the generation of fluorescently labelled cellular subsets and zebrafish knockout 56 lineages, allowing us to explore specific determinants of host immunity [8]. In particular, the 57 zebrafish embryo is optically transparent and possesses a functional innate immune response 58 from approximately 30 hours post-fertilisation, allowing the visualisation of early host-59 pathogen interactions at the cellular level [7, 8]. Therefore, we explored whether zebrafish 60 61 embryos could be used as a model for *M. kansasii* infection and examined initial host-pathogen interactions. 62

63 Methods

64 Bacterial culture and generation of fluorescent M. kansasii

M. kansasii ATCC 12478 and *M. marinum* M strain (ATCC BAA-535) were grown in Middlebrook 7H9 broth (BD Difco) and supplemented with 0.025% Tyloxapol and 10% oleic acid, albumin, dextrose, catalase (OADC enrichment) (7H9^{OADC/T}) or on Middlebrook 7H10 agar (BD Difco) containing 10% OADC enrichment (7H10^{OADC}). Bacteria were cultured at 37°C unless otherwise indicated. Antibiotics were included where required for fluorescent strains.

Fluorescent *M. kansasii* was generated using pTEC15 and pTEC27 (a gift from Lalita Ramakrishnan) as previously described [9], however 7H9^{OADC/T} was used instead of Sauton's and all incubation and wash steps were completed at 37°C and not 4°C. For selection of fluorescent colonies, *M. kansasii* was plated on 7H10^{OADC} containing 50 μ g/mL hygromycin (Euromedex). Positive colonies were selected based on fluorescence and maintained in 7H9^{OADC/T} supplemented with 50 μ g/mL hygromycin. Fluorescent *M. marinum* M strain has been previously described [10].

78

79 Zebrafish maintenance and infection

Zebrafish experiments were completed in accordance with the Comité d'Ethique pour l'Expérimentation Animale de la Région Languedoc Roussillon under the reference CEEALR36-1145. Experiments were performed using the *golden* mutant and macrophage reporter Tg(*mpeg1:mCherry*) line, as previously described [6]. Single cell bacterial preparations were completed as previously described [6] and used immediately and not from frozen stocks. Zebrafish caudal vein injections and lipoclodronate macrophage depletion were conducted, as reported [6]. Following infection, embryo age is expressed as days post-infection (dpi). Embryo survival was monitored daily based on the presence of a heartbeat. For live
imaging, zebrafish embryos were anaesthetised in 0.02% tricaine solution [6, 10], mounted on
3% (w/v) methylcellulose and imaged using a Zeiss Axio Zoom.V16 coupled with an Axiocam
503 mono (Zeiss). Fluorescence Pixel Count (FPC) measurements were determined using the
'Analyse particles' function in ImageJ and normalised for each experiment against the
corresponding control. All experiments were completed at least three times independently.

93

94 Statistical analysis

Survival curve analysis was completed using the log-rank (Mantel-Cox) statistical test. Cord,
granuloma and bacterial burden (FPC) analysis were completed using unpaired Student's *t*test. All statistical tests were completed using Graphpad Prism (Version 8.0.1).

98

99 **Results**

100 Mycobacterium kansasii infection causes a chronic infection in zebrafish embryos

A critical characteristic of zebrafish manipulation is the requirement for lower ambient temperatures of 28.5°C. As such, we first determined whether *M. kansasii* can replicate at lower temperatures compared to traditional 37°C incubation. We found that *M. kansasii* is able to grow and replicate at 30°C, albeit slower than at 37°C which was to be expected as for other

105 NTM species [6] (Supplementary Figure 1).

Previous research has demonstrated that the zebrafish embryo is an attractive model organism for mycobacterial pathogens such as *M. marinum* and *M. abscessus* and recapitulating important pathophysiological hallmarks of human mycobacterial infection and leading to larval death [6, 7]. With this in mind, we wanted to determine whether *M. kansasii* was able to promote larval death in a dose-dependent manner. We found that *M. kansasii* resulted in ≤10% embryo mortality in a 12-day period, irrespective of infectious dose.
Comparatively, *M. marinum*, which is an important fish and human pathogen, led to 100%
embryo mortality within 9 dpi (Figure 1A), as previously reported [7].

We next wanted to explore whether M. kansasii ATCC 12478 is rapidly cleared 114 following infection, or whether it causes a persistent infection. Using live whole embryo 115 116 fluorescent imaging, we observed the formation of granulomas in 50-60% of embryos from 3 dpi onwards (Figure 1B-D). Moreover, when we examined the average number of granulomas 117 per infected embryo over time, we observed a trend for the increasing number of granulomas 118 between 3 and 9 dpi, however this was not significant (P = 0.06) (Supplementary figure 2). 119 120 Overall, these results highlight that the zebrafish embryo is permissive to *M. kansasii* infection, 121 leading to a chronic infection characterised by granuloma formation.

122

Mycobacterium kansasii is able to form large extracellular cords and resist phagocytosis 123 Macrophages are important immune cells recruited to an infection and are the predominant cell 124 subset within the mycobacterial granuloma [7, 11]. Previous work showed that macrophage 125 ablation results in rapid larval death within several days of infection, underpinning the crucial 126 127 role of these phagocytes in containing mycobacterial infections [6, 12]. Thus, we explored whether macrophages are responsible for *M. kansasii* infection control. Following liposomal 128 129 clodronate treatment, which depletes macrophages until approximately 6 days post-treatment 130 (Supplementary figure 3), we observed progressive larval death resulting in~70% of dead embryos at 12 dpi as compared to standard M. kansasii infection or following liposomal PBS 131 injection prior to infection (Figure 2A). When we examined the bacterial burden in clodronate-132 133 treated embryos, there was a significant increase from 2 dpi onwards when compared to wild-134 type infections (Figure 2B).

135 Using whole embryo live fluorescent imaging in clodronate-treated embryos, we identified the presence of large extracellular cords in 20% of embryos from 3 dpi, which 136 increased to >80% by 9 dpi (Figure 2C). Comparatively, extracellular cord formation in wild-137 type embryos was a rare event, occurring in <10% of infected embryos and only at 9 dpi 138 (Figure 2C). When we further categorised the proportion of cords per embryo, we identified a 139 progressive increase in the number of moderate (6-10 cord/embryo) and heavy (>10 140 cords/embryo) number of cords up to 9 dpi, as compared to 3 dpi in which 100% of corded 141 embryos had a low number of cords (1-5 cords/embryo) (Figure 2C). Following macrophage 142 143 repopulation at 6 days post clodronate-treatment, we observed the gradual recruitment of macrophages to extracellular cords at approximately 9 dpi (Figure 2D). However, based on the 144 sheer size of these observed cords in comparison to the size of the recruited macrophages, they 145 146 were unable to degrade or control cord formation, in agreement with the increase in the number 147 of embryos with cords and the average number of cords (Figure 2C). Finally, to determine if cord formation is an acquired in vivo phenotype of M. kansasii, we examined stationary-phase 148 149 liquid cultures for the presence of cords. Using fluorescent microscopy, we observed the presence of large bacterial aggregates resembling cords, demonstrating that this is an intrinsic 150 *M. kansasii* phenotype (Supplementary Figure 4). 151

152 Collectively, these findings highlight the critical role of macrophages in the control of 153 *M. kansasii* infection and preventing the formation of large extracellular cords. Importantly, to 154 our knowledge this is the first reported observation of *M. kansasii* cord formation both *in vitro* 155 and *in vivo*, which has significant implications for our understanding of *M. kansasii* 156 pathogenesis.

157

158

160 **Discussion**

In this study, we report for the first time the applicability of the zebrafish embryo to study the pathogenesis of *M. kansasii*. Using this model organism, we demonstrate that under standard infection conditions, a type I *M. kansasii* strain causes a persistent and replicating infection typified by the presence of mycobacterial granulomas. Most importantly, we report here that *M. kansasii* forms large extracellular cords, representing a significant pathogenic trait of this NTM that was previously unexplored.

In recent years, the zebrafish model has established itself as a revolutionary platform to 167 examine host-pathogen interactions in real-time [7, 8]. Using this to our advantage in the 168 current study, we were able to monitor the kinetics of granuloma formation following M. 169 170 kansasii infection which revealed a chronic in vivo phenotype, representative of other slow-171 growing mycobacteria, including *M. tuberculosis*. It is notable that the zebrafish embryo does not possess adaptive immunity until later developmental stages and as such, the role of adaptive 172 immunity in *M. kansasii* remains unexplored. However, adult zebrafish have fully functional 173 innate and adaptive immunity and may be a feasible model organism to examine the role of 174 complete immunity of *M. kansasii* infection in subsequent studies. 175

Cord formation is a pathophysiological hallmark of acute mycobacterial disease [6, 13-176 15]. Following the successful escape from the phagosome, mycobacteria such as *M. abscessus*, 177 M. marinum and M. tuberculosis [4], are able to grow extracellularly as large cords which 178 vastly exceed the size of recruited immune cells and ultimately resist phagocytosis [6]. To date, 179 this phenotypic characteristic has only been identified within a few pathogenic mycobacteria 180 181 species and has often been reserved for those that have been considered most virulent. We 182 report here, the formation of large extracellular cords in *M. kansasii* both in vitro and in vivo. That *M. kansasii* shares this phenotypic trait with *M. tuberculosis* further supports the recent 183

184 view that M. kansasii is an environmental ancestor of M. tuberculosis and can be used as a model organism to investigate the transition from an environmental opportunistic pathogen to 185 a host-restricted pathogen [2]. Following macrophage depletion, we observed a large increase 186 187 in the presence and abundance of cords, highlighting the importance of macrophages in M. kansasii infection. Severe infections with M. kansasii are often associated with co-existing 188 infections such as HIV, suggesting that immunodeficiency is an important risk factor in the 189 190 severity of *M. kansasii* infection [3]. Consequently, our observations of extensive cord formation following macrophage ablation may be viewed as clinically relevant and may 191 192 provide an explanation for the higher mortality rate associated with M. kansasii co-infection in immunocompromised patients. As this is the first report of cord formation in *M. kansasii*, it is 193 important that subsequent studies examine the role of cord formation in clinical isolates and 194 195 screen patient sputum for the presence of M. kansasii extracellular cords to support these 196 observations.

While cording in wild-type embryos can be considered a rare event in the context of the 197 current study, it is not entirely absent suggesting that *M. kansasii* can eventually escape from 198 the phagosome. This observation fits with previous work which has identified that M. kansasii 199 200 is able to rupture the phagosomal membrane, indicative of phagosomal escape [2, 4]. Moreover, 201 M. kansasii has been demonstrated to secrete esxA and esxB; two effector proteins of the well-202 characterised ESX-1 locus, which are known to assist in phagosomal escape for M. marinum 203 and M. tuberculosis [2]. However, to date, the role of ESX-1 in M. kansasii pathogenesis remains to be explored. 204

In summary, our findings highlight the applicability of the zebrafish embryo to examine host-pathogen interactions and pathogenesis for the study of *M. kansasii*. Furthermore, using our new animal model, we have demonstrated that *M. kansasii* causes a chronic infection which is largely controlled by patrolling macrophages in a wild-type infection. Most importantly, we 209 have shown that *M. kansasii* forms large cords both *in vitro* and *in vivo*, ultimately representing

a previously unexplored virulence trait of this clinically significant NTM species.

211

212 Funding

MDJ received a post-doctoral fellowship granted by Labex EpiGenMed, an «Investissements
d'avenir» program (ANR-10-LABX-12-01).

215

216 **Conflicts of interest**

217 The authors have no conflict of interest to declare.

218

219 **Figure legends**

Figure 1: Mycobacterium kansasii induces a chronic infection in zebrafish. (A) Zebrafish 220 221 embryos at approximately 30 hours post-fertilisation were infected with increasing doses of M. kansasii ATCC 12478, or 100 CFU of M. marinum M strain via caudal vein injection and 222 survival was monitored over a 12-day period. Each group consists of approximately 20 223 embryos, with each experiment completed at least three independent times. Statistical 224 significance was determined using the log-rank (Mantel Cox) test. (B) The proportion of 225 infected zebrafish embryos with granulomas detected via fluorescent microscopy at 3, 6 and 9 226 days post-infection (dpi). Each experiment was completed three independent times, with 227 approximately 20 embryos per group. Error bars represent standard deviation. Statistical 228 229 analysis was completed using the Student's *t*-test. (C) Example image of granulomas located in the trunk region of zebrafish embryos infected with *M. kansasii* at 9 dpi. Macrophages are 230 231 labelled in red, while *M. kansasii* is labelled in green. Granulomas are highlighted with white 232 arrows. Scale bar represents 1 mm. (D) Representative image of several macrophages (red)

infected with *M. kansasii* (green) forming a granulomatous infection foci at 9 dpi. Scale bar
represents 100 µm.

235

Figure 2: Macrophages are critical determinants of host immunity to *M. kansasii* infection 236 and prevent extracellular cord formation. (A) Zebrafish embryos at 24 hours post-237 fertilisation were injected with lipoclodronate *via* caudal vein injection to deplete macrophages. 238 Following 6 hours treatment, embryos were infected with 300 CFU M. kansasii ATCC 12478 239 via caudal vein injection and survival was monitored daily over a 12-day period. Statistical 240 significance was determined using the log-rank (Mantel Cox) test. (B) Bacterial burden was 241 quantified by fluorescent pixel counts (FPC) using ImageJ, with each point representing an 242 individual embryo. Error bars represent standard deviation. Statistical analysis was completed 243 244 using the Student's *t*-test. (C) The proportion of infected zebrafish embryos with extracellular cords detected (left panel) and the average number of cords per embryo split into low (1-5 245 cords/embryo), moderate (6-10 cords/embryo) and high (>10 cords/embryo) number of cords 246 per embryo (right panel) that were detected using fluorescent microscopy at 3, 6 and 9 days 247 post-infection (dpi). The number of embryos analysed is indicated into brackets. Error bars 248 249 represent standard deviation. Statistical analysis was completed using Student's t-test. (D) 250 Representative images of extracellular cords identified using whole embryo live fluorescent 251 microscopy on the trunk region at 6 dpi (left panel) and 9 dpi (right panel). Macrophages are 252 labelled in red and *M. kansasii* is labelled in green. Cords are highlighted with white arrows. Scale bars represent 1 mm and 100 μ m, respectively. *** $P \le 0.001$, **** $P \le 0.0001$. ns, non-253 significant. 254

255

Supplementary Figure 1: *M. kansasii* is able to replicate at lower temperatures. *M. kansasii* ATCC 12478 was inoculated in $7H9^{OADC/T}$ at an optical density of 0.05 (OD₆₂₀) and

incubated at 30°C and 37°C under shaking at 100 rpm. Growth measurements were taken daily
over a 16-day period until cultures reached stationary phase.

260

Supplementary Figure 2: Aaverage numbers of granulomas per embryo over time. Macrophage reporter embryos Tg(*mpeg1:mCherry*) at approximately 30 hours postfertilisation were infected with 300 CFU of *M. kansasii* ATCC 12478 *via* caudal vein injection. The average number of granulomas per infected zebrafish embryos was detected *via* fluorescent microscopy at 3, 6 and 9 days post-infection (dpi). Each experiment was completed three independent times, with approximately 20 embryos per group. Error bars represent standard deviation. Statistical analysis was completed using the Student's *t*-test.

268

Supplementary figure 3: Efficacy of liposomal clodronate macrophage depletion and restoration kinetics. Macrophage reporter embryos Tg(*mpeg1:mCherry*) at 24 hours postfertilisation were injected with either PBS clodronate or liposomal clodronate *via* caudal vein injection to deplete macrophages (in red). Following 6 hours treatment, embryos were infected with 300 CFU *M. kansasii* ATCC 12478 (in green). Embryos were monitored at 3 and 6 dpi to determine the rate of regeneration of macrophages in treated embryos. Scale bars represent 1mm.

276

Supplementary Figure 4: *M. kansasii* forms large bacterial cords under standard *in vitro* culture conditions. *M. kansasii* ATCC 12478 was inoculated in $7H9^{OADC/T}$ at an optical density of 0.05 (OD₆₂₀) and incubated at 37°C with no agitation for approximately 2 weeks, after which bacteria were mounted onto a glass slide and visualised using fluorescent microscopy. Scale bars presents 50 µm.

282 **References**

283	1. Marras TK, Morris A, Gonzalez LC, Daley CL. Mortality prediction in pulmonary
284	Mycobacterium kansasii infection and human immunodeficiency virus. Am J Respir Crit Care
285	Med 2004 ; 170:793-8.
286	
287	2. Wang J, McIntosh F, Radomski N, et al. Insights on the emergence of Mycobacterium
288	tuberculosis from the analysis of Mycobacterium kansasii. Genome Biol Evol 2015; 7:856-70.
289	
290	3. Lovell JP, Zerbe CS, Olivier KN, et al. Mediastinal and disseminated Mycobacterium
291	kansasii disease in GATA2 deficiency. Ann Am Thorac Soc 2016; 13:2169-73.
292	
293	4. Houben D, Demangel C, van Ingen J, et al. ESX-1-mediated translocation to the cytosol
294	controls virulence of mycobacteria. Cell Microbiol 2012; 14:1287-98.
295	
296	5. Madigan CA, Cameron J, Ramakrishnan L. A zebrafish model of Mycobacterium leprae
297	granulomatous infection. J Infect Dis 2017; 216:776-9.
298	
299	6. Bernut A, Herrmann J-L, Kissa K, et al. Mycobacterium abscessus cording prevents
300	phagocytosis and promotes abscess formation. Proc Nat Acad Sci USA 2014; 111:943-52.
301	
302	7. Davis JM, Clay H, Lewis JL, Ghori N, Herbomel P, Ramakrishnan L. Real-time visualization
303	of mycobacterium-macrophage interactions leading to initiation of granuloma formation in
304	zebrafish embryos. Immunity 2002 ; 17:693-702.
305	
306	8. Cronan MR, Tobin DM. Fit for consumption: zebrafish as a model for tuberculosis. Dis
307	Model Mech 2014 ; 7:777-84.
	13

309	9. Viljoen A, Gutierrez AV, Dupont C, Ghigo E, Kremer L. A simple and rapid gene disruption
310	strategy in Mycobacterium abscessus: On the design and application of glycopeptidolipid
311	mutants. Front Cell Infect Microbiol 2018; 8:69.
312	
313	10. Takaki K, Davis JM, Winglee K, Ramakrishnan L. Evaluation of the pathogenesis and
314	treatment of <i>Mycobacterium marinum</i> infection in zebrafish. Nat Protoc 2013 ; 8:1114-24.
315	
316	11. Ramakrishnan L. Revisiting the role of the granuloma in tuberculosis. Nature Reviews
317	Immunology 2012 ; 12:352-66.
318	
319	12. Clay H, Davis JM, Beery D, Huttenlocher A, Lyons SE, Ramakrishnan L. Dichotomous
320	role of the macrophage in early Mycobacterium marinum infection of the zebrafish. Cell Host
321	Microbe 2007 ; 2:29-39.
322	
323	13. Julian E, Roldan M, Sanchez-Chardi A, Astola O, Agusti G, Luquin M. Microscopic cords,
324	a virulence-related characteristic of Mycobacterium tuberculosis, are also present in
325	nonpathogenic mycobacteria. J Bacteriol 2010; 192:1751-60.
326	
327	14. Sanchez-Chardi A, Olivares F, Byrd TF, Julian E, Brambilla C, Luquin M. Demonstration
328	of cord formation by rough Mycobacterium abscessus variants: implications for the clinical
329	microbiology laboratory. J Clin Microbiol 2011; 49:2293-5.
330	
331	15. Middlebrook G, Dubos RJ, Pierce C. Virulence and morphological characteristics of
332	mammalian tubercle bacilli. J Exp Med 1947 ; 86:175-84.



