# Images of the month: Intrahepatic multiple low-signal lesions from *Aeromonas salmonicida* infection

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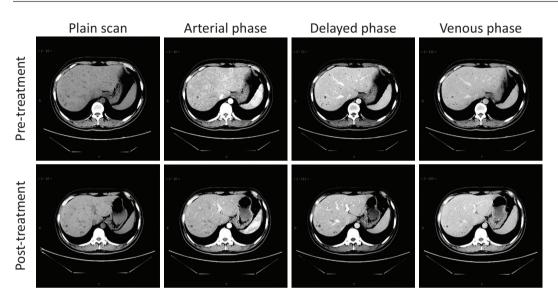


Fig 1. Computed tomography of the liver indicating multiple low-signal lesions in the liver before and after anti-Aeromonas salmonicida treatment.

**KEYWORDS:** Aeromonas salmonicida, liver infection, liver image, AIDS

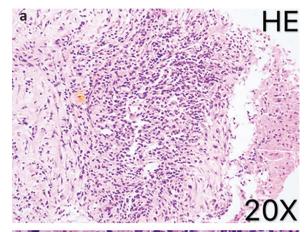
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# Case presentation

A 28-year-old male, with HIV infection for 1 year, initially presented with 5 days of fever (38.6°C) and mild abdomenal pain. He had been irregularly receiving antiretroviral therapy including lamivudine, tenofovir disoproxil fumarate and lopinavir/ritonavir over the past year. The physical examination on admission was notable for tenderness in the epigastrium and pain on percussion in liver area. Laboratory tests indicated physical infections with white blood cell counts at  $2.88\times10^9$ /L and neutrophils counts at  $1.58\times10^9$ /L, but multiple blood cultures did not reveal bacterial, funqi or mycobacterial growth. CD4 cell

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counts were 292 cells/µL. Multiple low-signal lesions in the liver area were shown by computed tomography (CT; Fig 1, upper row). A lumbar puncture and cerebrospinal fluid analysis were performed to exclude the possibility of intracranial infection. The patient was suspected as having a liver abscess and empirical antibiotics were administrated including cefotaxime sodium plus sulbactam sodium and metronidazole, which could cover the common *Escherichia coli* and amoeba. But the symptoms of the patient did not alleviate. In order to find the cause of the disease, we performed a percutaneous liver needle biopsy and the histopathology revealed considerable neutrophil infiltration, which were consistent with inflammation in the liver tissue (Fig 2). But no pathogenic microorganism was found in liver tissue with specific stain and culture. Further metagenomic sequencing (MGS) detected Aeromonas salmonicida in the liver tissue as standard protocol, although the detected A salmonicida reads were only four (Fig 3), and phylogenetic reconstruction showed each of the four sequences had a good homology with reference sequences (Fig 4). The patient then confessed eating half cooked fish before he was sick. Based on this finding, empirical antibiotics were conversed to moxifloxacin that targets A salmonicida and the patient rapidly recovered to his premorbid status, although liver lesions did not completely disappear in CT (Fig 1, lower row). One month later, second liver needle biopsy and MGS did not find A salmonicida infection again.



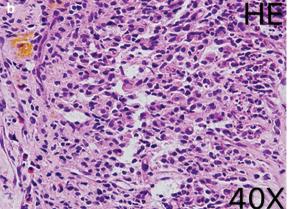


Fig 2. Haematoxylin and eosin stain of the liver biopsy specimen revealing considerable neutrophil infiltration consistent with inflammation.

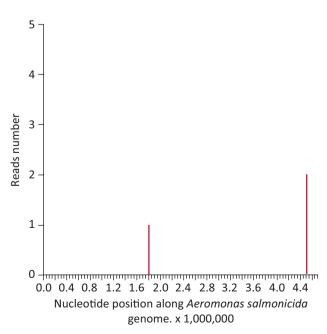


Fig 3. Nucleotide position chromosome coverage of metagenomic sequencing detecting *Aeromonas salmonicida* genome. *Aeromonas salmonicida* 0.0115 % total coverage.

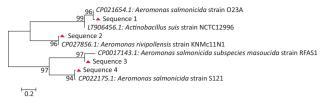


Fig 4. Phylogenetic tree constructed by the neighbour-joining method and Kimura two-parameter model and evaluated by the bootstrap resampling method using 1,000 replicates as implemented in Mega 5.0 software. Five reference sequences were obtained from GenBank. The numbers of the phylogenetic tree denoted the reliability of phylogenetic analysis. The scale bar represents genetic distance (0.2 substitution per site).

#### Discussion

A salmonicida is a member of vibrio family and is a rare multi-host pathogen which can infect immunocompromised individuals. mainly resulting in aastroenteritis and sepsis. Human liver infection of A salmonicida has rarely been reported, so its characteristic image performance is not clear. MGS is a type of DNA sequencing technology which mainly targets a person's genome for exceptional genetic disorders and detects rare pathogen infection when traditional tests are negative or quicker result judgement could increase patients' survival rate.<sup>3,4</sup> However, the high sensitivity of MGS sometimes results in false-positive results, which requires the clinician to accurately judge the detection report according to the clinical diagnosis pathway for specific disease.<sup>5</sup> It is reasonable that an AIDS patient with fever of unknown origin and liver low-signal lesions are conventionally suspected as having liver infection even if no pathogen diagnosis evidence. As traditional blood pathogen culture and empirical antibacterial therapy did not benefit this patient, MGS played a pivotal role in providing a valuable diagnostic support.

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### References

- 1 Janda JM, Abbott SL. The genus Aeromonas: taxonomy, pathogenicity, and infection. *Clin Microbiol Rev* 2010;23:35–73.
- 2 Tsai MS, Kuo CY, Wang MC et al. Clinical features and risk factors for mortality in Aeromonas bacteremic adults with hematologic malignancies. J Microbiol Immunol Infect 2006:39:150–4.
- 3 Human Microbiome Project Consortium. A framework for human microbiome research. *Nature* 2012;486:215–21.
- 4 Pfeiffer F, Zamora-Lagos MA, Blettinger M et al. The complete and fully assembled genome sequence of Aeromonas salmonicida subsp. pectinolytica and its comparative analysis with other Aeromonas species: investigation of the mobilome in environmental and pathogenic strains. BMC Genomics 2018;19:20.
- 5 Taylor JC, Martin HC, Lise S *et al*. Factors influencing success of clinical genome sequencing across a broad spectrum of disorders. *Nat Genet* 2015;47:717–26.

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