

# Deep Vein Thrombosis in Rickettsiosis of the Typhus Group

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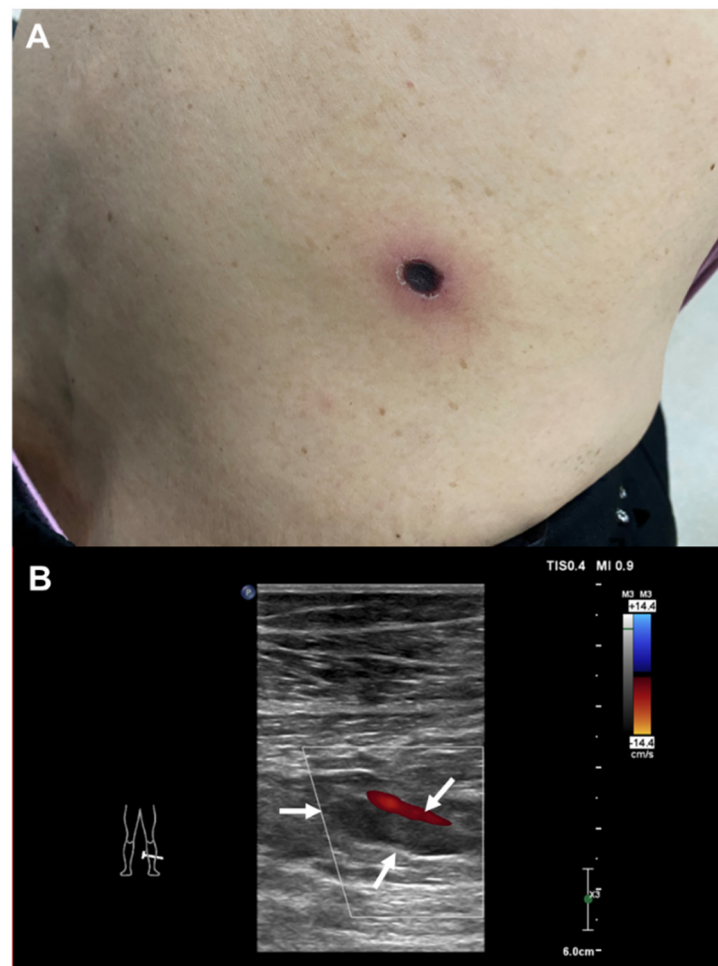
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**Figure 1.** A 58-year-old previously healthy Chinese female sanitation worker was admitted with a one-week history of fever, headache, nausea, and vomiting. Although oseltamivir had been prescribed 6 days before admission, she was unresponsive. On the day of admission, physical examination revealed an eschar (Panel A) on the right lower back and mild swelling of the left lower leg, which was non-pitting edema. Complete blood count showed

lymphopenia and thrombocytopenia. Liver function tests indicated a marked increase in transaminase levels. Coagulation function tests revealed a high D-dimer (11.02 µg/mL) with normal prothrombin time and activated partial thromboplastin time. Rickettsiosis was clinically diagnosed, and doxycycline was subsequently started. Given the mild swelling of the left lower extremity and abnormal D-dimer, suggesting a risk of deep vein thrombosis (DVT), a subsequent doppler ultrasound examination was performed. Color Doppler flow imaging for the left lower leg (Panel **B**) revealed a hypoechoic clot (arrow) within the intermuscular vein lumen, measuring 27 mm × 7 mm, not fully compressible under probe pressure, with no obvious blood flow signal in the local intermuscular vein. The lower-limb DVT of the left lower leg was confirmed, which was treated with low-molecular-weight heparin and rivaroxaban. Although fever and DVT responded promptly to treatment, the initial Weil-Felix test by Shenzhen Center for Disease Control and Prevention was negative, and the genes of rickettsiae were not detected in serum by metagenomic next-generation sequencing (mNGS) [1]. Owing to the negative serological and NGS results, a repeat Weil-Felix test was conducted on the residual sample, which showed an OX-19 titer of 1:160. The final diagnosis was rickettsiosis of the typhus group. Notably, DVT develops in only a tiny minority of foci of vascular infection and injury in rickettsiosis, making this presentation exceptional [2,3]. Although NGS is an advanced powerful tool for laboratory diagnosis of infectious diseases, it has to be used in conjunction with traditional laboratory tests [4,5].

### Author Contributions

F.X.: conceptualization, methodology, visualization, writing-original draft preparation; C.D.: investigation, visualization, writing-reviewing and editing. All authors have read and agreed to the published version of the manuscript.

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### Institutional Review Board Statement

The retrospective study was approved by the Medical Ethics Committee of The University of Hong Kong-Shenzhen Hospital ([2022]120).

### Informed Consent Statement

Patient consent was exempted as this is a retrospective study and no additional testing was performed.

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### Conflicts of Interest

The authors declare no conflict of interest.

### Use of AI and AI-Assisted Technologies

No AI tools were utilised for this paper.

### References

1. Xing, F.; Deng, C.; Huang, J.; et al. Usefulness of Next-Generation Sequencing for Laboratory Diagnosis of Rickettsiosis. *PLoS Negl. Trop. Dis.* **2024**, *18*, e0012546. <https://doi.org/10.1371/journal.pntd.0012546>.
2. Del Prete, E.; Pizzanelli, C.; Moretti, P.; et al. Mediterranean Spotted Fever: An Unusual Clinical and Neuroradiological Presentation. *Neurol. Sci.* **2015**, *36*, 2141–2143. <https://doi.org/10.1007/s10072-015-2313-z>.

3. Nagaki, Y.; Hayasaka, S.; Kadoi, C.; et al. Branch Retinal Vein Occlusion in the Right Eye and Retinal Hemorrhage in the Left in a Patient with Classical Tsutsugamushi Disease. *Jpn. J. Ophthalmol.* **2001**, *45*, 108–110. [https://doi.org/10.1016/s0021-5155\(00\)00293-8](https://doi.org/10.1016/s0021-5155(00)00293-8).
4. Xing, F.; Yang, Q.; Deng, C.; et al. Clinical Impact of Next-Generation Sequencing on Laboratory Diagnosis of Suspected Culture-Negative Meningitis and Encephalitis. *J. Infect.* **2022**, *85*, 573–607. <https://doi.org/10.1016/j.jinf.2022.08.026>.
5. Deng, C.; Yang, Q.; Li, L.; et al. Yield of Metagenomics in Suspected Central Nervous System Infections with Negative Cerebrospinal Fluid Cultures. *eMicrobe* **2025**, *1*, 6. <https://doi.org/10.53941/emicrobe.2025.100006>.