

A serological assay using Tropheryma whipplei antigens for the presumptive exclusion of Whipple disease

Author

Liew, Kwee Chin, Nguyen, Chelsea, Waidyatillake, Nilakshi T, Nguyen, Trang, Walton, Aaron, Harris, Owen, Athan, Eugene, Stenos, John, Graves, Stephen R

Published

2023

Journal Title

Pathology

Version

Accepted Manuscript (AM)

DOI

[10.1016/j.pathol.2023.09.010](https://doi.org/10.1016/j.pathol.2023.09.010)

Rights statement

© 2023. This manuscript version is made available under the CC-BY-NC-ND 4.0 license <https://creativecommons.org/licenses/by-nc-nd/4.0/>

Downloaded from

<http://hdl.handle.net/10072/428337>

Griffith Research Online

<https://research-repository.griffith.edu.au>

1 A serological assay using *Tropheryma whipplei* antigens for the presumptive exclusion of
2 Whipple Disease

3

4 Kwee Chin Liew^{1,2,4*}, Chelsea Nguyen², Nilakshi T Waidyatillake^{3,5}, Trang Nguyen⁶, Aaron
5 Walton^{2,4}, Owen Harris^{1,3,4}, Eugene Athan^{2,3,4}, John Stenos² and Stephen R. Graves^{2*}

6

7 ¹Department of Microbiology, Australian Clinical Labs, Geelong 3220, VIC, Australia.

8 ²Australian Rickettsial Reference Laboratory, University Hospital Geelong, Geelong 3220, VIC,
9 Australia.

10 ³Deakin University, School of Medicine, Geelong 3220, VIC, Australia.

11 ⁴Barwon Health, University Hospital Geelong 3220, VIC, Australia.

12 ⁵Allergy and Lung Health Unit, Centre for Epidemiology and Bio statistics, Melbourne School of
13 Population and Global Health, The University of Melbourne 3010, VIC, Australia.

14 ⁶Centre for Infectious Diseases and Microbiology Laboratory Services, ICPMR-Pathology West,
15 Sydney, Australia.

16

17 Running Head: Whipple's Disease exclusion serological assay

18

19 *Address correspondence to Kwee Chin Liew (K.C), lwkwel@yahoo.com.au.

20 *Alternative Corresponding author: Prof Stephen R. Graves, graves.rickettsia@gmail.com

21 Author order was determined based on seniority and contribution to this work.

22 This work was funded by the Australian Rickettsial Reference Laboratory Foundation Limited.

23

24

25
26
27
28
29
30
31
32
33
34
35
36
37
38
39
40
41
42
43
44
45
46
47

SUMMARY

Whipple’s disease (WD) is a rare infection in genetically susceptible persons, caused by the bacterium *Tropheryma whipplei*. An indirect-immunofluorescence serological assay (IFA), detecting patient antibodies to the bacterium, was developed using *T. whipplei* as antigen. We hypothesised that this assay could be used to rule out WD in patients in whom the diagnosis was being considered, based on a high immunoglobulin (Ig) G antibody titre to *T. whipplei*. In this study 16 confirmed WD patients and 156 age-matched controls from across Australia were compared serologically. WD patients mostly under-produce IgG antibody to *T. whipplei*, with titres of $\leq 1:32$ being common. While at an antibody titre of $< 1:64$ antibody titre the assay sensitivity for WD was only 69% (95% CI 41-89%), its specificity for not being WD was 91% (95% CI 85-95%). This specificity increased to 95% (95% CI 90-98%) at an antibody titre of $< 1:16$. Patients with antibody titres of $> 1:64$ were unlikely to have WD. At this titre, seroprevalence of *T. whipplei* IgG antibody was 92% (223/242) in Australian blood donors. Unlike other serological assays which are used to confirm a specific infection, this novel assay is designed to rule out WD infection with a specificity in Australia of 91%. Further validation of this assay, by trialing in other countries, should now be undertaken, as its usefulness is dependent on there being a high background seropositivity to *T. whipplei* in the general population at the location in which the assay is being used to rule out WD.

48 INTRODUCTION

49

50 Whipple disease (WD) is a rare infection due to *Tropheryma whipplei* with an estimated
51 prevalence of 1.1-9.8 cases per million persons and an annual incidence of 1-6 per 10 million
52 persons¹⁻³. In a small case series from Germany, Spain and north-western Italy, the maximum
53 incidence of WD was in persons between 40-60 years of age, of whom 72-96% were middle-aged
54 Caucasian men²⁻⁷. In contrast to these findings, a large retrospective population-based study
55 carried out between 2012 and 2017 in the United States (US) found that men and women were
56 affected at similar rates and that WD was more common in people aged greater than 65 years³.

57

58 The clinical spectrum of WD is wide^{1,5,8,9}. In classical WD, infection with *T. whipplei* probably
59 occurs years before the first clinical manifestations, which are most often chronic diarrhea, due to
60 gastroenteritis, or migratory arthralgia^{1,9}. In localised WD *T. whipplei* can infect the joints, bones,
61 brain, heart, skin, eyes, lymph nodes, lung and/or kidney^{1,8,9}. A broad spectrum of clinical severity
62 has been observed, ranging from asymptomatic carriage^{5,8} to organ failure and death. Fatal
63 outcome is associated with delayed diagnosis as well as misdiagnosis as an
64 autoimmune/inflammatory disorder with consequent immunosuppressive treatment⁹.

65

66 It has been observed that patients with confirmed WD frequently have reduced or undetectable *T.*
67 *whipplei*-specific (Ig) G antibody levels. WD patients may not be able to mount an adequate IgG
68 antibody response, which may be associated with a subtle immune defect. In contrast, individuals
69 who do not go on to develop clinical WD appear to mount a strong IgG antibody response when
70 exposed to *T. whipplei*. While healthy individuals not yet exposed to the bacterium (mostly

71 children and young adults) are necessarily IgG seronegative, they respond normally when exposed.
72 The aim of this study was to compare the sera of patients with confirmed WD to individuals
73 without WD and to evaluate the diagnostic utility of a *T. whipplei* screening IFA serology assay
74 as a tool to help distinguish patients with WD from those without WD. Seropositivity to *T. whipplei*
75 in healthy, age and sex matched blood donors in Australia was measured as the control for this
76 study of WD patients.

77
78

79 **PATIENTS, MATERIALS AND METHODS**

80

81

82 **Study Design**

83 A case-controlled comparative study of *T. whipplei* IFA was conducted from November 2019 to
84 October 2022 at the Australian Rickettsial Reference Laboratory (ARRL), Geelong, Victoria,
85 Australia, according to the *National Statement on Ethical Conduct in Human Research (2007)*
86 guideline (Barwon Health ethics reference number: 19/135). All sera tested in this study were de-
87 identified.

88

89 Adult patients aged more than 18 years were included. Grounds for exclusion were pregnancy,
90 active malignancy, primary immunodeficiency, and HIV with CD4 lymphocyte count < 200.
91 Subjects were considered WD cases if they displayed consistent symptoms with supportive
92 diagnostic test results, such as positive periodic acid-Schiff (PAS) staining, positive *T. whipplei*-
93 specific immunohistochemistry staining or positive *T. whipplei* PCR or 16s rRNA PCR of a tissue
94 biopsy.¹⁰⁻¹³

95

96 **Sample size and collection**

97 Given the limited number of WD patients in Australia, we employed a convenience sampling
98 strategy. Patients (n=16) with confirmed WD as described above were recruited from across
99 Australia. Of the 16 WD cases, eleven had sera already in storage at ARRL. The remaining five
100 cases had serum sample collection organized through their local doctors and pathology services.
101 In addition, 242 sera were obtained from healthy individuals via the Australian Red Cross
102 Lifeblood; 156 of these sera were age and sex matched to WD patients to act as controls and all
103 sera were tested for antibodies to *T. whipplei*. Data such as gender, age, clinical manifestations,
104 IgM, IgG and IgA titres to *T. whipplei*, 16S rRNA, 13 *T. whipplei* PCR and histology results were
105 obtained from WD patients and controls.

106 **Diagnostic techniques**

107

108 **Culture and IFA slide preparation**

109 A French strain of *T. whipplei* was cultivated in liquid axenic medium at 35°C for several weeks.¹⁴
110 The *T. whipplei* cells were pelleted and resuspended in phosphate-buffered saline (PBS). To
111 optimise antigen concentrations for the final IFA, slides were prepared using doubling dilutions of
112 *T. whipplei* antigen and subjected to sandwich IFA after incubation with a strongly reacting
113 positive serum sample. Antigen aliquots were stored at -70°C until required. When thawed, they
114 were spotted onto clean 3×10-well glass slides. After air drying, antigens were fixed with acetone
115 for 5 min. Slides could then be stored at -70°C for later use.

116

117 **Indirect-immunofluorescence assay (IFA)**

118 All sera from the WD patients and controls were tested by in-house *T. whipplei* IFA. Known
119 reactive and non-reactive sera were tested on the same slide as the assay samples. Initial 1:8

120 dilutions of test and control sera were prepared using a 2% casein PBS buffer. Serial doubling
121 dilutions from 1:8 to 1:4096 were prepared and tested by IFA. Each serum dilution was spotted in
122 duplicate onto pre-prepared *T. whipplei* slides. Slides were incubated for 30 minutes under humid
123 conditions at 35°C and then washed for 5 minutes in a 1:10 dilution of PBS and allowed to dry.
124 Fluorescein-conjugated goat-anti-human antibodies, [1:100 anti-human IgA (KPL, Cat # 02-10-
125 01), 1:100 anti-human IgM (KPL, Cat # 02-10-03) and 1:100 anti-human IgG (KPL, Cat # 02-10-
126 02)] were applied, and slides were incubated, washed and dried as above. Fluorescence mounting
127 fluid (Dako North America, Inc., USA) and a coverslip were applied. Microscopic evaluations of
128 the IFA were carried out under 400x magnification using an LED Leica DMLS fluorescence
129 microscope. The assay was repeated in duplicate for all sera. Reading of the slides were blinded.
130 Antibody titres for IgM, IgG and IgA were defined as the highest serum dilution giving strong
131 immunofluorescence (like the reactive control serum).

132

133 **Preparation of positive control sera for IFA**

134 IFA-positive *T.whipplei* human sera were kept in frozen storage as 5 uL aliquots until required.
135 They had been previously titrated and diluted in PBS to reach a fluorescence endpoint of 1:512
136 and used as controls on each IFA slide.

137

138 **Statistical analysis**

139 Statistical comparisons of groups, by 2-sided Fisher exact test was done on the quantitative data.
140 Sensitivity was defined as the proportion of WD patients correctly identified at a particular
141 antibody cutoff¹⁵. Specificity was defined as the proportion of healthy controls correctly identified
142 as not having WD¹⁵. Significance was set at $p < 0.05$.

143
144
145
146
147
148

RESULTS

Case demographics and clinical presentations

149 Fifteen male and one female WD patients were recruited to the study. They ranged in age from
150 37–82 years, with a mean age of 57 years [95% confidence interval (CI) 50–65 years] (Table 1).
151 The clinical presentations of these 16 WD cases varied: seven had infective endocarditis; three
152 displayed central nervous system manifestations; four had classic WD with intestinal
153 malabsorption; and two had joint manifestations.

154

Demographics of the controls

156 The 156 age-matched controls ranged from 31-69 years old (Table 1). The mean age of age-
157 matched controls was 50 years (95% CI 48-52). There was no obvious difference between the WD
158 and age-matched controls in terms of gender and age.

159

IFA serology: IgG

161 WD patients' antibody titres are shown in Table 2 and compared with controls in Figure 1. WD
162 patients were less able to produce IgG antibodies to *T. whipplei* than healthy age-matched controls.
163 This difference was statistically significant when comparing antibody titres at 1:16, 1: 32 and 1:64
164 dilutions (Table 3).

165

166 With an IFA IgG titre cutoff set at <1:64, sensitivity (for diagnosing WD) and specificity (for
167 diagnosing not having WD) were 69% (95% CI 41-89%) and 91% (95% CI 85-95%) respectively
168 (Table 3).

169

170 **IgM and IgA**

171 No difference in *T. whipplei* IgM or IgA IFA antibody titres was observed between WD cases and
172 age-matched controls ($p > 0.05$). Titres of this antibody isotype did not discriminate between these
173 two groups (data not shown for controls).

174

175 **DISCUSSION**

176

177 This study appears to confirm the hypothesis that patients with WD under-produce IgG antibody
178 compared with age and sex-matched controls exposed to *T. whipplei*. This finding is consistent
179 with previous reports¹⁶⁻²⁰. We were able to establish an antibody titre cutoff (<1:64) and arrive at
180 a potentially useful serological assay to rule out WD (specificity 91%). In patients who had been
181 naturally exposed to *T. whipplei*, and who had IgG antibody titres $\geq 1:64$, there was a reasonable
182 likelihood that they did not have WD. Those without WD rarely have such low titers (<1:64). In a
183 previous French study, when the cutoff titre was set at 1:100, IgG did not discriminate between
184 subjects with and without WD²¹. The discrepancy between these results might be explained by the
185 different antibody cutoff titres used as well as possible technical differences in the assays. In an
186 Australian context, this assay has the potential to be useful to screen for the absence of WD in
187 patients with WD-compatible signs and symptoms.

188

189 In contrast to a previous study, we did not find serum IgA to *T. whipplei* to be useful in
190 discriminating between patients with and without WD. Secretory IgA in intestinal mucus
191 might correlate with recent exposure to *T. whipplei*, but it would be challenging to measure
192 routinely. The usual early serological response to a primary bacterial or viral infection would

193 normally involve a conversion from IgM to IgG isotypes modulated by T-cell cytokines.
194 Seroconversion or a four-fold rise in antibody titre would be expected after about a 10–14-
195 day interval.²² Our serological findings in WD patients were contrary to this. This may be due
196 to a specific genetic predisposition to WD associated with human leukocyte antigen alleles
197 DRB1*13 and DQB1*06, which may disrupt optimum antigen presentation.²³ Interleukin
198 (IL)-16 gene polymorphisms and other polymorphisms polarise cytokine production towards
199 T-helper (Th) 2 cell activity.^{24,25} This specific genetic predisposition probably explains the
200 inability to switch from IgM to IgG antibody isotypes. In addition, there appears to be a
201 decrease in Th1 and Th17 cell activity, poor synthesis of transforming growth factor-beta 1
202 and excess synthesis of IL-10 by WD patients,^{17,26–28} a biological mechanism that changes B-
203 cell production of antibody from one class to another.^{16,19,20}

204

205 This is the first study to test for the seroprevalence of *T. whipplei* IgG antibody in Australia.
206 The finding that 223/ 242 (92%) Australian blood donors had *T. whipplei* IgG antibody
207 detected by IFA at $\geq 1:64$ serum dilution suggests that the microbe is widespread in the
208 Australian environment, with most adult Australians having been exposed to it. *Tropheryma*
209 *whipplei* has been detected in 37–66% of waste from sewage plants^{4,28–32} and in stool samples
210 of 48% children in Laos,³³ and 12–26% of sewage plant workers³⁰ and 1–11% of healthy
211 individuals in France.²⁹ The viable bacterium can also be detected in human faeces and
212 saliva,¹² supporting the notion of human colonisation, probably through faecal-oral
213 transmission. Further work with larger sample populations will be required to fully
214 characterise rates of exposure to *T. whipplei* in relevant communities and different settings.

215

216 There are several limitations to this study. Only a single strain of *T. whipplei* was used as
217 antigens for the IFA. Thus, if there were strain-specific differences in antibody affinities, it
218 might have affected our results. Antigenic shift and loss of antigen specificity after a few
219 subcultures is also a possibility. Ideally, a few robust antigenic targets could be identified
220 through a proteomic approach for the development of more specific assays. Testing patients
221 with diseases other than WD should now be undertaken. Finally, the absent/low antibody
222 responses we observed in some healthy controls complicate the reliance on an absence of *T.*
223 *whipplei* IgG responses as a marker for active WD. We assume that the absence of antibodies
224 in healthy individuals reflects a lack of exposure to the bacterium. Additional studies to
225 differentiate asymptomatic carriers could be considered by undertaking PCR testing on saliva
226 and/or stool. The control sera were obtained from Australian Red Cross Lifeblood, which
227 does not collect blood from persons older than 70, meaning this is an inherent bias in the
228 study. However, the study findings remain the same even if the three WD patients over 70
229 years old are removed from the analysis. This study demonstrates that the detection of *T.*
230 *whipplei* IgG antibody on IFA serology is a promising screening assay to rule out WD (Fig.
231 2). The usefulness of this assay (or the confidence a doctor may have in ruling out WD in a
232 patient) will only be shown by the passage of time, as the assay is used more widely. Given
233 the rarity of WD and the invasive nature of existing diagnostic tests, an inexpensive,
234 noninvasive, screening serological assay with sufficient accuracy to reliably exclude the
235 diagnosis may be a worthwhile addition to the diagnostic armamentarium of the physician.

236

237 **Acknowledgements:** The authors thank Professor Didier Raoult and Professor Pierre-
238 Edouard Fournier for providing the French *Tropheryma whipplei* isolate. They also thank all

239 the patients involved in this study and Ms Manidipa Majumber for assisting with the
240 seroprevalence study. Thanks are also due to the following medical colleagues: Ashley
241 Watson, Brian Chong, Callum Maggs, Cameron Jeremiah, Carly Hughes, Caroline Bartolo,
242 Christopher Swan, David Andresen, David Foley, David Sheffield, David Sowden, Denis
243 Spelman, Freya Langham, Hao Yu, Harsha Sheorey, Gabrielle O’Kane, Grace Butel-Simoes,
244 Hui Yi Ng, James Branley, Jenny Robson, Katy Lai, Lucy Crawford, Mike Catton, Renjy
245 Nelson, Rob Pickles, Tristan Gibbs, Sandra Jones, Smathi Chong, Su Ann Ho, Tony Korman,
246 Vitali Sintchenko and Zaal Meher-Homji. Dr Iain Gosbell, Dr David Irving and Dr Rena
247 Hirani of Australian Red Cross Lifeblood are thanked for providing the age- and sex-matched
248 serum samples.

249

250 **Ethics approval:** This study was conducted according to the National Statement on Ethical
251 Conduct in Human Research (2007) of the Australian National Health and Medical Research
252 Council (Barwon Health ethics reference number: 19/135).

253

254 **Conflicts of interest and sources of funding:** This study was supported by the Australian
255 Rickettsial Reference Laboratory Foundation Limited. The authors state that there are no
256 conflicts of interest to disclose.

257

258 Address for correspondence: Dr Kwee Chin Liew, Department of Microbiology, Australian
259 Clinical Labs, Geelong, Vic 3220, Australia. E-mail: lwkweel@yahoo.com.au

260

261

262 **REFERENCES**

263

264 1. Schneider T, Moos V, Loddenkemper C, et al. Whipple's disease: new aspects of
265 pathogenesis and treatment. *Lancet Infect Dis.* 2008;8(3):179-190.

266 [https://doi:10.1016/S1473-3099\(08\)70042-2](https://doi:10.1016/S1473-3099(08)70042-2)

267

268 2. von Herbay A, Otto HF, Stolte M, et al. Epidemiology of Whipple's disease in Germany.
269 Analysis of 110 patients diagnosed in 1965-95. *Scand J Gastroenterol.* 1997;32(1):52-57.

270 <https://doi:10.3109/00365529709025063>

271

272 3. Elchert JA, Mansoor E, Abou-Saleh M, Cooper GS. Epidemiology of Whipple's Disease
273 in the USA Between 2012 and 2017: A Population-Based National Study. *Dig Dis Sci.*

274 2019;64(5):1305-1311. <https://doi:10.1007/s10620-018-5393-9>

275

276 4. Dutly F, Altwegg M. Whipple's disease and "*Tropheryma whippelii*". *Clin Microbiol Rev.*
277 2001;14(3):561-583. <https://doi:10.1128/CMR.14.3.561-583.2001>

278

279 5. Lagier JC, Lepidi H, Raoult D, et al. Systemic *Tropheryma whippelii*: clinical presentation
280 of 142 patients with infections diagnosed or confirmed in a reference center. *Medicine*

281 (Baltimore). 2010; 89(5): 337–345. <https://doi:10.1097/MD.0b013e3181f204a8>. PMID:

282 20827111.

283

- 284 6. Biagi F, Balduzzi D, Delvino P, Schiepatti A, Klersy C, Corazza GR. Prevalence of
285 Whipple's disease in north-western Italy. *Eur J Clin Microbiol Infect Dis.*
286 2015;34(7):1347-1348. <https://doi:10.1007/s10096-015-2357-2>
- 287
288 7. Ojeda E, Cosme A, Lapaza J, Torrado J, Arruabarrena I, Alzate L. Whipple's disease in
289 Spain: a clinical review of 91 patients diagnosed between 1947 and 2001. *Rev Esp Enferm*
290 *Dig.* 2010;102(2):108-123. <https://doi:10.4321/s1130-01082010000200006>
- 291
292 8. Günther U, Moos V, Offenmüller G, et al. Gastrointestinal diagnosis of classical Whipple
293 disease: clinical, endoscopic, and histopathologic features in 191 patients. *Medicine*
294 *(Baltimore)*. 2015;94(15): e714. <https://doi:10.1097/MD.0000000000000714>
- 295
296 9. Marth T, Schneider S. Whipple's disease. In: *Principles and practice of infectious disease,*
297 8th edn. Mandell G, Bennett J, Dolin R, eds. Philadelphia: Elsevier, 2015; 2418–2425.
- 298
299 10. Fenollar F, Puéchal X, Raoult D. Whipple's disease. *N Engl J Med.* 2007;356(1):55-66.
300 <https://doi:10.1056/NEJMra062477>
- 301
302 11. Lepidi H, Fenollar F, Gerolami R, et al. Whipple's disease: immunospecific and
303 quantitative immunohistochemical study of intestinal biopsy specimens. *Hum Pathol.*
304 2003;34(6):589-596. [https://doi:10.1016/s0046-8177\(03\)00126-6](https://doi:10.1016/s0046-8177(03)00126-6)
- 305
306 12. Fenollar F, Laouira S, Lepidi H, Rolain JM, Raoult D. Value of *Tropheryma whipplei*
307 quantitative polymerase chain reaction assay for the diagnosis of Whipple disease:

- 308 usefulness of saliva and stool specimens for first-line screening. Clin Infect Dis.
309 2008;47(5):659-667. <https://doi:10.1086/590559>
- 310
311 13. Relman DA. Universal bacterial 16S rRNA amplification and sequencing. In: Persing DH,
312 Smith TF, Tenover FC, White TJ, editors. Diagnostic Molecular Microbiology: Principles
313 and Applications. Washington DC: ASM Press, 1993; 489–95.
- 314
315 14. Renesto P, Crapoulet N, Ogata H, et al. Genome-based design of a cell-free culture
316 medium for *Tropheryma whipplei*. Lancet. 2003;362(9382):447-449.
317 [https://doi:10.1016/S0140-6736\(03\)14071-8](https://doi:10.1016/S0140-6736(03)14071-8)
- 318
319 15. Empson MB. Statistics in the pathology laboratory: Characteristics of diagnostic tests,
320 Pathology 2001; 33:1, 93-95. <https://doi:10.1080/00313020120034966>
- 321
322 16. Black-Schaffer B. The Tinctoral Demonstration of a Glycoprotein in Whipple's
323 Disease. Proceedings of the Society for Experimental Biology and Medicine.
324 1949;72(1):225-227. <https://doi:10.3181/00379727-72-17388>
- 325
326 17. Marth T, Strober W. Whipple's disease. Semin Gastrointest Dis 1996; 7:41-8.
- 327
328 18. Mahnel R, Kalt A, Ring S, Stallmach A, Strober W, Marth T. Immunosuppressive therapy
329 in Whipple's disease patients is associated with the appearance of gastrointestinal
330 manifestations. Am J Gastroenterol. 2005;100(5):1167-1173. [https://doi:10.1111/j.1572-](https://doi:10.1111/j.1572-0241.2005.40128.x)
331 [0241.2005.40128.x](https://doi:10.1111/j.1572-0241.2005.40128.x)

332
333
334
335
336
337
338
339
340
341
342
343
344
345
346
347
348
349
350
351
352
353
354

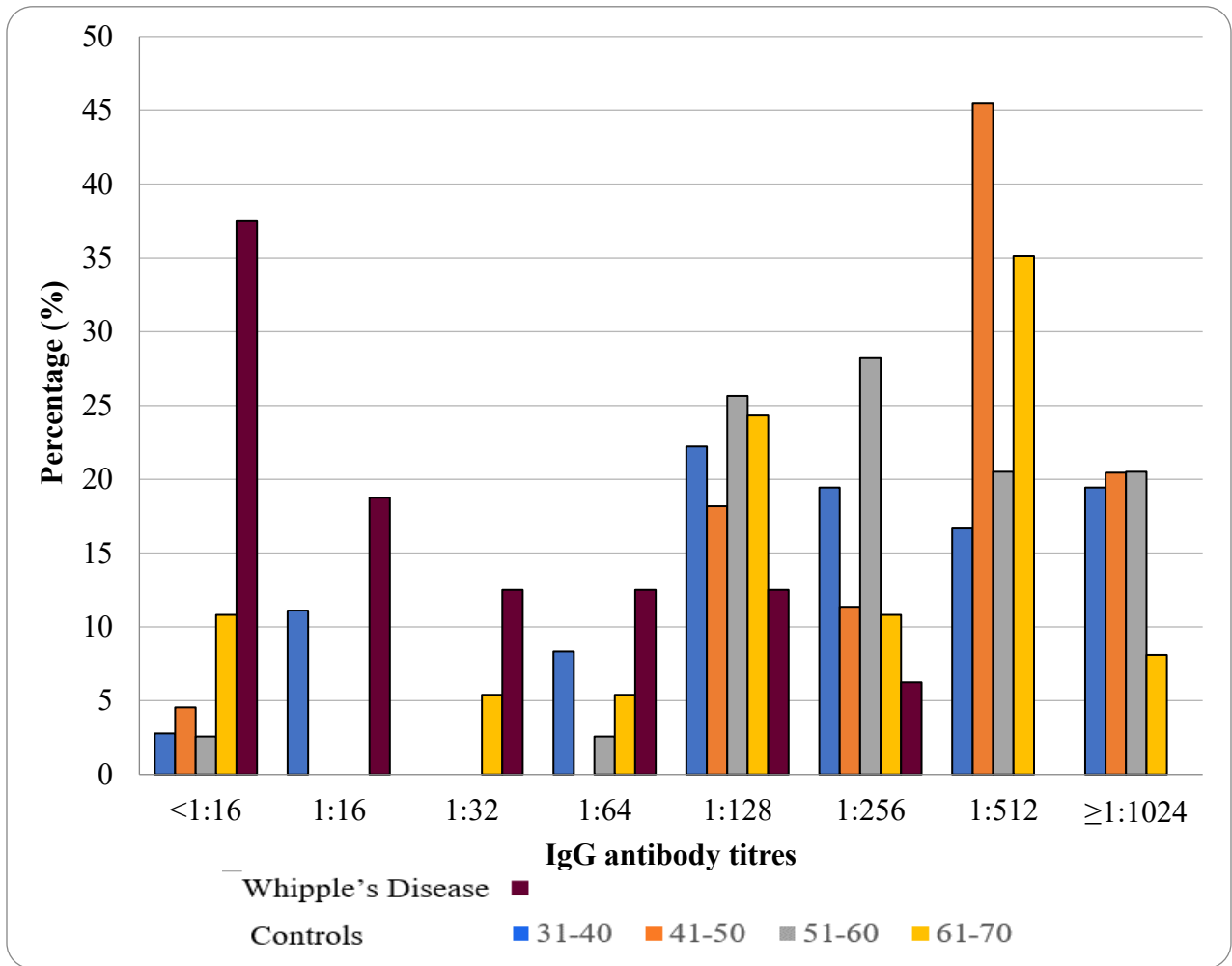
19. Bonhomme CJ, Renesto P, Nandi S, Lynn AM, Raoult D. Serological microarray for a paradoxical diagnostic of Whipple's disease. *Eur J Clin Microbiol Infect Dis*. 2008; 27(10):959-968. <https://doi:10.1007/s10096-008-0528-0>
20. Bonhomme CJ, Renesto P, Desnues B, et al. *Tropheryma whippelii* glycosylation in the pathophysiologic profile of Whipple's disease. *J Infect Dis*. 2009;199(7):1043-1052. <https://doi:10.1086/597277>
21. Raoult D, Birg ML, La Scola B, et al. Cultivation of the bacillus of Whipple's disease. *N Engl J Med* 2000; 342: 620–5 [correction *N Engl J Med* 2000; 342: 1538].
22. Carroll KC, Pfaller MA, Landry ML, et al., editors. *Manual of Clinical Microbiology*. 12th ed. Washington, DC: ASM Press, 2019.
23. Martinetti M, Biagi F, Badulli C, et al. The HLA alleles DRB1*13 and DQB1*06 are associated to Whipple's disease. *Gastroenterology*. 2009; 136 (7):2289-2294. <https://doi:10.1053/j.gastro.2009.01.051>.
24. Biagi F, Schieppatti A, Badulli C, et al. -295 T-to-C promoter region IL-16 gene polymorphism is associated with Whipple's disease. *Eur J Clin Microbiol Infect Dis*. 2015;34(9):1919-1921. <https://doi:10.1007/s10096-015-2433-7>

- 355 25. Biagi F, Badulli C, Feurle GE, et al. Cytokine genetic profile in Whipple's disease. Eur J
356 Clin Microbiol Infect Dis. 2012;31(11):3145-3150. [https://doi:10.1007/s10096-012-1677-](https://doi:10.1007/s10096-012-1677-8)
357 8
358
359
- 360 26. Relman DA, Schmidt TM, MacDermott RP, Falkow S. Identification of the uncultured
361 bacillus of Whipple's disease. N Engl J Med. 1992;327(5):293-301.
362 <https://doi:10.1056/NEJM199207303270501>
363
364
- 365 27. Moter A, Janneck M, Wolters M, et al. Potential Role for Urine Polymerase Chain
366 Reaction in the Diagnosis of Whipple's Disease. Clin Infect Dis. 2019;68(7):1089-1097.
367 <https://doi:10.1093/cid/ciy664>
368
- 369 28. Morgenegg S, Dutly F, Altwegg M. Cloning and sequencing of a part of the heat shock
370 protein 65 gene (hsp65) of "*Tropheryma whippelii*" and its use for detection of "*T.*
371 "*whippelii*" in clinical specimens by PCR [published correction appears in J Clin Microbiol
372 2000 Oct;38(10):3914]. J Clin Microbiol. 2000;38(6):2248-2253.
373
- 374 29. Fenollar F, Trani M, Davoust B, et al. Prevalence of asymptomatic *Tropheryma whippelii*
375 carriage among humans and nonhuman primates. J Infect Dis. 2008;197(6):880-887.
376 <https://doi:10.1086/528693>
377

- 378 30. Schöniger-Hekele M, Petermann D, Weber B, Müller C. *Tropheryma whipplei* in the
379 environment: survey of sewage plant influxes and sewage plant workers. Appl Environ
380 Microbiol. 2007;73(6):2033-2035. <https://doi:10.1128/AEM.02335-06>
381
- 382 31. Maiwald M, Schuhmacher F, Ditton HJ, von Herbay A. Environmental occurrence of the
383 Whipple's disease bacterium (*Tropheryma whippelii*). Appl Environ Microbiol.
384 1998;64(2):760-762. <https://doi:10.1128/AEM.64.2.760-762.1998>
385
- 386 32. Marth T, Raoult D. Whipple's disease. Lancet. 2003;361(9353):239-246.
387 [https://doi:10.1016/S0140-6736\(03\)12274-X](https://doi:10.1016/S0140-6736(03)12274-X)
388
- 389 33. Keita AK, Dubot-Pérès A, Phommasone K, et al. High prevalence of *Tropheryma whipplei*
390 in Lao kindergarten children. PLoS Negl Trop Dis. 2015;9(2):e0003538. Published 2015
391 Feb 20. <https://doi:10.1371/journal.pntd.0003538>

392 **FIG 1:** Distribution of immunoglobulin G (IgG) titres to *Tropheryma whipplei* in Whipple
393 disease patients and age-matched controls.

394



395

396

397 **TABLE 1:** Gender and age comparison of Whipple Disease patients (n=16) and age-matched
 398 controls (n=156)

399

Gender	Whipple's Disease	Age-matched controls
Female (%)	1 (6)	2 (1)
Male (%)	15 (94)	154 (99)
Age Group		
31-40 (%)	4 (25)	36 (23)
41-50 (%)	1 (6)	44 (28)
51-60 (%)	5 (31)	39 (25)
61-70 (%)	3 (19)	35 (24)
71-80 (%)	2 (13)	0 (0)
81-90 (%)	1 (6)	0 (0)
Mean age	57	50

400

401 **TABLE 2:** Indirect immunofluorescence assay antibody titres in 16 patients with Whipple
 402 disease (WD)

403

Study patients	Gender	Age Range	Clinical Manifestation & Diagnostic method	IgA	IgM	IgG
Patient 1	M	41-50	Endocarditis Dx Cardiac valve 16s rRNA +	<32	256	≤8
Patient 2	M	51-60	Endocarditis Dx Cardiac valve 16s rRNA +	<32	≤32	≤8
Patient 3	M	51-60	Classic WD + Joint Dx Duodenal and knee tissues PCR +	<32	128	≤8
Patient 4	M	51-60	CNS WD + Joint Dx Small bowel biopsy PAS +; PCR + CSF PCR +	<32	256	≤8
Patient 5	M	61-70	CNS WD Dx CSF PCR +	<32	≤32	≤8
Patient 6	M	71-80	CNS WD Dx: Vitreous PAS + Gastric PCR +	<32	≤32	≤8
Patient 7	M	31-40	Classic WD	<32	≤32	≤8

			Dx: Duodenum and Lymph node PAS+; PCR +			
Patient 8	M	51-60	Endocarditis Dx Cardiac valve 16s rRNA +	32	1024	16
Patient 9	F	61-70	Classic WD + Joint Lymph node PAS +; PCR +	<32	512	16
Patient 10	M	31-40	Endocarditis Dx Cardiac valve 16s rRNA +	<32	128	32
Patient 11	M	31-40	Classic WD Dx Duodenum tissue PAS +	<32	256	32
Patient 12	M	61-70	Classic WD Dx Gastric tissue PAS+; PCR +	64	≤32	64
Patient 13	M	81-90	Classic WD Dx Small bowel tissue PCR +	<32	≤32	64
Patient 14	M	51-60	Endocarditis Dx Cardiac valve 16s rRNA +	<32	512	128

Patient 15	M	71-80	Classic WD → Endocarditis (15 years later) Dx Cardiac valve 16s rRNA +	<32	256	128
Patient 16	M	31-40	Endocarditis Dx Cardiac valve 16s rRNA +	<32	256	256

404

405 CNS, central nervous system; CSF, cerebrospinal fluid; Dx, diagnosis; PAS, periodic acid–Schiff
406 stain; PCR+, *Tropheryma whipplei* DNA was detected by polymerase chain reaction (PCR)
407 amplification; 16s rRNA+, at least 99% homology to *T. whipplei* TW08/27 in National Center
408 for Biotechnology Information taxonomy database; WD, Whipple disease.

409

410

411 **TABLE 3:** Role of IgG titre in excluding Whipple disease (WD) in patients: serological
 412 comparison of IgG antibody titres in WD patients (n=16) and age-matched controls (n=156)

413
 414

Antibody Titres	WD	Age-matched controls	Sensitivity & Specificity	p-value [@]
IgG <64 (Test +)	11	14	Sn: 69% (95% CI 41% to 89%) Sp: 91% (95% CI 85% to 95%)	<0.001
IgG ≥ 64 (Test -)	5	142		
IgG < 32 (Test +)	9	12	Sn: 56 % (95% CI 30% to 80%) Sp: 92% (95% CI 87% to 96%)	<0.001
IgG ≥ 32 (Test -)	7	144		
IgG < 16 (Test +)	6	8	Sn: 38% (95% CI 15% to 65%) Sp: 95% (95% CI 90% to 98%)	<0.005
IgG ≥ 16 (Test -)	10	148		

415

416 [@] p-values are determined by Fisher's exact probability test; Sn: Sensitivity; Sp: Specificity

417 **FIG 2:** Algorithm for ruling out Whipple's Disease using a *Tropheryma whipplei* Indirect
 418 Immunofluorescence Assay (IFA).

