

1 **Title:**

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3 **The Accuracy of Histopathological and Cytopathological Techniques in the**
4 **Identification of the Mycetoma Causative Agents**

5

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23 **Abstract:**

24 Mycetoma is a devastating neglected tropical disease, caused by various fungal
25 and bacterial pathogens. Correct diagnosis to the species level is mandatory for
26 proper treatment. In endemic areas, various diagnostic tests and techniques are
27 in use to achieve that, and that includes grain culture, surgical biopsy
28 histopathological examination, fine needle aspiration cytological (FNAC)
29 examination and in certain centres molecular diagnosis such as PCR.

30 In this retrospective study, the sensitivity, specificity and diagnostic accuracy of
31 grain culture, surgical biopsy histopathological examination and FNAC to identify
32 the mycetoma causative organisms were determined. The histopathological
33 examination appeared to have better sensitivity and specificity.

34 The histological examination results were correct in 714 (97.5%) out of 750
35 patients infected with *Madurella mycetomatis*, in 133 (93.6%) out of 142 patients
36 infected with *Streptomyces somaliensis*, in 53 (74.6%) out of 71 patients infected
37 with *Actinoadura madurae* and in 12 (75%) out of 16 patients infected with
38 *Actinoadura pelletierii*.

39 FNAC results were correct in 604 (80.5%) out of 750 patients with *Madurella*
40 *mycetomatis* eumycetoma, in 50 (37.5%) out of 133 *Streptomyces somaliensis*
41 patients, 43 (60.5%) out of 71 *Actinoadura madurae* patients and 11 (68.7%) out
42 of 16 *Actinoadura pelletierii*. The mean time required to obtain the FNAC result
43 was one day, and for the histopathological examinations results it was 3.5 days,
44 and for grain, it was a mean of 16 days.

45 In conclusion, histopathological examination and FNAC are more practical
46 techniques for rapid species identification than grain culture in many endemic
47 regions.

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50 **Author summary**

51 In mycetoma endemic regions, the medical and health settings are commonly
52 suboptimal, and only a few diagnostic tests and techniques are available. That had
53 badly affected the patients' proper diagnosis and management and thus the late

54 presentation of patients with advanced disease. In this retrospective study, the
55 experience of the MRC on the common in use diagnostic tests in the period
56 between 1991 and 2018 is presented.

57 In this study, the sensitivity, specificity rates and diagnostic accuracy of grain
58 culture, surgical biopsy histopathological examination and FNAC to identify the
59 mycetoma causative organisms were determined. The histopathological
60 examination appeared to have better sensitivity and specificity. Furthermore, the
61 grain culture identification needs high experience, it is the tedious procedure, and
62 cross-contamination is common hence misdiagnosis is frequent. It can be
63 concluded that histopathological examination and FNAC are more practical
64 techniques for rapid species identification than grain culture in many endemic
65 regions with poor diagnostic setting.

66

67

68 **Introduction:**

69 Mycetoma is a chronic granulomatous subcutaneous inflammatory infection
70 endemic in subtropical and tropical regions, but it is reported globally [1,2]. It is
71 characterised by a painless subcutaneous swelling, multiple sinuses formation and
72 a discharge that contain grains [3,4]. The clinical presentation can give a clue to
73 the diagnosis, but without further diagnostic testing it will to misdiagnosis and
74 inaccurate treatment [5]. Mycetoma can be caused by different bacteria
75 (actinomycetoma) or fungi (eumycetoma) [6,7]. More than 70 different micro-
76 organisms were reported to cause this infection, and hence it is essential to identify
77 the causative agents to the species level for appropriate treatment [8,9]. In
78 endemic regions, the most commonly used tools are culturing of the grains,
79 surgical biopsy followed by histopathological examination and fine needle
80 aspiration cytological (FNAC) examination [10,11].

81 Currently, culturing the grains culture is still considered to be the golden standard
82 for species identification in many centres [12,13]. However, this technique is
83 tedious, time-consuming due to the slow growth rate and it needs expert
84 microbiologists to identify the causative agents based on the macroscopic

85 appearance of the isolates. Furthermore contamination is common. Patients on
86 medical treatment may have non-viable gains, and hence it is difficult to identify
87 the causative organism [14,15].

88 To overcome these difficulties, histological examination is often used
89 complementary to culture. In a histopathological examination, it is easy to
90 discriminate between fungal and bacterial causative agents [16,17]. However,
91 identification to the species level is more challenging and considered far from
92 reliable [18,19]. At the Mycetoma Research Centre (MRC), University of Khartoum,
93 Khartoum, Sudan FNAC is a common tool to identify the causative organisms. It is
94 less invasive, and time-consuming compared to the histopathological and culture
95 techniques [20,21]. However, to the best of our knowledge there was no study
96 performed in which the sensitivity and specificity of the three techniques for the
97 identification of the mycetoma causative organisms were compared. With this
98 background, this study was conducted at the Mycetoma Research Centre were
99 8500 confirmed mycetoma patients were seen and treated. In this retrospective
100 study, the records of these patients were reviewed, and patients who undergone
101 the three diagnostic tests were included.

102

103

104 **Materials and Methods**

105 **Study cohort**

106 Following the Mycetoma Research Centre Institutional Review Board ethical
107 approval, all the histopathological, cytological and microbiological reports of the
108 patients seen in the Mycetoma Research Centre over a 27-year period (January
109 1991 to January 2018) were reviewed. The data were collected in the pre-designed
110 data collection sheet. The data analysis was anonymised The patient demographic
111 characteristics, results of the three techniques were collected.

112 .

113 In this study, only patients in whom the causative organisms were identified by
114 culture and had undergone both a fine needle aspirate for cytological examination

115 and deep-seated excisional biopsy for histopathological examination were
116 included. (Fig. 1).

117

118 The diagnostic accuracy of these techniques for identification of the causative
119 agents was calculated as the percentage of cases with culture identification divided
120 by the number of cases with species identified by histopathological or cytological
121 techniques.

122

123

124 **Grains isolation**

125 The grains were obtained by surgical biopsy and/or FNAC. For the latter, a 25-
126 gauge needle was inserted into the lesion, and aspirates were taken. The yield of
127 grains was assessed visually by the number and size of grains obtained. If the
128 yield was low, a second aspiration was taken with a 23-gauge needle. When
129 excessive bleeding from the lesions was encountered, a 27-gauge needle was
130 used. The obtained sample usual spiltted into two parts; one was transported
131 immediately to the microbiology department for culturing, the other part was sent
132 to the histology department for histopathological and cytological.

133

134 **Grains culture**

135 The mycetoma grains were washed three times in sterile normal saline and cultured
136 on Sabaroud dextrose agar with gentamicin for fungal grains. The actinomycetoma
137 gains were cultured in Yeast Extract agar. Grains were incubated at 37°C and growth
138 was checked daily. The isolates were identified by their microscopical appearance
139 and biochemical testing.

140

141

142 **Cytological Examination**

143 The aspirate was allowed to air dry and was stained using Diff-Quick stain. The
144 stained aspirates were examined by an expert histopathologist for the presence of
145 the following cytomorphological features: smears cellularity, the host inflammatory

146 tissue reaction, the presence and types of the causative organisms' grains. Species
147 identification was based on species-specific criteria. In general, *M. mycetomatis*
148 grains can be either small or large, are light to dark brown in colour and have
149 irregular outlines and a crushing artefact when stained with hematoxylin and eosin
150 (H&E) (Fig. 2A).

151 *S. somaliensis* grains are difficult to see in H&E stained sections, they stain bright
152 pink to hazy pink in colour, are often oval to irregular shaped and can be as
153 aggregates (Fig. 2C).

154 *A. madurae* grains are small oval shaped, and it stained pink to red colour in H&E
155 and tend to be as one mass without any fractures. *A. pelletierii* grains are small
156 rounded to oval shaped, and they stained deep blue in H&E stained sections and
157 tend to be fractured.

158

159 **Histopathological Examination**

160 All patients underwent surgical biopsy under anaesthesia, which was fixed in 10%
161 formalin and processed further into paraffin blocks. 3-5- μ m sections were obtained
162 and stained with H&E. Species identification was made based on species-specific
163 criteria. *M. mycetomatis* grains tend to be large, light to dark brown in colour with
164 irregular outlines. They tend to fracture when sections are cut. *M. mycetomatis* has
165 two different types of grains, and these are the filamentous and vesicular. The
166 filamentous type, is the most common type and consists of brown septated and
167 branched hyphae that may be slightly more swollen towards the edges (Fig. 3D).

168

169 *S. somaliensis* grains are rounded to oval in shape, with homogenous appearance
170 in tissue sections. They appear faint yellow in unstained sections, and the grains are
171 not well stained with H&E. Moreover, as a result of sectioning they may show
172 longitudinal cracks, the filaments are fine (measured between 0.5 – 2 μ m in
173 diameter), closely packaged and embedded in cement matrix (Fig. 3B).

174

175 *A. pelletierii* grains are small, round to oval in shape and semicircular and sickle like
176 shapes have been observed as well. The filamentous structures are pretty difficult

177 to be detected. However, a careful and meticulous examination of the periphery of
178 the grains may show some of them. *A. pelletierii* grains stain deep violet with H&E,
179 which is very characteristic and allows the definitive diagnosis without a need for
180 culturing techniques (Fig. 3C).

181

182 *A. madurae* grains ranged from yellow to white. Therefore, it can be difficult to
183 discriminate them from the surrounding fat. Histologically the grain size ranges from
184 small to large. The large grains have a characteristic variegated pattern. The
185 periphery of the grain is opaque, homogenous and deep purple when stained with
186 H&E stain, while the centre is less densely stained. Additionally, the periphery of the
187 grains shows an eosinophilic material (Fig. 3A). Smaller grains are more
188 homogeneous and are difficult to distinguish from *A. pelletierii*. However, even the
189 small grains of *A. madurae* have a more deeply stained purple fringe, which is not
190 seen in *A. pelletierii*.

191

192

193 **Results:**

194 In this study, 991 patients out of 7940 patients were eligible and were included in
195 the analysis. Their ages ranged between 5 and 75 years old. The majority were
196 males 737 (74.3%), and most of them were students 327 (32.9%) and farmers 167
197 (16.8%). The majority of the patients (837 out of 991), gave a history of discharge
198 that contained grains and the majority of these grains were black (565; 57%)
199 followed by yellow (104;10.5%), white (60; 6.1%) and red grains (14; 1.4%). In this
200 cohort, the majority of patients, (72.6%) had no history of local trauma, only
201 (19.3%) patients did recall a local trauma and the remaining 73 (7.4%) patients were
202 not certain.

203

204 Based on the culture reports of the grains, in 750/991 (75.6%) of the patients the
205 mycetoma was caused by *M. mycetomatis*, in 142/991 (14.4%) it was caused by *S.*
206 *somaliensis*, in 71/991 (7.16%) it was caused by *A. madurae* and in 16/991 (1.6%)
207 it was caused by *A. pelletieri*. In 11 patients no growth was reported from the grains

208 obtained during the sample collection. The time to growth differed case by case and
209 ranged between 5 and 28 days.

210

211 In this study, out of the 991 mycetoma cases, the correct species identification was
212 obtained for 912 cases using histopathological examination. Using FNAC, the
213 correct diagnosis was obtained in 708 cases. The histopathological examination
214 confirmed the diagnosis of *M. mycetomatis* in 714 of 750 cases with 95.2%
215 sensitivity, 95.4% specificity and diagnostic accuracy of 95.3%. For FNAC only 604
216 out of 750 *M. mycetomatis* cases were identified, resulting in a sensitivity of 80.5%,
217 a specificity of 88.4% and a diagnostic accuracy of 82.4%.

218 Out of 142 *S. somaliensis* cases, 133 were also identified with histopathological
219 examination with 93.7% sensitivity, 98.9% specificity and diagnostic accuracy of
220 98.2%. With FNAC only 50 out of 133 *S. somaliensis* cases were identified, resulting
221 in a sensitivity of 35.2%, a specificity of 99.3% and a diagnostic accuracy of 90.1%.
222 53 out of 71 cases with *A. madurae* identification were identified by
223 histopathological examination, with a sensitivity of 74.7%, 99.5 % specificity and
224 diagnostic accuracy of 97.7%. FNAC identified 43 out of 71 cases with a sensitivity
225 of 60.6%, specificity of 94.4 % and diagnostic accuracy of 91.9%.

226

227 For *A. pelletierii* out of 16 cases; 12 were also identified with histological examination
228 with 75.0% sensitivity, 100% specificity, and diagnostic accuracy of 99.6%. For
229 FNAC a sensitivity of 68.8%, a specificity of 99.7% and a diagnostic accuracy of
230 99.2% were obtained.

231

232 With the histopathological examination, false negative result was reported in 36/750
233 *M. mycetomatis* cases, 9/142 *S. somaliensis* cases, 18/71 *A. madurae* cases and
234 4/16 *A. pelletierii* cases. To determine the false negative results reasons, the
235 histopathological slides were re-examined. There were various reasons for the false
236 negative, and that included the absence of mycetoma histopathological architecture
237 resulted in overlooking the causative agent (Fig 4). Furthermore, in some blocks, the
238 grains were absent, either because the tissue was not homogenously infected by

239 the causative agent and that the part which was taken for histology or the section
240 contained no grains.

241 This latter might be overcome by examining multiple sections at different depths of
242 the histology blocks especially when inflammation and necrosis are noted.

243

244 False positive results were obtained in 28 of the cases. This was attributed to the
245 presences of numerous structures that can mimic the appearance of *M. mycetomatis*
246 and that included vegetables, synthetic fibres and algae (Fig 5) which can resemble
247 fungal hyphae and calcification (Fig 5.). In overall, using histology correct species
248 identification was obtained in the majority of cases.

249

250 The mean time to identify the culture isolates was 16 days (range 5 to 28 days), for
251 histology it was 3.5 days (range 2 to 5 days), and for cytology, it was one day (range
252 1 to 2 days). This demonstrated that reliable species identification using histology
253 was obtained in 92.0% of cases within an average time reduction of 13.5 days, for
254 cytology this was 71.4% of cases with time reduction of 15 days, indicating that
255 adding histology or cytology to the diagnostic techniques used for species
256 identification resulted in an earlier start of treatment.

257

258 **Discussion:**

259 The accurate identification of mycetoma causative agents is considered the
260 cornerstone for the initiation of appropriate therapy. Hence a rapid and accurate
261 diagnostic tool to achieve the definitive species identification is considered a critical
262 part in patient treatment and management [21-23]. Different laboratory techniques
263 for species identification are in use, including culture, histopathology, [7, 25], FNAC
264 [8, 24], serological assays and imaging [26 - 29] as well as different molecular
265 diagnostic tools [30 - 35]. However, not all these assays are available in endemic
266 regions. In the Mycetoma Research Centre, culturing of the grains, histopathology
267 and FNAC are routinely performed and have been used for the past 27 years. In this
268 communication we have used the data collected for the last 27 years to assess the
269 sensitivity, specificity and diagnostic accuracy of histopathology and cytology in the

270 identification of mycetoma causative agents in comparison to the current golden
271 standard: culturing.

272 This study showed that the histopathology was more accurate to FNAC in terms of
273 species identification. Our results are in line with that reported previously by Yousif
274 and colleagues [36]. They reported 90.9% agreement when histopathology was
275 compared to FNAC for the diagnosis of *M. mycetomatis* (90.9%) while for
276 actinomycetoma causative agents it was only 60%. The lower diagnostic agreement
277 of actinomycetoma causative agents could have been caused by morphological
278 similarities of these microorganisms. Furthermore, both techniques are operator
279 dependent and need intensive training and experience which could have its
280 reflections on the accuracy.

281

282 Mycetoma can be caused by more than 70 different causative agents [37], but the
283 distribution of these species is not everywhere the same which could cause
284 differences in diagnostic accuracy in different regions. In some of the mycetoma
285 endemic regions, mycetoma is caused by closely related species. Morphologically
286 these organisms may look similar which could cause a challenge in the identification
287 of these organisms. In Mexico, the most common causative agents are *Nocardia*
288 *brasiliensis* and *Nocardia asteroides* [37], two closely related species which are
289 difficult to differentiate from each other based on histopathology [38, 39]. In Senegal,
290 the most common causative agents of eumycetoma are *M. mycetomatis* and
291 *Falciformispora senegalensis* which both can cause black grain mycetoma [37, 40].
292 In the black grains of *F. senegalensis*, the centre is non-pigmented, and the cement
293 is absent, whereas at the peripheries the grains are dark coloured and brown cement
294 is present. However, this is also seen in black grains of *Trematosphaeria grisea* and
295 certain grains of *M. mycetomatis*. Hence an expert pathologist is needed to
296 differentiate between these organisms [41].

297

298 The study performed here was a retrospective study, looking back at the records of
299 the Mycetoma Research Centre for the past 27 years. During that time molecular
300 identification of the causative agents was not performed and culture was considered

301 the golden standard. Recently in the study conducted by Borman and colleagues
302 demonstrated that using morphological identification, misidentifications occurred in
303 many cases [42]. Out of 28 previously identified *Trematosphaeria grisea* isolates, 22
304 were, other fungal species [42]. For actinomycetoma causative organisms,
305 misidentifications also have been described. In 2008, Quintana and associates
306 demonstrated that half of the *S. somaliensis* isolates obtained from Sudan appeared
307 to be *Streptomyces sudanensis* [43]. Furthermore, next to *A. madurae* and *A.*
308 *pelletieri* also *Actinomadura latina* was described [44].

309

310 In this study, the sensitivity of histopathological technique was superior to that FNAC
311 for all species tested. In that study they studied the performance of FNAC in
312 comparison to histology in 19 different mycetoma patients. Out of these 19 patients,
313 five patients had to be excluded due to inadequate aspirated materials. From the 14
314 remaining patients, 10 were diagnosed as *M. mycetomatis* with histopathology, and
315 4 were actinomycetoma. With this limited number of patients, they could conclude
316 that FNAC could identify the causative agent in 9 out of 10 *M. mycetomatis* patients.
317 One patient identified by histology could not be identified with cytology, again
318 confirming that histology was superior to FNAC in respect to species identification
319 [15]. A result confirmed in our current study, as in our study 146 patients with *M.*
320 *mycetomatis* mycetoma were missed with FNAC. However, of the three different
321 identification methods used, FNAC was the most rapid and resulted in a species
322 identification within 1 day, instead of 3.5 days for histology or 16 days for culture.
323 FNAC is a simple and rapid diagnostic technique which can be used at the one-stop
324 diagnosis clinic and in epidemiological and field surveys. However, it has many
325 limitations: it is an operator dependent technique, can be painful and can lead to
326 deep-seated bacterial infection. At the moment the fine needle aspirate is often taken
327 blindly without guidance of ultrasound imaging which creates a risk that the operator
328 might miss the pockets which contains grains. With the use of the ultrasound-guided
329 aspiration, the diagnostic yield of the technique will improve which in its turn could
330 enhance the number of cases in which a positive species identification might be
331 obtained.

332

333 The grains culture remains in many centres the cornerstone for the diagnosis of
334 mycetoma. However it is a time-consuming procedure and an experienced
335 microbiologist is needed to identify the organisms to the species level. Cross-
336 contamination is a common problem. However, by complementing culturing with
337 histology or FNAC a preliminary identification might be obtained earlier.

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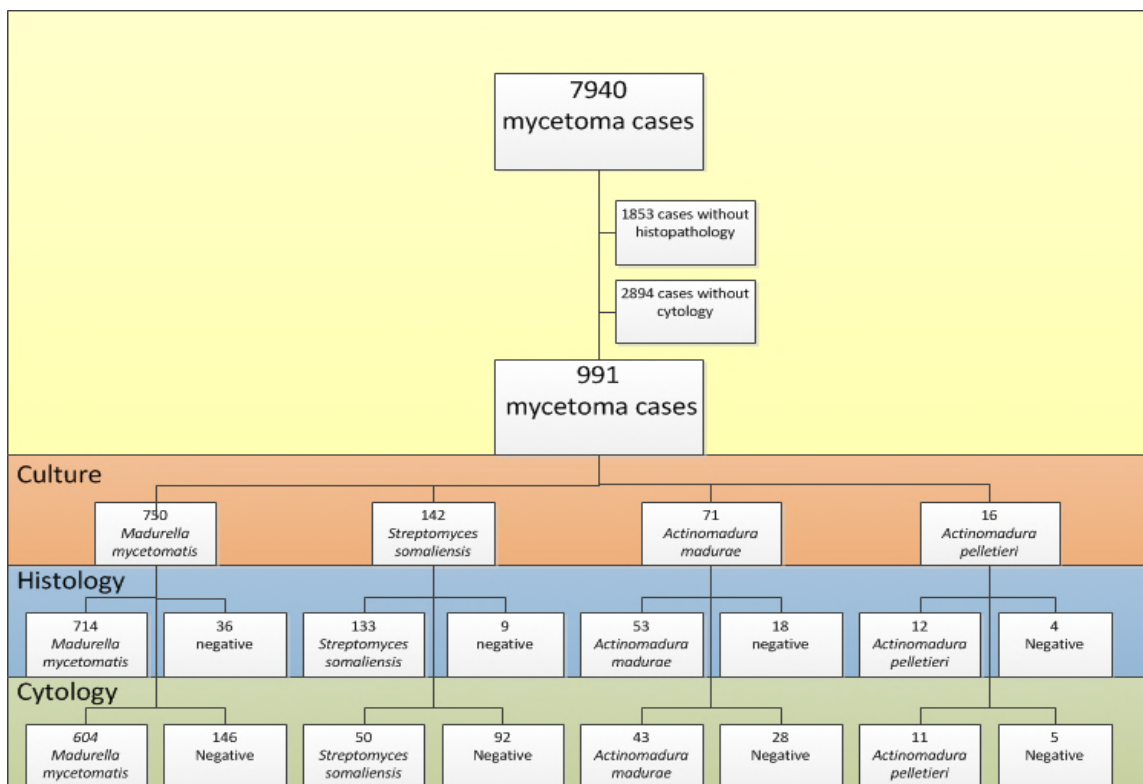
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481 Fig 1: the study data collection method

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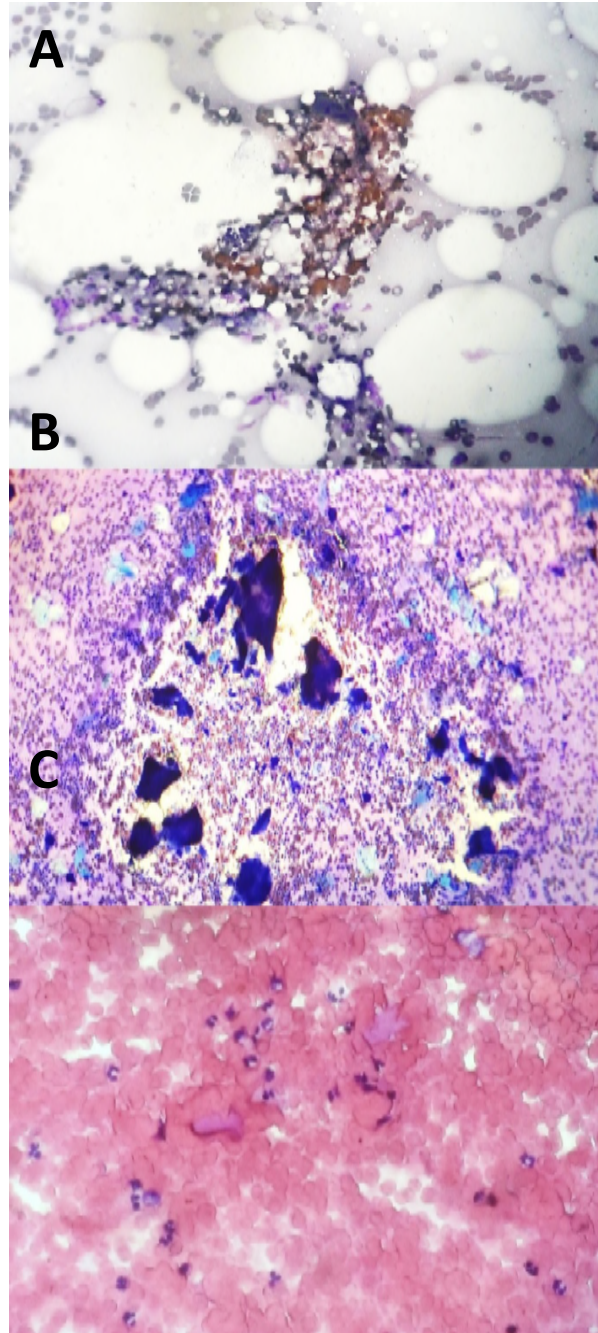
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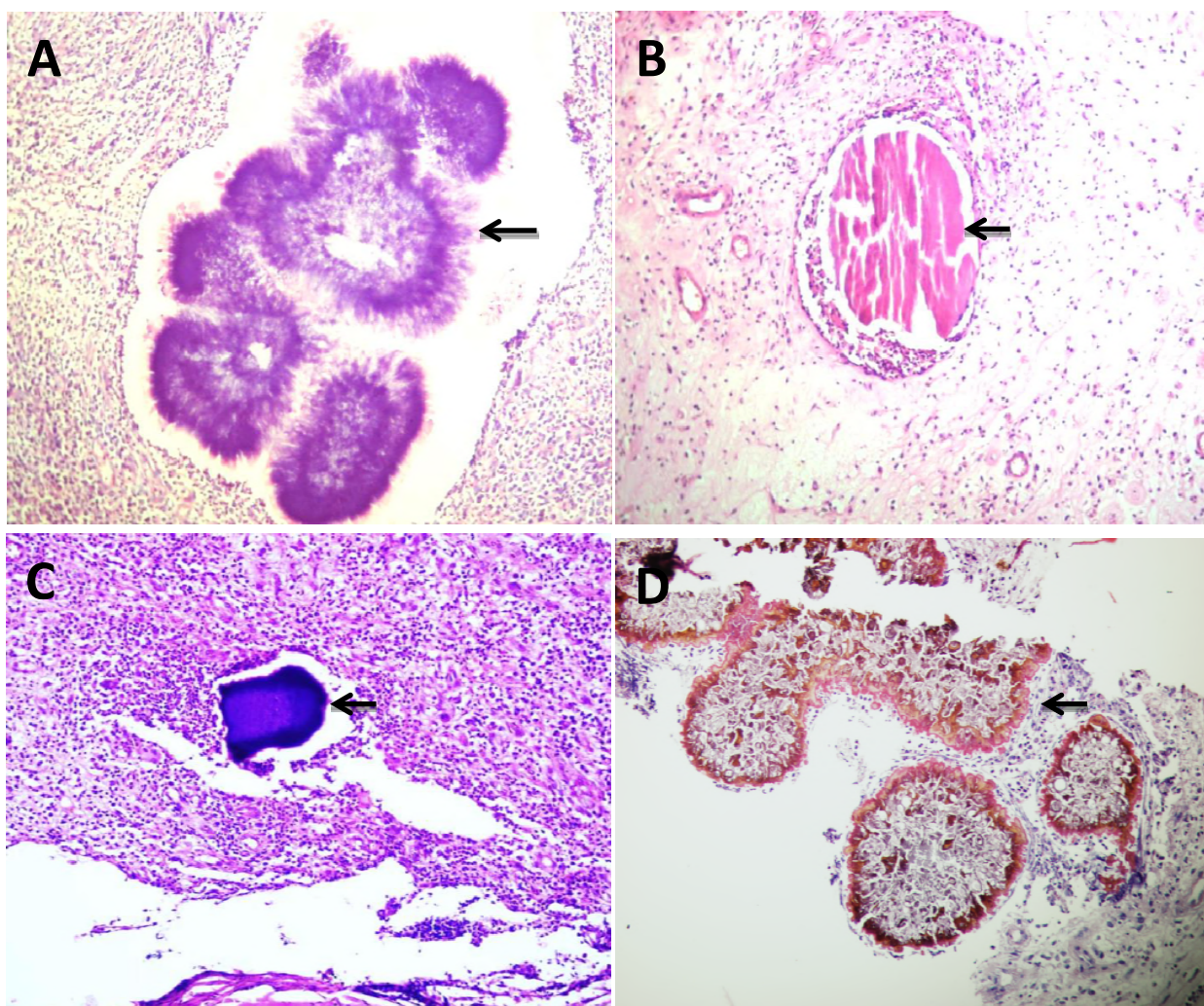
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519 Fig 2: photomicroscopy of the cytopathological appearance of some mycetoma
520 causative agents. *M. mycetomatis* (A), *A. pelletierii* (B), *S. somaliensis* in fine (H&E
521 X10)

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525 Fig 3. Histopathological appearance of mycetoma causative agents; *A. madurae*

526 (*A*), *S. somaliensis* (*B*), *A. pelletierii* (*C*) and *M. mycetomatis* (*D*). (H&E X10).

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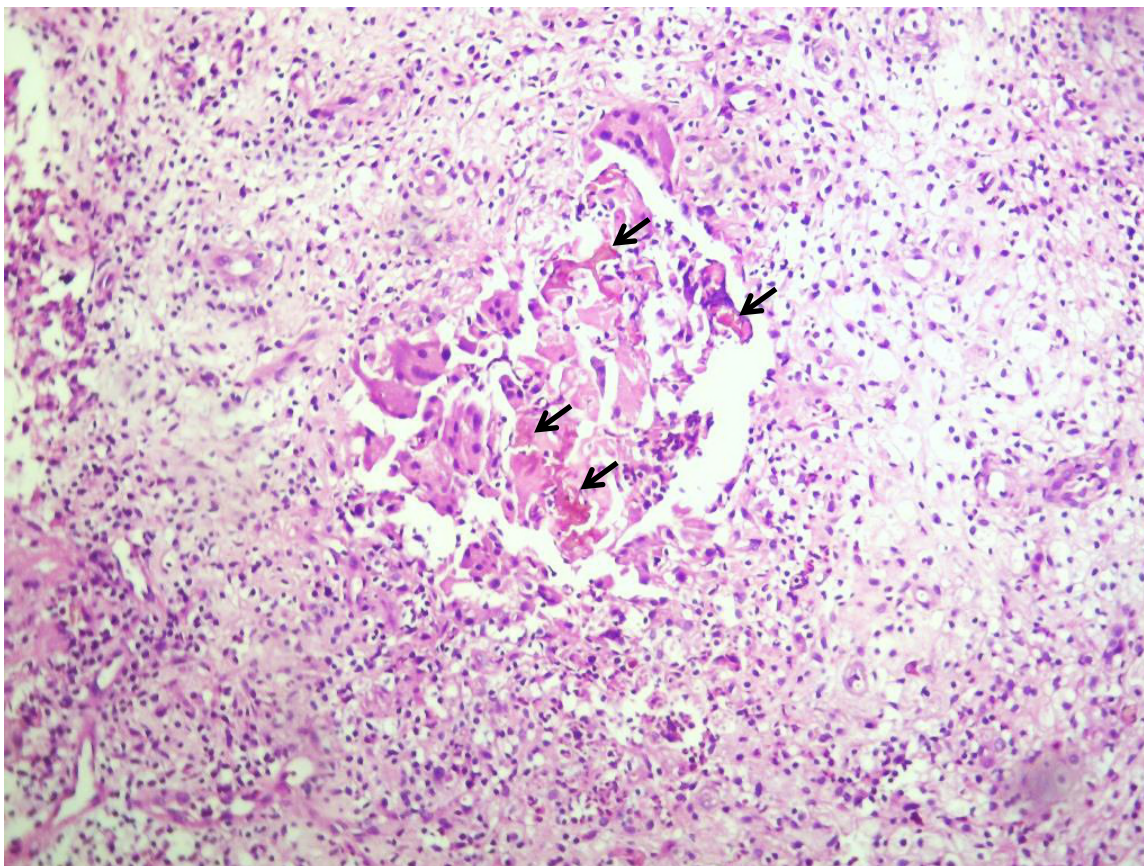
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535 Fig 4: photomicroscopy showing histopathological section with histiocytes at the
536 centre with sparse amount of *M. mycetomatis* the grains (arrows). It was reported
537 first as false negative. (H&E X 10).

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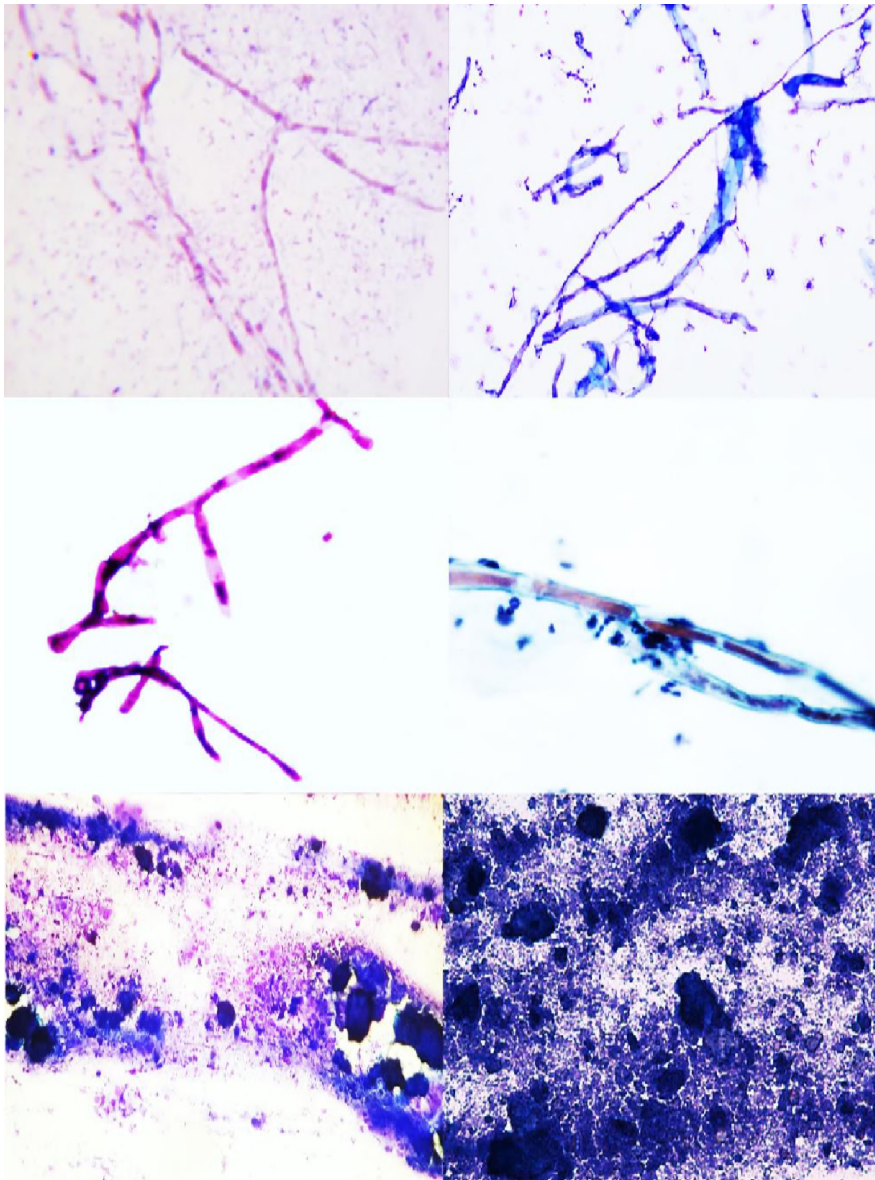
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567 Fig 5: false positive identifications in cytology.

568 Showing cytological smears of (A) *M. mycetomatis* hyphae stained with Giemsa

569 stain, (B) Synthetic Fibers stained Wright-Giemsa stain. (Giemsa stain, X40). (C)

570 Smear showing hyphae of *M. mycetomatis*. (D) Smear showing elongated

571 structures of the Oedogoniales order. The chloroplasts form a chain interrupted by

572 clear zones (X40). (E) Smear showing *M. mycetomatis* grains after being crushed

573 on the smear. (F) smear with abundant calcific debris without intact cells taken

574 from patient with tumoral calcinosis (Diff Quick, X10).

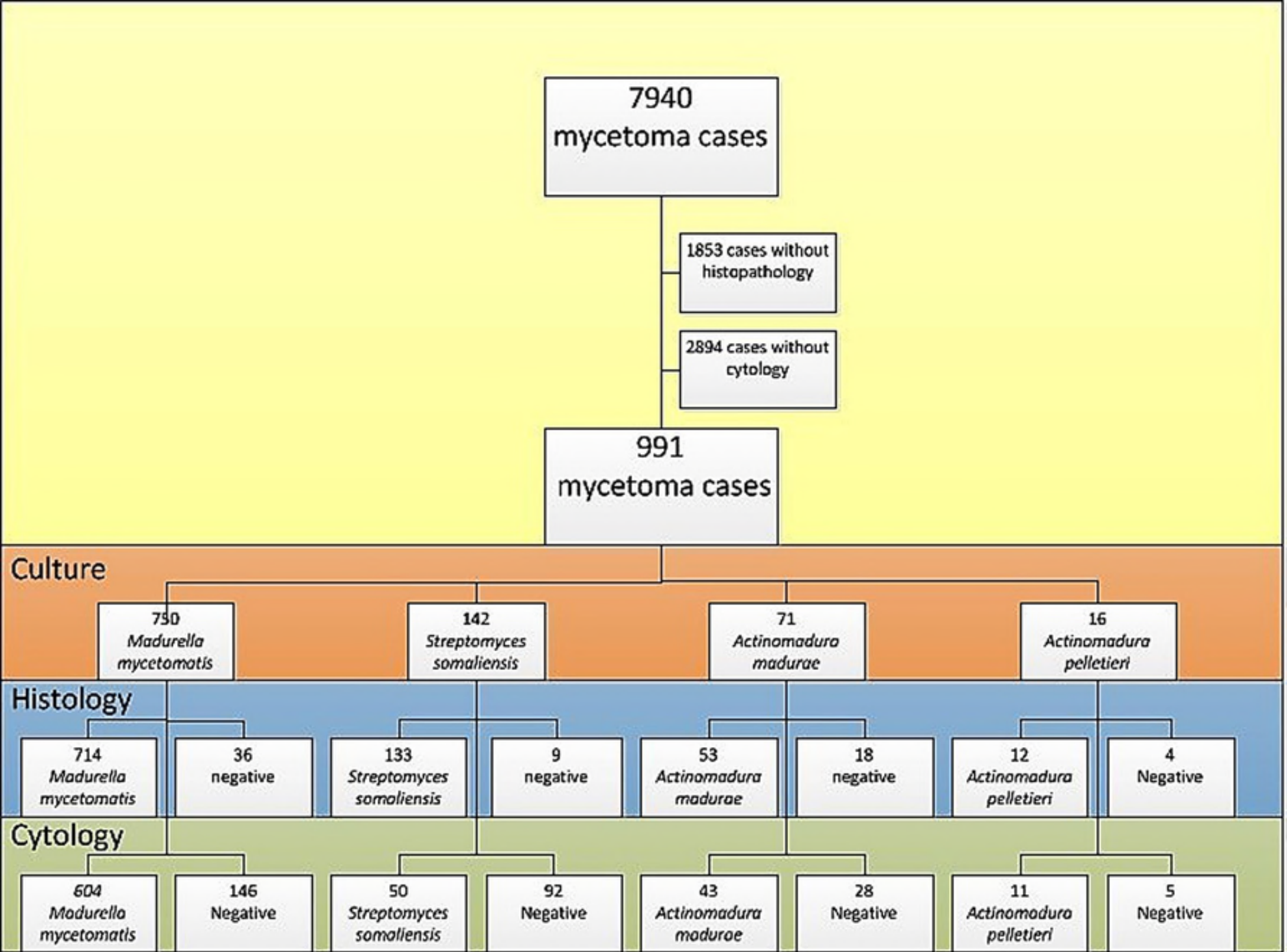


Fig 1

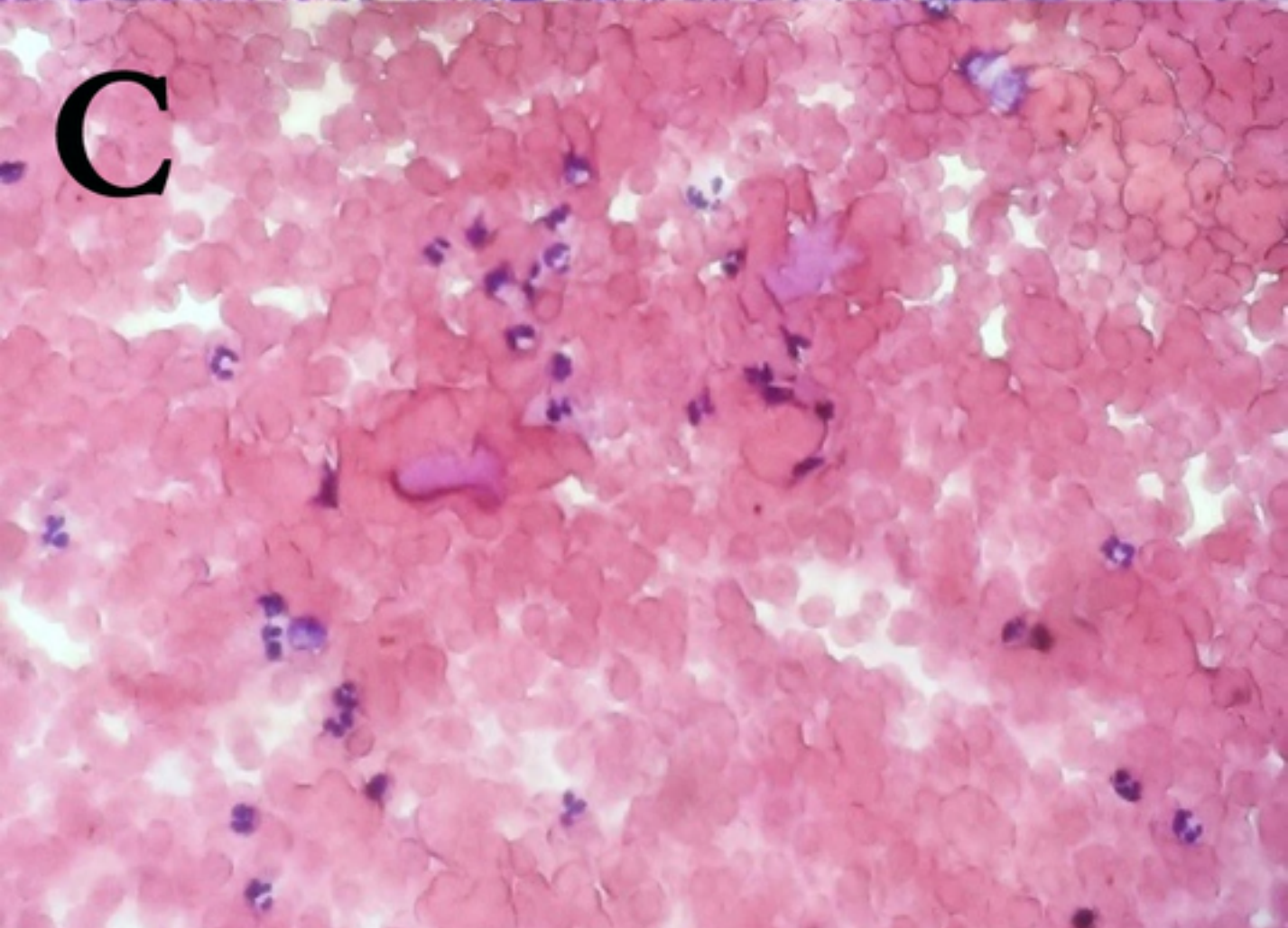
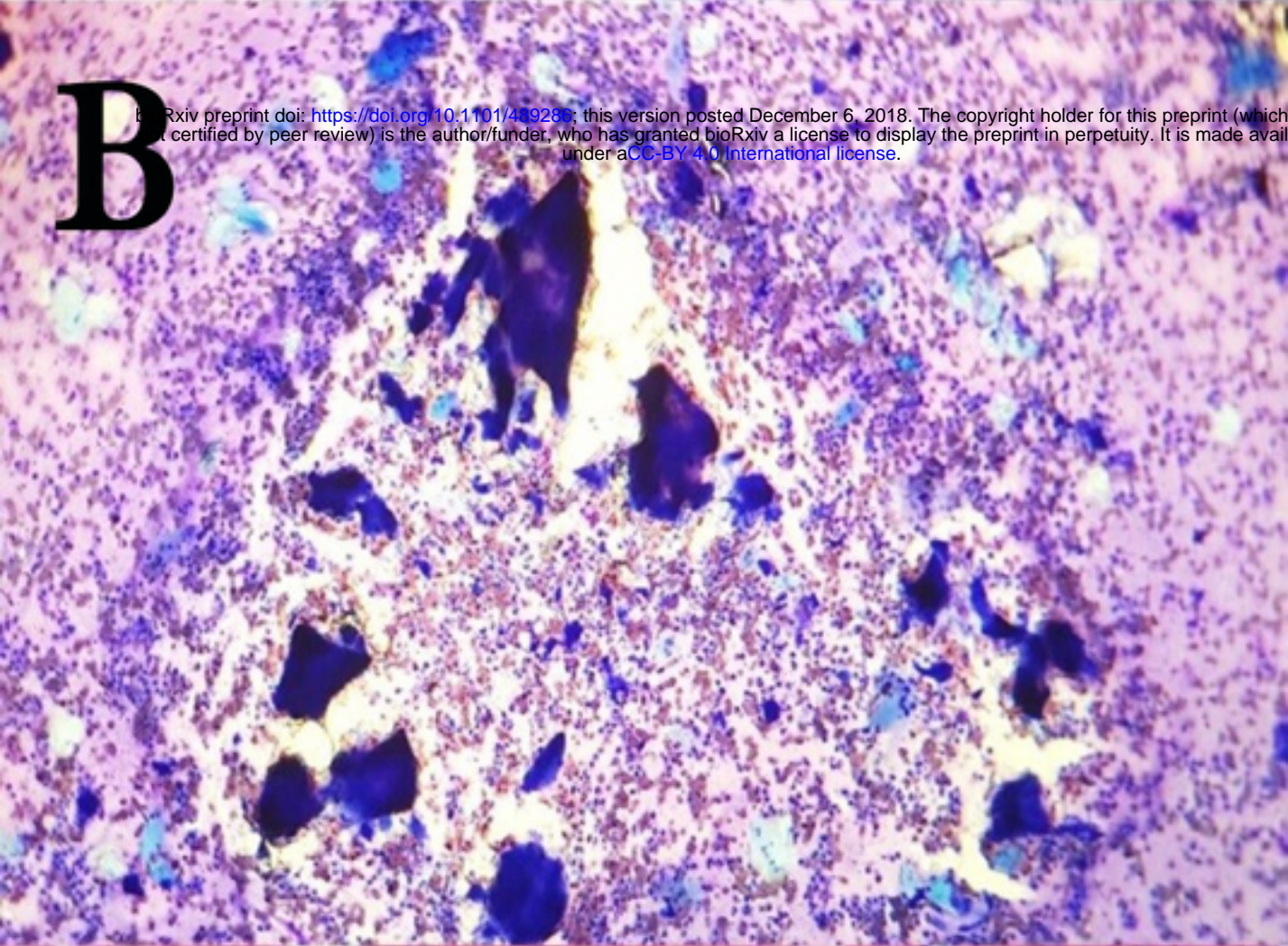
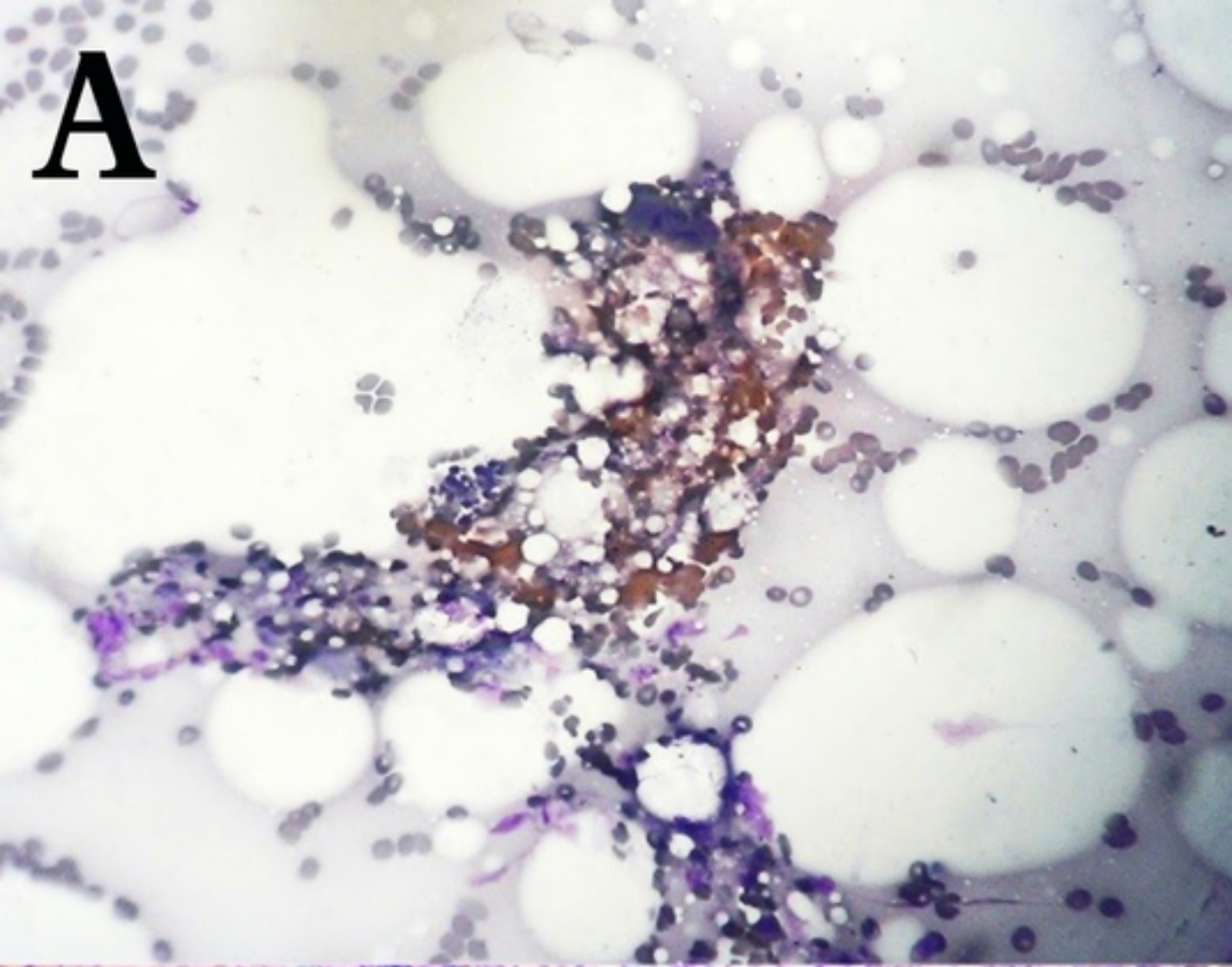


Fig.2

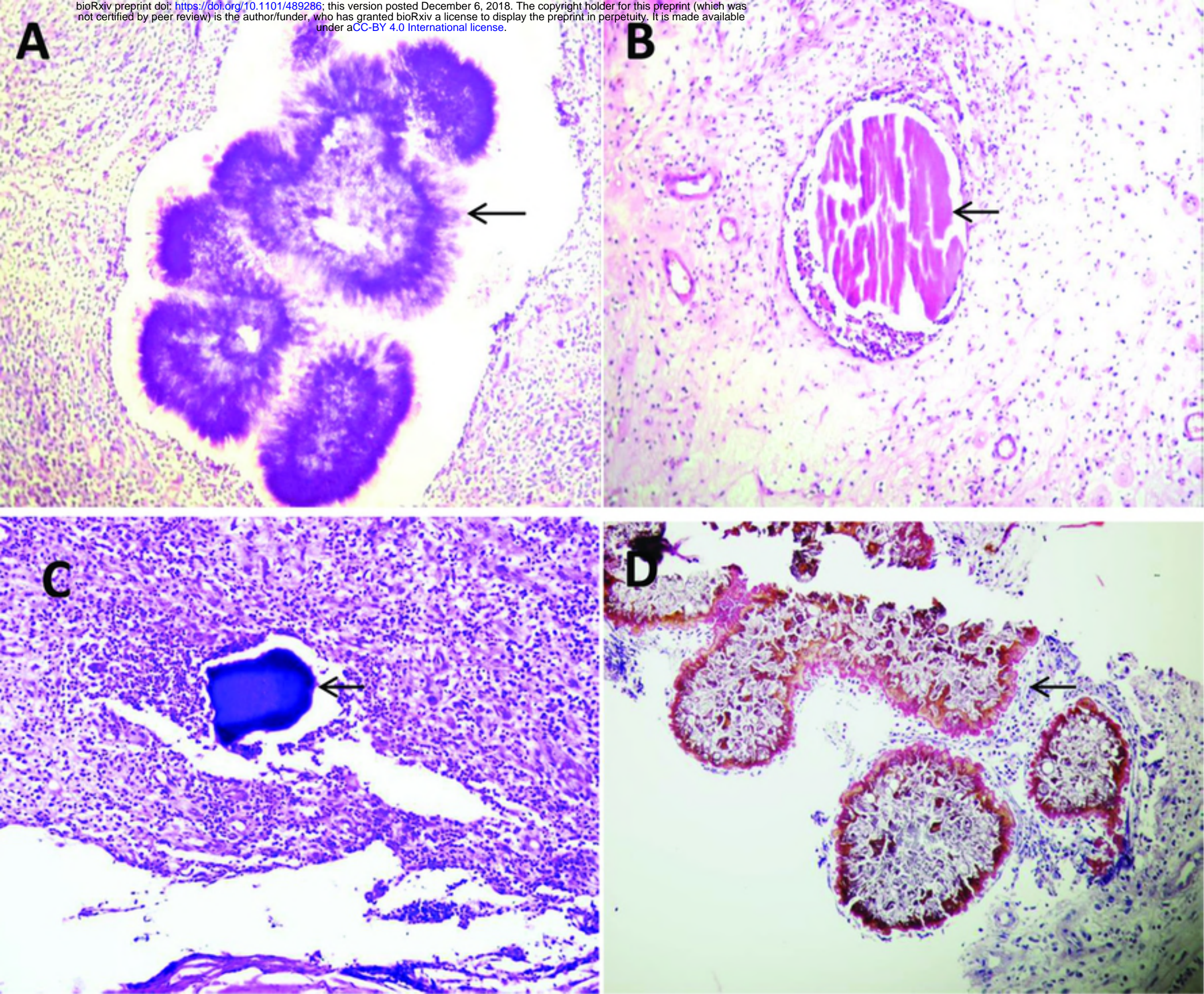


Fig. 3

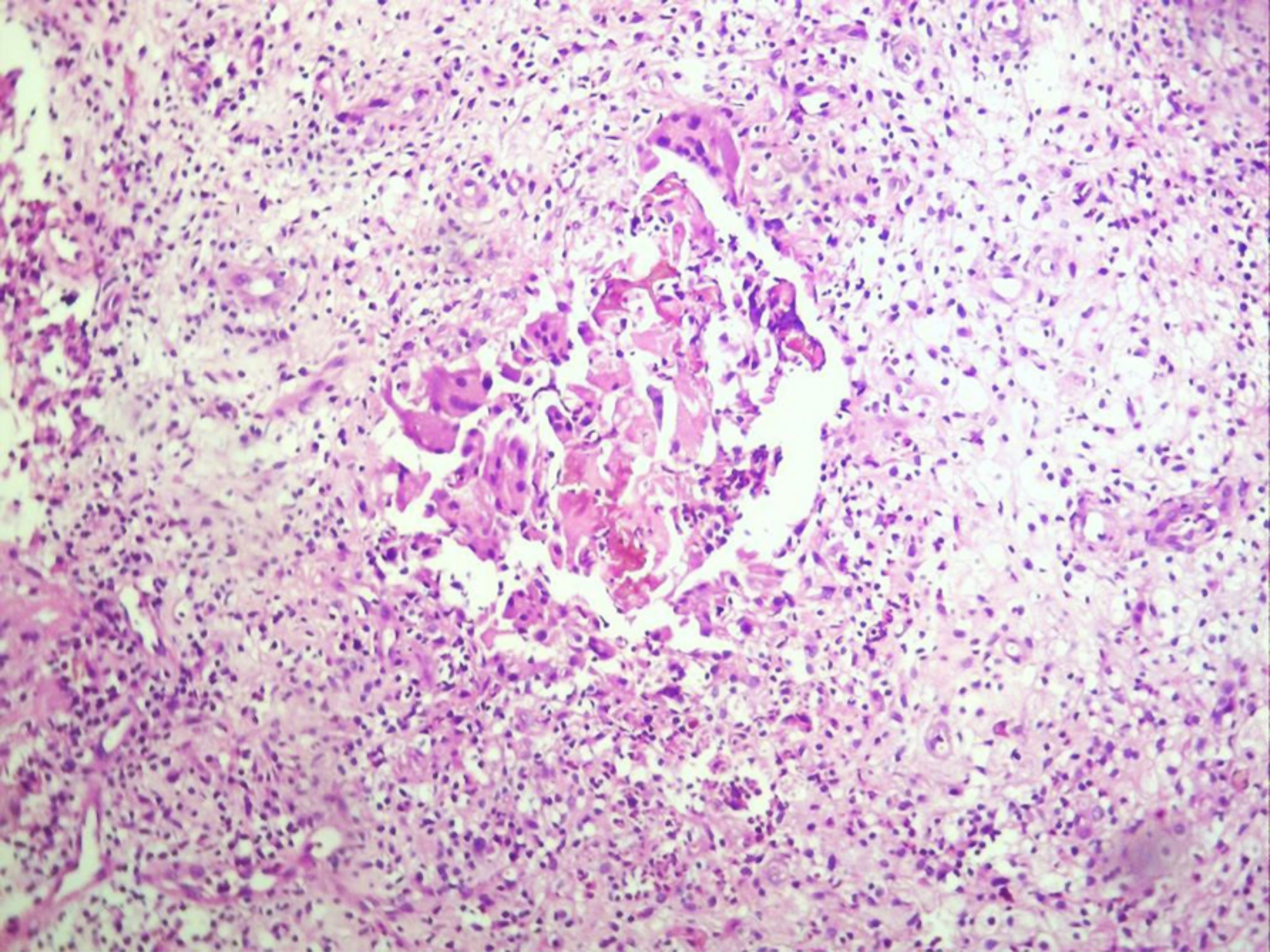
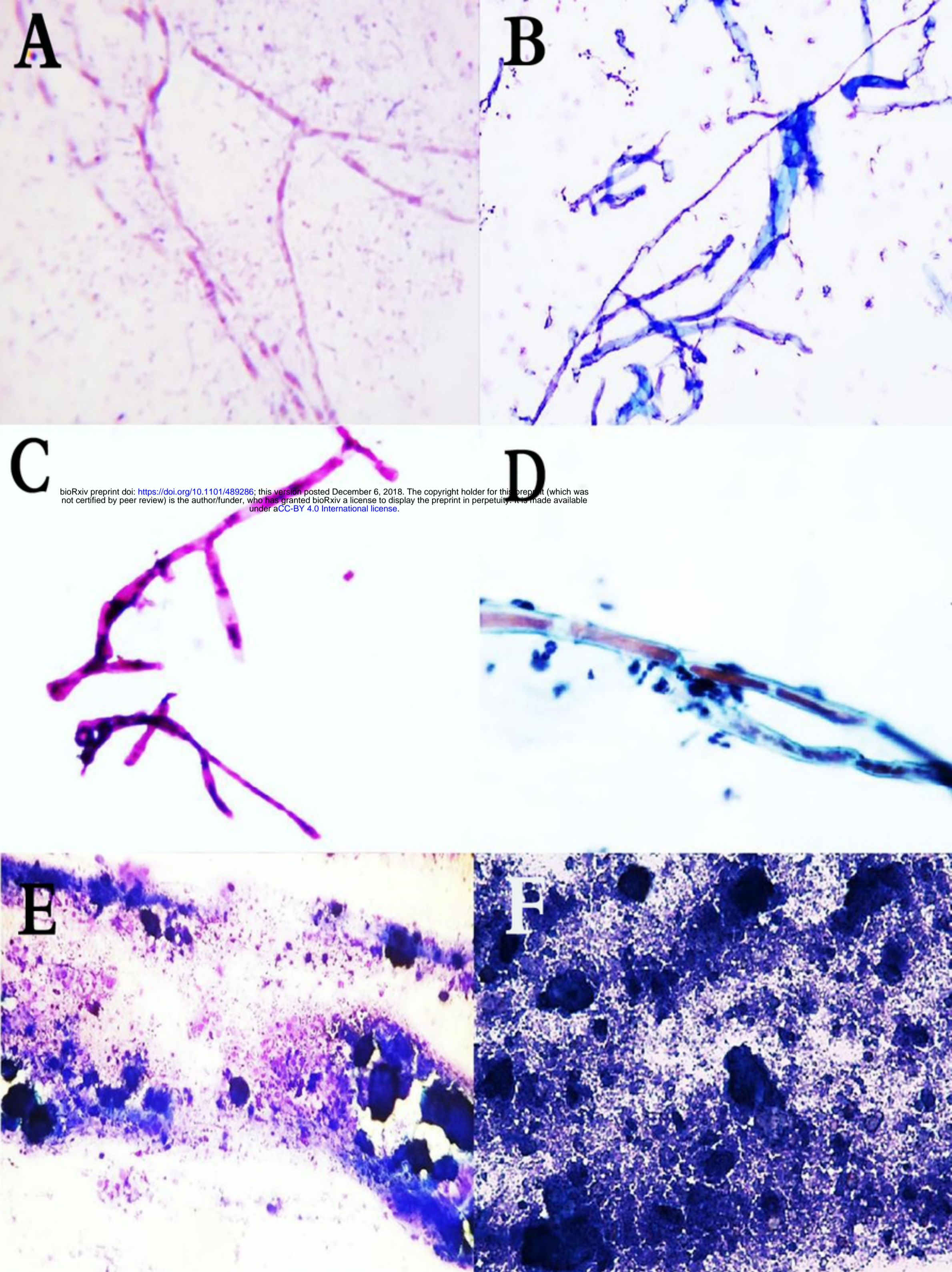


Fig.4



Figure