1 High prevalence of the digenean *Plagiorchis* sp. in the wood mouse *Apodemus*

2 sylvaticus in a periaquatic ecosystem.

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21 SUMMARY

The prevalence of the digenean *Plagiorchis* sp. was investigated in a natural wood 22 mouse population (Apodemus sylvaticus) in a periaquatic environment. Classical 23 24 identification was complemented with the use of molecular differentiation to determine prevalence and verify species identity. Use of the complete ITS1-5.8S rDNA-ITS2 and 25 partial 28S rDNA gene sequences have confirmed that the species reported at this 26 location was *Plagiorchis elegans* and not *Plagiorchis muris* as reported previously, 27 illustrating the difficulties in identification of these morphologically similar parasites. P. 28 elegans is typically a gastrointestinal parasite of avian species but has also been reported 29 from small mammal populations. The occurrence of this digenean in A. sylvaticus in the 30 UK is rare however in the area immediately surrounding Malham Tarn, Yorkshire, it had a 31 high prevalence of 23% and a mean worm burden of 26.6±61.5. The distribution of P. 32 elegans followed a typically overdispersed pattern and both mouse age and sex were 33 determined to be two main factors to be associated with prevalence. Male mice harboured 34 the majority of worms carrying 688 of 717 recovered during the study and had a higher 35 prevalence of 32.4% in comparison to only 8.7% in the small intestine of female mice. A 36 higher prevalence of 43% was also observed in adult mice compared to 14% for young 37 adults. No infection was observed in juvenile mice. These significant differences are likely 38 to be due to differences in the foraging behaviour between the sexes and age cohorts of 39 wood mice. 40

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Key words: Plagiorchiidae; *Plagiorchis muris*; *Plagiorchis elegans*; wood mouse; *Apodemus sylvaticus*; DNA; aquatic environment.

Wild rodent populations are commonly examined for their helminth assemblages 47 48 and studies conducted worldwide have revealed a plethora of nematode, cestode and trematode species being harboured by various rodent fauna. In the UK, several commonly 49 identified helminth species have been documented in rodents from different sites. The 50 most frequently reported digeneans of the wood mouse Apodemus sylvaticus tending to 51 be Corrigia vitta and Brachylaemus recurvum typically found infecting the pancreatic ducts 52 and the small intestine of their host respectively (Elton et al., 1931; Lewis, 1968; Lewis and 53 Twigg, 1972; Behnke et al., 1999; Abu-Madi 2000). The parasites of A. sylvaticus and 54 other small mammals have been extensively studied in populations located in a 55 periaquatic environment at the Malham Tarn Nature Reserve, North Yorkshire, UK, 56 including the discovery of a new parasite species (Allan et al 1999; Hughes et al 2006, 57 2008; Rogan et al 2007; Hide et al 2009; Thomasson et al 2011; Boyce et al 2012). 58 Specifically, Rogan et al., (2007) reported the occurrence of the intestinal digenean 59 Plagiorchis muris in the wood mouse A. sylvaticus over a 13-year period with an overall 60 prevalence of 16.9%. 61

The occurrence of *P. muris* within the UK is rare. As far as can be determined, the 62 first report of *P. muris* in the UK occurred in Oxford during a study on the health of a wild 63 mouse population which took place from September 1925 to January 1928 (Elton et al., 64 1931). During this study the digenean Lepoderma muris (syn. P. muris) was described 65 from the small intestine of the wood mouse A. sylvaticus at a very low prevalence of 0.1%. 66 67 L. muris was furthermore reported in the brown rat Rattus norvegicus in Cambridgeshire, UK (Baylis, 1939) and later an occurrence of *P. muris* was recorded in Scotland in 1963 68 when the digenean was unexpectedly recovered from the intestine of a Scottish Hill sheep 69 during a parasitological necropsy (Fahmy and Rayski, 1963). These studies, including 70

those of Rogan *et al.*, (2007) were all based on classical parasite identification using morphology and have not benefited from the greater precision available for DNA sequencing analysis.

74 Members of the genus Plagiorchis are cosmopolitan and tend to demonstrate low definitive host specificity. Species of this genus have been previously described from the 75 intestines of reptiles and birds in addition to mammals (Janssen and Bock, 1990; Biserkov 76 and Kostadinova, 1998; Ito and Itagaki, 2003). P. muris is no exception. This species was 77 originally described from the small intestine of the black rat, Rattus rattus and the brown 78 rat, Rattus norvegicus by Tanabe in Kyoto, Japan (Tanabe, 1922) and has since been 79 considered to be predominantly a digenean of wild rodents (Elton et al., 1931; Seo et al., 80 1964: Ito and Itagaki, 2003; Chai et al., 2007; Rogan et al., 2007). Definitive host variability 81 for *P. muris* has, however been, frequently recorded including that of the domestic dog, 82 Canis familiaris in Japan (Saito et al., 1995), the feral Japanese raccoon, Procyon lotor 83 (Yamada, 2000; Sato and Suzuki, 2006) and the Mexican Greater funnel-eared bat; 84 Natalus mexicanus (Perez-Ponce de Leon et al., 1996) in addition to several cases of 85 natural avian infection in the USA (McMullen, 1937; Cort and Ameel, 1944; Secord and 86 Canaris, 1993). Typically, P. elegans has been considered foremost in the genus 87 Plagiorchis for infecting birds (Shimalov, 2002) however this species has also been 88 reported to parasitize several species of rodent, including A. sylvaticus (Montgomery and 89 Montgomery, 1990a), the yellow necked mouse, Apodemus flavicollis (Hildebrand and 90 Zaleśny, 2009), the striped field mouse, Apodemus agrarius (Shimalov, 2002; Hildebrand 91 and Zaleśny, 2009), and the bank vole, Myodes glareolus (Hildebrand and Zaleśny, 2009). 92

Little is known about the exact life cycle of *Plagiorchis spp.* that infect these rodents and, in particular, the identity and role of intermediate hosts. The high prevalence reported in this periaquatic site (Rogan *et al.,* 2007) suggests that the presence of water and aquatic organisms might be key factors. The unique nature of this site and the

development of molecular tools for detection and identification of these parasites presents
an opportunity to dissect the ecology of this parasite and its hosts.

In the current study we investigate the rare occurrence of *Plagiorchis muris* at Malham Tarn, Yorkshire, UK (Rogan *et al.*, 2007) by examining prevalence, intensity and seasonality of adult stages collected from rodents trapped at defined woodland sites around this upland lake. We furthermore highlight the difficulty in distinguishing *P. muris* and *P. elegans* in the absence of molecular tools and investigate life cycle indicators within this periaquatic environment.

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106 MATERIALS AND METHODS

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The study was carried out at Malham Tarn Nature Reserve located in North Yorkshire, Northwest England at an altitude of 375m above sea level and is an area that has previously been investigated for a range of host parasite systems (Kennedy and Burrough, 1978; Allan *et al.*, 1999; Hughes *et al.*, 2006, 2008; Rogan *et al.*, 2007; Behnke *et al.*, 2009; Hide *et al.*, 2009; Thomasson *et al.*, 2011; Boyce *et al.*, 2012).

Rodent trapping and examination was conducted according to the methods 113 described by Boyce et al., (2012). In total 117 wood mice (Apodemus sylvaticus) and 63 114 voles (54 bank voles Myodes glareolus and 9 field voles Microtus agrestis) were trapped 115 from four sampling sites around Malham Tarn: Tarn Woods, Tarn Fen, Spiggot Hill and Ha 116 Mire Plantation between January 2010 and October 2011 (Figure 1). A permit was 117 granted by the National Trust to allow sampling within the boundaries of the reserve. 118 119 Plantations within these boundaries are represented by Tarn Woods, Ha Mire Plantation and Spiggot Hill. Ownership boundaries form part of the reserve perimeter restricting 120 sampling in part. Tarn Fen is an area of raised bog covered by deciduous woodland 121 located within the reserve boundaries. As water is considered to be an important aspect in 122

the life cycle of *Plagiorchis* these four trapping sites were selected in order to cover a vast
 area of woodland throughout the reserve, which is located in close proximity to each of the
 tarn's borders.

126 Morphological examination of the digenean specimens recovered from the small intestine of A. sylvaticus was carried out according to the methods described by Boyce et 127 al., (2012). The majority of the specimens were fixed in 70% ethanol suitable for molecular 128 analysis. DNA was extracted from 12 individual worms collected from different host 129 specimens of A. sylvaticus when feasible using a phenol: chloroform method modified from 130 Thomasson et al., (2011) which encompassed halving the amount of reagent at each 131 stage of the protocol. DNA was extracted from three individual worms isolated from three 132 different wood mice trapped from each of the four sampling sites. Only one wood mouse 133 however was found to be infected from Spiggot Hill therefore three individual worms from 134 the same host specimen needed to be used in this instance. 135

The Internal transcribed spacer (ITS), including the ITS1, 5.8S, ITS2 and flanking 136 regions of the 3' end of the 18S and 5' end of the 28S were amplified using the forward 137 universal primer BR (^{5'}GTAGGTGAACCTGCGGA^{3'}) and reverse digenean specific primer 138 dig11 (^{5'}GTGATATGCTTAAGTTCAGC^{3'}) according to Tkach et al., (2000a). The partial 139 28S rDNA gene region was amplified using the forward digenean specific primer dig12 140 (⁵'AAGCATATCACTAAGCGG³') and the reverse universal primer Lo 141 (⁵GCTATCCTGAGRGAAACTTCG³) according to Tkach et al., (2000b). 142

Each 50µl PCR reaction contained 5µl 10X DreamTaq buffer including 2mM MgCl₂ (Fermentas, Life Sciences), 0.05µmol dNTPS (100mM, Bioline), 2.5µM forward primer, 2.5µM reverse primer, 5U DreamTaq DNA polymerase and 2µl DNA template. All PCR reactions were performed using a Robocycler 96 PCR machine (Stratagene, CA) and visualised on a 1% (w/v) Tris-acetate-EDTA (TAE) agarose gel stained with gel red using a G: Box gel imaging system (Syngene, UK). The amplification profile consisted of 1 cycle at

94°C for 10 minutes, followed by 35 cycles of 1 minute at 94°C, 1 minute at 54°C and 1 149 minute at 72°C and one final cycle at 72°C for 10 minutes. The target bands were excised 150 from the gel using a UV transilluminator and purified using a PCR purification kit 151 152 (Geneflow) according to the manufacturer's instructions. Samples were commercially sequenced in both directions (Source Bioscience, Nottingham, UK). The 12 DNA 153 sequences for the ITS gene were primarily aligned using the multiple sequence alignment 154 program ClustalW (www.genome.jp/tools/clustalw/) to check for sequence homology 155 between specimens from each of the four sampling sites. This procedure was repeated 156 with the 28S rDNA data. In both instances FinchTV trace viewer (Geospiza, Seattle, WA) 157 was utilised in order to verify any regions of ambiguity. The two DNA sequences generated 158 for the ITS and 28S rDNA from Malham Tarn were compared with those held in the 159 National Center for Biotechnology Information (NCBI) database using the BLAST program 160 (www.blast.ncbi.nlm.nih.gov/Blast.cgi). 161

Differences in prevalence observed between sampling sites and seasons were 162 investigated using Chi-squared test for heterogeneity. Host sex and host age were 163 statistically analysed using 2 x 2 contingency tables using Fisher's exact test 164 (http://www.graphpad.com/quickcalcs/ConfInterval2.cfm). For determination of 165 age, rodents were split into three age cohorts according to Behnke et al., (1999). Associations 166 between prevalence and host length (cm), weight (g) and rainfall data (mm) were analysed 167 using Spearman's rank of correlation. Monthly rainfall data was provided by Malham Tarn 168 Field Centre. Prevalence calculated during each season over a two-year period was 169 analysed in relation to the previous three months rainfall (mm) adapted from Rogan et al., 170 171 (2007).

In an attempt to identify life cycle indicators of *Plagiorchis* sp. at Malham Tarn, molluscan species were also collected quarterly between January 2010 and October 2011 in order to examine for intramolluscan stages. Several water bodies located within close

proximity to the rodent trapping sites including the tarn margin were selected for analysis. 175 Snails were collected using a D-frame aquatic dip net and kick sample technique and were 176 also hand-picked from the stems of vegetation and underlying surfaces of rocks. Snails 177 178 were speciated according to Macan and Cooper (1960) and housed in the laboratory in 4 litre glass covered tanks containing pond water. Snails were maintained at 4°C and fed 179 washed lettuce ad libitum according to Voutilainen et al., (2009). For the examination of 180 Plagiorchis sp. intramolluscan larval stages, a snail crushing method was employed 181 according to Caron et al., (2008). Binomial confidence intervals on parasite prevalences 182 two-tailed test), were calculated (P = 0.05,based on standard methods 183 (http://statpages.org/confint.html). 184

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186 RESULTS

Five helminth species were recorded from the sampled wood mouse population; *Heligmosomoides polygyrus, Syphacea oblevata, Capillaria murissylvatici, Brachylaemus recurvum* and *Plagiorchis* sp. with species richness varying from 0 to 5 in individual animals. *Plagiorchis* sp. was not detected from any of the vole species despite careful observations.

In total, 717 *Plagiorchis* worms were successfully recovered from the small intestine 192 of 27 of 117 examined wood mice between January 2010 and October 2011. An overall 193 prevalence rate of 23% was determined from all four sampling locations: Tarn Woods, 194 Tarn Fen, Spiggot Hill and Ha Mire Plantation indicating this digenean species to be 195 dispersed throughout the Malham Tarn area. Worm burden ranged from 0 to 275 with a 196 197 mean of 26.6±61.5 (717/27) appearing to be typically overdispersed in distribution (Variance to mean ratio (VMR): σ^2/μ = 158.89) with 84.6% of wood mice being infected 198 with zero or just a single worm in comparison to just 1.7% harbouring the majority of 199 parasites (>100 worms). A goodness of fit indicated a good agreement with the negative 200

binomial distribution ($G_{calculated} = 21.17$, df = a - 3 (12 - 3) is 21.67, p = 0.01). *Plagiorchis* sp. was identified following the microscopical analysis of 10 fixed and unstained specimens.

The adult worm (Figure 2) measured 1.66 to 2.93mm (mean 2.64mm) in length by 204 0.46 to 0.76mm (mean 0.58mm) in width at the widest part (across the region of the most 205 anterior testis). The tegument possesses minute spines covering the entire surface. The 206 ventral surface appears to be obscured by a vast array of vitelline glands that fail to 207 maintain confluency within the anterior region. The oral sucker is roughly spherical in 208 shape and measures 200 to 300µm in length (mean 256µm) by 200 to 300µm (mean 209 247µm) in width. The oral sucker lies anterior to the pharynx from which the intestinal 210 bifurcation occurs. The intestine appears short and indistinct and thereafter extends into 211 two very long blindly ending caeca that are often difficult to observe, commonly being 212 masked by a copious mass of vitelline glands, but reaching the near posterior extremity of 213 the body. The ventral gland is smaller in size than that of the oral sucker measuring 130 to 214 200µm (mean 163µm) in length by 130 to 200µm (mean 163µm) in width. These 215 measurements indicate an oral to ventral sucker ratio of 1.52: 1 (width ratio) - 1.57: 1 216 (length ratio). Two oval shaped testes are situated posterior to a single ovary. The testes 217 are obliquely positioned with the anterior testis situated slightly right of the median line and 218 219 the most posterior testis to the left. The ovary lies just posterior to the ventral sucker, separated by the cirrus sac, which curves in a posterior direction along the left hand side 220 of ventral sucker. The metraterm can be visualised to curve posteriorly along the right 221 hand side of the ventral gland adjoining the anterior region of the uterus when not 222 obscured by the vitelline glands. The uterus extends to the posterior extremity of the body 223 presenting a characteristic s-shape that reaches from the region of the ovary and 224 continues intertesticularly towards a posterior vitellarian commissure. The vitelline glands 225 continue along both lateral sides, from the far extremity of the hind body and into the 226

forebody surpassing the ventral sucker and creating confluency often up to the posterior border of the pharynx. The morphological features described in this section hold a sufficient similarity for both *P. muris* and *P. elegans* therefore the use of morphology alone could be considered ambiguous and the use of molecular differentiation be regarded an important means to complement species classification in this case.

DNA was successfully extracted and amplified from 12 individual Plagiorchis worms 232 recovered from the small intestine of 10 A. sylvaticus mice trapped from the four different 233 sites. Amplification of the internal transcribed spacers (ITS) including the ITS1, 5.8S and 234 ITS2 generated a sequence of 1213bp (GenBank accession: JX522536). The 12 ITS 235 sequences generated from all four sampling sites were 100% identical. The ITS sequence 236 generated from Malham Tarn was compared against five DNA sequences generated from 237 Plagiorchis adults that were held in the NCBI database: P. maculosus (AF316152) 238 collected from the Chaffinch, Fringilla coelebs (Snyder and Tkach, 2001), P. elegans 239 (AF151952) collected from The Red-Backed Shrike, Lanius collurio, P. koreanus 240 (AF151944) collected from Kuhl's pipistrelle, Pipistrellus kuhli, the common noctule, 241 Nyctalus noctula and Daubenton's bat Myotis daubentoni, P. vespertilionis (AF151949) 242 from *M. daubentoni* and *P. muelleri* (AF151947) from the serotine bat *Eptesicus serotinus* 243 all obtained within the Ukraine (Tkach et al., 2000a). The ITS sequence from Malham Tarn 244 shared a 100% sequence homology with that of P. elegans with only one omission of 245 adenosine at site 571. This omission was observed in all 12 generated DNA sequences. 246 The Malham Tarn sequence shared only 94% sequence homology with P. maculosus, 247 89% with P. koreanus, 91% with P. vespertilionis, and 90% with P. muelleri. 248

Amplification of the 28S rDNA gene generated a partial sequence of 1263bp (GenBank accession: JX522535). The 12 28S rDNA sequences generated from all four sampling sites were also 100% identical. This sequence was also compared against five available DNA sequences from the NCBI database: *P. elegans* (AF151911) (Tkach *et al.*,

1999), P. muris (AF096222) obtained from the intestine of a rat in the Republic of Korea 253 (Lee et al., 2004), P. muelleri (AF184250) (Tkach et al., 2001), P. koreanus (AF151930) 254 and P. vespertilionis (AF151931) collected from N. noctula in the Sumy region of the 255 256 Ukraine and *M. daubentoni* in the vicinity of Kiev, Ukraine (Tkach et al., 2000b). The partial 28S rDNA sequence available for P. muris (AF096222) was only 304bp in length (Lee et 257 al., 2004). All other available 28S sequences for Plagiorchis species including the 258 sequence generated from Malham Tarn were therefore trimmed and aligned with this 259 sequence for P. muris. The 28S sequence generated for specimens collected from 260 Malham Tarn again shared a 100% sequence homology with that of *P. elegans*, 98% with 261 P. muelleri, P. koreanus and P. vespertilionis and only a 95% match with P. muris (Figure 262 **3**). This data in combination with the 100% sequence homology match demonstrated by 263 the ITS region, questions the identity of P. muris at Malham Tarn and infers that the 264 species present at this location is in fact that of *P. elegans*. For the remainder of this 265 paper, references to *Plagiorchis* sp. in this study should be read as *P. elegans*. The 266 prevalence of *P. elegans* at this location was examined. 267

Prevalence was analysed in relation to both extrinsic and intrinsic factors. During the study a comprehensive data set was established which recorded trapping location, date, host sex, host weight and host length. All prevalence data, 95% confidence limits and mean intensities have been summarised in **Table 1**.

To determine whether there was an association with prevalence and a particular trapping site, the rate of prevalence was examined between the four sampling sites using chi squared test for heterogeneity. The greatest prevalence was observed at Ha Mire Plantation in which 37.03% (n = 27) of sampled wood mice carried a mean intensity of 19±24.2 worms (187/10). The prevalence at Tarn Woods which is the original sampling site reported by Rogan *et al.*, (2007) was less with only 23.07% (n = 52) of wood mice being infected despite a slightly higher mean intensity of 21±50.33 (246/12). At Tarn Fen, only

four of the 21 examined wood mice were infected with *P. elegans* giving a prevalence of 19.05%. A very low worm burden of just 2±2.5 (9/4) was also observed at this site. Only a single wood mouse was infected with *P. elegans* at Spiggot Hill providing the lowest prevalence of the study at 5.88% (n = 17). This mouse however harboured 275 worms, the highest number recorded during the study. No significant heterogeneity was found between the prevalence of *P. elegans* and any of the four sites ($X^2 = 3.79$, p = 0.05, v = 3).

As studies of *Plagiorchis* infection in natural rodent populations have demonstrated 285 seasonal variation in prevalence and intensity, we examined this hypothesis at Malham 286 Tarn. Mean prevalence and mean intensity was calculated each season over a two year 287 period. Both mean prevalence and mean intensity were zero during the winter (January) 288 sampling periods (n = 8). Only one mouse was infected during the spring (April) giving a 289 prevalence of 12.5% (n = 8). This mouse however harboured 179 worms. Peak prevalence 290 occurred during the summer (July) when 27.3% (n = 22) of mice carried a mean intensity 291 of 3 ± 2.94 worms (20/6). Prevalence thereafter decreased slightly to 25.3% (n = 79) during 292 the autumn (September/October) however mean intensity increased considerably to 293 26±61.46 worms (518/20) for this period. Despite observable differences in prevalence no 294 significant heterogeneity was found between the prevalence of *P. elegans* and season (X^2) 295 = 3.38, p = 0.05, v = 3). A Spearman's rank of correlation also indicated a very weak but 296 significant correlation (r = 0.095, $P = \langle 0.05 \rangle$) between the prevalence of *P. elegans* and 297 rainfall for the three months preceding each sampling session (Table 2). 298

In total, 71 male mice and 46 female mice were examined. Prevalence in male mice was greater at 32.4% in comparison to only 8.7% for female mice. Male mice furthermore harboured the majority of worms carrying 688 of 717 recovered with a mean intensity of 30 ± 66.09 (688/23) against only 7±12.5 (29/4) for female mice. A highly significant difference between *P. elegans* infection and *A. sylvaticus* sex was identified (*p* = 0.003).

The data was furthermore analysed to determine whether there was an age-304 dependent prevalence occurring at this location. P. elegans was recorded in a total of 19 305 out of 44 adult wood mice (43.2%) which was greater than that observed for young adult 306 307 mice (14.3%, n = 56). No juvenile mice were infected during the present study (n = 17). A similar pattern was observed for mean intensity with adult wood mice harbouring the vast 308 majority of infection carrying 677 of 717 worms recovered. Adult males (n = 34) however 309 carried 650 of these worms in comparison to just 27 worms carried by adult female mice (n 310 = 10). Mean worm burden in the adult group was 36 ± 71.72 (677/19) in comparison to only 311 5±7.86 (40/8) for young adults and zero for juvenile mice. No significant difference was 312 observed between the juvenile and young adult age categories (p = 0.185) however the 313 difference between adult wood mice and young adults (p = 0.002) and adult and juvenile 314 (p = 0.001) age cohorts was found to be highly statistically significant. Wood mice ranged 315 in weight from 4.0 to 29g and length from 4.5 to 9.8cm. A Spearman's rank of correlation 316 using these data against prevalence gave a very strong correlation in both cases (r = 317 0.929 $P = \langle 0.05 \text{ and } r = 0.955, P = \langle 0.05 \text{ respectively} \rangle$ indicating an age-dependent 318 prevalence to be evident at this location. 319

In an attempt to identify life cycle indicators of *P. elegans* at Malham Tarn, a total of 320 2021 snails consisting of 11 species were examined for intramolluscan stages by crushing 321 (Table 3). No larval stages of *P*. elegans were found in any of the snails despite careful 322 observation and the frequent detection of other trematode larvae such as Notocotylus. 323 Binomial confidence intervals were calculated (P = 0.05, two-tailed test) to establish 324 whether zero prevalence was a significant result using these samples. The results 325 326 however indicate that none of the examined snail species can be currently ruled out as a potential intermediate host for *P. elegans* with zero prevalence not being significant given 327 the small sample sizes. Despite this for snail species where sample sizes were high, very 328 low prevalences (of less than 1%) are likely. Further investigation is required. 329

331 DISCUSSION

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333 In the present study we employed the use of molecular differentiation to investigate further the occurrence of Plagiorchis muris at Malham Tarn. Results indicate that the 334 currently identified specimens of *P. muris* at this location are *Plagiorchis elegans*. As far as 335 we can determine, our previous report (Rogan et al., 2007) was the fourth known report of 336 P. muris in British wildlife (Elton et al., 1931; Baylis 1939; Fahmy and Rayski, 1963) 337 although the conclusion from this study indicates that this trematode was, in fact, P. 338 elegans, and also raises the question as to whether other previous studies have correctly 339 identified the species. Elton et al., (1931) reported only a prevalence and Baylis, (1939) 340 simply listed an occurrence of the digenean. Neither author described the morphology of 341 the parasite involved nor was DNA sequencing an aspect of biological surveys at that time. 342 The third report by Fahmy and Rayski (1963) was a short report which provided no written 343 account, but rather incorporated a diagram as a means of description. Unfortunately, the 344 diagram contained insufficient detail to clarify its status as *P. muris* from that of *P. elegans*. 345 Previously, Fahmy (1954) described a new species of *Plagiorchis*, *P. lutrae* from the 346 otter (Lutra lutra) in Scotland. This new species closely resembled the description of P. 347 muris but was differentiated on the basis of size. The measurements of this new species 348 did however coincide with those provided in the original description of *P. muris* described 349 by Tanabe in Kyoto, Japan in 1922; despite such P. lutrae was compared with P. muris 350 described by McMullen (1937) from Douglas Lake in Michigan State, USA which was 351 much larger in size. 352

The description of *P. muris* from various locations appears ambiguous. The majority of reports describing *P. muris* from outside of Southeast Asia indicate both a greater range in length and oral to ventral sucker ratio. The description of *P. muris* by McMullen (1937)

was based upon adult digeneans recovered from a range of experimental hosts in addition
to a range of naturally infected avian fauna. The average length and oral to ventral sucker
ratio of these specimens was however beyond the maximum dimensions reported for
Southeast Asian *P. muris* (Tanabe, 1922; Seo *et al.*, 1964).

Interestingly, the measurements provided for *P. muris* by McMullen (1937), Rogan 360 et al., (2007) and during the current study overlap with the description generated for P. 361 elegans from a PhD study conducted by Gorman at Leeds University in 1980. 362 Experimental evidence provided by Gorman (1980) did however identify intraspecific 363 variation within a pure strain of *P. elegans*. The study identified various manifestations in 364 several anatomical structures, not only between different definitive host species but 365 furthermore within the same definitive host including differences in the extent of the 366 vitellaria and aperture shape of both the oral and ventral suckers (Gorman, 1980). 367 Confluency of the vitelline glands have been used on several occasions as a means of 368 differentiation for *Plagiorchis* species (Fahmy, 1954; Tkach et al., 2000a) however as 369 pointed out by Blankespoor (1974) and Gorman (1980) use of the glands for diagnosis 370 may not be appropriate due to the intraspecific variation observed in this feature. For 371 instance the vitelline glands of *P. muris* have been reported to extend to either the 372 posterior border of the pharynx (Tanabe, 1922; Hong et al., 1996) or the level of the oral 373 sucker (Fahmy and Rayski, 1963; Seo et al., 1964; Hong et al., 1998). Hong et al., (1996) 374 nonetheless described *P. muris* from a human case of plagiorchiasis using the positioning 375 of the vitelline glands to morphologically differentiate *P. muris* from both *P. vespertilionis* 376 and *P. koreanus*. There was however no mention of *P. elegans* in this report despite this 377 species appearing to display the most morphological similarity to P. muris in the 378 distribution of the vitellaria. As far as can be determined there have currently been no 379 reported cases of *P. elegans* infection in either Korea or Japan where *P. muris* appears to 380 be considered the typical dominant *Plagiorchis* species found in rodents. 381

Currently, there appears to be ambiguity in the criteria used to morphologically 382 differentiate P. muris and P. elegans. Fortunately, the use of DNA sequencing could be 383 employed in the current study to confirm the identity of the Malham Tarn specimens. The 384 385 use of the internal transcribed spacer regions and the 28S rDNA gene indicate the specimens recovered from Malham Tarn to be P. elegans. Based on these results and 386 taking into consideration the unreliability of morphological differentiation for the two 387 species in guestion (Blankespoor, 1974; Gorman, 1980; Hong et al., 1998), it could be 388 speculated that other reports describing P. muris based purely on morphology have also 389 misidentified the species involved. For example, in his report McMullen (1937) commented 390 that the cercariae used for experimental infection possessed seven or eight pairs of 391 penetration glands on either side of the stylet which is a combination typical of *P. elegans* 392 (Faltýnková et al., 2007) as opposed to the four pairs originally described for P. muris by 393 Tanabe (1922). This morphological description provided for *P. muris* by McMullen (1937) 394 has since been a basis for morphological comparison made by some European authors 395 (Fahmy, 1954; Rogan et al., 2007). 396

Despite the questionable identity over *Plagiorchis* at Malham Tarn, the occurrence 397 of this digenean at this location within the UK is nonetheless considered rare, in particular 398 with a consistent prevalence recorded since 1993 (Rogan et al., 2007). Furthermore, the 399 overall prevalence rate of 23% recorded during this study appears to be much greater than 400 that reported in the literature. Other UK reports involving either P. muris or P. elegans 401 have encompassed very low prevalence rates of 0.1% and 0.05% respectively (Elton et 402 al., 1931; Montgomery and Montgomery, 1990a). A further two reports of P. muris in the 403 wood mouse A. sylvaticus in Ireland have also indicated very low prevalence rates of 1% 404 or less (Langley and Fairley, 1982; O'Sullivan et al., 1984). Further afield, Ito and Itagaki, 405 (2003) reported a prevalence of just 1.7% in the large Japanese field mouse Apodemus 406 speciosus in Japan and Chai et al., (2007) recorded an overall prevalence of 5.3% in the 407

striped field mouse *Apodemus agrarius* in Korea. *P. elegans* does however appear to be the species reported most often from small mammals within Europe. Hildebrand and Zaleśny (2009) reported a prevalence of 1.3% in the bank vole *Myodes glareolus* trapped in Poland. A single specimen of *P. elegans* recovered from *M. glareolus* in Pallasjärvi, Finland, gave a prevalence of just 0.5% (Tenora *et al.*, 1983) and a slightly higher prevalence rate of 3.1% was reported by Shimalov (2002) from *A. agrarius* in Belarus.

The reasons for the occurrence of *P. elegans* at such a high prevalence at Malham 414 Tarn are unclear. Malham Tarn is a 'Site of Special Scientific Interest' (SSSI) boasting a 415 vast array of plant and animal species. The surface area of the tarn is approximately 150 416 acres with an average depth of 2.4m and a maximum depth of 4.4m in various regions 417 (Woof and Jackson, 1988). Similarly to this study, previous studies have observed the 418 occurrence of *Plagiorchis* in regions of close proximity to significant water bodies (Cort and 419 Olivier, 1943; Cort and Ameel, 1944; Bock, 1984; Hong et al., 1999; Hildebrand and 420 Zaleśny, 2009). Being the only upland marl lake of its kind in Britain (Rogan et al., 2007), it 421 could be speculated that Malham Tarn itself may play an important role by providing 422 important breeding sites for intermediate host species. Molluscs of the genus Lymnaea are 423 the dominant snails acting as the first intermediate host for *Plagiorchis* species worldwide 424 (Tanabe, 1922; Velasquez, 1964; Bock, 1984; Manga-Gonzalez et al., 1994; Zakikhani 425 and Rau, 1999; Väyrynen et al., 2000; Faltýnková et al., 2007). Four species of Lymnaea 426 have been recorded at Malham Tarn including L. stagnalis, L. peregra, L. palustris and L. 427 truncatula (Norris, 2003) although currently none of these hosts have been positively 428 implicated in the life cycle of *P. elegans* at this location. Negative infections, in all of the 429 1603 Lymnaea specimens examined, suggest that these are unlikely to be intermediate 430 hosts, but this is difficult to absolutely rule out. Insects act as the second intermediate host 431 and dragonflies are the most commonly reported insects found to be naturally infected 432 (Hong et al., 1999). Malham Tarn is home to several species of dragonfly (Shorrock and 433

434 Sutton, 2010) however further study is required in order to identify first and second 435 intermediate host species involved in transmission at this location.

In the present study, *P. elegans* was recorded from all four trapping sites and no 436 437 association between prevalence and rainfall amount was identified suggesting that temporary water bodies may not be an important determinant for transmission and that 438 rather infection is related to the presence of intermediate host species that breed within the 439 tarn body itself. Each trapping site however is separated from the tarn by a narrow shingle 440 beach and earth ridge, terrain that wood mice are unlikely to cross. A crude morphological 441 examination of the stomach contents of A. sylvaticus (n = 117) at Malham Tarn 442 demonstrated the presence of adult insect remains suggesting the main source of infection 443 for A. sylvaticus to be adult insects infected with metacercariae that may migrate into the 444 home range of the wood mouse following emergence from the tarn body. Other studies 445 have also indicated the diet of A. sylvaticus to include various adult insects (Montgomery 446 and Montgomery, 1990b; Khammes and Aulagnier, 2007). 447

The distribution of *P. elegans* at Malham Tarn demonstrated a typical pattern of 448 over-dispersion with rodent age and sex being the two main factors associated with 449 prevalence. Khammes and Aulagnier (2007) used three age categories to examine the 450 differential diet of A. sylvaticus; juvenile, sub-adults and adults and indicated arthropod 451 remains to be more abundant in the stomach contents of adult mice. In the present study, 452 the prevalence of *P. elegans* was significantly greater in adult mice than younger age 453 cohorts. This is likely due to differences in exposure to infective stage parasites through 454 differences in diet as a result of adult mice demonstrating greater foraging behaviour 455 456 (Lewis, 1968; Lewis and Twigg, 1972).

457 Several studies have indicated a change in the feeding habits of *A. sylvaticus* from 458 a granivorous diet to one that consumes animal material during periods when seeds are 459 scarce during the spring and early summer months (Lewis, 1987; Montgomery and

Montgomery, 1990b). A study by Montgomery and Montgomery (1990b) in County Down, 460 Northern Ireland, compared the stomach contents of A. sylvaticus at two locations. In both 461 sample sets animal material was seen to rapidly decline after September. The typical 462 463 consumption of insect material by A. sylvaticus between spring and autumn coincides with the detectable prevalence period of *P. elegans* in the present study. Other studies have 464 furthermore identified seasonal patterns of infection in natural rodent populations (Chai et 465 al., 2007). Seasonality in prevalence is likely due to the developmental cycle of the 466 intermediate host species involved in transmission. Hong et al., (1998) demonstrated that 467 up to 96% of *Plagiorchis* worms were expelled from the intestine of albino laboratory rats 468 within 28 days post infection indicating the likelihood of finding adult digeneans from the 469 previous season to be low. 470

Male mice carried both a statistically higher prevalence and a higher worm burden 471 than their female counterparts. It is likely that the differences observed between male and 472 female and adult and younger adult mice are due to variation in behaviour between the 473 various groups. It has been well documented that the home range of male rodents is much 474 greater than that for female rodents (Langley and Fairley, 1982; Wolton, 1985; 475 Attuquayefio et al., 1986), particularly during the breeding season between April and 476 October (Bueshing et al., 2007) during which male wood mice have been observed to 477 increase their home range by as much as five times (Corp et al., 1997). Male adult wood 478 mice are also much more arboreal than their younger counterparts as well as female wood 479 mice (Buesching, 2007). Arboreality in wood mice is considered to be due to their 480 insectivorous nature, with insects and other small invertebrates often inhabiting the tree 481 482 canopy. Climbing as a means to acquire food may however be energetically expensive (Bueshing et al., 2007) and female wood mice that have a greater dependency on 483 resources have been observed establishing mutually exclusive breeding territories 484 (Flowerdew, 1993) and as such are less likely to become infected due to a reduced 485

exposure to metacercarial infected insects than adult males that appear more prone towanderlust.

This study sheds light on the detailed parasite-multihost interactions within this 488 489 complex periaquatic ecosystem but demonstrates the difficulties associated with understanding these interactions. A key issue is clearly the need for accurate identification 490 of all stages of the parasite, an issue that molecular tools can significantly enhance in 491 particular for species such as *P. muris* and *P. elegans* that are difficult to distinguish by 492 classical morphology alone. The greatest challenges lie in linking the life cycle stages 493 together and to link transmission dynamics to the ecology of the parasites and hosts within 494 the ecosystem. Further studies are required to fully understand these complex 495 interactions. 496

497

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499

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Table 1. Summarised prevalence and mean intensity data for *P. elegans* in *A. sylvaticus*

⁷⁹⁴ from Malham Tarn. * Figure derived from a single host infection.

	Prevalence	95% Confidence limits		Mean intensity
	(%)	Lower	Upper	± S.E.M
<u>Overall</u>	23.0	16.33	31.54	27 ± 61.5
Location Tarn Woods Tarn Fen Spiggot Hill Ha Mire	23.07 19.05 5.88 37.03	13.58 7.80 0.01 21.47	36.28 40.59 28.92 55.84	21± 50.33 2 ± 2.5 275* 19 ± 24.2
<u>Season</u> Winter Spring Summer Autumn	0 12.5 27.3 25.3	0 0.11 12.88 16.96	37.22 49.22 48.43 35.96	0 179* 3 ± 2.94 26 ± 61.46
<u>Sex</u> Male Female	32.4 8.7	22.62 2.90	43.98 20.86	30 ± 66.09 7 ± 12.5
<u>Age cohort</u> Adult Young adult Juvenile	43.2 14.3 0	29.67 7.16 0	57.79 26.00 21.63	36 ± 71.72 5 ± 7.86 0

Table 2. Correlation of rainfall data and *P. elegans* prevalence (%). Total rainfall over a three monthly period was analysed, using Spearman's rank of correlation, with parasite prevalence Rainfall data was supplied by Malham Tarn Field centre. Numbers in bold are the figures that have been used in the statistical analysis.

Month	Total monthly	Total three	Trapping	P. elegans
	rainfall (mm)	months	month/ year	prevalence (%)
		rainfall (mm)		
<u>2009</u>				
October	128.3			
November	378.9			
December	123.6	630.8	January 2010	0
<u>2010</u>				
January	074.4			
February	064.4			
March	098.3	237.1	April 2010	0
April	027.2			
Мау	021.2			
June	033.5	081.9	July 2010	11
July	151.4			
August	116.4			
September	296.1	559.9	September 2010	35
October	121.4			
November	189.1			
December	041.5	352.0	January 2011	0

<u>2011</u>

January	150.5			
February	221.4			
March	036.2	408.3	April 2011	20
April	031.4			
Мау	126.2			
June	094.1	251.7	July 2011	38
July	090.6			
August	170.8			
September	157.7	419.1	October 2011	18
October	NA			

Table 3. Prevalence of intramolluscan stages found in snail species. Confidence intervals
(95%, P = 0.05, two tailed test) were calculated using an online Binomial Confidence
Interval calculator (http://statpages.org/confint.html).

Molluscan species	Number	Prevalence	95% Confidence interval	
	examined (n)	(%)	Lower	Upper
Lymnaea stagnalis	32	0	0.00	10.89
Lymnaea palustris	58	0	0.00	06.16
Lymnaea peregra	1270	0	0.00	00.29
Lymnaea truncatula	243	0	0.00	01.51
Anisus leucostoma	35	0	0.00	10.00
Bithynia tentaculata	10	0	0.00	30.85
Physa fontinalis	8	0	0.00	36.94
<i>Planorbis</i> sp.	7	0	0.00	40.96
Potamopyrgus antipodarum	188	0	0.00	01.94
Sphaerium corneum	33	0	0.00	10.58
Valvata cristata	137	0	0.00	02.66

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Figure 1. Map of Malham Tarn Nature Reserve and adjacent woodlands. Redrawn from
Shorrock and Sutton (2010). TW: Tarn Woods; TF Tarn Fen; SP Spiggot Hill; HM Ha Mire
Plantation. The dotted line represents the reserve boundary.

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Figure 2. *Plagiorchis* specimen recovered from the small intestine of *Apodemus sylvaticus*at Malham Tarn. The drawing was made from a photograph taken with a Leica ICC50
digital camera attached to a Leica DM500 microscope. Abbreviations: OS, oral sucker; P,
pharynx; C, cirrus; CS, cirrus sac; VS, ventral sucker; M, metraterm; O, ovary; T, testis;
Ca, caecum; U, uterus; V, vitellaria; Vc, vitellarian commissure. Scale bar = 500µm.

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Figure 3. Comparison of the partial 28S rDNA sequence of *P. muris* collected from Malham Tarn with other *Plagiorchis* species retrieved from the NCBI database: *P. elegans* (AF151911); *P. koreanus* (AF151930); *P. muelleri* (AF184250); *P. vespertilionis* (AF151931); *P. muris* (AF096222). The comparison was made using the GeneDoc alignment tool. Black shading indicates regions of conserved homology. Grey indicates regions of conservation between four or more species.