

RESEARCH ARTICLE

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Sperm parameters and anti-Müllerian hormone remain stable with *Helicobacter pylori* infection: a cross-sectional study

Chun Feng^{1†}, Ping-Ping Lv^{2†}, Chang-Chang Huang¹, Song-Qing Yang¹, Qiu-Ping Yao¹, Jin-Ming Shen^{3*} and Min Jin^{1*}

Abstract

Background and aims: It has been reported that *Helicobacter pylori* (HP) infection was more prevalent in infertile populations. HP infection could lead to decreased sperm parameters, and treating the HP infection could improve the quality of sperm. However, studies investigating the relationship between infertility and HP infection are still limited, and more evidence is required. Therefore, we performed the present study to investigate the impact of HP infection on sperm quality in males and on ovarian reserve in females.

Methods: A total of 16,522 patients who visited the Second Hospital of Zhejiang University from January 2016 to June 2019 due to abdominal discomfort and underwent a ^{13/14}C-urea breath HP test were included in this retrospective cross-sectional study. Among them, 565 had performed sperm analysis or ovarian reserve tests in the past three months and were involved for further analyses. Sperm parameters were examined with a computer-assisted sperm analysis system, and serum anti-Müllerian hormone (AMH) and sex hormones were tested with an electrochemiluminescence method.

Results: Among 363 patients who underwent the sperm test, 136 (37.47%) had HP infection. Among 202 patients who underwent the AMH test, 55 (27.23%) had HP infection. There was no difference in sperm concentration and motility between the HP+ and HP- groups ($P > 0.05$). Further subgroup analyses stratified into 5-year age groups confirmed that there was no significant difference in sperm parameters ($P > 0.05$). When pooled with previously published data, no significant difference in sperm concentration or motility was found ($P > 0.05$). Meanwhile, this study found that the serum AMH level was similar between the HP+ and HP- groups ($P > 0.05$). Further subgroup analyses confirmed that there was no significant difference in serum AMH level ($P > 0.05$).

Conclusions: There were no differences in sperm parameters and AMH levels based on history of HP infection among Chinese patients.

Keywords: *Helicobacter pylori* (HP), Anti-müllerian hormone (AMH), Sperm parameters, Progressive motility

Background

Currently, it is widely accepted that *Helicobacter pylori* (HP) may be related to a series of extragastric diseases, including cardiovascular, neurologic, respiratory, hematologic, metabolic, dermatologic, obstetric, autoimmune, and kidney diseases [1, 2]. Among them, the impact of HP on fertility has attracted much attention. As early as

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*Correspondence: shenjinmg@gmail.com; min_jin@zju.edu.cn

†Chun Feng and Ping-Ping Lv contributed equally to this work

¹ Department of Reproductive Medicine, The Second Affiliated Hospital of Zhejiang University School of Medicine, 88 Jiefang Road, Hangzhou 310009, Zhejiang, China

³ Department of Orthopedics, The First Affiliated Hospital of Zhejiang Chinese Medicine University, 54 Youdian Road, Hangzhou 310006, Zhejiang, China

Full list of author information is available at the end of the article



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Journal : BMCTwo 12894

Dispatch : 26-9-2020

Pages : 9

Article No : 725

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TYPESET

MS Code :

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20 years ago, a study in Italy suggested that the prevalence of HP infection was significantly higher in an infertile population than in controls, and antibodies against HP could be found in follicular fluids, semen, and vaginal secretions [3]. Ten years ago, a study in Japan found that the seropositive rate of HP in an infertile population with unknown etiology was higher than that in a population with known infertility factors, indicating that HP infection could be the cause of infertility [4]. In a cytotoxin-associated gene A (CagA)-positive population, the incidence of early pregnancy loss (EPL) after assisted reproductive technology increased significantly [5]. Recently, studies about infertility have focused on the impact of HP infection on sperm quality.

The first study from Italy reported a lower sperm quality in HP-infected patients with idiopathic infertility than in HP-uninfected patients. In CagA-positive patients, both sperm motility and fertility index are reduced [6, 7]. It has been suggested that anti-CagA antibodies might block spermatozoa acrosomes and disturb fertilization [8]. Further study found that compared with HP- patients, HP+ patients showed reduced sperm concentration, motility, and fertility index [9]. All the above studies indicate that HP infection may be a deteriorating factor for sperm quality, which deserves further investigation and treatment. However, most studies are from Italy, and additional data from different ethnicities may provide more robust evidence.

In females, there is a possible association between HP infection and polycystic ovarian syndrome (PCOS). A study from Turkey reported that the proportion of HP seropositivity was almost doubled in the PCOS population [10]. It is speculated that HP infection may lead to the release of certain substances or stimulate the immune response of the host, leading to the occurrence of PCOS. PCOS is manifested by increased ovarian reserve, while decreased ovarian reserve is an even worse problem that is difficult to treat. Since anti-Müllerian hormone (AMH) is an excellent indicator of ovarian reserve, we plan to investigate the association between HP infection and AMH.

Overall, there appears to be an association between HP infection and infertility, but available support is not sufficient and thus requires further validation. The purpose of the present study is as follows: (1) to investigate the correlation between HP infection and sperm quality in males and (2) to explore the association between HP infection and ovarian reserve in females.

Methods

Population of study

From January 2016 to June 2019, patients aged 20–50 years who came to the Second Hospital of

Zhejiang University School of Medicine due to abdominal discomfort and underwent HP testing were included in this study. Among them, 565 had plans for pregnancy and had performed sperm analysis or ovarian reserve tests in the past three months, who were involved for further analyses (Fig. 1).

Detection of HP infection

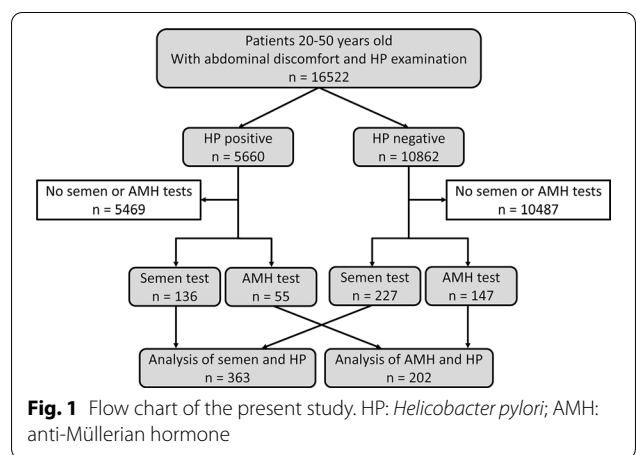
The ¹³C-urea breath test (UBT) or ¹⁴C-UBT was used to examine HP infection. Two breath samples were collected before and after ingestion of a ¹³C-urea (Richen-Force, Beijing, China) or ¹⁴C-urea (Xinke, Shanghai, China) reagent dissolved in water. For the ¹³C-urea breath test, a change over baseline value greater than 4.0 delta over baseline (DOB) was taken as a positive result (HP+). For ¹⁴C-UBT, a result greater than 100 DPM was taken as a positive result (HP+).

Detection of sperm parameters

Sperm samples were collected with sterile containers by masturbation after 2–7 days of sexual abstinence. After liquefaction at 37 °C for 30 min, routine parameters including sperm concentration and motility were examined with a computer-assisted sperm analysis (CASA) system (WLJY-9000, Beijing, China) according to World Health Organization guidelines [11]. Sperm morphology was assessed by the Papanicolaou staining modified for spermatozoa according to World Health Organization guidelines [11].

Detection of serum AMH and sex hormones

Serum AMH was tested with the electrochemiluminescence method by an Elecsys® AMH from Roche Diagnostics on a Roche Cobas e602 analyzer. The total imprecision for the assays was 1.2% at a level of 1.19 ng/mL with a measuring range of 0.01–23 ng/mL. Serum



sex hormone levels were detected with the electrochemiluminescence method by kits from Siemens Healthcare Diagnostics Inc.. The total imprecision for the assays was 3.0% at a level of 10,585 pmol/L for estradiol (E₂), 12.6% at a level of 0.37 nmol/L for testosterone (T), 2.7% at a level of 4.2 IU/L for luteinizing hormone (LH), 3.9% at a level of 6.9 IU/L for follicle-stimulating hormone (FSH), 4.8% at a level of 69.9 mIU/L for prolactin (PRL), and 12.7% at a level of 3.8 nmol/L for progesterone (P). The measuring ranges were 43.6–11,010 pmol/L for E₂, 0.24–52.05 nmol/L for T, 0.07–200 IU/L for LH, 0.3–200 IU/L for FSH, 6.4–4240 mIU/L for PRL, and 0.67–190.8 nmol/L for P.

Search strategy and data extraction

To search for studies investigating the correlation between HP and sperm parameters, two reviewers independently searched the studies published in English via three databases, including PubMed, Embase, and Cochrane CENTRAL, until June 30, 2019. Articles were identified through computerized searches using the keywords as follows: ("semen analysis" OR "sperm count" OR "sperm motility") AND ("*Helicobacter pylori*" OR "*Campylobacter pylori*"). Meanwhile, we hand-searched the references listed in the achieved papers to obtain additional studies.

Two reviewers extracted the common characteristics and outcome parameters of the searched manuscripts independently. The common characteristics included the name of the first author, publication year, country, and number of patients. The clinical outcomes included sperm concentration and progressive motility percentage (PR).

Statistical analysis

Analyses were performed by using the SPSS 19.0 statistics package (SPSS, Chicago, IL, USA). Continuous variables are expressed as the mean values ± standard deviation (SD). Student's t test was used for comparisons between two groups. Pearson correlation analysis was performed to analyze the relationship between two variables. A P value of <0.05 was considered statistically significant.

Data from our hospital and previously published results were pooled and calculated together by Review Manager Software (RevMan Version 5.3). When the mean and SD were not provided in the published article, we used formulas to estimate them [12–14]. The results were presented as the mean difference (MD) and 95% confidence interval (CI), and statistical significance was calculated by the Z test. If there was no serious heterogeneity (P value ≥ 0.1 by the Q test), a fixed-effects model (FEM) was applied for calculation, and if there was

serious heterogeneity, a random-effects model (REM) was applied [15].

Results

Baseline characteristics of the involved population.

As shown in Fig. 1, a total of 16,522 patients who underwent the HP test were included in this study. Among these patients, 34.26% (5660) were HP positive. Among the patients with HP infection, 136 underwent the sperm test, and 55 underwent the AMH test. Among the patients without HP infection, 227 underwent the sperm test, and 147 underwent the AMH test. Finally, 363 were involved in the analysis between sperm and HP and 202 between AMH and HP.

As shown in Table 1, the baseline characteristics were similar in both the sperm and AMH analyses. In the analysis of sperm and HP, there was no significant difference in age, weight, height, or body mass index (BMI) between the HP+ and HP- groups (P>0.05). Similarly, in the analysis of AMH and HP, no significant difference was found in age, weight, height, or BMI between the HP+ and HP- groups (P>0.05).

Comparison of sperm parameters between groups with or without HP infection

As shown in Table 1, the mean sperm concentration was 53.00 × 10⁶ Sp/mL and 53.90 × 10⁶ Sp/mL in the HP+ and HP- groups, respectively, with no significant difference (P>0.05). Sperm PR was also similar between the HP+ and HP- groups (39.39% vs. 39.92%), with no significant difference (P>0.05). There was no difference in either normal sperm morphology percentage or sperm head defects (P>0.05).

To further exclude the impact of age, we divided the population into subgroups of 20–24, 25–29, 30–34, 35–39, 40–44, and 45–50 years of age. As shown in Fig. 2a, c, there was no significant difference in sperm concentration or PR between the HP+ and HP- groups for any age group (P>0.05). As shown in Fig. 2b, d, in both the HP+ and HP- groups, there was no significant correlation between sperm concentration and age or between PR and age (P>0.05).

Comparison of AMH and sex hormones between groups with or without HP infection

As shown in Table 1, the mean serum AMH level was 3.49 in the HP+ group and 3.25 in the HP- group, with no significant difference (P>0.05). No difference was found between HP+ and HP- groups in serum E₂, T, LH, FSH, PRL, or P levels (P>0.05).

As shown in Fig. 2e, there was no significant difference in AMH level between the HP+ and HP- groups in every age span (P>0.05). Meanwhile, AMH correlated

Table 1 Characteristics of the present study

Characteristics	Sperm and HP			AMH and HP		
	HP+	HP-	P	HP+	HP-	P
	n = 136	n = 227		n = 55	n = 147	
Age (y)	31.08 ± 4.28	31.19 ± 4.52	0.822	32.89 ± 6.82	34.27 ± 6.83	0.202
Weight (kg)	69.72 ± 10.05	69.58 ± 9.82	0.898	52.91 ± 8.57	52.85 ± 7.01	0.961
Height (m)	1.74 ± 0.06	1.74 ± 0.06	0.838	1.61 ± 0.05	1.61 ± 0.04	0.546
BMI (kg/m ²)	22.98 ± 2.78	22.97 ± 2.78	0.967	20.26 ± 2.72	20.39 ± 2.50	0.757
Conc. (Sp/ml × 10 ⁶)	53.00 ± 42.36	53.90 ± 46.95	0.855	NA	NA	NA
PR (%)	39.39 ± 18.61	39.92 ± 18.81	0.793	NA	NA	NA
Normal (%)	6.73 ± 3.97	6.63 ± 4.43	0.865	NA	NA	NA
Head (%)	86.61 ± 8.71	84.77 ± 13.44	0.248	NA	NA	NA
AMH (ng/ml)	NA	NA	NA	3.49 ± 3.03	3.25 ± 2.82	0.605
E2 (pmol/L)	NA	NA	NA	194.46 ± 78.53	192.56 ± 68.24	0.906
T (nmol/L)	NA	NA	NA	0.94 ± 0.71	1.12 ± 0.78	0.118
LH (IU/L)	NA	NA	NA	6.31 ± 4.32	4.98 ± 3.32	0.106
FSH (IU/L)	NA	NA	NA	8.36 ± 3.84	8.38 ± 3.43	0.978
PRL (mIU/L)	NA	NA	NA	256.19 ± 154.03	234.81 ± 129.36	0.489
P (nmol/L)	NA	NA	NA	1.99 ± 1.03	1.65 ± 0.97	0.128

BMI: body mass index; AMH: anti-Müllerian hormone; NA: not available; Conc.: concentration; PR: progressive motility; Normal: normal sperm morphology percentage; Head: sperm head defects; DFI: sperm DNA fragmentation index; E₂: estradiol; T: testosterone; LH: luteinizing hormone; FSH: follicle-stimulating hormone; PRL: prolactin; P: progesterone

221 significantly negatively with age (Fig. 2f, for HP-, Pear-
 222 son correlation coefficient = -0.482, P=0.000; for HP+,
 223 Pearson correlation coefficient = -0.431, P=0.001).

224 **Pooled analysis of the association between sperm**
 225 **parameters and HP infection**

226 Since 2010, six studies investigated the correlation
 227 between HP infection and sperm parameters, as listed
 228 in Table 2. Most studies found that sperm motility was
 229 reduced significantly in CagA+ patients [6, 7, 16, 17]. The
 230 latest study found that sperm concentration and PR were
 231 reduced in the HP+ population, and in the CagA+ popu-
 232 lation PR was reduced further than in the CagA- popu-
 233 lation [9].

234 Five studies including 703 participants were pooled to
 235 compare HP+ and HP- groups, and five studies includ-
 236 ing 210 participants were pooled to compare CagA+ and
 237 CagA- groups. As shown in Fig. 3a, c, FEM analysis
 238 showed that there was no significant difference in sperm
 239 concentration between the HP+ and HP- groups or
 240 between the CagA+ and CagA- groups (P>0.05 for
 241 both). In the sperm motility analysis between HP+
 242 and HP-, since serious heterogeneity (P<0.01) was
 243 found, a REM was applied and suggested no signifi-
 244 cant difference in PR (Fig. 3b, 95% CI -11.44 to 1.87,
 245 P=0.16). FEM analysis was applied to compare sperm
 246 PR between CagA+ and CagA- groups, which suggested
 247 that PR was 16.18% lower in the CagA+ group than in

the CagA- group (Fig. 3d, 95% CI -18.86 to -13.50, P<0.01).

250 **Discussion**

251 In the present study, no difference was found in sperm
 252 concentration or sperm motility between HP+ and
 253 HP- groups. Further subgroup analyses confirmed that
 254 there was no significant difference in sperm parameters
 255 between HP+ and HP- groups. Furthermore, we pooled
 256 our data and those of previous studies and found no sig-
 257 nificant difference in sperm concentration or motility,
 258 indicating that in the Chinese population, HP infection
 259 does not disturb spermatogenesis.

260 The results of previous studies were not consistent.
 261 Some suggested that sperm concentration and motil-
 262 ity were reduced in HP+ patients [9] and that treating
 263 HP could improve the quality of sperm [18], while some
 264 suggested no significant difference in sperm parameters
 265 between HP+ and HP- groups [6, 7, 16, 17]. The incon-
 266 sistent results may be due to different test methods and
 267 ethnicities investigated.

268 This is the first study that used UBT to detect HP infec-
 269 tion and to investigate its relationship with sperm qual-
 270 ity. In previous studies, HP infection was detected with
 271 a serology test by enzyme-linked immunosorbent assay
 272 (ELISA) and confirmed with western blotting (WB) [6,
 273 7, 9, 16-18], whereas in the present study, ¹³C- and ¹⁴C-
 274 UBT were used to detect HP infection. UBT is the best

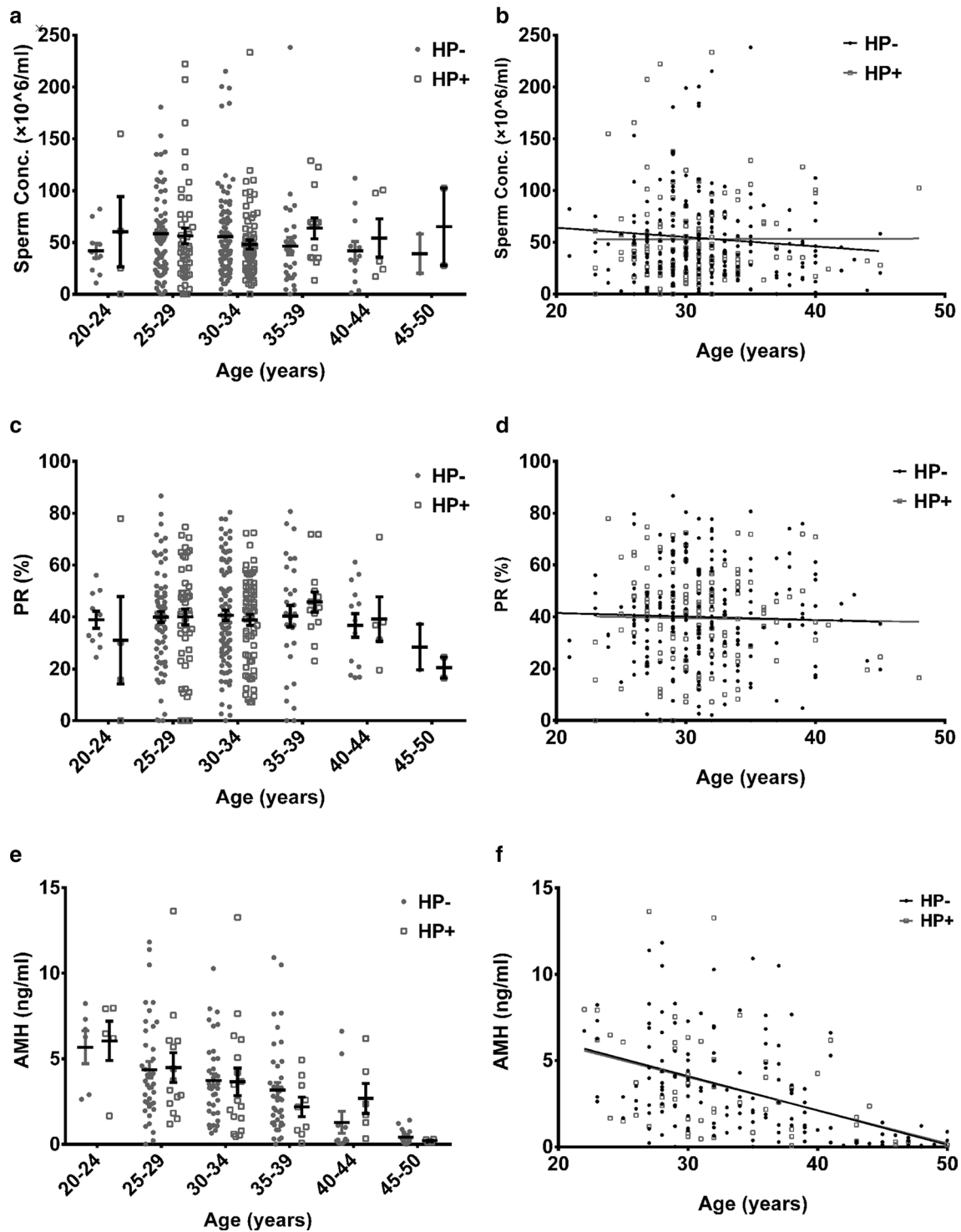


Fig. 2 Correlation between sperm parameters and anti-Müllerian hormone (AMH) and *Helicobacter pylori* (HP) infection. **a, b** Correlation between AMH and HP. **c, d** Correlation between sperm concentration (Conc.) and HP. **e, f** Correlation between progressive motility percentage (PR) and HP



Table 2 Previous studies investigating the impact of HP infection on sperm parameters

Study	Public year	Country	HP (+ vs. -)			CagA (+ vs. -)			Results
			Population	Conc	PR (%)	Population	Conc	PR (%)	
Moretti [9]	2017	Italy	32 vs. 41	38.0 vs. 55.0*	17.0 vs. 34.0***	20 vs. 12	33.4 vs. 42.5	10.5 vs. 22.5***	HP+ reduced conc. and PR
Moretti [17]	2015	Italy	28 vs.81	61 vs. 72	32 vs. 32	12 vs. 16	61 vs. 61.5	24 vs. 36.5*	CagA+ reduce PR
El-Garem [18]	2014	Egypt	NA	NA	NA	22 vs. 201	NA	NA	HP treatment improved PR
Moretti [16]	2013	Italy	NA	NA	NA	37 vs. 50	58 vs. 63	18 vs. 32**	CagA+ reduced PR
Moretti [7]	2012	Italy	27 vs. 51	94 vs. 72	32 vs. 30	11 vs. 16	65 vs. 94	24 vs. 38**	CagA+ reduced PR
Collodel [6]	2010	Italy	36 vs.44	24.5 vs. 23.5	22 vs. 28.5	17 vs. 19	25.5 vs. 23	18 vs. 29*	CagA+ reduced PR

NA: not available; Conc.: sperm concentration; PR: progressive motility; CagA: cytotoxin-associated gene A

* $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$

275 noninvasive method for patients without gastric resec- 312
 276 tion or proton pump inhibitor (PPI) treatment, with both 313
 277 high positive predictive value and negative predictive 314
 278 value [19–21]. A meta-analysis suggested that UBT had 315
 279 high diagnostic accuracy for detecting HP infection in 316
 280 patients with dyspepsia, with a pooled sensitivity of UBT 317
 281 in adult patients of 96% and a pooled specificity of 93% 318
 282 [22, 23]. Another meta-analysis involving 34 studies with 319
 283 serology evaluation and 57 studies with UBT detection 320
 284 reported that the sensitivity of HP diagnosis was 0.94 for 321
 285 ¹³C-UBT, 0.92 for ¹⁴C-UBT, and 0.84 for serology tests. 322
 286 UBT showed a higher diagnostic accuracy than the serology 323
 287 test for HP infection diagnosis [24, 25]. Therefore, in 324
 288 this study, UBT was used, as it provides a more accurate 325
 289 HP diagnosis than serology tests. 326

290 Moreover, serology tests cannot distinguish between 327
 291 active and inactive infections [26]. In a letter from Cav- 328
 292 iglia et al., the authors emphasized that the presence of 329
 293 serological HP antibodies could only indicate previous 330
 294 exposure, not necessarily a current infection, and based 331
 295 on this, they recommended UBT as a direct diagnostic 332
 296 test [27]. Similarly, in the present study, UBT examina- 333
 297 tion represented the status of current HP infection better 334
 298 than serology tests. 335

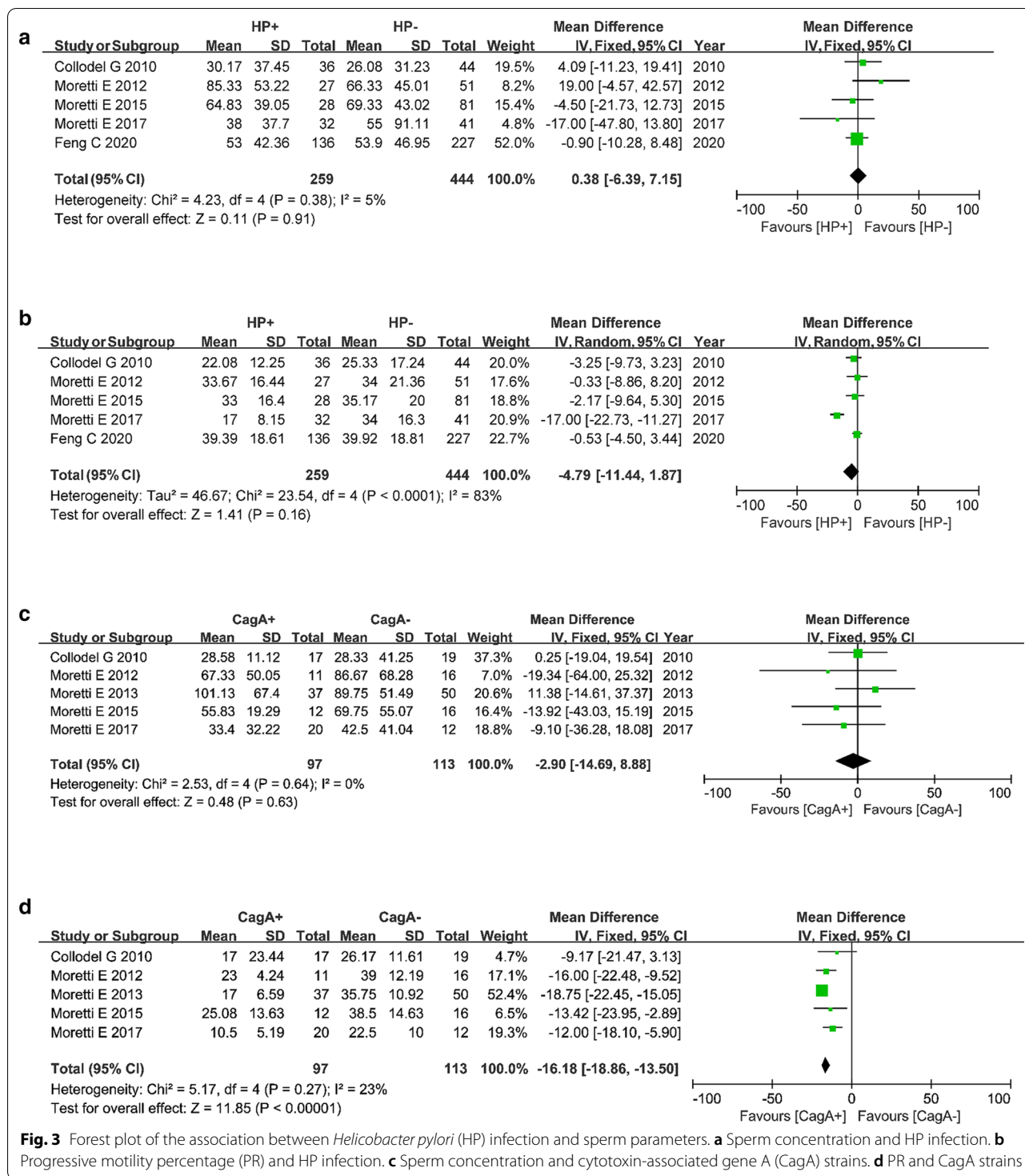
299 CagA is the major virulence factor in HP, encoding 336
 300 the CagA protein in the cag pathogenicity island [28]. 337
 301 HP infection can be divided into two isolates: CagA- 338
 302 producing strains (CagA+) and CagA-nonproducing 339
 303 strains (CagA-). Our meta-analysis of sperm motility 340
 304 and CagA-producing/nonproducing strain infection sug- 341
 305 gested that PR was 16.18% lower in the CagA+ group 342
 306 than in the CagA- group. The underlying mechanism 343
 307 may be that CagA+ HP infection induces overexpression 344
 308 of miR-543 and downregulation of the p14ARF tumor 345
 309 suppressor to inhibit autophagy and increase cytokine 346
 310 production, which induces inflammatory responses of 347
 311 HP accordingly [29–31]. Anti-CagA antibodies may 348

block spermatozoa acrosomes and disturb fertilization [8].

The prevalence of the CagA genotype in HP infection varies significantly among different regions. In Western countries, CagA+ strains comprise 50–60% of the HP+ population, and in the Chinese population, CagA+ strains occupy nearly 100% of the HP+ population [32, 33]. Studies investigating the CagA status of Chinese HP strains with polymerase chain reaction (PCR) detected CagA genotypes in nearly all strains [34, 35]. Considering the high CagA positivity in the Chinese HP+ population, the sperm concentration and motility should be weakened in HP+ patients, but the present study showed stable parameters. Further study investigating CagA antibody status should be performed to clarify the role of CagA in sperm quality.

The variability of sperm parameters after HP treatment is an interesting question. It was reported that after the treatment of HP, seminal HP IgA level decreased significantly, and meanwhile progressive sperm motility, nonprogressive sperm motility, and sperm normal forms increased significantly ($P = 0.001$) [18]. In the present study, sperm analyses were performed before HP test, and most patients with HP+ suspended their plans of pregnancy after HP treatment. Therefore, we did not follow the sperm parameters.

In the present study, there was no difference in serum AMH level between HP+ and HP- groups, which was confirmed with further age-divided subgroup analyses. Published results of the relationship between PCOS and HP infection are inconsistent. Yavasoglu et al. found that HP antibody positivity was significantly more common in the PCOS group than in the age-matched control group [10]. The possible explanation may be that the antigenic mimicry to HP antigens leads to an immune cross-reaction between HP antigens and the ovaries, inducing the onset of PCOS [36]. Nevertheless,



349 Tokmak et al. found no significant difference in HP
 350 IgG positivity between PCOS and non-PCOS groups
 351 [37]. AMH is a potential future substitute for detect-
 352 ing polycystic ovarian morphology (PCOM) and a

useful biomarker for predicting the risk of PCOS [38–
 40]. Our data indicated no correlation between PCOS
 and HP infection. Meanwhile, AMH is considered the
 best serum biomarker of ovarian reserve, reflecting
 the number of primordial follicles and its response to

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358 exogenous gonadotropins [41]. The present study indi-
 359 cates that ovarian reserve is stable with HP infection.

360 **Conclusion**

361 This is the first observation investigating the impact of
 362 HP infection on ovarian reserve, which found that HP
 363 infection was not related to the serum ovarian reserve
 364 biomarker AMH. In general, HP infection is not a
 365 crucial factor affecting sperm parameters or ovarian
 366 reserve.
 367

368 **Abbreviations**

369 DOB: Delta over baseline; E₂: Estradiol; EPL: Early pregnancy loss; FEM: Fixed-
 370 effects model; FSH: Follicle-stimulating hormone; HP: *Helicobacter pylori*; LH:
 371 Luteinizing hormone; MD: Mean difference; P: Progesterone; PCOM: Polycystic
 372 ovarian morphology; PCOS: Polycystic ovarian syndrome; PCR: Polymerase
 373 chain reaction; PPI: Proton pump inhibitor; PR: Progressive motility percent-
 374 age; PRL: Prolactin; REM: Random-effects model; SD: Standard deviation; T:
 375 Testosterone; UBT: Urea breath test; WB: Western blotting.

376 **Acknowledgements**

377 We are especially grateful to Ya-Wen Xing for his kind help in data collection.

378 **Authors' contributions**

379 CF, MJ, and JMS conceived and designed the study; CF, MJ and CCH enrolled
 380 the patients and collected the data; CF, PPL, SQY, and QPY analyzed the data;
 381 PPL and MJ reviewed the references and extracted the data; CF, MJ, and JMS
 382 organized and wrote the manuscript. All authors read and approved the final
 383 manuscript.

384 **Funding**

385 This work was financially supported by the following funding sources: the,
 386 including National Key R&D Program of China (2018YFC1004900 to CF),
 387 National Natural Science Foundation of China (81871176 to CF, 81701461 to
 388 PPL, and 81671487 to MJ), and the Medical and Health Science and Technol-
 389 ogy Plan of Zhejiang Province (2018251579 to PPL) for the English language
 390 editing service costs, and article-processing charge. No funders had any role in
 391 the study design, data analysis of data, or writing manuscript.

392 **Availability of data and materials**

393 The data analyzed in this study are available from the corresponding author
 394 upon request.

395 **Ethics approval and consent to participate**

396 The present study was approved by the Ethics Committee of the Second
 397 Affiliated Hospital of Zhejiang University School of Medicine (IR2019001059).
 398 Since this was a retrospective observational study and no intervention was
 399 needed, no formal ethical approval or written consent was required, which
 400 was approved by the Ethics Committee of the Second Affiliated Hospital of
 401 Zhejiang University School of Medicine.

402 **Consent for publication**

403 Not applicable.

404 **Competing interests**

405 The authors declare that they have no competing interests.

406 **Author details**

407 ¹ Department of Reproductive Medicine, The Second Affiliated Hospital
 408 of Zhejiang University School of Medicine, 88 Jiefang Road, Hangzhou 310009,
 409 Zhejiang, China. ² The Women's Hospital of Zhejiang University School
 410 of Medicine, Hangzhou 310006, Zhejiang, China. ³ Department of Orthope-
 411 dics, The First Affiliated Hospital of Zhejiang Chinese Medicine University, 54
 412 Youdian Road, Hangzhou 310006, Zhejiang, China.

Received: 21 December 2019 Accepted: 24 September 2020

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