AVIAN GASTRIC YEAST (AGY) INFECTION (MACRORHABDIOsis OR MEGABACTERIOSIS)

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Summary


The review on avian gastric yeast (AGY) aimed to collect and update available information about this infection from clinical point of view which can help for better understanding of the disease. Macrorhabdus ornithogaster (M. ornithogaster) is the causative agent of proventriculitis in birds. The organism is large, rod-like, classified as anamorphic ascomycetous yeast. Clinical signs may vary from acute with sudden death or chronic wasting. Diarrhoea or enteritis has also been reported in infected birds. Birds can have other concurrent infections (enteric parasites, bacteria or viruses). Gross lesions may include proventricular oedema, hyperaemia, or haemorrhage, with overproduction of mucus accumulated in the proventricular lumen. The proventriculus may be dilated with or without ulceration in mucosa. The organism can be cultured on De Man, Rogosa and Sharpe (MRS) agar, but it is easily detected in Gram’s, periodic acid-Schiff (PAS) and Giemsa stained proventricular/ventricular junction, or isthmus sections. Histological changes are more prominent in the ventriculus. Affected birds have marked disruption of the koilin layer with disorganisation and degeneration, and demonstrate large numbers of yeasts with matchstick or logjam appearance. The organism can be detected by polymerase chain reaction (PCR). The disease is common in budgerigars, canaries, finches, and parrotlets. Organisms are identified retrospectively in approximately one-fourth of canaries and budgerigars. The infection has also been reported in chickens, partridges, and ostriches but mildly in chickens. The affected birds can be successfully treated with amphotericin B.

Key words: budgerigars, histologic changes, M. ornithogaster, proventriculitis, yeast

INTRODUCTION

M. ornithogaster is the causative organism of avian gastric yeast (AGY). M. ornithogaster is an anamorphic Ascomycota yeast (Tomaszewski et al., 2003). It infects a wide range of bird species as chickens, turkeys, ostriches, several species of parrots, passerine species, in addition to captive-bred and wild finches (Phalen, 2014). The organism has a worldwide prevalence and differs extensively in pathogenicity. It was first described in canaries by Van Herck et al.
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(1984). Under some conditions, *M. ornithogaster* infection does not show any signs. Certain individual birds will show signs as a result of infection with *M. ornithogaster*. In some species of birds, the prevalence of the disease may be high. This can be attributed to variation in the pathogenicity of *M. ornithogaster* strains or variation in susceptibility of the affected birds.

AGY is a highly contagious disease where *M. ornithogaster* colonises the proventriculus close to the proventricular-ventricular isthmus of birds and has not been recognised in other parts of the body or in the environment (Tomaszewski et al., 2003; Phalen, 2005; Hannafusa et al., 2007). Frequently affecting budgerigars, the disease appears without any symptoms in a chronic course and even the whole flock can become infected without showing any symptoms for several weeks or months. The chronic course can be interrupted by intermittent acute episodes. The disease has a negative influence on the digestion; therefore the affected bird loses weight in spite of excessive feeding.

SYNONYMS

*M. ornithogaster* name is derived from Greek words macrorhabdus (meaning long rod), and ornithogaster (meaning stomach of bird). *M. ornithogaster* (Megabacterium) disease has many synonyms as megabacteriosis, macrorhabdosis, going light (GL) (Baker, 1985), wasting disease, budgie wasting disease (Henderson et al., 1988), bacteria giganticus, Megabacteria Associated Disease (MAD) (Perry, 1993), or Proventricular/Ventricular Disease (PVD) (Filippich & Parker, 1994). The commonest name, however, is avian gastric yeast (AGY) infection.

HISTORY

*M. ornithogaster* was recognised as a bacterium for a long time; however latest research studies affirmed that the organism was a yeast (Tomaszewski et al., 2003; Hannafusa et al., 2007). The organism is a large Gram-positive bacillus. It was generally noticed in the proventriculus of budgerigars like normal flora (Scanlan & Graham, 1990).

Since 2000, *M. ornithogaster* "megabacteriosis" was distinguished clinically by emaciation, prostration, loss of appetite, cachexia and death. Normal chronic course was observed in chickens, turkeys, guinea fowls while acute disease was seen in the canary (*Serinus canarius*), the zebra finch (*Taeniopygia guttata*) and budgerigars (*Melopsittacus undulatus*) (Martins et al., 2006).

ECONOMIC IMPORTANCE

The infection can be subclinical and birds appear normal. Variable mortality rates were reported in infected avian species: 50% mortality in rhea, 100% in a free-range flock of guinea fowl and 40% to 80% mortality in affected ostrich flocks (Huchzermeyer et al., 1993). In chickens, experimental infection with *M. ornithogaster* showed only decrease in the feed conversion rate without any clinical signs (Phalen & Moore, 2003).

CHARACTERISATION OF *M. ORNITHOGASTER*

**Scientific classification**

*M. ornithogaster* is the only member of genus *Macrorhabdus*, Order Saccharomycocetales, Class Saccharomycetes, Division
Ascomycota, Kingdom Fungi (Tomaszewski et al., 2001).

Morphology

M. ornithogaster is an ascomycetous yeast (Tomaszewski et al., 2003). The vegetative cells are motile, elongated (2 to 20 µm) and divided by fission. The cells are single or in short chains of two to four cells. It is Gram positive, but only the cytoplasm stains with Gram stain (Hannafusa et al., 2007). In mucosal scrapings and in the faeces of infected birds, the organism appears as a stiff, straight rod, 20 to 80 µm long and 2 to 3 µm wide, with rounded ends. The long rods may bend slightly in a gentle curve. On a wet mount, the organisms are viewed directly as small oblong refractile structures while the nuclei, found at regular intervals, are readily seen. The nucleus was demonstrated in M. ornithogaster isolates from budgerigars by electron microscopy and in situ hybridization with a pan-eukaryote rRNA probe was positive (Ravelhofer-Rotheneder et al., 2000). On electron microscopy, it had a nucleus that contained eukaryotic ribosomal DNA as reported by Ravelhofer-Rotheneder et al. (2000). Motility pattern of M. ornithogaster was seen as cellular lateral swinging or linear forward movements (Martins et al., 2006).

Staining

M. ornithogaster stained weakly basophilic by haematoxylin and eosin (H&E) and positive periodic acid reaction (Scanlan & Graham, 1990). It stained with silver stains and the periodic acid-Schiff stain (PAS) (Dorrestein et al., 1980; Har- greaves, 1981). The organism stained strongly with stains that bind the polysaccharide chitin as blanchopeur BA (Ravelhofer et al., 1998) and calcofluor white M2R (Moore et al., 2001). The organism was Gram-positive, PAS-positive and acidophilic in Gram’s, PAS and Giemsa stained sections, respectively (Kheirandish & Salehi, 2011).

Biochemical features

The isolates were catalase-negative and oxidase-negative and did not reduce nitrate. All isolates failed to utilise arginine, lysine, ornithine or tryptophan but produced acid from glucose, galactose, levulose, maltose, melibiose, starch, and sucrose. All isolates produced acetoin from glucose and hydrolysed esculin (Scanlan & Graham, 1990).

Culture media

The bacterium is extremely pleomorphic, facultatively anaerobic and capnophilic. When subcultured on agar media, it changes particularly in both diameter and length. After incubation on blood agar for several days, the bacterium is haemolytic, forming flat colonies with irregular edges. All isolates were reported to grow on sodium azide agar but not on MacConkey agar (Scanlan & Graham, 1990). Isolation of this organism on De Man, Rogosa and Sharpe (MRS) agar medium was reported (Huchzermeyer et al., 1993; Gerlach, 2001). The organism appeared to grow gradually in cell culture media enhanced with dextrose, foetal calf serum, in addition to antibiotics (Ravelhofer-Rotheneder et al., 2000).

NATURAL HOSTS

AGY was reported in several avian species including canaries (van Herck et al.,
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1984), budgerigars (Scanlan & Graham, 1990; Henderson et al., 1988; Baker et al., 1992; Filippich & Hendrikz, 1998; Pennycott et al., 1998), ostriches (Huchzermeier et al., 1993; Baker, 1997; Martins et al., 2004), chickens (Mutlu, 1997; Schulze & Heidrich, 2001) and turkeys (Schulze & Heidrich, 2001). Also, Martins et al. (2006) reported the disease in ostrich, rhea, canary, budgerigar, zebra finch, industrial broiler, free range chicken, turkey, guinea-fowl, domestic pigeon, ruddy ground-dove, toucan, chukar partridge. Macrorhabdus-like agents were reported also in mammals as dogs, cats and laboratory mice (Cooke, 2000; Rossi, 2000).

EXPERIMENTAL HOSTS

In chickens and Japanese quails, the attempts for experimental infection by using chicken isolate revealed the presence of M. ornithogaster in apparently normal birds in two organs (proventriculus and liver) (Martins et al., 2006). Experimental intraperitoneal infection of mice with 10^3 CFU/dose was resulted in 100% mortality (Martins et al., 2006), while Hanafusa et al. (2013) was failed to induce infection in mice.

INCIDENTE AND DISTRIBUTION

The large rod-shaped organism could be identified in both infected and normal chickens, turkeys, quails and pigeons, rheas (Rhea americana), ostriches (Struthio camelus), canaries, zebra-finches, guinea-fowl (Numida meleagris) and budgerigars (Martins et al., 2006). The prevalence of infection in budgerigar, parrotlet, canary, and finch aviaries was often high (Dorrestein et al., 1980; Simpson, 1992; Ravelhofer et al., 1998; Baker, 1985; Filippich & Herdrikz, 1998). The prevalence of M. ornithogaster among cockatiels, budgerigars, lovebirds and barred parakeets was: 42.4% (14/19), 29.0% (9/22), 50.0% (2/4) and 0% (0/3) respectively in Uberaba, state of Minas Gerais (Paula et al., 2018). The infection was also reported in wild parrots including the galah and sulphur-crested cockatoo (Filippich & Parker, 1993; Phalen et al., 2007; Doneley, 2012). In Germany, Hanka et al. (2010) reported the infection in birds of 16 orders through Giemsa stained impression smears of the glandular stomach surface and M. ornithogaster was found in different frequencies in birds of the orders Psitaciformes, Gruiformes, Galliformes and Columbiformes.

M. ornithogaster has been described in chickens in Europe, North and South America, and Australia (Mutlu et al., 1997; Schulze & Heidrich, 2001; Hanka et al., 2010; Behnke & Fletcher, 2011; Martins et al., 2006; Phalen et al., 2007). Also, Japanese quails, partridges and turkeys have been reported to be infected with M. ornithogaster (Martins et al., 2006; Jansson et al., 2008; Hanka et al., 2010). Hanka et al. (2010) reported the infection in domestic ducks.

The infection with M. ornithogaster does not result in disease under most circumstances. Therefore, the recognition of the organism in a sick or dead bird does not confirm that it was the cause of the bird's sickness. Predisposing factors to M. ornithogaster--associated disease comprise variation in M. ornithogaster strains, poor management in addition to genetic factors of the host (Filippich & Perry, 1993; Speer et al., 2004; Filippich & Hendrikz, 1998).
TRANSMISSION

M. ornithogaster, agent of megabacteriosis, of yeast form, is frequently encountered in the normal flora of bird’s proventriculus. Under nonspecific circumstances, the host-fungus balance is disturbed, and the so-called “going light syndrome” clinical signs are manifested (Queirós et al., 2011). M. ornithogaster colonises the isthmus (narrow junction) of the proventriculus (glandular stomach) and the ventriculus (grinding stomach) of birds (van Herck et al., 1984). Healthy appearing normal birds can shed the organism with faeces while diseased birds may not continuously shed. Likely, most infections result from faecal-oral contamination from sick or subclinical birds shedding M. ornithogaster in their faeces (Kheirandish & Salehi, 2011; Lanzarot et al., 2013). If one bird feeds another, the pathogen can be spread. A chronically infected bird that seems to be healthy can introduce and spread the disease when introduced into a new flock. Birds suffering from a megabacterial infection often show secondary bacterial infections. Detection of the infection could occur in some avian species along with other infectious or disease problems, for example, endoparasites (helminths, coccidia) and ectoparasitism (order Mallophaga and/or order Acarina) (Martins et al., 2006). Transmission may happen among species and is supported by the absence of a traditional biosecurity policy (Martins et al., 2006).

It was suggested the implication of genetic factors as the M. ornithogaster prevalence was significantly greater in chicks from parents positive for M. ornithogaster than from M. ornithogaster-negative parents, even when reared by M. ornithogaster-negative parents (Filippich & Hendrikz, 1998; Hoppes, 2013). Infection was found to pass from experimentally infected chicks to uninfected chicks housed with them (Phalen & Moore, 2003). Wild birds were considered as possible source of infection. M. ornithogaster infections were identified in 13 laying hens and 4 cocks located in 14 different flocks in addition to one turkey. As all infected birds were reared under circumstances permitting contact to wild birds and M. ornithogaster infections have been detected earlier in wild-living green finches, wild birds were considered a potential source of infection (Schulze & Heidrich, 2001). Various conditions such as recent shipping, crowded housing, reproductive activities, mixed species avaiaries are commonly reported as stressors.

CLINICAL SIGNS

Generally, signs are non-specific varying from asymptomatic carriedness to dangerous illness with high mortality. Birds become emaciated and suffer from anorexia, vomiting, cachexia and death. Dropping consistency varies from slight softness to severe diarrhoea. In some cases, birds eat frenziedly, but in fact are just grinding seeds without ingestion. In the acute stage, birds die quickly within few days. In the chronic stage, they become gradually more emaciated and weakened then die within weeks or months, or recover but with retrogression after weeks or months (Gerlach, 2001; Moore et al., 2001; Martins et al., 2006).

Budgerigars and parrotlets

Two clinical forms of the infection were reported in budgerigars.

- Acute form

The apparently healthy birds suddenly reduce the food intake than normal, regur-
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Gnitate ingesta (which may be blood stained), then die within 2 days (Baker, 1985; Filippich et al., 1993; Moore et al., 2001; Phalen et al., 2002). In the green-rumped parrotlet, infection and disease seems to be widespread. Also, clinical signs are widespread in middle-aged budgerigars, the infection begins very early and large numbers of organisms were detected in the isthmus of nestling birds aged 12 days (Phalen et al., 2002; Phalen, 2006).

• Chronic form

Affected birds typically seem to be hungry and settle long time at the food dish. Rather than eating, the birds are grinding the food without ingesting it. Regurgitation is frequent, so the tops of affected birds’ heads usually contain fresh or dried saliva. Undigested seeds may be present in the droppings. Diarrhoea with or without melena may also be present. These birds go through a long period of weight loss (going light) looking unhealthy and finally, they die (Baker, 1992; Filippich et al., 1993; Antinof, 2004; Phalen, 2006). Few birds with the chronic form show clinical signs in the large affected budgerigar colony. Most commonly, affected birds are mature birds with an average age of 2.7 years (Filippich et al., 1993).

Canaries and finches

Initially, owners perceive that there is a problem when a skinny bird is discovered dead (van Herck et al., 1984; Phalen et al., 2002). The infection in canaries and finches likely resembles the chronic form shown in budgerigars (Dorrestein et al., 1980; van Herck et al., 1984; Filippich & Parker, 1994).

The avian gastric yeast (megabacteriosis) recognised by high morbidity and mortality was reported in a flock of zebra finches, the adults being exclusively affected. Clinical signs comprised puffed and ruffled feathers, feather loss, lethargy, and raised respiratory stress (Snyder et al., 2013).

Endoventricular mycosis results from yeast proliferation within the koilin of the ventriculus in finch and finch-like birds. The clinical signs vary from sudden death to weight loss and presence of undigested seeds in the droppings.

Chickens

Schulze & Heidrich (2001) demonstrated a megabacteriosis-associated proventriculitis in 13 laying hens and 4 cocks located in 14 different flocks in addition to one turkey. Birds suffered from progressive runting, high mortality as well as poor laying performance. At necropsy, the proventriculus was enlarged with thickened wall and the mucosa filled with cloudy, grey-white mucus. At the proventricular-ventricular junction mainly, petechial haemorrhages and ulcerations were observed accompanied with irregular sloughing of the necrotic koilin layer of the ventriculus. In outbreak of mortality in chickens and Japanese quails sharing the same airspace, megabacteria were observed in examined 8 out of 11 chickens and 16 out of 24 quails (Pennycott et al., 2003).

White Leghorn chickens experimentally infected with M. ornithogaster did not show any clinical signs, only the feed conversion rate was decreased in infected birds in comparison to non-infected controls (Phalen & Moore, 2003).

Ostriches

In ostriches, cases of M. ornithogaster infection were reported in 10-day to 12-week-old chicks. Birds appeared normal
but stopped growing and lost weight. Finally, they became weak and died. Birds were anaemic and had soiled vents. Some birds suffered from diarrhoea, while others excreted dry pelleted droppings. Mortality rates in affected flocks ranged from 40% to 80% (Huchzermeyer et al., 1993).

Pigeons

Hanka et al. (2010) examined a total of 1,137 birds in 16 orders for presence of *M. ornithogaster*. Most of the findings were accomplished on preparations from glandular stomach surface. *M. ornithogaster* was detected in birds of the orders Psittaci-, Passed-, Anseri-, Galli- and Columbiformes in different frequencies. The authors presented the first proof of *M. ornithogaster* infection in two diseased feral pigeons.

GROSS LESIONS

At necropsy, atrophy of pectoral muscle, accompanied with proventricular dilation was observed. Proventricular and ventricular walls became thickened containing thick white mucus of alkaline pH 7–7.3 in addition to koilin layer loosening. Haemorrhage and ulceration in the proventricular-ventricular junction are reported (Baker, 1992; Werther et al., 2000; Schulze & Heidrich, 2001; Marlier et al., 2006; Martins et al., 2006).

HISTOPATHOLOGY

Histologically, examined Gram-, PAS-, or silver-stained proventriculus sections showed pale eosinophilic organisms present at the tips of the glands of the isthmus lined up in parallel, like logs in a logjam. As the number of organisms increased, they moved into the spaces between the glands and could be seen on the surface of the koilin of the gizzard, and, at times, invade it. Atrophy of the glands of the isthmus and ulceration of the isthmus and koilin developed in the most severe cases. Before ulceration, little or no inflammation was present. When inflammation occurred, it was typically a lymphoplasmacytic infiltration of the lamina propria of the glands of the isthmus. A moderate thickening of the lamina propria of the isthmus glands was noted in chickens infected experimentally with *M. ornithogaster*. Dilation of the proventricular glands and disruption of the normal structure of the koilin are other lesions that may be detected (Dorrestein et al., 1980; Hargreaves, 1981; van Herck et al., 1984; Baker, 1985; Filippich et al., 1993). *M. ornithogaster* was histologically diagnosed in the mucosal isthmus of the proventriculus and ventriculus of adult hobby chickens showing intermittent signs of enteritis (Behnke & Fletcher, 2011). The most severe histopathological lesions in budgerigars were observed in the proventriculus and gizzard, especially at the proventricular-ventricular junction, including penetration of an organism to the lumen of the superficial proventricular crypts and occasionally to deeper parts of the glands; lymphocytic plasmacytic proventriculitis, lymphocytic plasmacytic ventriculitis and disruption of the koilin layer (Kheirandish & Salehi, 2011; Powers et al., 2019).

CLINICAL PATHOLOGY

Henderson et al. (1988) reported that budgerigars showing clinical signs of *M. ornithogaster* infection presented some haematologic changes as reduced packed cell volumes (anaemia), leukocytosis, heterophilia, monocytosis, lymphocytosis, basophilia, and thrombocytosis as well as
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low phosphate, sodium, chloride, glucose, cholesterol, and aspartate transference values.

DIAGNOSIS OF M. ORNITHOGASTER INFECTION

The characteristics of the disease and shape of the organisms found in the ulcerated gastric mucosa suggested megabacteriosis (Martins et al., 2006).

Diagnosis of M. ornithogaster infection is based on the evaluation of clinical signs, gross lesions and the microscopic examination of Gram or Giesma-stained direct smears from gastric mucosa, fresh faeces, or using mini-Flotac technique and detection of long rod-shaped organisms at 400 and 1000 times magnification (Martins et al., 2006; Cringoli et al., 2013; Borrelli et al., 2015). Birds may shed the organism intermittently, so a negative faecal examination does not exclude infection. Wet mount, modified Wright’s, or Gram stain of a faecal sample often reveal M. ornithogaster appearing as a large, Gram-positive rod, with mottling or stippling throughout its length (Hoppes, 2013). Examining five droppings from each bird will increase the chance of finding M. ornithogaster. Definitive diagnosis of megabacteriosis is most consistently demonstrated by histopathology and fresh smear of the proventricular mucus (Kheirandish & Salehi, 2011). Sullivan et al. (2017) compared polymerase chain reaction (PCR) of cloacal swab samples and faecal Gram’s stain (FGS) for diagnosis of active M. ornithogaster shedding in a captive flock of budgerigars, where 57% sampled birds were positive by PCR and 24 – by FGS. All FGS-tested birds were positive on PCR, but the overall percent agreement for the two methods was only 67%.

The experimental infection of mice using the chicken isolate may be an indication of the pathogenicity of the isolate (Martins et al., 2006).

Direct detection of M. ornithogaster cells in faeces by PCR (Phalen, 2014) and by culture methods using cloacal cotton-tipped swabs (Lanzarot et al., 2013). In budgerigars, PCR was reported to be more preferable than FGS in diagnosis of M. ornithogaster (Sullivan et al., 2017).

Isolation and identification

M. ornithogaster lives at the junction between proventriculus and ventriculus. Isolation of organism occurs mainly from crop wash, fresh faecal samples by sterile cotton-tipped swab or scraping of the gut lining. In case of crop wash and fresh faeces, samples are mixed with 20 times of their volume of saline. After 10 seconds, a wet preparation of the surface film was ready for preparation. Organisms can be seen by the 10× objective on the microscope. Fresh impression smears of gut mucosa are prepared onto glass slides for observation after staining by Giemsa and Gram’s methods under light microscopy (Hannafusa et al., 2007; Lanzarot et al., 2013).

Molecular diagnosis

Genetic characterisation for M. ornithogaster is considered the proper method for diagnosis due to difficulty in its culturing. So, PCR application on tissue lesions of proventriculus and its contents from dead birds lead to a final diagnosis of M. ornithogaster.

PCR by pan-fungal DNA primer sets was applied for amplification of extracted rDNA from purified cells. Specific primer sets amplified only rDNA extracted from isthmus scrapings of an infected bird, but not rDNA extracted from a non-infected
bird or other control DNA. The sequence was assured to be obtained from the purified organism by *in situ* rRNA hybridization using a specific probe. Phylogenetic analysis of sequences of the 18S rDNA and domain D1/D2 of 26S rDNA demonstrated the organism to be previously undescribed anamorphic ascomycetous yeast appointing a new genus (Tomaszewski *et al.*., 2003).

There are two sets of the PCR primers targeting 18S rDNA of *M. ornithogaster*. The first PCR primer set (AGY1/SM2) is selected as a specific PCR test for *M. ornithogaster*; the forward primer being AGY1 5’-GGACTTATATTACTAGTCAGATGG-3’ (positions 620–643). AGY1 does not match with any other reported fungi sequences and the reverse primer: Sm2 5’-CAATACGCCTGCTTTGAACACTC-3’ (positions 761–783) (Razmyar *et al.*, 2016).

The second PCR set (SM1/SM2) used for sequencing analysis of *M. ornithogaster* comprised the forward primer, Sm1 5’-ATCTGGTT¬GATCCTGCCAGTAGTC-3’ (positions 2–25) and the reverse primer, Sm2 (5’-CAATACGCCTGCTTTGAACACTC-3’ (positions 761–783) (Tomaszewski *et al.*, 2003).

**DIFFERENTIAL DIAGNOSIS**

The signs associated with *M. ornithogaster* infection are not specific and can occur with many other diseases, including trichomoniasis and giardiasis, bacterial and other fungal infections of the crop and stomach, helminth infections of the digestive tract, *Bornavirus* infection, crop and gastric foreign bodies and heavy metal poisoning.

**DISEASE CONTROL AND TREATMENT**

*M. ornithogaster* control and treatment are very difficult for avian clinicians, as subclinical infections without clear clinical signs can happen in many birds. It is practically difficult to keep extensive aviaries free from the disease (Filippich *et al.*, 2004). Numerous antimicrobial and antifungal medications are not efficient in treating the disease as iodine arrangements, luphenuron, nystatin, fluconazole, ketoconazole, itraconazole, and terbinafine (Filippich *et al.*, 1993; Phalen *et al.*, 2002; Phalen, 2005). Amphotericin B is considered the best drug for treatment. It is given orally two times every day for a month at a dose of 100 mg/kg (Phalen *et al.*, 2002; Phalen, 2005). As megabacteria develop in an alkaline environment, another treatment technique based on increasing the proventricular fluid acidity by oral administration of apple vinegar or grapefruit juice can be used (Gerlach, 2001). On the other hand, Püstow & Krautwald-Junghanns (2017) applied a medication with amphotericin B (100 mg/kg PO q12 h) for 4 weeks in positive infected budgerigars. The reported results were unacceptable and stressful for the birds because of the handling and the long treatment duration.

In a pilot study, amphotericin B was administered at 100 mg/kg twice daily for 30 days in two naturally infected bird species (*Melopsittacus undulatus* and *Agapornis roseicollis*) and at 25 mg/kg and 100 mg/kg twice daily for 10 days in experimentally infected chickens. The retrospective analysis indicated treatment failure in 80.4% of 36 cases, but significantly decreased *M. ornithogaster* burden, followed by profound rebound effect of the number of organisms shed in the faeces. The findings proved treatment failure in 3
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scenarios and indicated that treatment efficacy of amphotericin B against *M. ornithogaster* was poor (Baron et al., 2019).

In a colony of zebra finches, correction of the light cycle treatment with amphotericin B improved dramatically the health of the birds and baseline mortality returned to normal (Snyder et al., 2013).

Strains of *M. ornithogaster* were reported to be amphotericin B resistant in Australia (Filippich & Perry, 1993).

For moderate improvement of the flock mortality, oral application of nystatin medication in the feed in addition to vinegar administration in the drinking water for 3 weeks was recommended, followed by 500 mg/L sodium benzoate in the drinking water is administrated orally for 4 weeks (Madani et al., 2014).

Successful treatment of megabacteriosis in a canary was obtained with nystatin (Scullion & Scullion, 2004). The rate of mortality decreased and reached zero in budgerigars treated with nystatin (Kheirandish & Salehi, 2011). Probiotics are also thought to be of use through promoting health generally and by helping lower digestive tract pH through the production of lactic acid.

The mini-Flotac system enables the counting of yeast cells under microscope which is more obvious than the wet mount or Gram stain. So, evaluation of the effectiveness of the applied treatment can be based on yeast counting by this quantitative examination (Borrelli et al., 2015).

PREVENTION

It is believed that as *M. ornithogaster* is spread faecally, it would be prudent to maintain good hygiene in bird’s cages or house by thorough cleaning and disinfection with daily droppings elimination.

Common utilisation of waterers may be another source of disease transmission that can be controlled by good thorough house cleaning. It is also thought that disease transmission can occur through natural behaviour of the birds feeding each other. The *M. ornithogaster* infection was reported in both captive and wild psittacine and passerine species. Therefore, it is important to prevent cross infection between species. Strict hygiene measures should be undertaken when dealing with any sick bird.

Many AGY-infected birds are asymptomatic but still are shedding the organism (Phalen et al., 2002), therefore it is important to screen all new birds especially parrots admitted into aviaries from wildlife so that infected birds can be isolated from AGY negative birds.

Proper and hygienic hand raising chicks from incubator-hatched chicks will also break the infection cycle (Moore et al., 2001). Experimentally, it appeared that the disease did not happen if budgerigar eggs are collected from their parents, cleaned and the produced chicks are kept away from contact with the egg or infected birds (Moore et al., 2001).

CONCLUSION

*M. ornithogaster* infection is prevalent worldwide in aviculture. In Egypt, there are no available data about the impact of this yeast on aviculture. Studies are required to illustrate the role of this yeast in avian disease and, eventually, the factors which render avian hosts susceptible to megabacteriosis.

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