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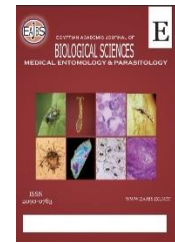
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Virulence of *Entamoeba histolytica*

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ABSTRACT

Entamoeba histolytica is an invasive protozoan, *E. histolytica* can adapt to the host's gut environment due to the virulence of the parasite. Ca²⁺, Acetylcholine (ACh), Fe, and EhRho1 are believed to be important for parasite motility and adhesion. Cytolysis, phagocytosis, and microcytosis cause parasitic invasion. Amebiasis is treated with metronidazole (MTZ), although strains resistant to metronidazole have emerged. With the help of studying the virulence of the parasite, it was possible to search for other drugs.

This review aims to present the key enzymes and the latest studies on promising drugs.

INTRODUCTION

Amoebiasis is an infectious parasitic disease in humans and one of the deadliest parasites in the world with a strong pathogenic possibility (Betanzos *et al.*, 2019). The life cycle of *E. histolytica* begins when infection cysts are ingested by swallowing polluted water or food and travel through the digestive system until they reach the large intestine, Infection occurs indirectly there are multiple stages and mechanisms for the onset of the disease (Kantor *et al.*, 2018). *E. histolytica* can invade and destroy human tissues, using many molecules and biological properties related to virulence. The amoeba utilizes three main virulence factors: Gal/GalNAC lectin, amebapore, and proteases. scientists work hard to understand the pathogenesis of the disease, however, the details about the sub- pathways involved in tissue invasion and damage are unclear (Babuta *et al.*, 2020). An understanding of the virulence of the parasites and the signals that lead to invasion is needed to design a better treatment. several virulence factors have been identified, and an unclear association between parasite genotype and invasive disease has been observed (Debnath *et al.*, 2019a). It is believed that the gut environment and the parasite's genotype, along with the host's genotype, all interact to create an environment conducive to invasion by *E. histolytica* ("Hunter's Tropical Medicine and Emerging Infectious Diseases," 2020).

1. Morphology

E. histolytica occurs in four forms, trophozoite, precyclic, cystic, and metacyst stages. (Chakraborty, 2004; Paniker's, 2018).

1.1. The Trophozoite (Fig 1):

It's the vegetative feeding phase of the parasite (Chakraborty, 2004). It is a facultative amoebozoan that metabolizes glucose as the main energy source. The trophozoite is measured in 20-30 μm . In the cytoplasm, there's a central nucleus called the karyosome. Some soluble lysosomes and

organelles perform the same function as mitochondria called mitosomes. Researchers thought that mitosomes are debris resulting from the breakdown of the mitochondria organelles (Despommier *et al.*, 2017). Endoplasm contains food vacuoles containing erythrocytes sometimes leukocytes and tissue residues that the parasite swallowed from the host's body (Paniker's, 2018).

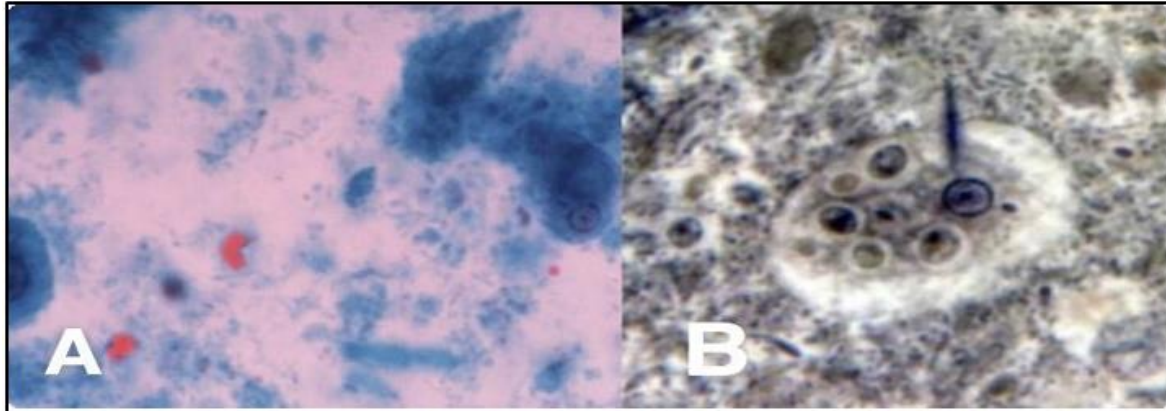


Fig. 1: (A) Trophozoite contains one nucleus and diameter is 10-60 μm

(B) Trophozoite of *E. histolytica* in stool of patient suffering from amoebic (Guerrant *et al.*, 2011)

1.2. Cystic Stage (Fig 2):

The diameter of the cyst is 10-15 μm . *Entamoeba histolytica* infection occurs when ingesting cysts (Guerrant *et al.*, 2011). It has a cyst wall to protect it from the harsh environment, it's resistant to dehydration, even some chemicals such as (chlorinated compounds, fluorides) and can live in water for a month, while those in feces can live in drylands for more than 12 days, and can

withstand temperatures of up to 50 C (Bogitsh *et al.*, 2012). It contains four or fewer nuclei. Initially, it has one nucleus in the precyst stage, then it multiplies by binary fission and becomes two nuclei, then four nuclei, and the adult cyst contains four nuclei. Immature cysts may supplement their growth outside when they're excreted through the feces (Chakraborty, 2004; Guerrant *et al.*, 2011; Bogitsh *et al.*, 2012).



Fig2: Cysts (the infective stage) contain four nuclei, the shape of the cyst 10-15 μm (Guerrant *et al.*, 2011)

2. Life Cycle (Fig 3):

Entamoeba histolytica passes its life cycle in one host, humans. The infection is transmitted by swallowing cysts in contaminated food or water, which pass through the stomach without damage due to the cyst wall, is resistant to gastric juice (Paniker's 2018), it reaches the ileum where its excystation occurs, the cyst wall shattered due of the alkaline medium of the caecum or lower part of the ileum (Guerrant *et al.*, 2011). This results in the metacystic where

the mitotic divided produces it causes the nucleus division first, followed by the cytoplasm then there will be eight small amoebulae. Each amoebulae can evolve into trophozoites. There are small size trophozoites that feed on bacteria. The large ones, feed on erythrocytes and attack the epithelial cells of the mucosal epithelium of the large intestine, causing amoebiasis. Some develop into cysts in the bowel lumen that passed into the feces to repeat the cycle (Baker 2017; Chakraborty 2004; Bogitsh *et al.*, 2012).

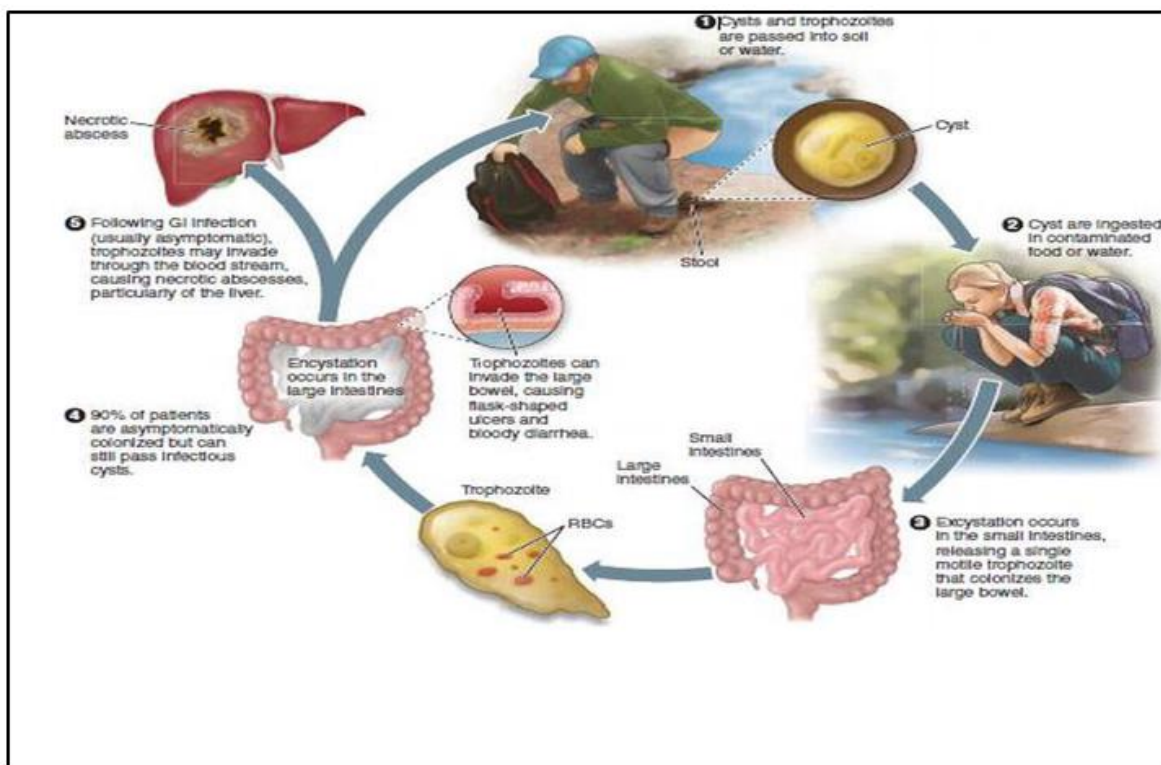


Fig3: The life cycle of *Entamoeba histolytica*

3. Amebic Dysentery

In most injuries, the inflammation resides on the mucous membrane, where the process of repairing and repairing the damaged tissue equals. It affects the gut and also the liver, lungs, and brain (Chakraborty, 2004; Pritt & Graham Clark, 2008). the *E. histolytica* is nourished by cellular debris and bacteria. The injury occurs when cysteine Proteases from *E. histolytica* digest the Muc2 mucin that surrounds the inner and outer layers and eliminates collagen and

fibronectin, causing the trophozoite to invade the deep layers of the gut. normally this process is following the regeneration of damaged tissue of the intestinal wall for a period, leading to the deposition of fibrous connective tissue (Nozaki & Bhattacharya, 2015; Rogier *et al.*, 2014; Song *et al.*, 2020). Severe ulceration follows a bacterial infection, which increases tissue breakdown, this infection is called amoebic colitis (Fig 4) (A. Singh *et al.*, 2009). Due to the high activity and high mobility of trophozoites,

when they traverse the epithelium and invade the tissues, they can reach the liver through the lymphatic and mesenteric veins to the hepatic portal vein, resulting in amoebic hepatitis and amoebic liver abscess (Fig 5) (Abubakar *et al.*, 2020). Usually, this leads to a single abscess in the right upper lobe. The parasite multiplies rapidly in the liver, causing blockage of the blood circulation leading to ischemic necrosis of the surrounding liver cells, containing the formation of small abscesses, it becomes one large abscess when the small abscesses merge (Bogitsh *et al.*, 2012; Samie *et al.*, 2012). The abscess contains a pus-like (not pus) mixture of liver

tissue and blood and appears brown and is called anchovy-sauce pus. Liver abscesses are grave given the fact that they release toxic substances due to ulceration (Chakraborty, 2004). There is a rare infection of the lungs and brain, and pulmonary amebiasis usually occurs as a complication of amoebic abscess and adheres to the diaphragm when trophozoite invades the portal circulation to the pulmonary capillaries and the appearance of abscesses like those in the liver. Cerebral amebiasis is an infection caused by a liver or lung abscess through the spread of trophozoites in the blood (Chakraborty, 2004; Samie *et al.*, 2012).

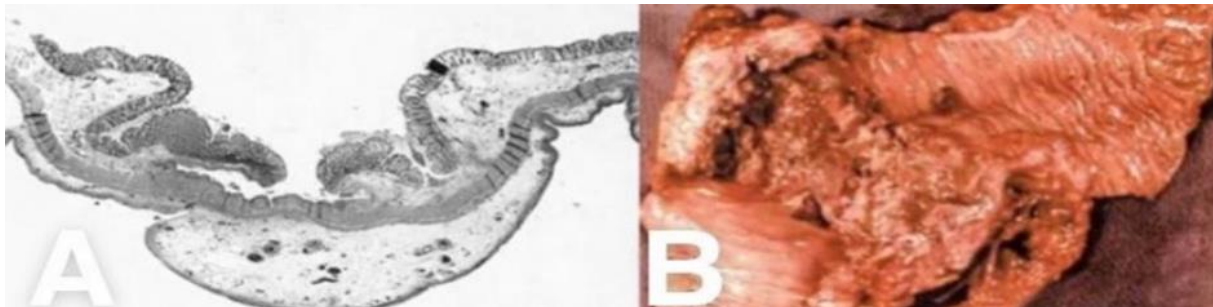


Fig 4: (A) Section of the human colon showing chronic amoebic ulcer (Bogitsh *et al.*, 2012). (B) a portion of the transverse colon showing extensive ulceration due to intestinal infection with *E. histolytica* (Baker 2017).

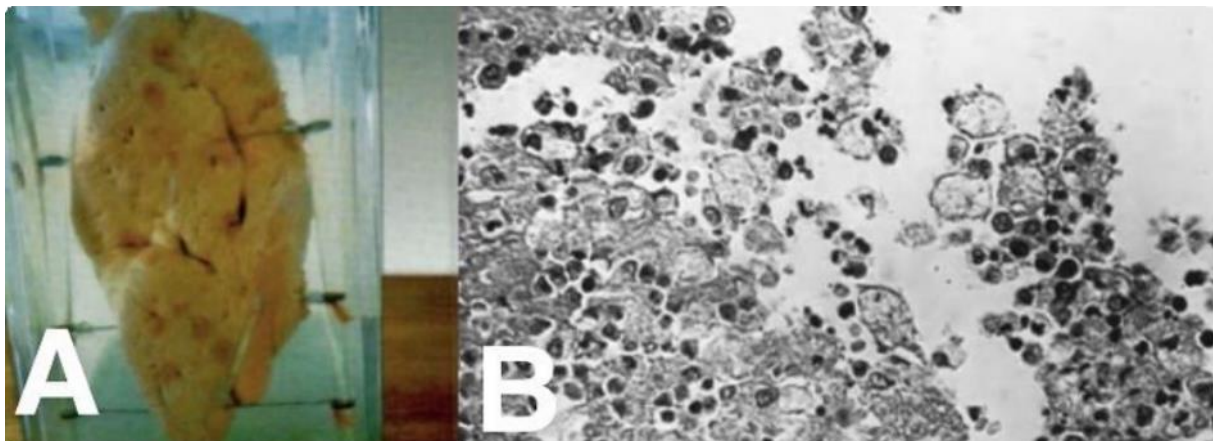


Fig 5: (A) Abscess in the human liver due to *Entamoeba histolytica*. (B) Specimen showing amoebic liver abscess (Paniker's 2018).

4. Virulence Factors of *E. histolytica*

4.1. Iron (Fe) Virulence Factors of *E. histolytica*

Iron is dependent on many organisms, including parasites and their hosts. *E. histolytica* depends on iron for metabolism, reproduction, and survival in the

host body. Iron is toxic and insoluble in the host cells mammalian host cells avoid iron toxicity by isolating certain proteins such as transferrin serum and mucosal lactoferrin as part of the prosthetic head group in hemoglobin (Hb) and cytochromes in iron storage protein ferritin (ft) (Hernandez-Flores

et al., 2016). These strategies prevent the parasite from growing in the host's body, since the parasite needs iron to survive and complete its life cycle, and to cause disease the host must have access to a cation at the site of infection, the parasite evolves special mechanisms to gain iron, namely reduction of the Fe + 3 to Fe + 2 complex by ferric reductase, which releases iron, and also generates specific receptors for iron-containing host protein and the secretion of a protease which separates iron from the protein cleavage (Gastelum-Martínez et al., 2018). *E. histolytica* can harness hemoglobin as a source of iron and externalize this protein across the cell membrane of erythrocytes through hemolysins and phospholipases (Cruz-Castañeda et al., 2009). Also, trophozoites have evolved two blood-binding proteins, Ehhmbp45 “EHI_096540” and Ehhmbp26 “EHI_022250.” These proteins are expressed by iron deficiency and can bind to the protoporphyrin ring in heme, suggesting that they have the same function as homophones. The mechanisms of hemoglobin endocytosis to obtain iron in *E. histolytica* are not yet clear, however, *E. histolytica* can phagocytose red blood cells and obtain hemoglobin through hemoglobinase, which has optimal protein vitality at acidic pH (Cruz-Castañeda et al., 2009, 2011). *E. histolytica* has evolved many strategies to exploit host resources and evade its defenses (Anaya-Velázquez & Padilla-Vaca, 2011). Parasite virulence depends on gene expression and variables in the host environment, including iron. Iron regulation was studied in *E. histolytica* by assessing different gene expressions in the medium, either deficient or excessive in iron (J. Lee et al., 2008). The mRNA differential display technique has been used in the study of cultures grown under iron-deficiency conditions. The researchers identified several genes involved in translation including those that encode the EF-1 α surveillance factor and five ribosomal proteins. These data indicate that low levels of aberration increase the need for translational mechanisms. what's more, a gene was found that codes for CP

(Hernández-Cuevas et al., 2014). Similarly, under iron-restricted conditions gene expression was significantly increased in 6 of the CPs (EhCP1 to EhCP6) (Gastelum-Martínez et al., 2018). CPs were implicated in *E. histolytica* adherence to enterocytes, which are important for host defense suppression. Cruz-Castañeda identified two types of proteins associated with *E. histolytica*, Hb-binding proteins of 45 and 26kDa (Ehhmbp 45 and Ehhmbp26 respectively). Both are involved in extracting iron from the blood. It is possible that they are hemophores that scavenge the heme from Hb or other heme proteins and can therefore be considered virulence factors of the parasite (Cruz-Castañeda et al., 2009; Gastelum-Martínez et al., 2018).

4.2. EhGEF virulence factors of *E. histolytica*:

GEF proteins help and facilitate the associated protein activation of Rho (small GTPase) by exchanging GDP with GTP. Direct activation of small GTPases with their master regulators (GEF) mediates many essential cellular activities in high eukaryotic cells, including cell morphology, actin dynamics, gene transcription, cell cycle progression, apoptosis, and malignancy progression in human cells (Toma-Fukai & Shimizu, 2019). Trophozoites interact with various host cells and tissues during invasive processes and metastasis (Cui et al., 2019), Generate signals for phagocytosis and trogocytosis to begin along with motility. Small GTPases are crucial regulators of cytoskeletal dynamics, hence they play a significant function. The Rho GTPases family plays a key role in actin dynamics by modulating the activity of actin-regulatory proteins (Mao & Finnemann, 2015; Xu et al., 2018), since GEFs act as the main regulator of the activity of Rho routines act. A study through structural bioinformatics analysis of EhGEF revealed that it is a PT-barrel structure capable of binding GTP and is not like any Dbl-family GEF structure the peptide sequence also revealed that EhGEF differs from the GEF family in length and sequence. Secondary structure predictions from

different software revealed that EhGEF is an essential part of α helices and β -sheets, whereas the conventional Dbl-family GEF structures are formed exclusively of α helices. The main function of GEF is to bind exchanges with GDP in the Rho protein pocket with GTP (Bharadwaj et al., 2017). Neither the T34N nor the EhRho1 mutant Q63L support this exchange due to a mutation in their regulatory domain (Bharadwaj et al., 2018). MANT-GTP binding was tested in a study and the results showed that EhGEF showed almost no activity towards EhRab1a, while EhRho1 was well activated. However, EhGEF showed a slow association with GTP, which was confirmed by the tryptophan quenching assay. However, MANT-GTP in a reaction containing EhGEF and EhRho1 showed a significant increase in rate and density, suggesting that EhGEF may swap nucleotides on EhRho1 by yet unknown mechanisms (Bharadwaj et al., 2021; Bosch et al., 2012). A study showed that EhGEF is one of the most important activators and specific regulators of EhRho1 in vivo. Sequence analysis showed that EhRho1 is identical to human HsRhoA except that the amoeba molecule lacks a distinct and conserved feature of the Rho GTPases "Rho insert domain", which is required for activation of the downstream Rho-associated kinase. However, both human and amoebic molecules appear to be similar in that they function in which overexpression of EhRho1 can be generating stress fibers in mammals (Bharadwaj et al., 2021; Hall et al., 2006). The GDP-bound form of EhRho1 lies in the cytosol, while the active form (GTP-bound) transports toward the membrane and attaches to it through the prenylation domain. Like EhRh1, EhGEF localization has also been observed in the cytosol and membrane of trophozoites. EhGEF can bind to the P3 "PtfIns (3,4,5)" in the membrane through the PH domain and activate EhRho1 in its way. participation in Rho-GEF in endocytosis has been well studied. The Rho-GEF has been found with Fc receptors during phagocytosis (Buchsbaum, 2007; García-Mata & Burridge, 2007). Therefore, a study examined the

participation of EhRho1 in phagocytosis, they found EhGEF in phagocytic cups next to EhRho1 and a reduced formation of phagocytic cups on their down-regulation of expression. The threshold concentration of active EhRho1 (GTP bound) is Required at the RBC attachment site, for the initiation and progression of the phagocytic cup. It has been observed that phagocytosis proceeds slowly in those trophozoites down-regulated for EhGEF expression where there is sufficient EhGEF down-regulation by antisense RNA, thereby prolonging the time it takes for EhRho1 activity to reach the critical level reached on spot. Therefore, the rate of formation and development of the phagocytic cup may be directly proportional to the concentration of some key molecules, such as EhRho1 and its regulators on the phagocytic cups (Dong et al., 2010; Hall et al., 2006; W. L. Lee et al., 2007)

4.3. Ca²⁺ virulence factors of *E.histolytica*:

E.histolytica contains ironically released Ca²⁺ totaling 70% Ca²⁺ which can be divided into two parts:

The first part is impelled by the second messenger inositol 1,4,5-triphosphate (Ins (1,4,5) P3), which releases endogenous Ca²⁺ from the endoplasmic reticulum-like structures. The second part is sensible to ins (1,3,4, 5) P4 (Babuta et al., 2020). Although these two messengers' affairs in two disparate Ca²⁺ stores, it is unclear whether there is a connection between them in this organism *E.histolytica* cipher calpain-like protein, as well as many nuclear components that require Ca²⁺, including Ca²⁺-dependent ATPase/ADPase, Ca²⁺-dependent thiamine pyrophosphatase, and acid phosphatase. Calpain-like protein is expected to be interrelated with apoptosis in parasites, as its level increases during programmed cell death. It is also found in the cytoplasm and near the cell nucleus (Domínguez-Fernández et al., 2018; Monroy et al., 2015), The inner layer of cytoplasmic vacuoles, which may or may not be phagocytic, contains a small number of nucleotide enzymes. unclear whether these enzymes are into in calcium homeostasis in this parasite, genomic analysis identified 27

multi-EF-hand-containing CaBPs in *E. histolytica* (Bhattacharya et al., 2006). Various of these proteins are thought to be Ca²⁺ buffers and are accordingly involved in governing Ca²⁺ concentration in different cell compartments. The first step: The target cells' adhesion is significant during the invasion process; after contact is made, some molecules involved in this process have been identified, including glucose and N-acetyl-d-galactosamine (Gal/GalNAc) lectin, a 260-kDa heterodimeric cell-surface protein consisting of a 170-kDa heavy chain (hgl) linked to a 35/31-kDa light chain (lgl) (Babuta et al., 2020). The light subunit is believed to be attached to the membranes by the glycosphosphoinositol stabilizers and the lectin-260 kDa is complexed with an intermediate subunit 150 kDa. The heavy chain consists of the carbohydrate identification domain (CRD), which is introduced on the cell surface. Overexpression of the mutant heavy chain subunit is permanently absent of the N-terminal i.e., In animal models, CRDs frequently confer a dominant-negative phenotype with reduced adhesion and pathogenicity. Furthermore, light subunits play a role in pathogenicity. The expression of the mutant from lgl (Part of the C-terminal) appears a negative phenotype (Babuta et al., 2020; Katz et al., 2002). The heterogeneous junction of mutant lgl is formed with hgl, but this junction is functionally inactive. Added to the Gal/GalNAc lectin, some other cell surface molecules involved in the adhesion action have been discovered (Daniela et al., 2014). In one study, the Ca²⁺ binding site for interaction with the ligand was identified (Chadee et al., 1988). Another study was alike determined the Ca²⁺ binding location, and a mutant was created that lost the ability to bind Ca²⁺. Although the mutant protein retained its carbohydrate-binding function, it lost its ability to agglutinate erythrocytes, suggesting that some properties of the Gal/GalNAc lectin are modulated by Ca²⁺ ions, similar to the Ca²⁺-binding protein calreticulin (CRT) on the surface of a cell protein (Yadav et al., 2016). *E. histolytica* attaches to complement

component 1q (C1q), which is involved in phagocytosis of apoptotic immune cells but does not attack or kill normal cells such as Chinese Hamster Ovary (CHO) cells (Vaithilingam et al., 2012).

4.3.1. Relationship of Cytolysis Of Target Cells and Ca²⁺:

Amoebic cells express many different genes that exhibit proteolytic activity (Siqueira-Neto et al., 2018). The cysteine protein 5 (Ehcp5) is gaining attention because it is localized on the cell surface and because there is no active homolog in the non-pathogen *E. dispar* (D. Singh et al., 2004). Porin-like proteins from *E. histolytica* are also involved in cytolysis performed by amoeba cells. Interaction of the interaction of *E. histolytica* with target cells is the dramatic increase in Ca²⁺ levels at the end after contact. Preventing Ca²⁺ Chan-Nelson target cell death induced by Gal/GalNAc lectin due to purified protein itself is believed to increase Ca²⁺ levels in target cells (Babuta et al., 2020; Ravdin et al., 1985a), but the pathway to release Ca²⁺ from the target cell upon contact with *E. histolytica* is unclear. Multifold studies have displayed that Ca²⁺ signaling shares the ability of *E. histolytica* to kill the target cell. Avert the increase of Ca²⁺ between the parasitic cells avert the start of the cytolytic process (Makioka et al., 2001; Ravdin et al., 1982, 1985b). The direct involvement of Ca²⁺ in amoebae wildness was observed when the upstream factor of regulatory element 3 was found to regulate gene expression of the virulence-associated genes. UREBP binds the promoter element of Gal/GalNAc lectin hgl5. It contains 2 Ca²⁺-binding EF-hand booster and regulates negative transcription in the presence of Ca²⁺, meaning it only binds the DNA promoter in the deficiency of Ca²⁺. (Gilchrlst et al., 2003). The Ca²⁺ binding flaw dialogue showed a dominant phenotype and the cells expressing the mutated protein were further virulent. There is a possibility that URE3BP plays a major role in virulence since a large proportion of amoeba genes contain the URE3 form which is recognized by URE3BP and thus controls the expression of a variety

of genes (Gilchrist et al., 2010). An unusual endemic number of URE3BP appears in the plasma membrane of trophozoites regardless of which one is in the nucleus (Moreno et al., 2010)

4.4. Acetylcholine (ACh) Virulence Factors of *E.histolytica*:

ACh is released from the gut nervous system in response to enteritis and regulates digestive functions, including the transfer of epithelial ions on its membrane (Wood, 2008). Trophozoites *E.histolytica* can bind ACh and thus modify parasite virulence by remodeling cell structure, parasite movement, phagocytosis, and increased expression and secretion of virulence factors, thus promoting amoebic cytotoxic (Medina-Rosales et al., 2021). The trophozoites' invasion of the gut is promoted by causing damage to the gut system and inflammation of the gut during infection, thence ACh may conceivably be involved in triggering amoebiasis. One study showed that ACh binds to trophozoites membranes, thereby increasing parasite proliferation and chemical transformation resulting in cell structure remodeling, regulation of the expression, and virulence factors such as Gal/GalNA lectin, L220, Amebapore C, and CP, which improved the Amoeba's ability to destroy and invade host tissues (Diamond et al., 1978). There is a study that showed that the activities increase the expression of actin and increase the polymerization of actin (F-actin) in response to the physiological levels of ACh. The dynamics of the cellular structure of the *E. histolytica* cytoskeleton are indispensable for cellular processes like adhesion, migration, phagocytosis, and killing of host cells. The host cells of *E.histolytica* are killed in a contact-dependent and contact-independent manner (Manich et al., 2018; Tavares et al., 2005). ACh upregulation of virulence factors, particularly secreted soluble components, resulted in a great increase in cellular toxicity of ACh-treated trophozoites against HepG2 cell monolayers, as evidenced by increased expression of amebapore and CP, as well as their will activities, through contact-independent processes. ACh stimuli also

segmented cytopathic damage to the HpG2 monolayer (contact-dependent), a process closely related to frequent trogocytosis and phagocytosis (Talamás-Lara et al., 2014). Erythrocytosis is certainly the characteristic toxicity of pathogenic amoebas, which involves adhesion, cytoskeletal reorganization, and secretion of soluble agents that mediate host cell killing and ultimately ingest cellular debris. Ach supports erythrocytic activity in treated activities by increasing Gal/GalNAc and protein expression L220 (Manich et al., 2018; Medina-Rosales et al., 2021)

5. The Latest Drugs for Amebiasis:

E. histolytica may withstand current metronidazole (MTZ) treatment, and enhanced production of iron-containing superoxide dismutase and peroxiredoxin has been linked to drug resistance. MTZ resistance has been discovered in some clinical strains of *E.histolytica*, showing that the generation of drug-resistant strains is linked to enhanced production of iron-containing superoxide dismutase and peroxiredoxin. Molecular resistance to MTZ has been described in some clinical strains (Debnath et al., 2019). These observations led to the search for new drugs, and studies based on drug testing were published. The first study identified 1H-1,2,3-triazole-linked isatin-metronidazole conjugates that are most potent against *E.histolytica* and *Giardia lamblia* of MTZ (Kumar et al., 2018). The second study, E, targeting the primary biosynthetic pathway of the parasite cysteine, described the fungal metabolite as a compound that inhibits cysteine and amoeba growth in a cysteine-dependent manner with low mammalian cytotoxicity (Mori et al., 2018). A third study identified drugs capable of killing Riptaceae and MNZ resistant *E.histolytica* (Ehrenkauffer et al., 2018). In another study, farnesyltransferase (FT), the final co-enzyme of the mevalonate pathway-derived product function, was targeted. This enzyme is critical for a variety of functions including cell differentiation and growth (Probst et al., 2019). A promising therapy option for amebiasis is a synergistic

combination of metronidazole and the FT inhibitor lonafarnib. In many regions of the world, plants and their extracts are used to treat gastrointestinal problems. (Kelber et al., 2017).

Conclusions

E.histolytica virulence factors are essential for host-parasite interaction due to the severity of infection and tissue invasion. These factors can be modified by the parasite and the host to create a suitable environment for that parasite to live. Ca²⁺ is an important virulent factor. Calcium is known to be involved in many cellular processes in all eukaryotic systems. It regulates a large number of *E.histolytica* signaling pathways that affect the virulence of the parasite. In addition to iron, *E.histolytica* has an iron response. Two proteins (Ehhmb25 and Ehhmbp45) are involved in extracting iron from the blood, so it can be considered one of the virulences. The enteric nervous system releases ACh in response to intestinal inflammation, and ACh can modulate pathogenesis. EhGEF is involved in phagocytosis and plays an important role in

endocytosis dependent on these proteins. EhGEF is a protein that binds to EhRho1 and interacts with it. Most virulence factors are up or downregulated by these factors, and their functions in *E. histolytica* are not yet clear. We encourage researchers to study their functions to understand more about the virulence of the parasite to develop a treatment for this disease.

Declarations:

Ethics approval and consent to participate:

No application

Availability of data and materials: data available, Complete experimental details and data sets are available at the ‘‘ArrayExpress’’ MIAME-based database (www.ebi.ac.uk/arrayexpress/) with the accession number E-MTAB-1158.

Competing interests: The author declare that she has no competing interests

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List of abbreviations

Abbreviation	Definition	Aberration	Definition
Fe	Iron	Hb	Hemoglobin
Ft	Ferritin	CPs	Carbamoyl Phosphate synthase II
GEF	Guanine nucleotide exchange factors	Rho	Ras homologous
Ca ²⁺	Calcium	MTZ	Metronidazole
hgl	Heavy chain-linked	Lgl	Light subunit
CRD	Carbohydrate recognition	Ach	Acetylcholine

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