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RESEARCH ARTICLE

GC-MS ANALYSIS OF BIOACTIVE COMPOUNDS AND ANTIMICROBIAL ACTIVITY OF CRYPTOCOCCUS RAJASTHANENSIS KY627764 ISOLATED FROM BOMBYX MORI GUT MICROFLORA.

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Abstract

The insect gut microflora acts as a large reservoir of bioactive natural products as a diversity of resident microflora is found to be symbiotically associated with the insect. The present study reports the bioactive chemical compounds and the antimicrobial nature of the yeast isolated from the insect *Bombyx mori* gut. The yeast is identified as *Cryptococcus rajasthanensis* by Molecular characterization. The Fourier transform infrared spectroscopy (FT-IR) studies of the chloroform and ethyl acetate crude yeast extracts indicated a number of functional groups like alcoholic, phenolic, ester, aldehydic, etc. that accounts for the bioactive nature of the extracts. The Gas chromatography-Mass spectroscopy (GC-MS) analysis of the crude extracts revealed a large number of bioactive compounds of high and low molecular weight that are considered biologically active. Some of the biologically active molecules like phenol 2,4-bis(1,1-dimethylethyl), 1,2-benzenedicarboxylic acid dibutyl ester, celidoniol deoxy, nonadecane, tetratetracontane, 2-methyloctacosane and pentadecane bearing antimicrobial property were detected in both chloroform and ethyl acetate extracts. Furthermore, the crude extracts were subjected to antimicrobial assay by agar well diffusion method and are found to be antimicrobial against four pathogenic bacteria *Staphylococcus aureus*, *Bacillus subtilis*, *Escherichia coli* and *Pseudomonas aeruginosa* and two fungi *Candida albicans* and *Aspergillus flavus*. The chloroform extract showed maximum inhibition for *E. coli* and minimum for *A. flavus*.

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Introduction:-

Insects being one of the oldest group of animals on our planet that are ubiquitous in nature and have been adapted to several environmental conditions represent the most diverse group of animals on earth (Chapman, 2007). Thus, a wide range of microbes are found associated with these insects and are found to thrive on the insect's exoskeleton and in the hemocoel and gut of insects (Douglas et al., 2015). The prevalence of microbes in the insect gut is high as compared to the other parts of the insect anatomy as the gut tends to provide a hostile environment for the microorganisms. It is studied that the microbial load in the insect gut is 10 times the total cells of the insect (Rajagopal, 2009). The gut microbiota represent both positive as well as negative interaction

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with the host ranging from pathogenic to mutualistic relationships. Most of the microbes that live in association with the insect play beneficial roles for the host and hence play symbiotic or mutualistic role (Dillon and Dillon, 2004). The resident microbial flora in insects thus benefits the insect by producing essential compounds like vitamins, digesting and metabolizing food, nutrient absorption, etc. (Douglas et al., 2001; Nasir and Noda, 2003; Calderon-Cortes et al., 2012) and pheromones production and chemical communication (Ezenwa et al., 2012). The commensal flora are found to play crucial part in survival of the insects by protecting their insect hosts against the natural enemies and pathogens through varied mechanisms that include colonial resistance, production of toxins and insects immune system activation to fight the invader organism (Boulanger et al., 2001; Dillon and Charnley, 2002; 1995).

Although most studies in the past have focused on the bacterial endosymbionts, certain recent studies have now been conducted on the use of fungi, the yeasts being one of the predominant group as the beneficial microbes in the insect health (Gonzalez, 2014). Most of the yeasts insect endosymbionts are known to belong to the genus *Candida*, *Metschnikowia*, *Pichia*, *Saccharomyces*, *Cryptococcus* and *Pseudozyma* (Urubschurov and Janczyk, 2011). The yeast-insect symbiotic relationship benefits the yeasts as the insect provides hostile environment for the yeast to thrive in the gut, and facilitates dispersal of yeast spores and outbreeding (Vega and Dowd, 2005). On the contrary the insect is benefitted from the yeasts as the yeasts provide nutrients (vitamins like B3 and B5, trace metals, proteins, amino acids, etc.), detoxify harmful substances (usually by producing detoxifying enzymes like hydrolase, glucosidase, etc.), provides protection from biotic and abiotic stress (by providing protection from invader pathogens) and can also aid in chemical communication (Gibson and Hunter, 2010; Engel and Moran, 2013; Pozo et al., 2012; Davis et al., 2011; Christensen, 2010). This insect-microbe association directs the research onto a wide range of novel chemical moieties that can be obtained from the microorganism isolated from the insect gut that can be used in human therapy (Beemelmanns et al., 2016; Brachmann and Bode, 2013). The research on the secondary metabolites produced by the gut microflora can be justified in 2 ways: 1) to identify the natural products produced by the microflora that can be beneficial to humans and 2) to identify the known as well as novel microorganisms involved in production of these natural compounds so as to exploit these microbes as they may produce these secondary metabolites to combat pathogenic microbes (Soria-Mercado et al., 2012).

The insect *Bombyx mori* (*B. mori*) belongs to the Lepidopteran class of insects wherein the yeast association with the insect has been less well studied (Gonzalez, 2014). Although the yeasts are known to produce volatile compounds that can be used as biocontrol agents especially in post harvest crop management (Witzgall et al., 2012), very less research has been conducted for use of the bioactive compounds produced by the yeasts isolated from the insect gut that can be used in human therapy (Bode, 2011). Therefore, the present study was focused on the investigation of antimicrobial nature of the yeast isolated from the *B. mori* gut and evaluation of its bioactive chemical constituents by the GC-MS analysis. Our studies have successively proved that the yeast *Cryptococcus rajasthanensis* isolated from the *B. mori* gut bears antimicrobial potential and the GC-MS profiling has shown a wide range of chemical compounds bearing therapeutic properties including antimicrobial, antioxidant, anticancer, etc.

Materials and Methods:-

Isolation of yeast from *Bombyx mori* gut:-

Fifth instar healthy *B. mori* larva was dissected under sterile conditions, the gut was isolated and washed in sterile distilled water to remove solid contaminants. The fluid was collected after slight maceration of the gut in sterile bacteriological saline. The extracted fluid containing microorganisms was swab inoculated onto the sterile potato dextrose agar (PDA) plate. The inoculated plate was incubated for 48-72 hours at 27-30 °C.

Identification of the yeast strain:-

The isolated yeast Bm5F5 strain was initially checked for its morphological characteristics and gram stained. It was further identified by molecular characterization using 18S rRNA gene sequencing and phylogenetic analyses.

Preparation of yeast extracts:-

The yeast extracts were prepared according to Cita et al., 2017. Briefly, the yeast from the PDA plate was inoculated into an Erlenmeyer flask containing 500 ml sterile sabouraud's broth. The inoculated flask was incubated at 27 °C for 5-7 days under shaker conditions. After incubation, the culture broth was filtered to remove the cell mass. The filtrate was extracted with chloroform as well as ethyl acetate (1:1 v/v). The solvent extract was concentrated by rota-evaporation. The chloroform and ethyl acetate crude extracts of the yeast was used for further evaluation.

Fourier transform infrared radiation (FTIR) analysis of yeast extracts:-

The various functional groups responsible for the biological activities of the crude yeast extracts were recorded (Poojary et al., 2015). KBr pellets of the dried yeast extracts were prepared and the spectra were recorded ranging from 4000–400 cm^{-1} at a resolution of 2 cm^{-1} using a FTIR spectrophotometer (NICOLET 6700, USA).

Gas chromatography-Mass spectroscopy analysis of yeast extracts:-

The GC-MS analysis of the crude yeast extracts was carried out to evaluate the various bioactive compounds present in the yeast extracts (Sivaraman et al., 2017). This was carried out by GC-MS spectroscopy (Shimadzu GCMS-QP 2010S system) and the compounds were predicted using the National Institute Standard and Technology 11.0 (NIST) library database and Wiley 8.0.

Evaluation of Antimicrobial activity by agar well diffusion assay:-

The antimicrobial activities of the yeast extracts were performed by agar well diffusion method (Bonev et al., 2008). Lag phase cultures of gram positive bacteria *Staphylococcus aureus* and *Bacillus subtilis*, gram negative bacteria *Escherichia coli* and *Pseudomonas aeruginosa* and fungi *Candida albicans* and *Aspergillus flavus* were used as the test microorganisms. 100 μl of the test microorganisms were swab inoculated onto the sterile nutrient agar and sterile PDA petriplates respectively for which then wells were bored. 50 $\mu\text{g}/\text{mL}$ of the yeast extracts were added into the respectively labeled well. Tetracycline and Streptomycin were used as the reference drugs for the bacteria and Nystatin and fluconazole for fungi. These plates were incubated at 37°C for 24 hours for bacteria and 27–30°C for 48 hours for fungi followed by observing and measuring the antimicrobial activity of the yeast extracts by measuring the zones of inhibition in millimeters using the Hi media Antibiotic scale.

Results:-**Isolation of yeast from *Bombyx mori* gut:-**

A cream colored glistening yeast colony was isolated on potato dextrose agar as shown in figure 1 and labeled as yeast Bm5F5 strain for reference.

Identification of the yeast strain:-

The yeast Bm5F5 strain depicted encapsulated oval shaped cells on gram staining as in figure 2. The strain was identified as *Cryptococcus rajasthanensis* by 18S rRNA gene sequencing and phylogenetic analyses and the gene sequence was deposited in the NCBI GenBank under the accession number KY627764.

Fourier transform infrared radiation (FTIR) analysis of yeast extracts:-

FT-IR spectral data of the chloroform and ethyl acetate crude extracts revealed presence of various functional groups in the yeast as shown in figure 3 and 4 respectively. Spectral data in table 1 gives the interpretation of the frequencies, vibration bonds and the functional groups present in the extracts. Both the extracts show hydrogen bonded –OH stretching, C-H stretching, N-H bending, C-C stretching in ring, C-O stretching and C-N stretching in common.

Gas chromatography-Mass spectroscopy analysis of yeast extracts:-

The bioactive compounds present in chloroform and ethyl acetate extracts obtained from the yeast are shown in tables 2 and 3. The GC-MS studies of the yeast extracts showed 18 compounds in the chloroform extract and 15 compounds in the ethyl acetate extract. Based on the relative abundance, the top four major compounds present in the chloroform extract were Celidoniol deoxy (23.71%), eicosane (22.26%), nonadecane (18.68%) and tetratetracontane (15.93%). The ethyl acetate extract contained 2-methyloctacosane (16.37%), nonadecane (15.03%), Celidoniol, deoxy (13.64%) and tetratetracontane (10.43%) as the top four major compounds. The GC chromatogram of the chloroform and ethyl acetate extracts are depicted in figures 5 and 6 respectively.

Evaluation of Antimicrobial activity by agar well method:-

A concentration of 50 $\mu\text{g}/\text{mL}$ of the yeast extracts was evaluated for its antimicrobial activity against 4 bacterial and 2 fungal strains. The results shown in table 4 and figure 7 determine the chloroform and ethyl acetate extracts of the yeast as antimicrobial agents against human pathogens. It has shown good antibacterial activity against the gram negative bacterial strains used *Escherichia coli* and *Pseudomonas aeruginosa* as well as to the gram positive ones *Staphylococcus aureus* and *Bacillus subtilis*. The antifungal activity of the yeast extracts against the yeast *Candida albicans* and the fungal strain *Aspergillus flavus* predicts the antifungal nature of the extracts.

Fig. 1:- Cream colored glistening yeast colony on Potato Dextrose agar isolated from *Bombyx mori* gut



Fig. 2:- Gram staining showing gram positive encapsulated oval shaped cells

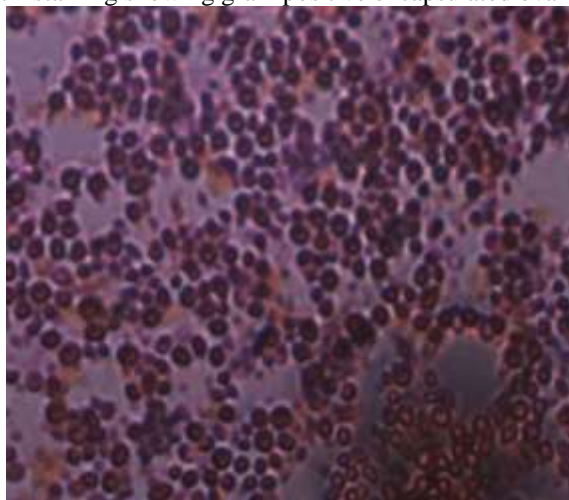


Fig. 3:- FT-IR spectrum of the yeast *Cryptococcus rajasthanensis* chloroform extract

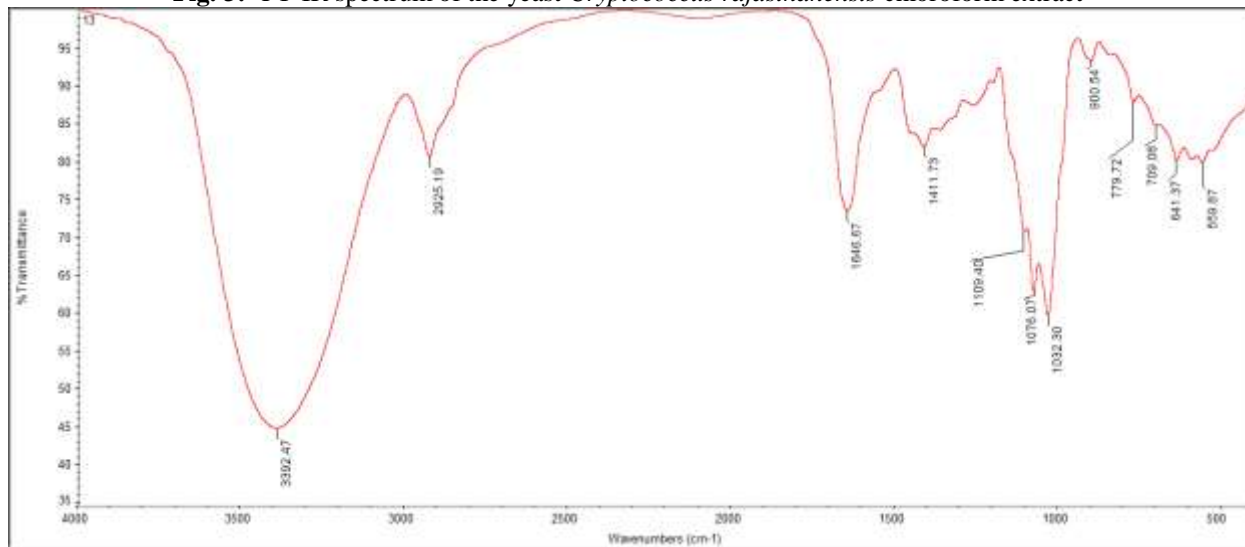


Fig. 4:- FT-IR spectrum of the yeast *Cryptococcus rajasthanensis* ethyl acetate extract

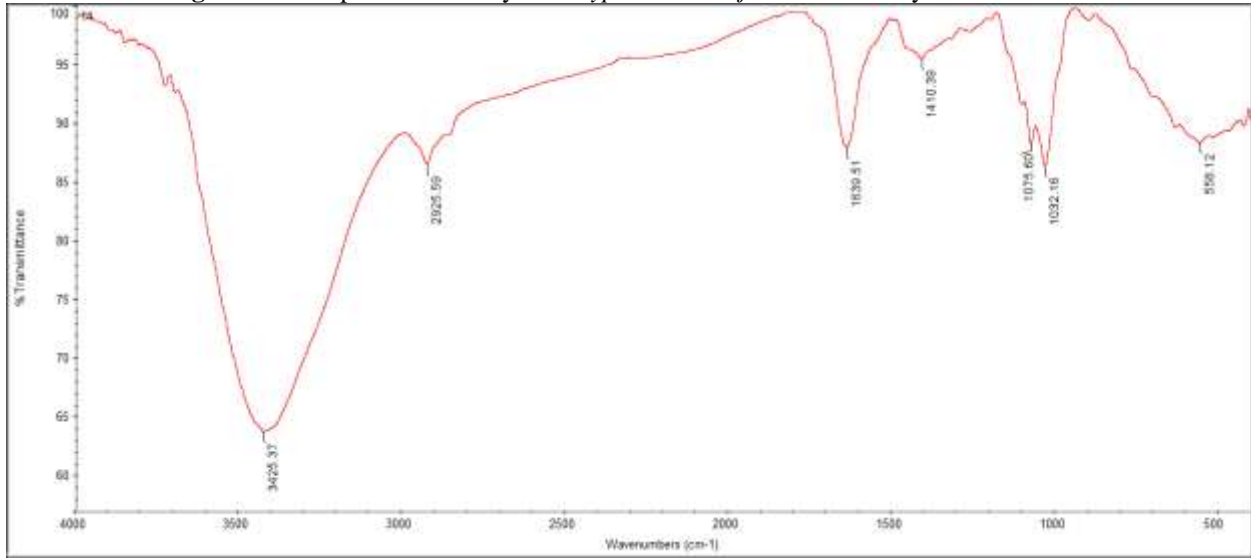
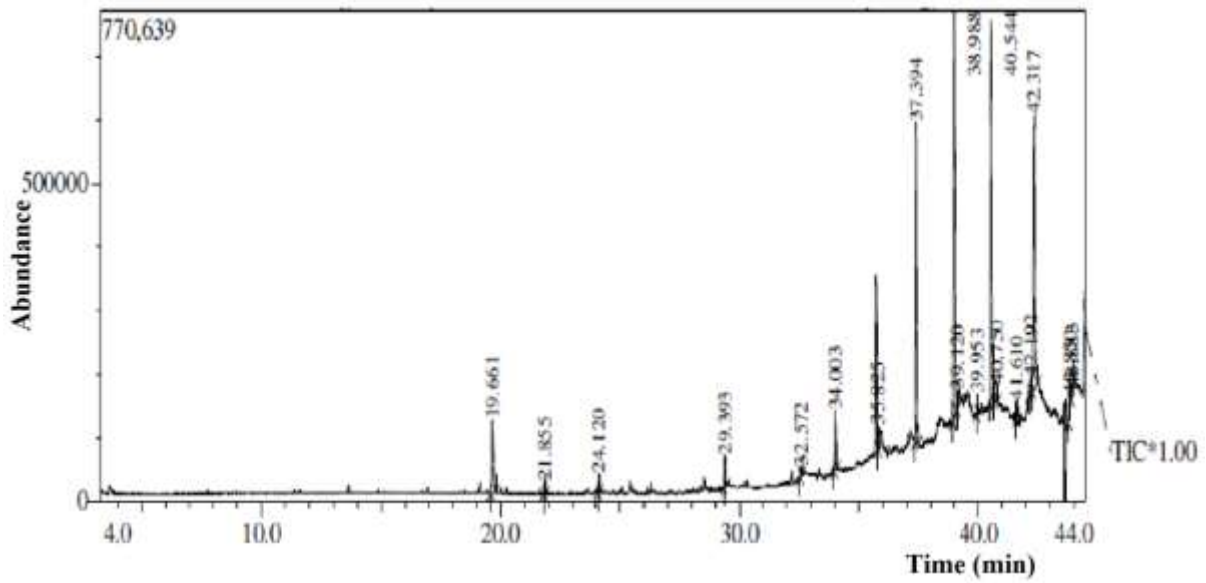


Fig.5:- GC-MS chromatogram of the yeast *Cryptococcus rajasthanensis* chloroform extract



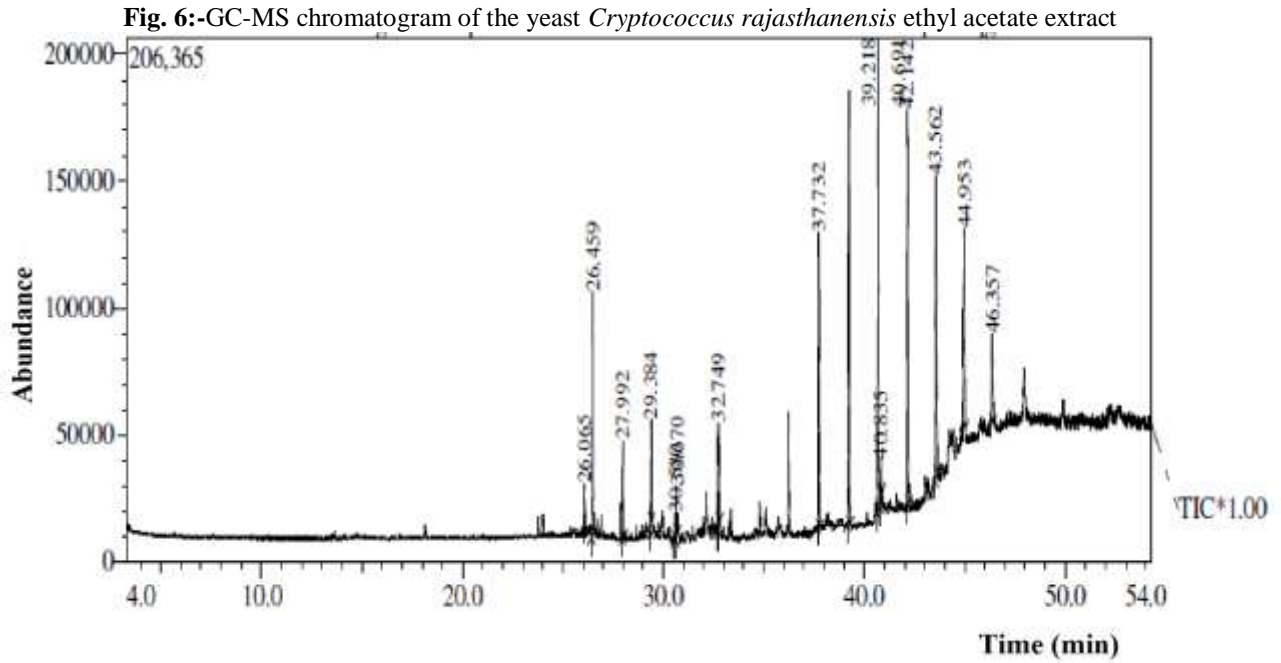
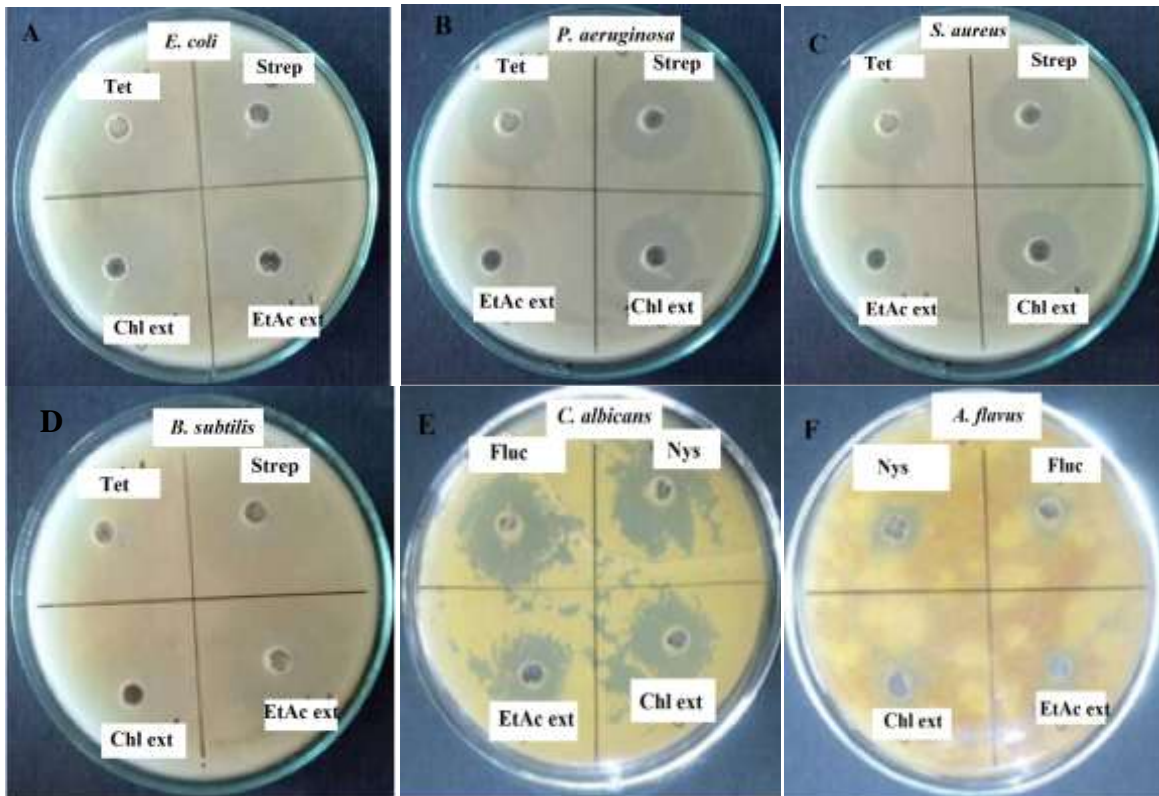


Fig. 7:- Antimicrobial activity of yeast *Cryptococcus rajasthanensis* extracts against human pathogens by agar well diffusion method



Note:- Tet-Tetracycline; Strep- Streptomycin; Fluc- Fluconazole; Nys- Nystatin; Chl ext- Chloroform extract; EtAc ext- Ethyl acetate extract

IR ν_{\max} (cm ⁻¹)	(Vibration mode)	Functional groups
Chloroform extract		
3392.47	H-bonded OH-stretching; -N-H stretching	Alcohols, phenols 1°, 2° amines, amides
2925.19	C-H stretching	Alkanes
1646.67	-C=C- stretching	alkenes
1411.73	C-C stretch (in-ring)	aromatics
1109.40, 1076.07, 1032.30	-C-O stretching; -C-N stretching	Alcohols, carboxylic acid, esters, ethers; Aliphatic amines
900.54	=C-H bending	alkenes
779.72; 709.08	N-H waging C-H "oop" C-Cl stretching	1°, 2° amines; Aromatics; Alkyl halides
641.37	-C≡C-H: C-H bending	alkynes
559.87	C-Br stretching	Alkyl halides
Ethyl acetate extract		
3425.37	H-bonded OH-stretching	Alcohols, phenols
2925.59	C-H stretching	Alkanes
1639.51	N-H bending	1°, 2° amines
1410.39	C-C stretch (in-ring)	aromatics
1075.60; 1032.16	C-O stretching; C-N stretching	Alcohols, carboxylic acids, esters, ethers; Aliphatic amines
558.12	C-Br stretching	Alkyl halides

Table 1:- FTIR data for the yeast *Cryptococcus rajasthanensis* extracts

Sl no	Compound name	Retention time	Peak Area (%)	Molecular formula and Molecular weight	Activity	Reference
1	Phenol-2,4-Bis(1,1-dimethylethyl)-	19.661	4.42	206 (C ₁₄ H ₂₂ O)	Antioxidant (Inhibits ROS) Antibacterial	Teresa et al., 2014 Manorenjitha et al., 2013
2	Pentadecane	21.855	0.49	212 (C ₁₅ H ₃₂)	Antibacterial	Yogeshwari et al., 2012
3	Heptane,3,3-dimethyl-	24.120	0.78	128.25(C ₉ H ₂₀)	Unknown	
4	1,2-Benzenedicarboxylic acid, dibutyl ester (Dibutyl phthalate)	29.393	1.51	278.344 (C ₁₅ H ₂₈ O)	Antibacterial Antifouling Pesticide	Khatiwora et al., 2012 Jenecius et al., 2012 Wanxi et al., 2014
5	3-Ethyl-3-methylheptane	32.572	0.40	142.28(C ₁₄ H ₂₂)	Unknown	
6	2-methyloctacosane	34.003	2.75	408.7867 (C ₂₉ H ₆₀)	Found in cuticles of various insects spp. (chemical communication)	Spikes et al., 2010
7	2-Cyclohexyl-3-isopropylpent-4-en-2-ol	35.825	0.94	210.356 (C ₁₄ H ₂₆ O)	unknown	
8	Nonadecane	37.394	16.37	268 (C ₁₉ H ₄₀)	Antimicrobial and cytotoxic A cuticular hydrocarbon of insects (chemical communication)	Hsouna et al., 2011 Colazza et al., 2007
9	Eicosane	40.544	22.11	282.547 (C ₂₀ H ₄₂)	Antitumor activity	Sivasubramanian and Brindha, 2013
10	2,5-Octadecanoic acid, methyl ester	39.120	1.35	408.7867 (C ₁₉ H ₃₀ O ₂)	Antiviral Antioxidant, antibacterial	Linton et al., 2013 Sudharshan et al., 2010
11	2-Methyltetracosane	39.953	0.87	352.680 (C ₂₅ H ₅₂)	Free radical scavenging activity	Ramya et al., 2015
12	Hexatriacontane	41.610	1.26	506.973 (C ₃₆ H ₇₄)	Unknown	
13	Celidoniol, deoxy (Nonacosane)	38.968	21.11	408 (C ₂₉ H ₆₀)	Antibacterial Antiinflammatory Chemical communication especially in Anopheles stephensi mosquito Pheromone of Orgyia leucostigma	Kose et al., 2016 Zakaria et al., 2014 Brei et al., 2004 Grant et al., 1987

14	2-Bromotetradecane	40.750	2.17	277.284 (C ₁₄ H ₂₉ Br)	Unknown	
15	Methenolone	42.192	1.25	302.451 (C ₂₀ H ₃₀ O ₂)	A naturally occurring drug for aplastic anemia treatment	Lockner, 1979
16	Tetratetracontane	42.317	21.18	619.185 (C ₄₄ H ₉₀)	Antibacterial	Gumgumjee and Hajar, 2015
17	1-Chlorononadecane	43.850	0.60	302.966 (C ₁₉ H ₃₉ Cl)	Unknown	
18	3-Phenoxypropylamine,2-allyl-,beta,hydroxyl-N-[3,3-dimethylpropargyl]-	43.883	0.43		Unknown	

Table 2:- Chemical constituents of the yeast *Cryptococcus rajasthanensis* chloroform extract obtained by GC-MS analysis

Sl no	Compound name	Retention time	Peak Area (%)	Molecular formula and Molecular weight	Activity	Reference
	Pentadecane	26.065	1.05	212 (C ₁₅ H ₃₂)	Antibacterial	Yogeshwari et al., 2012
	Phenol-2,4-Bis(1,1-Dimethylethyl)-	26.459	8.16	206 (C ₁₄ H ₂₂ O)	Antioxidant Antibacterial	Teresa et al., 2014 Manorenjitha et al., 2013
	Tetradecane	27.992	2.02		Unknown	
	Octadecane	29.384	1.99	254.494 (C ₁₈ H ₃₈)	Antifungal agent for plant and human pathogens	Abubacker and Devi, 2015
	Bis-(3,5,5-trimethylhexyl) ether	30.580	0.52		Unknown	
	Docosane	30.670	1.22	310.60064 (C ₂₂ H ₄₆)	Antibacterial activity Enhances host egg parasitization	Gumgumjee and Hajar, 2015 Paul et al., 2002
	1,2-Benzenedicarboxylic acid, dibutyl ester	32.749	3.10	278.344 (C ₁₅ H ₂₈ O)	Antibacterial Antifouling Pesticide	Khatiwora et al., 2012 Jenecius et al., 2012 Wanxi et al., 2014
	Tetracosane	37.732	9.65	338.66 (C ₂₄ H ₅₀)	Cytotoxic towards gastric cancer cells by induction of apoptosis	Uddin et al., 2012
	Nonadecane	39.218	15.64	268 (C ₁₉ H ₄₀)	Antimicrobial and cytotoxic A cuticular hydrocarbon of insects (chemical communication)	Hsouna et al., 2011 Colazza et al., 2007
	2-methyloctacosane	40.691	17.48	408.7867 (C ₂₉ H ₆₀)	Found in cuticles of various insects spp. (chemical	Spikes et al., 2010

					communication)	
	1-Heptanol,2,4-dimethyl-, (R,R)-(+)-	40.835	1.37		Unknown	
	Celidoniol, deoxy (Nonacosane)	42.142	15.06	408 (C ₂₉ H ₆₀)	Antibacterial Antiinflammatory Chemical communication especially in Anopheles stephensi mosquito Pheromone of <i>Orgyia leucostigma</i>	Kose et al., 2016 Zakaria et al., 2014 Brei et al., 2004 Grant et al., 1987
	Tetratetracontane	43.562	11.16	619.185 (C ₄₄ H ₉₀)	Antibacterial	Gumgumjee and Hajar, 2015
	Heptadecane, 2,6,10,15- tetramethyl-	44.953	8.16	296 (C ₂₁ H ₄₄)	Unknown	
	Decane,2,3,8-trimethyl-	46.357	3.41		Unknown	

Table 3:- Chemical constituents of the yeast *Cryptococcus rajasthanensis* ethyl acetate extract obtained by GC-MS analysis

Zones of inhibition in mm				
Test microorganism	Chloroform extract (50 µg/mL)	Ethyl acetate extract (50 µg/mL)	Tetracycline (50 µg/mL)	Streptomycin (50 µg/mL)
Antibacterial activity				
<i>Escherichia coli</i>	17	16	20	19
<i>Pseudomonas aeruginosa</i>	14	13	18	18
<i>Staphylococcus aureus</i>	13	12	22	20
<i>Bacillus subtilis</i>	15	14	23	21
Antifungal activity				
Test microorganism	Chloroform extract (50 µg/mL)	Ethyl acetate extract (50 µg/mL)	Nystatin (50 µg/mL)	Fluconazole (50 µg/mL)
<i>Candida albicans</i>	18	10	16	20
<i>Aspergillus flavus</i>	10	10	14	12

Table 4:- Antimicrobial activity of yeast *Cryptococcus rajasthanensis* extracts against pathogens by Agar well diffusion assay

Discussion:-

The insect gut microflora has become a focus of intense research in recent years owing to the useful natural products that can be derived from the insect-microbe association that can be used for human therapy (Douglas, 2015). Likewise in the present study, a potent yeast strain was isolated from the *Bombyx mori* gut and was identified as *Cryptococcus rajasthanensis*. A research by Saluja and Prasad (2007) has first reported *Cryptococcus rajasthanensis* strain from Rajasthan, India from the *Andrographis echioides* inflorescence. Later on, it has also been isolated and studied in different sources like the phylloplane of rice, sugarcane, etc. (Limtong et al., 2014). The *Cryptococcus* spp. have also been reported from many insect orders including the Lepidopteran larval gut (Vega and Dowd, 2005). The yeast grew as cream colored glistening colony on potato dextrose agar which on gram staining appeared as gram positive oval shaped cells with capsules surrounding the yeast cells which is the peculiar characteristics of the Cryptococcal strain (Botton, 1980).

The FT-IR spectral analysis of the chloroform and ethyl acetate yeast extract depicted that the yeast extracts contain various biologically active functional groups like the alcoholic, phenolic, ester, aldehydic, etc. and hence proves that

the yeast possesses bioactive chemical compounds (Poojary et al., 2015). FT-IR spectra of both the extracts showed peaks at 3392.47 cm^{-1} and 3425.37 cm^{-1} which could be due to the OH functional group present in the samples (Tejado et al., 2007). Although the functional groups present in the extracts responsible for the bioactive nature of the compounds can be predicted by the FT-IR analysis, it alone cannot justify the existence of the compounds especially when an extract contains a mixture of different molecules (Mak et al., 2013).

The GC-MS chromatographic analysis of the chloroform and ethyl acetate extracts revealed the presence of various bioactive compounds. The compounds like phenol 2,4-bis(1,1-dimethylethyl)-, 1,2-benzenedicarboxylic acid dibutyl ester, celidoniol deoxy, nonadecane, tetratetracontane, 2-methyloctacosane and pentadecane were present in both the extracts but in different quantities. The compounds phenol 2,4-bis(1,1-dimethylethyl)- is known for its antibacterial and antioxidant activity (Teresa et al., 2014; Manorenjitha et al., 2013); Pentadecane has been reported to bear antibacterial activity (Yogeshwari et al., 2012); 1,2-benzenedicarboxylic acid dibutyl ester is known to possess antibacterial, antifouling and pesticidal activity (Khatiwora et al., 2012; Jenecius et al., 2012; Wanxi et al., 2014); Nonadecane is antimicrobial and cytotoxic (Hsouna et al., 2011); 2,5-octadecanoic acid methyl ester is known for the antiviral, antibacterial and antioxidant nature (Linton et al., 2013; Sudharshan et al., 2010); 2-methyltetracosane is a free radical scavenger (Ramya et al., 2015); tetratetracontane is antibacterial (Gumgumjee and Hajar, 2015), tetracosane has shown cytotoxicity towards the gastric cancer cells by inducing apoptosis (Uddin et al., 2012); Celidoniol deoxy also known as nonacosane is antibacterial and anti-inflammatory (Kose et al., 2016; Zakaria et al., 2014). The biological activities of certain compounds like 3-Ethyl-3-methylheptane; 2-Cyclohexyl-3-isopropyl-pent-4-en-2-ol; 2-Bromotetradecane; hexatriacontane; 1-chlorononadecane; 3-Phenoxypropylamine,2-allyl-,beta,hydroxyl-N-[3.3-dimethylpropargyl]; 1-Heptanol,2,4-dimethyl-, (R,R)-(+)-; etc. have not been reported and hence marked as unknown. Based on the GC-MS studies, most of the chemical constituents appear as biologically active compounds and have shown to bear pharmacologic activities which may contribute to the therapeutic potential of the yeast. Apart from the chemical constituents with therapeutic potential certain compounds are found to help the insect in chemical communication. The compounds like Celidoniol deoxy is known to aid in chemical communication in the mosquito *Anopheles stephensi* and also act as a pheromone (Brei et al., 2004; Grant et al., 1987); nonadecane and 2-methyloctacosane and are the cuticular hydrocarbons of insects that also aids in chemical communication (Spikes et al., 2010; Colazza et al., 2007). Docosane is report to aid in host egg parasitization that can be used as biocontrol agent (Paul et al., 2002). Christensen (2010) has reported that the insect associated yeasts produce volatile compounds for intraspecific and interspecific communication in insects.

The yeast extracts has shown potent antimicrobial activity against the human pathogens used and hence proves its antimicrobial nature. The chloroform extract showed higher antimicrobial activity as compared to the ethyl acetate extract which may be attributed to the more number of bioactive chemical constituents in the chloroform extract as compared to the ethyl acetate extract evaluated by GC-MS analysis. Several research studies have previously reported the antimicrobial nature of yeasts especially in biopreservation of food and yeasts as biocontrol agents (Younis et al., 2017; Mewa-Ngongang et al., 2017; Knight and Witzgall, 2013) but the antimicrobial nature of yeasts in light of its chemical constituents has not been given much importance. Thus this study reports the probability of antimicrobial nature of the yeast isolated from insect gut with regards to its bioactive chemical constituents.

Conclusion:-

Insects display a diversity of microbial association owing to the diverse nature of insects on earth. These insect associated microbes act as a large reservoir of bioactive natural products that can be used in human therapy. The yeast *Cryptococcus rajasthanensis* reported in this study has been isolated from the insect *Bombyx mori* gut and has shown considerable antimicrobial activity against the pathogenic microbial forms. The GC-MS study has also analyzed diverse chemical constituents that are known to be biologically active. Thus, the study shows the bioactive potency of the yeast isolated from the insect gut. It also indicates that the yeasts associated with the insect gut act as the reservoir for bioactive natural products that can be explored for human therapy. However, this study needs to further purify and evaluate the main chemical constituents that play the biologically active role and also to explore the novel bioactive compounds from the insect-gut microflora.

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Conflict of interest:-

Authors do not have any conflict of interest related to the manuscript.

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