Center for Sustainable Development



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Isolation, cultivation, and characterization of novel local marine micro-algae for aquaculture feed supplement production

ABSTRACT

Aquaculture is considered as a promising alternative to support the food demands of the ever-increasing population. Currently, this sector faces several challenges such as using fishmeal, which is unsustainable and expensive. Therefore, it is necessary to identify an alternative feed component that is sustainable, cost-effective and can provide the essential nutrients required by the fish. In this context, microalgae are considered as a viable source of proteins, lipids, polysaccharides and high-value products (HVPs) such as essential fatty acids, amino acids and vitamins. They play a vital role in the marine food chain and hence can be easily assimilated by the fish. The current research targeted the isolation, identification and characterization of novel marine microalgae from Qatar coastline to produce aquaculture feed supplement.

For that, environmental samples have been subjected for microalgae isolation. The isolated microorganisms belong to mainly three classes: diatoms, cyanobacteria and green algae. Among them, three have been well characterized in terms of growth rate and metabolites profiling.

OBJECTIVES

- Isolation, Identification and Characterization of novel marine microalgae to produce aquaculture feed supplement.
- Cultivation and assessment of the growth parameters and the metabolites content.

LITERATURE REVIEW

Global total capture fisheries production reached 91 million tons in 2016, as per FAO's recent statistics. However, such production didn't meet the worldwide demand due to the drastic population increase. For these reasons, aquaculture becomes of particular interest worldwide as a viable alternative to produce a large quantity of seafood (Patrik et al., 2017). Marine microalgae are considered a very promising alternative for aqua feed production since they represent the main primary food producers for the aquatic food chain (Borowitzka 1998).

METHODOLOGY

1. Sample isolation & morphological characterization

Samples were isolated and purified using common plating method F/2 and BG11 mediums.



Light microscopy is used to determine the morphological characteristics.

2. Molecular identification

The Genomic DNA was extracted and PCR amplification was performed using the genomic markers (18SrDNA, 16S rDNA). The PCR products where purified then sequenced and bioinformatics analysis was done



3. Growth rate studies

Optical Density was measured daily using spectrophotometry at 750 nm. Growth rate (u) is: $\mu = \ln X2 - \ln X1/t2-t1$ where X1 and X2



4. Cell dry-weight determination

are optic densities at times t1 and t2.

10-ml of the culture was filtered through 0.47mm dry weight filters and kept at 80°C for 24 h, then transferred to desiccators for dehydration and dry weight calculations



5. Protein and lipid extraction

Lowry-BSA method is used to determine the protein content (%).

Total lipids was extracted using Folch Method. The total lipid content will be determined gravimetrically.



6. Crude extraction of algae carotenoid

Carotenoids is extracted via maceration using methanol.

Spectrophotometry was used at 480 nm to determine the carotenoid concentration.



CONCLUSION

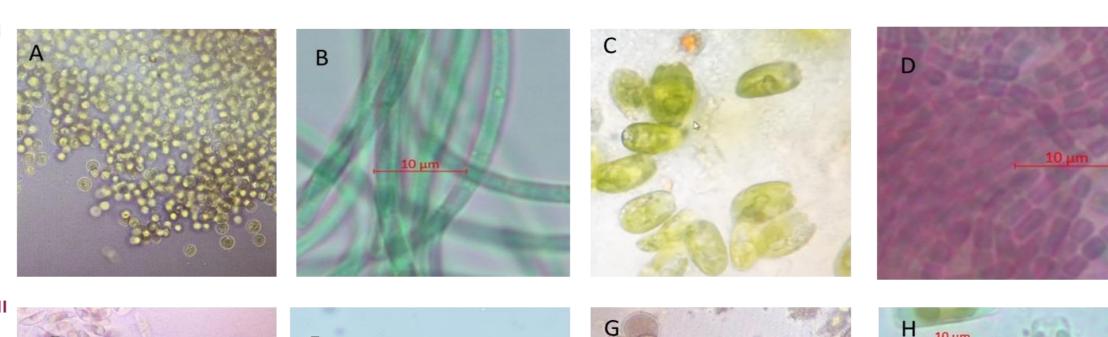
Based on the metabolites and he growth analysis, a rich feed supplement blending the three isolates can be considered as an alternative to fishmeal, well balanced and with high nutritional value.

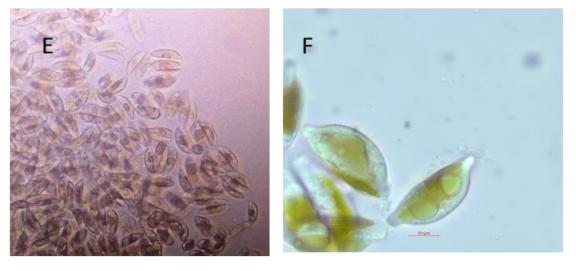
RESULTS and DISCUSSION

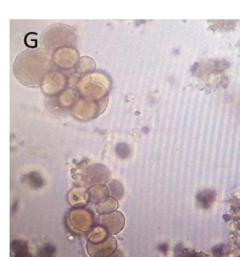
1. Isolation, purification and characterization of local marine microalgae isolates

Sample #	Location	GPS coordinates	Nature of sample	Date of isolation	T. (°C)	Salinity (ppt)	рН
1-3	Purple island	25°41′34″N, 51°32′56″E	seawater	29-9-2018	38	60	9.3
4	Purple island	25°41′34″N, 51°32′56″E	crab, leaf, shells and soil	29-9-2018	37	60	
5	Purple island	25°41′34″N, 51°32′56″E	soil sample (purple)	29-9-2018	37	60	8.7
6-7	Al-Ashat	24.74531N, 51.60062E	sea water	27-9-2018	35	47	8.9
8	Al-Ashat	24.74531N, 51.60062E	soil sample	27-9-2018	35	47	8.5

Table 1: Characteristics of the environmental samples collected







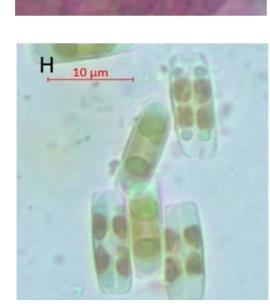


Figure 1: isolated strains under microscope

- > 20 strains were isolated, The main novel isolated strains were Diatoms, cyanobacteria and green microalgae. They were purified and characterized morphologically using light microscopy.
- > It is noted that samples collected from the purple island contain mainly diatoms and purple cyanobacteria. Accordingly, these two microorganisms might be at the origin of purple color of the soil.

2. Molecular identification of the new isolates

- DNA Extraction.
- PCR amplification of the genomic markers, such as 16s rDNA and the 18s rDNA for Microalgae and Cyanobacteria.
- PCR Product Purification to get Pure DNA
- Sequencing & bioinformatic analysis.

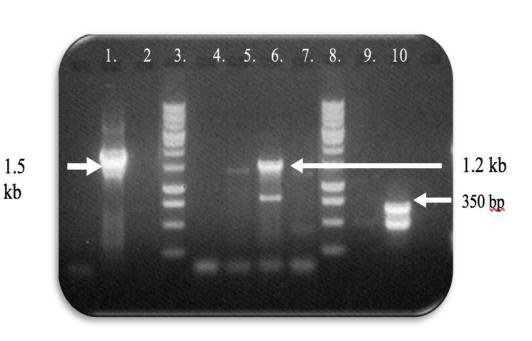


Figure 2: PCR amplification of the cellected genomic markers,

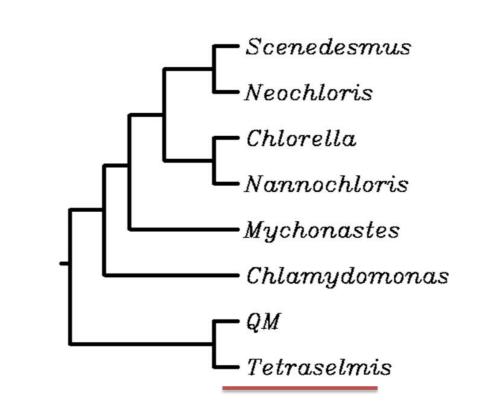
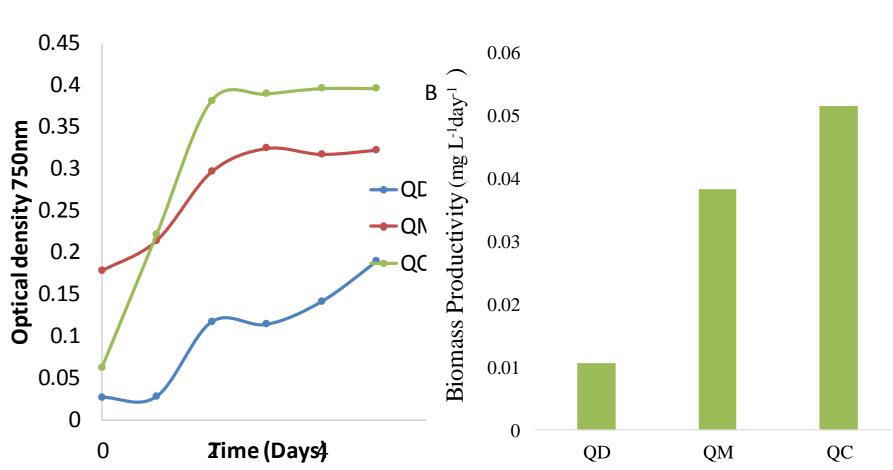


Figure 3: bioinformatics analysis., Phylogenetic tree of the strain QM

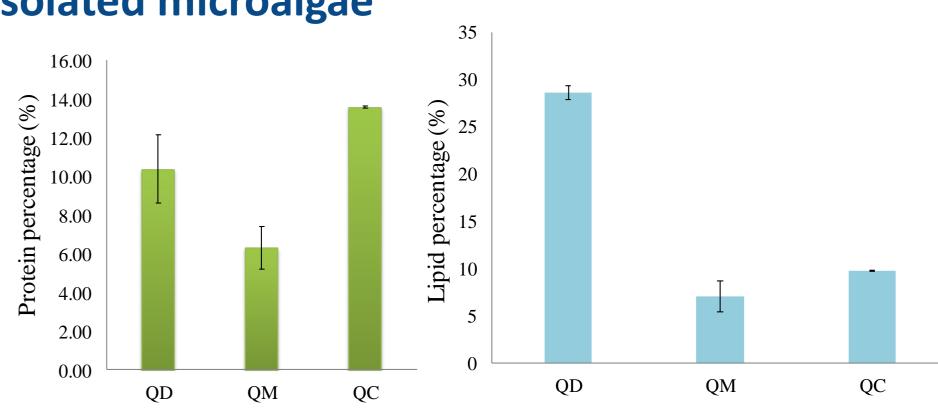
> The multiple alignment of the QM sequence proved that this strain is Tetraselmis subcoliformis (Figure 2)

3. Assessment of the Growth rate and biomass productivity

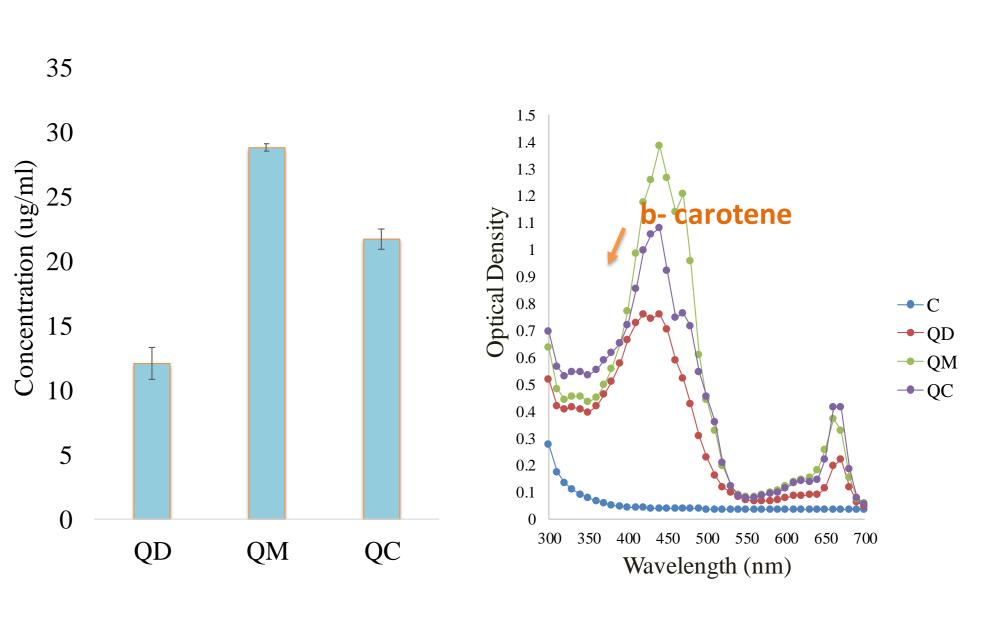


> QC presents the fastest growth and highest biomass productivity of (51.4 mgL-1day-1), followed by the Tetraselmis subcoliformis. (38.2 mgL-1day-1),

4. Metabolites profiling of the three newly isolated microalgae



- > The cyanobacteria strain (QC) presents the highest amount of protein 13.57%/g dry weight.
- > the Diatom (QD) was the highest in lipid concentration (28.62%).



- (Tetraselmis subcoliformis) present the highest amount of carotenoids.
- > The deep investigation of the nature of the carotenoids using spectral scan proved the presence of b- carotene.
- QM represents a very good candidate for producing high quality aquafeed enriched with high value added compounds.

REFERENCES