





Red algae of the genus *Laurencia* are known to produce structurally diverse terpenes that have been investigated for their natural products; most of them are halogenated compounds with important ecological functions and significant potential for the discovery of new biotechnological applications. The objective for this project is to obtain and identify molecular network from Laurencia species based on efficient methodologies, such as Vacuum Liquid Chromatography (VLC), Thin Layer Chromatography (TLC), and liquid chromatography–mass spectrometry (LCMS). **ODCTDOCT** 

# BACKGROUND AND INTRODUCTION

Marine organisms, specifically algae, are potentially prolific sources of highly bioactive secondary metabolites that might represent useful leads in the development of new pharmaceutical agents. Algae can be classified into two main groups: micro-algae and macro-algae. Macro-algae (seaweeds) which includes green, brown and red algae. Red algae are considered as the most important source of many biologically active metabolites in comparison to other algal classes.

The Red algal genus *Laurencia* is a prolific producer of halogenated secondary metabolites such as sesquiterpenes, diterpenes, and triterpenes. Biological properties of halogenated compounds have been researched for the past decades, with results showing antibacterial, antifungal, antiviral, anti-inflammatory, insecticidal activity. By obtaining Laurencia's molecular network, we can discover and characterize new halogenated compounds, along with a remarkable effort toward the evaluation of their possible biomedical and biotechnological applications.



## AQUIRED SAMPLE

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### EXTRACTION AND CHEMICAL ANALYSIS OF GENUS LAURENCIA SAMPLE

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### VLC & TLC

**Purpose:** Separation of different components within a mixture in a column based on each component's affinity (polarity and size).



Column development: separation of different strands.



Primary TLC plate









### TLC plate under UV light: useful to detect hidden spots.

### Chlorophyll



