Accelerating Biomolecular Nuclear Magnetic Resonance Assignment with A*

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Overview

- Introduction
 - Motivation
 - Nuclear Magnetic Resonance Spectroscopy
- NMR Assignment Overview
 - Data Collection and Manual Assignment
- Automation Algorithm
 - Preprocessing
 - Assignment
 - Goal State
- Conclusion
 - Results
 - Outlook

Introduction

Motivation

- Nuclear Magnetic Resonance Spectroscopy
 - Gain knowledge about protein structure
 - Study how mutations lead to diseases
- Problems
 - Generates large amounts of data
 - Data analysis is slow and error prone
- Goal
 - Automate the assignment process
 - Decrease human error
 - Increase productivity

Introduction •0

Nuclear Magnetic Resonance (NMR)

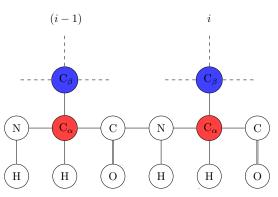
- Used to obtain structural information
 - Chemical shift values
- HNCACB experiment
 - Generates C_{α} and C_{β} residue i and i-1
- CBCA(CO) NH experiment
 - Generates C_{α} and C_{β} for residue i
 - Confirms residue data

Nuclear Magnetic Resonance Spectroscopy

Introduction 0

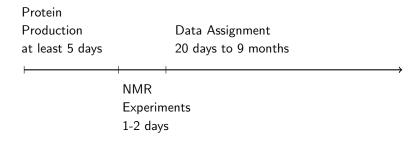
Chemical Shift Values

HNCACB



Data Collection and Manual Assignment

Time Line



Data Collection and Manual Assignment

Manual Methods

- Most time consuming part
- Prone to human error
- Missing and ambiguous data forces chunks to be skipped

Preprocessing

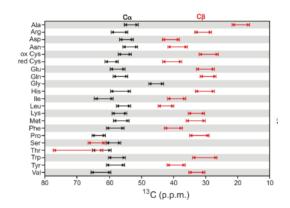
Initialization

- Input
 - Expected amino acid sequence
 - Covered to expectation chemical shift values
 - Stored as the protein chain
 - NMR chemical shift data
 - C_{α} and C_{β} for residue i and i-1
 - Stored in a tile
- Missing data
 - Place holder tile generation
- Grouping

00 0000 000

Preprocessing

Grouping



iew Automation Algorithm

Conclusior 0 0000

Assignment

Starting the assignment

Protein Chain	Tiles		
11.5	13 9 1.5	11 13 0.5	9 12 2.5
12.5			
9.6			

Assignment

Cost Calculation

- Accuracy matching the protein chain residue
- Accuracy matching the tile above current tile
- Cost of all tiles place before current tile

Automation Algorithm

OO
OO
OO

Conclusion 0 0000

Assignment

Generating child nodes

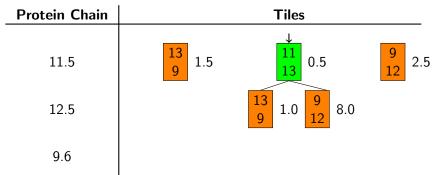
Tiles to assign: $\begin{bmatrix} 13 \\ 9 \end{bmatrix} \begin{bmatrix} 11 \\ 13 \end{bmatrix} \begin{bmatrix} 9 \\ 12 \end{bmatrix}$

Protein Chain		Tiles	
11.5	13 9 1.5	11 13 0.5	9 12 2.5
12.5			
9.6			

Assignment

Generating child nodes

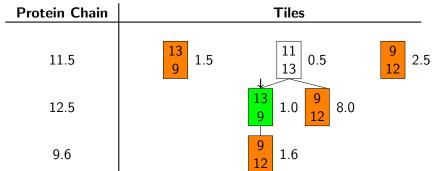




Goal State

Goal State

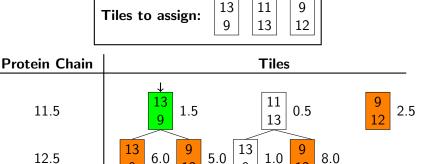




1.6

Goal State

Goal State



9.6

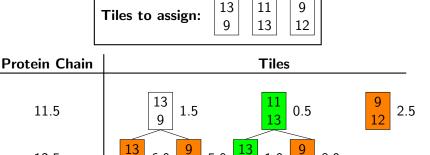
Goal State

Solution State

11.5

12.5

9.6



1.0

1.6

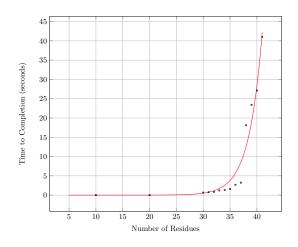
8.0

5.0

6.0

Results

Time of Assignment



Future Goals

- Parallelization
 - Decrease assignment time
 - Allow for larger data sets
- Machine learning
 - Increase accuracy of assignment
 - Optimize cost calculation

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Outlook

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- Leah Robison (research colleague)

Outlook

Bibliography



Sean Cahill and Mark Girvin. Introduction to 3d triple resonance experiments. 2012

Outlook

Thank You

