

PARAMETER ADVISING FOR  
MULTIPLE SEQUENCE ALIGNMENT

by

Daniel Frank DeBlasio

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THE UNIVERSITY OF ARIZONA  
GRADUATE COLLEGE

As members of the Dissertation Committee, we certify that we have read the dissertation prepared by Daniel Frank DeBlasio, entitled Parameter Advising for Multiple Sequence Alignment and recommend that it be accepted as fulfilling the dissertation requirement for the Degree of Doctor of Philosophy.

\_\_\_\_\_  
John Kececioglu

Date: 15 April 2016

\_\_\_\_\_  
Alon Efrat

Date: 15 April 2016

\_\_\_\_\_  
Stephen Kobourov

Date: 15 April 2016

\_\_\_\_\_  
Mike Sanderson

Date: 15 April 2016

Final approval and acceptance of this dissertation is contingent upon the candidate's submission of the final copies of the dissertation to the Graduate College.

I hereby certify that I have read this dissertation prepared under my direction and recommend that it be accepted as fulfilling the dissertation requirement.

\_\_\_\_\_  
Dissertation Director: John Kececioglu

Date: 15 April 2016

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SIGNED: Daniel Frank DeBlasio

## ACKNOWLEDGEMENTS

Insert your acknowledgements here.

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## ABSTRACT

Abstract text



## CHAPTER 1

### Introduction and Background

#### Overview

This gives some preview of what is contained in this chapter.

#### 1.1 Introduction

At this point you may want to include a figure, such as the one below:

#### 1.2 Background

Maybe this section should have a table:

#### 1.3 Plan of the dissertation

This may refer to chapters other than Chapter 1

This template was used to produce my dissertation (DeBlasio, 2016), and it was passed around by graduate students at the University of Arizona. At this point, I

<i>d1gvoa</i>	203	...	gsvenrarlvlevvdavcnewsad- <b>RIGIRVSP</b> igtfqnvndngpnee--adalyl---	...	255
<i>d2dora</i>	141	...	ydfeatekllke-----vftfftk- <b>PLGVKLPP</b> yf-----dlvhfdim	...	178
<i>d1oyb</i>	215	...	gsienrarftlevvdalveaighe- <b>KVGLRLSP</b> ygvfnsmsggaetgivaqyayvage	...	272
<i>d1o94a1</i>	193	...	gslenrarfwletlekvkhavgsdc <b>AIATRF</b> GV-----dtvygpgq	...	234
<i>d1ep3a</i>	147	...	tdpevaaalvka-----ckavskv- <b>PLYVKLSP</b> nvt-----divpiaka	...	185

(a) Higher Accuracy Alignment

<i>d1gvoa</i>	184	...	yl-lhqflspssnqrtdqyggsvenrarlvlevvdavcnewsad- <b>RIGIRVSP</b> igtfq	...	240
<i>d2dora</i>	159	...	k <b>P-LGVKLPP</b> yf--dlvhfdimaeilnqfplt <b>YVNSV</b> -nsig---nglfidpeaesv	...	209
<i>d1oyb</i>	196	...	yl-lnqfldphsnttrtdeyggsienrarftlevvdalveaighe- <b>KVGLRLSP</b> ygvf	...	252
<i>d1o94a1</i>	174	...	yl-plqflnpyynkrtkdyggslenrarfwletlekvkhavgsdc <b>AIATRF</b> --- <b>GV</b> dt	...	228
<i>d1ep3a</i>	164	...	kv <b>PLYVKLSP</b> nv-tdivpiakaveaagad <b>GLTMI</b> ntl-----mgvrfdlkrqp	...	212

(b) Lower Accuracy Alignment

Figure 1.1: This is the caption that ends up below the figure.

Table 1.1: *This is above the table itself*

Parameter choice ( $\sigma, \gamma_I, \gamma_E, \lambda_I, \lambda_E$ )	Advisor set			
	Default	Greedy	Exact	Oracle
$k = 2$				
(VTML200, 50, 17, 41, 40)	(2)	(2)		
(VTML200, 55, 30, 45, 42)		(2)	(3)	(1)
(BLSUM80, 60, 9, 43, 42)			(2)	
(BLSUM45, 65, 35, 44, 44)				(3)
$k = 3$				
(VTML200, 50, 17, 41, 40)	(2)	(2)		
(VTML200, 55, 30, 45, 42)		(3)	(5)	(1)
(BLSUM80, 60, 26, 43, 43)		(2)	(2)	
(VTML200, 55, 30, 41, 40)			(6)	
(VTML40, 45, 29, 40, 39)				(7)
(BLSUM62, 65, 16, 44, 42)				(8)
$k = 4$				
(VTML200, 50, 17, 41, 40)	(2)	(2)		
(VTML200, 55, 30, 45, 42)		(3)	(9)	(6)
(BLSUM80, 60, 26, 43, 43)		(2)		
(VTML200, 60, 15, 41, 40)		(1)		
(VTML200, 45, 6, 40, 40)			(8)	(1)
(VTML200, 55, 30, 41, 40)			(8)	
(BLSUM80, 55, 19, 43, 42)			(1)	
(VTML40, 45, 29, 40, 39)				(4)
(BLSUM62, 65, 35, 44, 42)				(3)

do not think they require a template, but some may still find it useful.

## REFERENCES

- DeBlasio, D. F. (2016). *Parameter Advising for Multiple Sequence Alignment*. PhD dissertation, Department of Computer Science, The University of Arizona.