FIRST PRINCIPLE CALCULATIONS OF DEFECT STRUCTURES IN ZINC OXIDE

By

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A thesis submitted in partial fulfillment of the requirements for the degree of

BACHELOR OF SCIENCE

UNIVERSITY OF THE PHILIPPINES - DILIMAN National Institute of Physics

MAY 2020

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To the Faculty of University of the Philippines Diliman National Institute of Physics:			
The members of the Committee appointed to examine the thesis of CHRISTIAN LOER			
T. LLEMIT find it satisfactory and recommend that it be accepted.			
	Roland V. Sarmago, Ph.D., Chair		
	Donald Trump, Ph.D.		
	Rodrigo Duterte, Ph.D.		

ACKNOWLEDGMENT

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FIRST PRINCIPLE CALCULATIONS OF DEFECT STRUCTURES IN ZINC OXIDE

Abstract

by Christian Loer T. Llemit, BS University of the Philippines - Diliman May 2020

: Roland V. Sarmago

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TABLE OF CONTENTS

		Pa	age
ACK	NOV	WLEDGMENT	iii
ABST	ΓRA	CT	iv
LIST	OF	TABLES	ix
LIST	OF	FIGURES	X
1	IN	ΓRODUCTION	1
	1.1	Semiconductors	1
	1.2	Applications of Semiconductors	1
	1.3	Defects in Semiconductors	1
	1.4	ZnO	1
		1.4.1 Subsection of section - double quotes	1
		1.4.2 Another subsection of section - citations	2
		1.4.2.1 Subsubsection of section - italic text	2
2	TH	EORETICAL FRAMEWORK	4
	2.1	Many-body Quantum Mechanics	4
		2.1.1 Simplifying Assumptions	6
		2.1.2 Time Independent Schrödinger Equation	6
		2.1.3 Use of Atomic Units	6
		2.1.4 Hamiltonian Operator	6
		2.1.5 Indistinguishability of electrons	6
	2.2	Early First Principle Calculations	6
		2.2.1 n-electron problem	6
		2.2.2 Hartree Method	6
		2.2.3 Hartree-Fock Method	6

	2.3	Density Functional Theory
		2.3.1 Electron Density
		2.3.2 Hohenberg-Kohn (HK) Formalism 6
		2.3.2.1 First HK Theorem
		2.3.2.2 Second HK Theorem
		2.3.3 Kohn Sham (KS) Formalism
		2.3.3.1 KS Equation
		2.3.3.2 Energy Terms
	2.4	Exchange-correlation Functional
		2.4.1 Local Density Approximation (LDA) 6
		2.4.2 Generalized Gradient Approximation (GGA) 6
3	\mathbf{DF}	T Calculation of Solids
	3.1	Pseudopotential Approach
		3.1.1 Freezing the core electrons
		3.1.2 Pseudizing the valence electrons
		3.1.3 Common Pseudopotentials
	3.2	Choosing the appropriate Calculation Size
		3.2.1 Use of Supercell
		3.2.1.1 Periodic Boundary Conditions (PBC)
		3.2.2 Use of Reciprocal Space
		3.2.2.1 Reciprocal Lattice
		3.2.2.2 First Brillouin Zone
		3.2.2.3 Irreducible Brillouin Zone
		3.2.3 k-point sampling
		3.2.3.1 Monkhorst-Pack method
		3.2.3.2 Gamma Point Sampling
	3.3	Bloch Representations
		3.3.1 Electrons in solid
		3.3.2 Bloch Theorem in periodic systems
		3.3.3 Fourier Expansion of Bloch representations
		3.3.3.1 Fourier Expansions

		3.3.3.2 Fast Fourier Transformation (FFT)
	3.4	3.3.3.3 Kohn-Sham Matrix Representations10Plane Wave (PW) Expansion103.4.1 Basis Set10
		3.4.1.1 Local Basis Set
		3.4.1.2 Plane Wave Basis Set
		3.4.2 Plane Wave Expansion for KS quantities
		3.4.2.1 Charge Density
		3.4.2.2 Kinetic Energy
		3.4.2.3 Effective Potential
	3.5	Electronic Structure
		3.5.1 Band Structure of free electrons
		3.5.2 Band Structure of electrons in solids
		3.5.3 Electronic Density of States
	3.6	Practical Aspects
		3.6.1 Energy Cutoffs
		3.6.1.1 Cutoff for Wavefunction
		3.6.1.2 Cutoff for Charge Density
		3.6.2 Smearing
		3.6.2.1 Gaussian Smearing
		3.6.2.2 Fermi Smearing
		3.6.2.3 Methfessel–Paxton Smearing
		3.6.2.4
4	MA	THEMATICS NOTATION
	4.1	Some Math Stuff
	4.2	Math equation
	4.3	Chapter section
	4.4	Chapter section
5	FIC	GURES AND TABLES
	5.1	Examples of a figure

	5.2	Example of a table
	5.3	Chapter section
REF	ERE	ICES
APP	END	\mathbf{X}
\mathbf{A}		
В		
\mathbf{C}		
D		
E		2

LIST OF TABLES

5.1	Whole-genome sequences	used in this study		16
-----	------------------------	--------------------	--	----

LIST OF FIGURES

5.1	Cost per raw megabase of DNA sequence from 2001 to 2015	15
A.1	Cost per raw megabase of DNA sequence from 2001 to 2015	20
A.2	Cost per raw megabase of DNA sequence from 2001 to 2015	21
B.1	Cost per raw megabase of DNA sequence from 2001 to 2015	22
C.1	Cost per raw megabase of DNA sequence from 2001 to 2015	23
D.1	Cost per raw megabase of DNA sequence from 2001 to 2015	24
E.1	Cost per raw megabase of DNA sequence from 2001 to 2015	25

Dedication

This dissertation/thesis is dedicated to my mother and father who provided both emotional and financial support ${\cal C}$

Chapter One

INTRODUCTION

- 1.1 Semiconductors
- 1.2 Applications of Semiconductors
- 1.3 Defects in Semiconductors
- 1.4 ZnO

This is an example of how to cite [1]

1.4.1 Subsection of section - double quotes

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Chapter Two

THEORETICAL FRAMEWORK

2.1 Many-body Quantum Mechanics

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2.1.1	Simplifying Assumptions		
2.1.2	Time Independent Schrödinger Equation		
2.1.3	3 Use of Atomic Units		
2.1.4	Hamiltonian Operator		
2.1.5	Indistinguishability of electrons		
2.2	Early First Principle Calculations		
2.2.1	n-electron problem		
2.2.2	Hartree Method		
2.2.3	Hartree-Fock Method		
2.3	Density Functional Theory		
2.3.1	Electron Density		
2.3.2	Hohenberg-Kohn (HK) Formalism		
2.3.2.1	First HK Theorem		
2.3.2.2	Second HK Theorem		
2.3.3	Kohn Sham (KS) Formalism		
2.3.3.1	KS Equation		
2.3.3.2	Energy Terms		

2.4 Exchange-correlation Functional

6

2.4.1 Local Density Approximation (LDA)

Chapter Three

DFT Calculation of Solids

3.1 Pseudopotential Approach

This is sample text

- 3.1.1 Freezing the core electrons
- 3.1.2 Pseudizing the valence electrons
- 3.1.3 Common Pseudopotentials
- 3.2 Choosing the appropriate Calculation Size
- 3.2.1 Use of Supercell
- 3.2.1.1 Periodic Boundary Conditions (PBC)
- 3.2.2 Use of Reciprocal Space
- 3.2.2.1 Reciprocal Lattice
- 3.2.2.2 First Brillouin Zone
- 3.2.2.3 Irreducible Brillouin Zone
- 3.2.3 k-point sampling
- 3.2.3.1 Monkhorst-Pack method
- 3.2.3.2 Gamma Point Sampling

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3.3 Bloch Representations

- 3.3.1 Electrons in solid
- 3.3.2 Bloch Theorem in periodic systems
- 3.3.3 Fourier Expansion of Bloch representations
- 3.3.3.1 Fourier Expansions
- 3.3.3.2 Fast Fourier Transformation (FFT)
- 3.3.3.3 Kohn-Sham Matrix Representations
- 3.4 Plane Wave (PW) Expansion
- 3.4.1 Basis Set
- 3.4.1.1 Local Basis Set
- 3.4.1.2 Plane Wave Basis Set
- 3.4.2 Plane Wave Expansion for KS quantities
- 3.4.2.1 Charge Density
- 3.4.2.2 Kinetic Energy
- 3.4.2.3 Effective Potential
- 3.5 Electronic Structure
- 3.5.1 Band Structure of free electrons
- 3.5.2 Band Structure of electrons in solids
- 3.5.3 Electronic Density of States
- 3.6 Practical Aspects

Chapter Four

MATHEMATICS NOTATION

4.1 Some Math Stuff

LaTeX has a special way to embed mathematical symbols and notations. Here are some of them. Also observe how a bullet list is made.

- greater than \geq
- less than <
- percent sign %
- multiply $N \times N$
- inline equation M = N(N-1)/2

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4.2 Math equation

Example of a mathematical formula:

$$ADD = \sum_{i=1}^{M} | \langle D(n+1, i) \rangle - \langle D(n, i) \rangle |$$
 (4.1)

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4.3 Chapter section

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Chapter Five

FIGURES AND TABLES

5.1 Examples of a figure

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Example of a figure. Example of reference to a figure in the text (Fig. 5.1). Phasellus dolor neque, vehicula vestibulum semper at, facilisis eget libero. Mauris interdum magna molestie, auctor felis a, condimentum odio. Pellentesque habitant morbi tristique senectus et netus et malesuada fames ac turpis egestas. Suspendisse maximus lacinia dignissim. Maecenas pharetra accumsan metus, sagittis dictum purus sollicitudin eget. Curabitur ut porttitor arcu, ut porttitor ipsum. Vestibulum porttitor finibus sapien, ac pharetra odio bibendum nec. Nullam tincidunt dignissim risus imperdiet dictum.

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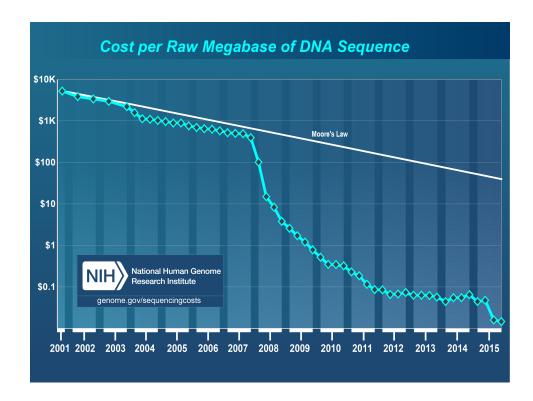


Figure 5.1 Cost per raw megabase of DNA sequence from 2001 to 2015. Straight line - Moore's Law, blue curve - cost in US dollars, Y-axis scale is logarithmic. Graph reproduced from [5]

5.2 Example of a table

Example of a table and here is the reference to Table 5.1. Tables in, my opinion, are the hardest thing to make.

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ORGANISM	Accession no.	GENOME SIZE (bp)	No. CDS
Mesorhizobium loti	NC_002678	7036071	6743
Sinorhizobium meliloti	NC_003047	3654135	3359
Bradyrhizobium japonicum	NC_004463	9105828	8317
Rhodopseudomonas palustris	NC_005296	5459213	4813
Bartonella quintana	NC_005955	1581384	1142
Bartonella henselae	NC_005956	1931047	1488
Rickettsia typhi	NC_006142	1111496	837
Beijerinckia indica	NC_010581	4170153	3569

Table 5.1 Whole-genome sequences used in this study

sapien, ac pharetra odio bibendum nec. Nullam tincidunt dignissim risus imperdiet dictum.

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5.3 Chapter section

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maximus lacinia dignissim. Maecenas pharetra accumsan metus, sagittis dictum purus sollicitudin eget. Curabitur ut porttitor arcu, ut porttitor ipsum. Vestibulum porttitor finibus sapien, ac pharetra odio bibendum nec. Nullam tincidunt dignissim risus imperdiet dictum.

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- 3. Altschul, S. F. *et al.* Gapped BLAST and PSI-BLAST: a new generation of protein database search programs. *Nucleic acids research* **25**, 3389–3402 (1997).
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- 5. Wetterstrand, K. A. DNA Sequencing Costs: Data from the NHGRI Genome Sequencing Program (GSP) www.genome.gov/sequencingcosts.



Appendix A

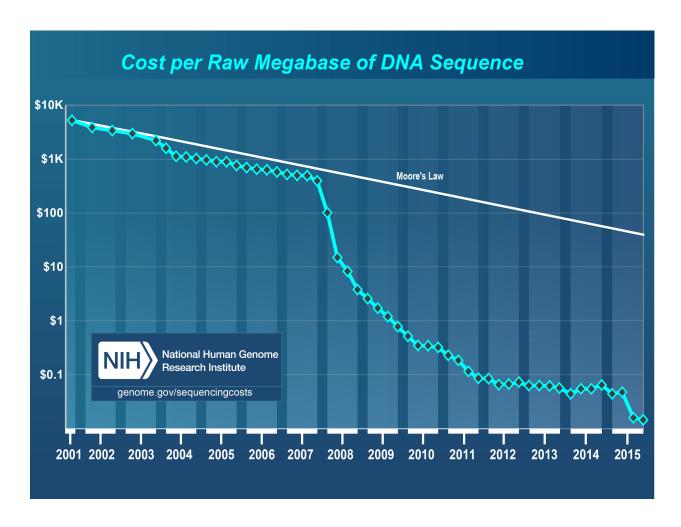


Figure A.1 Cost per raw megabase of DNA sequence from 2001 to 2015. Straight line - Moore's Law, blue curve - cost in US dollars, Y-axis scale is logarithmic. Graph reproduced from [5]

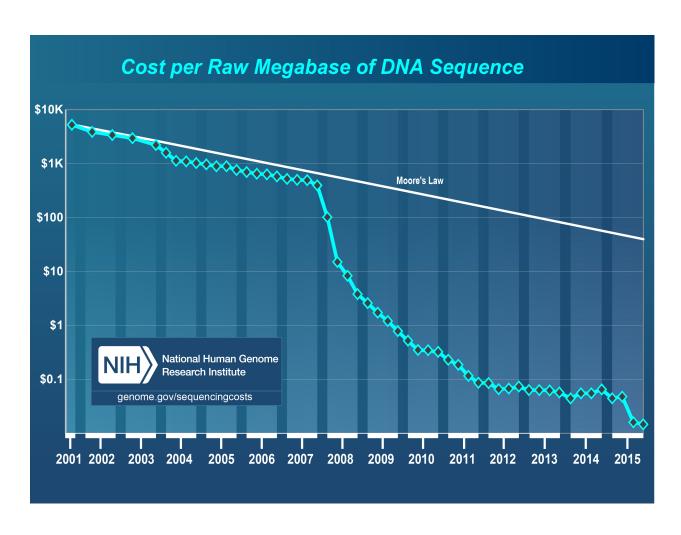


Figure A.2 Cost per raw megabase of DNA sequence from 2001 to 2015. Straight line - Moore's Law, blue curve - cost in US dollars, Y-axis scale is logarithmic. Graph reproduced from [5]

Appendix B

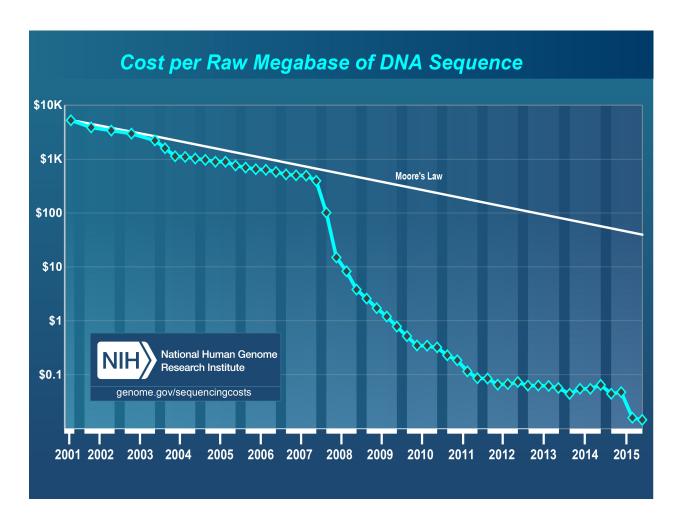


Figure B.1 Cost per raw megabase of DNA sequence from 2001 to 2015. Straight line - Moore's Law, blue curve - cost in US dollars, Y-axis scale is logarithmic. Graph reproduced from [5]

Appendix C



Figure C.1 Cost per raw megabase of DNA sequence from 2001 to 2015. Straight line - Moore's Law, blue curve - cost in US dollars, Y-axis scale is logarithmic. Graph reproduced from [5]

Appendix D

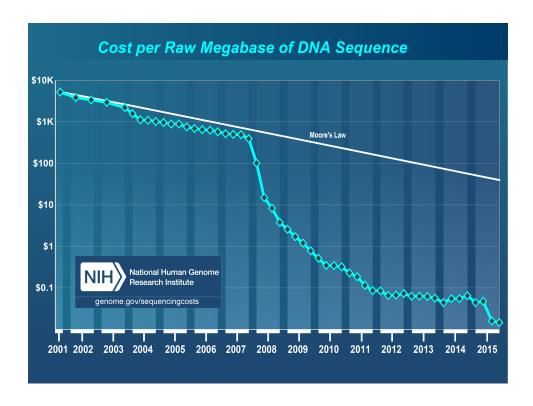


Figure D.1 Cost per raw megabase of DNA sequence from 2001 to 2015. Straight line - Moore's Law, blue curve - cost in US dollars, Y-axis scale is logarithmic. Graph reproduced from [5]

Appendix E



Figure E.1 Cost per raw megabase of DNA sequence from 2001 to 2015. Straight line - Moore's Law, blue curve - cost in US dollars, Y-axis scale is logarithmic. Graph reproduced from [5]