

BioDEG

Biodegradation and Corrosion Simulation using Finite Element

User Manual

Version 0.8

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[BioDeg website](#)

KU LEUVEN

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Contents

1	Introduction	3
1.1	Authors	3
1.2	Acknowledgments	3
1.3	Referencing BIODEG	3
2	Useful background information	3
3	Installation	4
3.1	Compiling and installing external libraries	4
3.1.1	Instructions for PETSc and Qt	4
3.1.2	Instructions for FreeFEM	4
3.2	Obtaining and Compiling BIODEG	4
4	Running BioDeg	4
4.1	Configuring the simulation	4
4.1.1	Example 1	4
4.1.2	Example 2	4
5	Future extensions to BioDeg	4
6	Finding answers to more questions	4
A	Run-time input parameters	4
A.1	Global parameters	5

1 Introduction

BiODEG is an open source software written in FreeFEM (a domain-specific language for finite element programming), C++, and Python for modeling the degradation of metallic biomaterials and simulating the biodegradation behavior of medical devices, implants and scaffolds in corrosion experiments. It can handle any geometry of desire and supports parallel computing to simulate models with a high level of complexity.

1.1 Authors

BiODEG is developed by the [Biomechanics Research group at KU Leuven and University of Liege](#). The code is currently maintained by its principal developer, who manage the development of the mathematical models and the core functionalities.

Principal developer

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1.2 Acknowledgments

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1.3 Referencing BioDeg

Please refer to [BiODEG repository](#), section "Publications and referencing" to properly cite the use of BiODEG in your scientific work.

2 Useful background information

You may refer to the following articles for a background of the methods and algorithms implemented in BiODEG.

In addition, below are some useful references on finite element method and some online resources that provide a background of finite elements and their application to the solution of partial differential equations.

3 Installation

All the underlying installation instructions assume a Linux operating system. We assume standard tools and libraries like CMake, compilers- (C, C++ and Fortran), and MPI libraries are pre-installed. Most high-performance computers would have the latest version of these libraries in the default environment.

3.1 Compiling and installing external libraries

3.1.1 Instructions for PETSc and Qt

3.1.2 Instructions for FreeFEM

3.2 Obtaining and Compiling BioDeg

Assuming that you have already installed the above external dependencies, next follow the steps below to obtain and compile BIODEG.

4 Running BioDeg

4.1 Configuring the simulation

4.1.1 Example 1

4.1.2 Example 2

5 Future extensions to BioDeg

The future versions of BioDeg will focus on the implementation of the following methodologies/features.

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6 Finding answers to more questions

If you have questions that go beyond this manual, there are a number of resources:

- For questions/suggestions about BIODEG installation, bugs, or similar stuff please use the [BIODEG issue tracker](#).
- BIODEG is primarily based on the [FreeFEM](#). If you have particular questions about FreeFEM, contact the community at <https://community.freefem.org/>.
- If you have specific questions about BIODEG that are not suitable for public and archived mailing lists, you can contact the primary developer and mentor:
 - Mojtaba Barzegari: mojtaba.barzegari@kuleuven.be.
 - Liesbet Geris: liesbet.geris@kuleuven.be (Mentor).

A Run-time input parameters

The underlying description of the input parameters also includes a “Standard/Advanced/Developer” label, which signifies whether an input parameter is a standard one, or an advanced level parameter, or a developer level one only meant for development purposes. The default values of the “Advanced” and “Developer” labelled parameters are good enough for almost all cases. However, in some cases user may need to use “Advanced” labelled parameters. For user convenience, all input parameters are also indexed at the end of this manual in Section ??.

A.1 Global parameters