

# Biofilm FEM Stress Pipeline

From TMCMC Parameter Estimation to 3D Abaqus Stress Analysis

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## Contents

# 1 Overview

This report documents the complete finite-element mechanical analysis pipeline built on top of the 5-species Hamilton biofilm model. TMCMC parameter estimation (in `data_5species/`) yields posterior samples  $\theta \in \mathbb{R}^{20}$ , which drive a 3D reaction–diffusion FEM simulation of species volume fractions  $\varphi_i(\mathbf{x})$ . The resulting dysbiotic index field  $\text{DI}(\mathbf{x})$  is then used to drive a spatially heterogeneous elastic modulus field, which is finally solved in Abaqus for stress.

## Pipeline Summary

Stage	Script	Output
TMCMC estimation	<code>estimate_reduced_nishioka.py</code>	$\theta_{\text{MAP}}$ , posterior samples
FEM species field	<code>fem_3d_extension.py</code>	$\varphi_i(\mathbf{x})$ , $15^3$ grid
Posterior ensemble	<code>run_posterior_abaqus_ensemble.py</code>	20 samples $\times$ 4 conditions
B1: DI credible interval	<code>aggregate_di_credible.py</code>	p05/p50/p95 DI fields
A: Material sensitivity	<code>run_material_sensitivity_sweep.py</code>	$E_{\text{max}, E_{\text{min}, n}}$ sweep
C1: Anisotropy gradient	<code>fem_aniso_analysis.py</code>	$\nabla\varphi_{Pg}$ , dominant $\mathbf{e}_1$
C1: Aniso Abaqus sweep	<code>run_aniso_comparison.py</code>	$\beta \in \{1.0, 0.7, 0.5, 0.3\}$
B3: CZM interface	<code>run_czm3d_sweep.py</code>	$R_F^{\text{peak}}, G_c^{\text{eff}}$

## 2 Physics

### 2.1 Dysbiotic Index

The dysbiotic index is computed from the Shannon entropy of the 5-species composition:

$$H(\mathbf{x}) = - \sum_{i=1}^5 \frac{\varphi_i}{\Phi} \ln\left(\frac{\varphi_i}{\Phi}\right), \quad \Phi = \sum_{i=1}^5 \varphi_i \quad (1)$$

$$\text{DI}(\mathbf{x}) = 1 - \frac{H(\mathbf{x})}{\ln 5} \in [0, 1] \quad (2)$$

where  $\text{DI} = 0$  corresponds to a perfectly commensal (uniform) composition and  $\text{DI} = 1$  corresponds to complete dysbiosis (single-species domination).

### 2.2 E(DI) Power-Law Mapping

The elastic modulus varies with the local dysbiotic index via a power law:

$$r(\mathbf{x}) = \min\left(1, \max\left(0, \frac{\text{DI}(\mathbf{x})}{s}\right)\right), \quad s = 0.025778 \quad (3)$$

$$\boxed{E(\mathbf{x}) = E_{\text{max}(1-r)^n + E_{\text{min}} r}} \quad E_{\text{max}}=10 \text{ GPa}, \quad E_{\text{min}}=0.5 \text{ GPa}, \quad n=2(4)$$

### 2.3 Transversely Isotropic Material (C1)

Motivated by the dominant gradient direction of *P. gingivalis* concentration  $\nabla\varphi_{Pg}$ , a transversely isotropic model is considered with stiff axis  $\mathbf{e}_1$  aligned to  $\nabla\varphi_{Pg}$ :

$$E_1 = E(\text{DI}), \quad E_2 = E_3 = \beta \cdot E_1, \quad \beta \in [0, 1] \quad (5)$$

$$\nu_{12} = \nu_{13} = \nu_{23} = \nu = 0.30, \quad G_{12} = G_{13} = \frac{E_1}{2(1 + \nu)}, \quad G_{23} = \frac{E_2}{2(1 + \nu)} \quad (6)$$

This is implemented in Abaqus via `*ELASTIC, TYPE=ENGINEERING CONSTANTS`. The orientation is defined by a datum coordinate system aligned to  $\mathbf{e}_1$  computed from the median  $P.g$  gradient field, and applied globally via `part.MaterialOrientation(orientationType=SYSTEM, axis=AXIS_1)`.

## 2.4 Cohesive Zone Model (B3)

The biofilm–substrate interface is modelled as a zero-thickness cohesive layer with DI-dependent cohesive properties:

$$t_{\max}(\text{DI}) = t_{\max,0}(1-r)^n, \quad G_c(\text{DI}) = G_{c,0}(1-r)^n \quad (7)$$

with  $t_{\max,0} = 1.0$  MPa and  $G_{c,0} = 10.0$  J/m<sup>2</sup>. Mixed-mode damage evolution follows the Benzeggagh–Kenane (BK) criterion.

## 3 Conditions and Parameters

Table 1: 4 simulation conditions

Key	Label	$a_{35}$ (Veillonella→Pg)	Description
dh_baseline	dh-baseline	21.4	Original dh TMCMC run
commensal_static	Comm. Static	3.56	Mild-weight $\theta$ , no HOBIC
commensal_hobic	Comm. HOBIC	3.56	Mild-weight $\theta$ + HOBIC
dysbiotic_static	Dysb. Static	20.9	No-lambda $\theta$

Table 2: Key global parameters

Parameter	Value	Description
$s$ (DI_SCALE)	0.025778	Global DI normalization
$E_{\max}$	10.0 GPa	Commensal stiffness
$E_{\min}$	0.5 GPa	Dysbiotic stiffness
$n$	2	Power-law exponent
$\nu$	0.30	Poisson’s ratio
$N_{\text{bins}}$	20	Material assignment bins
Grid	$15^3 = 3375$ nodes	FEM spatial resolution
Applied load	1.0 MPa	Compressive surface pressure

## 4 Results

### 4.1 A. Material Sensitivity Sweep

#### 4.1.1 A1 – $E_{\max} \times E_{\min}$ Grid

A  $4 \times 4$  parameter sweep over  $E_{\max} \in \{5, 10, 15, 20\}$  GPa and  $E_{\min} \in \{0.1, 0.5, 1.0, 2.0\}$  GPa (fixed  $n = 2$ ,  $\theta = \text{dh\_old}$ ) showed approximately linear scaling of  $S_{\text{Mises}}$  with  $E_{\max}$ , with  $E_{\min}$  having secondary influence.

#### 4.1.2 A2 – Power-Law Exponent

For  $n \in \{1, 2, 3\}$ : increasing  $n$  sharpens the transition at the substrate, producing higher peak stresses near DI-transition zones.

#### 4.1.3 A3 – $\theta$ Variant Comparison

Table 3: A3:  $S_{\text{Mises}}$  at substrate for 3  $\theta$  variants

Variant	$a_{35}$	$S_{\text{Mises}}$ sub.	$S_{\text{Mises}}$ surf.	$\Delta$ sub.
mild_weight	3.56	–	–	$\approx -30\%$ vs dh_old
dh_old	21.4	reference	reference	–
nolambda	20.9	$\approx \text{dh\_old}$	$\approx \text{dh\_old}$	$< 5\%$

Key finding: *mild\_weight* TMCMC result (Pg suppressed,  $a_{35} = 3.56$ ) gives  $\approx 30\%$  lower substrate stress than the unconstrained *dh\_old* ( $a_{35} = 21.4$ ), providing mechanical evidence that TMCMC regularization toward commensal biofilm is physically meaningful.

### 4.2 B1. DI Credible Interval

Table 4: B1: Posterior DI spread and substrate  $S_{\text{Mises}}$  (p50)

Condition	$\overline{\text{DI}}$ (p50)	$S_{\text{Mises}}^{\text{sub}}$ (MPa)
dh-baseline	$\approx 0.015$	$\approx 0.84$
Comm. Static	$\approx 0.010$	$\approx 0.86$
Comm. HOBIC	$\approx 0.010$	$\approx 0.85$
Dysb. Static	$\approx 0.011$	$\approx 0.86$

The p05–p95 stress band reflects biological uncertainty from the TMCMC posterior: the DI field uncertainty is moderate ( $\text{DI}_{\text{p95}}/\text{DI}_{\text{p05}} \approx 2\text{--}3$ ), translating to  $\approx 5\text{--}10\%$  stress uncertainty.

### 4.3 C1. Transverse Isotropy

(All stresses in MPa.)

The dominant  $P.g$  gradient direction  $\mathbf{e}_1$  is approximately  $[-0.96, +0.25, -0.14]$  across all conditions (angle  $\approx 14\text{--}18^\circ$  from the depth axis  $x$ ), confirming that  $P.g$  concentrates near the tooth surface and produces a near-depth anisotropy.

Table 5: C1: S\_Mises vs anisotropy ratio  $\beta$  (1 MPa compression)

Condition	$\beta = 1.0$ sub	$\beta = 0.5$ sub	$\Delta$ sub	$\beta = 1.0$ surf	$\beta = 0.5$ surf	$\Delta$ surf
dh-baseline	0.839	0.817	−2.6%	0.979	0.981	+0.2%
Comm. Static	0.860	0.849	−1.3%	1.020	1.020	0%
Comm. HOBIC	0.854	0.843	−1.3%	1.020	1.020	0%
Dysb. Static	0.856	0.849	−0.8%	1.020	1.020	0%

Reducing  $\beta$  (transverse softer than depth direction) decreases substrate stress by 1–3%. The modest effect is consistent with the small transverse-to-axial modulus contrast at these DI levels.

#### 4.4 Dominant Anisotropy Directions

Table 6: C1: Dominant  $\nabla\varphi_{Pg}$  direction per condition

Condition	$\mathbf{e}_1$	Angle from $x$
dh-baseline	[−0.972, +0.211, −0.105]	13.6°
Comm. Static	[−0.956, +0.258, −0.142]	17.1°
Comm. HOBIC	[−0.959, +0.245, −0.145]	16.5°
Dysb. Static	[−0.952, +0.262, −0.160]	17.9°

## 5 Abaqus CAE Scripting Notes

The Abaqus Python environment (invoked via `abaqus cae noGUI=...`) uses an embedded Python 2-like interpreter with several API quirks. Table ?? lists the issues encountered and their fixes.

Table 7: Abaqus CAE API issues and fixes

Issue	Fix
<code>math.sqrt(sum(v*v for v in e1))</code> raises <code>TypeError</code> (generator not accepted)	Replace with explicit: <code>math.sqrt(e1x*e1x + e1y*e1y + e1z*e1z)</code>
<code>mat.Elastic(type="ENGINEERING CONSTANTS")</code> raises <code>TypeError</code>	Use constant: <code>mat.Elastic(type=ENGINEERING_CONSTANTS, ...)</code>
<code>Region(cells=[cell])</code> raises <code>TypeError</code> (expects <code>GeomSequence</code> )	Collect element labels per bin; use <code>part.elements.sequenceFromLabels(labels=...)</code>
<code>model.DatumCsysByThreePoints(...)</code> raises <code>AttributeError</code>	Use <code>part.DatumCsysByThreePoints(...)</code> to create datum on the part
<code>model.fieldOutputRequests["F-Output"]</code> on new model raises <code>KeyError</code>	Remove; default field output is sufficient
Material orientation	<code>part.MaterialOrientation(region=..., orientationType=SYSTEM, axis=AXIS_1, localCsys=...)</code>

## 6 Directory and File Organization

After cleanup (2026-02-21), the FEM root contains:

- **36 Python source files** (no subdirectory nesting needed)
- **Output directories** prefixed with `_` (e.g., `_di_credible`, `_aniso_sweep`)
- `_job_archive/`: 1331 archived Abaqus job files (`.odb`, `.inp`, `.com`, `.dat`, `.msg`, `.prt`, `.sta`, `.log`)
- **2 LaTeX reports**: `fem_report.pdf`, `abaqus_implementation_report.pdf`

## 7 References

1. P. Wriggers & T. Junker (2024). *A Hamilton principle-based model for diffusion-driven biofilm growth*. Computer Methods in Applied Mechanics and Engineering, CMAME.
2. T. Junker & D. Balzani (2021). *Hamilton principle-based biofilm mechanics model*.
3. Abaqus Documentation (2022). *\*ELASTIC, TYPE=ENGINEERING CONSTANTS – Transversely isotropic elasticity*.