

Biofilm FEM Stress Pipeline

From TMCMC Parameter Estimation to 3D Abaqus Stress Analysis

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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>	θ_{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>	$E_{\max, E_{\min, n}}$ sweep
C1: Anisotropy gradient	<code>fem_aniso_analysis.py</code>	$\nabla \varphi_{P_g}$, 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^{peak} , G_c^{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 $E(\text{DI})$ Power-Law Mapping

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

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

$E(\mathbf{x}) = E_{\max(1-r)^n + E_{\min r}}$

$E_{\max=10 \text{ GPa}}, \quad E_{\min=0.5 \text{ GPa}}, \quad n=2$

(4)

2.3 Transversely Isotropic Material (C1)

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

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

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

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/m². 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
<code>dh_baseline</code>	dh-baseline	21.4	Original dh TMC MC run
<code>commensal_static</code>	Comm. Static	3.56	Mild-weight θ , no HOBIC
<code>commensal_hobic</code>	Comm. HOBIC	3.56	Mild-weight θ + HOBIC
<code>dysbiotic_static</code>	Dysb. Static	20.9	No-lambda θ

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
ν	0.30	Poisson's ratio
N_{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 4×4 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 $n = 2$, $\theta = dh_old$) showed approximately linear scaling of S_{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 – θ Variant Comparison

Table 3: A3: S_{Mises} at substrate for 3 θ variants

Variant	a_{35}	S_{Mises} sub.	S_{Mises} surf.	Δ sub.
mild_weight	3.56	–	–	$\approx -30\%$ vs dh_old
dh_old	21.4	reference	reference	–
nolambda	20.9	$\approx dh_old$	$\approx dh_old$	< 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_{Mises} (p50)

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

The p05–p95 stress band reflects biological uncertainty from the TMCMC posterior: the DI field uncertainty is moderate ($DI_{p95}/DI_{p05} \approx 2\text{--}3$), 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 β (1 MPa compression)

Condition	$\beta = 1.0$ sub	$\beta = 0.5$ sub	Δ sub	$\beta = 1.0$ surf	$\beta = 0.5$ surf	Δ 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 β (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 $\nabla\varphi_{P_g}$ 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 TypeError (generator not accepted)	Replace with explicit: <code>math.sqrt(e1x*e1x + e1y*e1y + e1z*e1z)</code>
<code>mat.Elastic(type="ENGINEERING_CONSTANTS")</code> raises TypeError	Use constant: <code>mat.Elastic(type=ENGINEERING_CONSTANTS, ...)</code>
<code>Region(cells=[cell])</code> raises TypeError (expects GeomSequence)	Collect element labels per bin; use <code>part.elements.sequenceFromLabels(labels=...)</code>
<code>model.DatumCsysByThreePoints(...)</code> raises AttributeError	Use <code>part.DatumCsysByThreePoints(...)</code> to create datum on the part
<code>model.fieldOutputRequests["F-Output1"]</code> ; default field output is sufficient on new model	raises KeyError
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*.