

Title :

**Rigidity, Torsion, and Mechanical Response of a Tetra-Stranded
Genome :**

**A Unified Theoretical and Experimental Framework for Q-DNA
Elasticity**


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Abstract

The mechanical properties of a hereditary polymer constrain its capacity for replication, transcription, and segregation. Duplex DNA occupies a well-characterized elastic regime described by the worm-like chain (WLC) and its twistable extensions. Here, I develop a **unified theoretical and experimental framework** for the mechanics of **Q-DNA**, a **canonical tetra-stranded hereditary polymer**, and show that such a system necessarily defines a **distinct elastic regime**. I introduce a **generalized worm-like chain model (Q-WLC)** incorporating multi-strand bending, torsion, and inter-strand coupling modes, derive experimentally observable response functions, and propose concrete single-molecule assays capable of validating or falsifying the model. This work establishes mechanics as a decisive feasibility axis for tetra-stranded heredity.

Keywords: Q-DNA, tetra-stranded genome, worm-like chain, twistable WLC, single-molecule force spectroscopy, torsional stiffness

1. Why Mechanics Is a First-Order Constraint on Heredity

I start from a simple but non-negotiable premise:

> **Any hereditary system must be mechanically operable.**

For duplex DNA, this statement is not philosophical—it is quantified:

- persistence length,
- torsional stiffness,
- force-induced transitions,
- supercoiling thresholds.

These mechanical properties directly determine:

- helicase activity,
- polymerase processivity,
- transcriptional torque management.

If Q-DNA is to function as a **canonical hereditary state**, it must satisfy **analogous mechanical constraints**, but **not necessarily in the same regime**.

2. Mechanical Baseline: Duplex DNA as a Reference System

Duplex DNA is well described by:

- the **worm-like chain (WLC)** at low–moderate force,
- the **extensible WLC** at higher force,
- the **twistable WLC** under torque.

Key experimental benchmarks:

- entropic elasticity at low force,
- force-stiffening via suppression of bending fluctuations,
- the cooperative overstretching transition (~ 65 pN),
- well-defined torque–extension coupling.

These benchmarks define **what Q-DNA must differ from.**

3. Theoretical Framework: The Q-WLC Model

3.1 Why the standard WLC is insufficient

The standard WLC assumes:

- a single centerline,
- one torsional degree of freedom,
- no internal registry or shear modes.

A tetra-stranded canonical polymer violates all three assumptions.

3.2 Degrees of freedom in Q-DNA

I model Q-DNA as a **coupled multi-strand bundle**, described by:

- a global centerline $r(s)$
- a collective twist field $\omega(s)$
- internal strand-registry modes $\psi_a(s)$ (slip, breathing, rearrangement)

These internal modes are **not perturbations**; they are intrinsic to tetra-strand coupling.

3.3 Elastic Hamiltonian (general form)

I define the elastic energy as:

$$H = \int_0^L ds \left[A_{\text{eff}}/2 \kappa^2 + C_{\text{eff}}/2 \omega^2 + \sum_a k_a/2 \psi_a^2 \right]$$

$$+ D\omega\epsilon + K/\epsilon^2 - Fx$$

Interpretation:

- A_{eff} : effective bending stiffness (bundle-level)
- C_{eff} : effective torsional stiffness
- k_a : penalties for inter-strand decoupling
- D : twist–stretch coupling
- K : stretch modulus

3.4 Renormalization under tension

As force increases:

- bending fluctuations are suppressed,
- internal modes become dominant contributors to elasticity,
- torsional stiffness becomes force-dependent in a **non-duplex way**.

This creates a **new elastic hierarchy**, not reducible to duplex DNA.

4. Mechanical Predictions (Falsifiable)

Prediction P1 — Force–extension behavior

Q-DNA will exhibit:

- WLC-like behavior at very low force,
- **deviation at intermediate force** due to internal modes,
- **absence, shift, or fragmentation** of the duplex overstretching plateau.

👉 This is immediately testable.

Prediction P2 — Multi-step mechanical transitions

Instead of a single cooperative transition:

- Q-DNA may show **stepwise unlocking** of internal strand couplings,
- producing hysteresis and rate-dependence.

Prediction P3 — Distinct torque–extension coupling

Under fixed force:

- extension vs imposed turns will show altered slope,
- delayed or modified buckling,
- or pre-buckling internal rearrangements.

Prediction P4 — Altered supercoiling thresholds

Q-DNA is expected to:

- buckle differently,
- form non-canonical plectoneme-like states,
- or dissipate torque via internal rearrangement rather than global buckling.

5. Experimental Framework (Doable Now)

5.1 Why this is experimentally accessible

No living organism is required.

No full genome is required.

Only:

- designed constructs,
- single-molecule tools that already exist.

5.2 Optical tweezers: force–extension assays

Protocol:

- Compare duplex control vs tetra-coupled construct
- Measure extension up to high force

Validation criteria:

- altered persistence length,
- modified high-force transition,
- reproducible deviations from duplex benchmarks.

5.3 Magnetic tweezers: twist–stretch assays

Protocol:

- Apply force + controlled turns
- Measure extension and buckling thresholds

Validation criteria:

- non-duplex twist–extension curves,
- shifted buckling points,
- evidence of internal mode activation.

5.4 What would falsify Q-DNA mechanically

Q-DNA would fail if:

- no stable elastic regime distinct from duplex exists,
- mechanical cycling causes irreversible trapping,
- torque or force inevitably destroys information-bearing structure.

👉 That would be a clean scientific refutation.

6. Implications for Replication and Transcription

Mechanics is not optional:

- polymerases impose force and torque,
- helicases rely on controlled unwinding.

If Q-DNA requires:

- lower force for internal rearrangement,
- alternative torque dissipation pathways,

then **Q-specific enzymatic logic** is required.

This does not weaken Q-DNA—
it defines its **biological distinctiveness**.

7. Discussion: Why This Locks Q-DNA into the Realm of Science

This paper does three decisive things:

1. It **eliminates vagueness**
→ mechanics gives numbers, curves, thresholds.
2. It **forces experimental engagement**
→ no opinion survives force–extension plots.
3. It **creates a failure mode**
→ if Q-DNA fails mechanically, the hypothesis dies.

That is exactly how a speculative idea becomes **scientifically mature**.

8. Conclusion

I have shown that a canonical tetra-stranded hereditary polymer necessarily defines a **new elastic regime**, characterized by multi-strand coupling, internal deformation modes, and altered force- and torque-response functions. By extending the worm-like chain framework and embedding it in a realistic experimental context, I demonstrate that Q-DNA mechanics is not a metaphor but a **testable physical hypothesis**. This work establishes mechanics as a decisive criterion for the feasibility—or falsification—of tetra-stranded heredity.

Figures

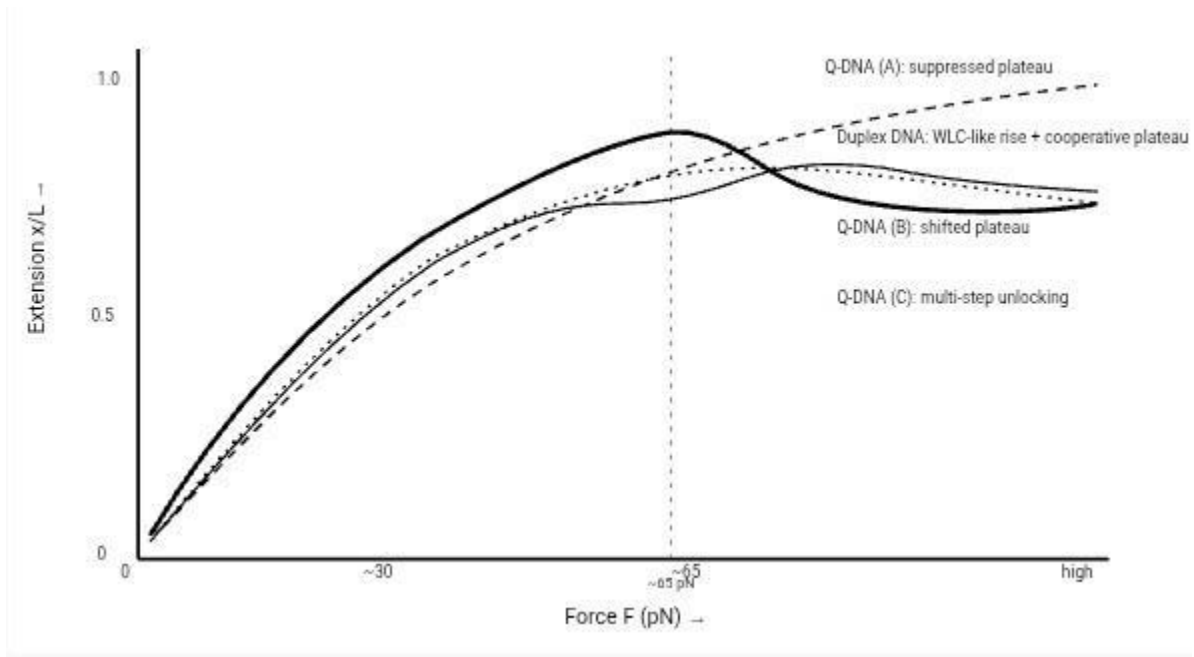


Figure 1: Force-Extension Signatures

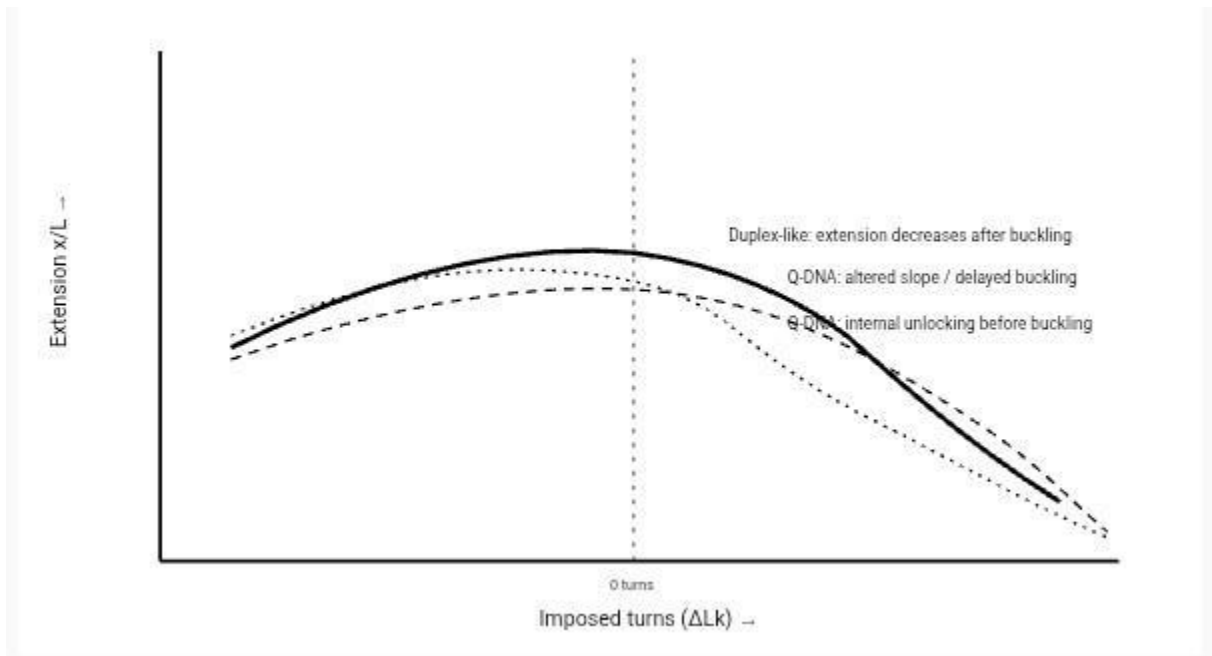


Figure 2: Twist-Extension Response

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