

## Cancer as Tau-Node Disruption

*Strand 1 Unbounded Growth · DNA Tau-Address Corruption vs Deletion · Warburg Effect as Tau-Energy Diversion · Tau-Restoration Therapy*

*Stephen Daubney*

*The Daubney Foundation*

### **P-CANC-1 The FOT Definition of Cancer**

In conventional oncology, cancer is defined as uncontrolled cell division driven by genetic mutation. The Force of Time framework gives a deeper structural definition: cancer is the uncoupling of Strand 1 (the replication programme) from Strand 2 (the Tau-regulatory signal). When Strand 2 H-bond tension is lost or disrupted at a specific Tau-address in the DNA, the Strand 1 programme continues executing without the Strand 2 brake — producing unlimited replication.

#### **P-CANC-1**

Cancer = Strand 1 replication programme running without Strand 2 Tau-regulation. The oncogenic mutation is not the cause but the symptom: the loss of Strand 2 coupling at a specific Tau-address in the DNA. Tau-address corruption is reversible; Tau-address deletion is not.

This distinction — corruption vs deletion — is clinically significant. A corrupted Tau-address retains the structural capacity to be restored by re-establishing Strand 2 coupling. A deleted Tau-address (terminal chromosomal damage) cannot be restored by any Tau-field intervention. FOT predicts that most early-stage cancers involve address corruption, while late-stage cancers increasingly involve deletion.

### **P-CANC-2 The Warburg Effect as Tau-Energy Diversion**

The Warburg effect — cancer cells preferentially use aerobic glycolysis rather than oxidative phosphorylation even in the presence of oxygen — is one of the most robust and puzzling observations in cancer biology. FOT provides the mechanism: a Tau-decoupled cell diverts its energy metabolism away from the mitochondrial Tau-generator (oxidative phosphorylation produces 36 ATP via the full Tau-cycle) toward the Strand-1-only glycolytic programme (2 ATP — the minimal {2} lattice energy output).

Full Tau-cycle (OXPHOS): 36 ATP =  $2^2 \times 3^2 = 4 \times 9$  Glycolysis (Strand 1 only): 2 ATP =  $2^1$  Ratio:  $36 / 2 = 18 = 2 \times 3^2$  Cancer chooses 2 over 36 — the Strand-1-only sub-lattice.

The factor  $18 = 2 \times 3^2$  is the same factor that appears in the water bond angle ( $18/\pi^2$  radians). A cancerous cell abandons the full {2,3} Tau-cycle in favour of the minimal {2} programme. The Warburg effect is not an inefficiency — it is the metabolic signature of Strand 2 decoupling.

### **P-CANC-3 Metastasis as Tau-Node Fragmentation**

Metastasis — the spread of cancer cells to secondary sites — represents Tau-node fragmentation: a cell that has lost its primary Tau-address coupling loses spatial registration within the tissue Tau-field and migrates to whichever register boundary it can couple with. The bloodstream is a Tau-medium (water-based, G1 register) that transports decoupled cells to secondary nodes.

FOT prediction on metastatic tropism: cancer cells from a given primary site preferentially colonise secondary sites whose Tau-address sub-lattice is closest to the corrupted primary address. This explains the non-random pattern of metastatic spread (e.g., breast cancer to bone, lung cancer to brain) without invoking organ-specific chemical signals: the pattern is geometric, set by Tau-address proximity in the DNA lattice.

### **P-CANC-4 Telomerase and the Strand 2 Bypass**

Cancer cells reactivate telomerase — the enzyme that extends telomere repeats — allowing unlimited replication. In FOT, telomerase is a Strand 2 bypass enzyme: it resets the Tau-counter (telomere length) without re-establishing Strand 2 regulation. The cell gains Tau-counter capacity (unlimited division capability) but remains Strand-2-decoupled (still cancerous). Telomerase inhibition without Tau-restoration addresses the counter but not the coupling deficit.

*FOT prediction: telomerase inhibition alone will not cure cancer because it does not restore Strand 2 Tau-coupling. A cancer cell with inhibited telomerase remains Strand-2-decoupled: it will find alternative Tau-counter bypass mechanisms (ALT pathways) or undergo crisis-and-recombination. Successful treatment requires Tau-address restoration — not just counter inhibition.*

### **P-CANC-5 Tau-Restoration Therapy Framework**

The FOT framework predicts that restoring Strand 2 Tau-coupling at the corrupted DNA address will reverse the cancerous state. The mechanism: externally applied coherent oscillations at the Tau-lattice frequencies corresponding to the disrupted address re-establish Strand 2 H-bond tension across the damaged region.

| <b>Therapy modality</b>    | <b>FOT mechanism</b>                        | <b>Proposed frequency</b>                  |
|----------------------------|---|--|
| Coherent EM radiation      | Re-couples Strand 2 at corrupted address    | Tau-lattice frequency of affected register |
| Coherent ultrasound        | Mechanical Tau-wave restores H-bond tension | Acoustic analogue of EM coupling frequency |
| 40 Hz gamma entrainment    | Brain Tau-lock (P-CONS) → immune Tau-boost  | 40 Hz = 40,075 km/1000 (P-SCHUM-2)         |
| Fasting / metabolic switch | Forces cell back to OXPHOS (36 ATP)         | Removes Warburg energy advantage           |

### **P-CANC-6 Testable Predictions**

| Prediction   | Measurement  | Distinguishes FOT from standard model                           |
|--|--|---|
| Metastatic tropism pattern encodes Tau-address proximity                                   | Map metastatic routes vs DNA lattice address distance  | Standard model predicts chemical signals; FOT predicts geometry |
| Early-stage cancer cells restore normal behaviour under coherent Tau-frequency EM exposure | In vitro cell line trials with lattice-frequency EM    | Standard model has no mechanism for frequency-specific reversal |
| Warburg ratio (OXPHOS/glycolysis) recovers toward 18:1 under Tau-restoration conditions    | Metabolic flux analysis during Tau-frequency treatment | $18 = 2 \times 3^2$ is a specific FOT prediction, not 17 or 19  |

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