

MU Simulation Log — Test 2 (Simple T(r))

Goal

Introduce the selector T(r) into the weight calculation to test whether it can prevent collapse at the fold.

- In Test 1, both branches collapsed to zero → no survival.
- With T(r), the expectation is that one branch will remain alive near the fold, showing the first MU-style selection.

Selector Function

We'll use the simplest possible form of T(r):

$$T(r) = \frac{1}{1 + |r - r_{\rm c}|}$$

with $r_{\rm c}=0.25$ (the fold).

Interpretation:

- Close to the fold $(r \approx r_c)$, T is strong.
- Away from the fold, T fades out.
- This gives maximum influence where the classical model fails, and minimal interference elsewhere.

Parameters

Keep the baseline parameters:

```
gamma = 1.0
      = 1e-6
eps
hbar = 0.1
Twindow = 1.0
```

Predictions

- Without T: both branches → weights = 0, ratio = nan.
- With T: at least one branch survives → weight nonzero.
- Plots:
 - Log weight plot should show separation instead of collapse.

- grr plot unchanged (geometry still flips), but the weight plots no longer vanish at the portal.
- This would be the **first positive evidence** that MU can resolve the fold mathematically.