Cross-Validation Consistency Across Patients
(Each fold tests on different patient)

E' mean: 0.542 ± 0.019
(CV = 3.5%)

Legend:
- E' (Poisson)
- D' (MSE)

Y-axis: Test SSIM (2μm)

X-axis: Cross-Validation Fold

Fold 1 (Train: P1+P2):
- P5: 0.527
- P5: 0.214

Fold 2 (Train: P1+P5):
- P2: 0.530
- P2: 0.200

Fold 3 (Train: P2+P5):
- P1: 0.568
- P1: 0.186