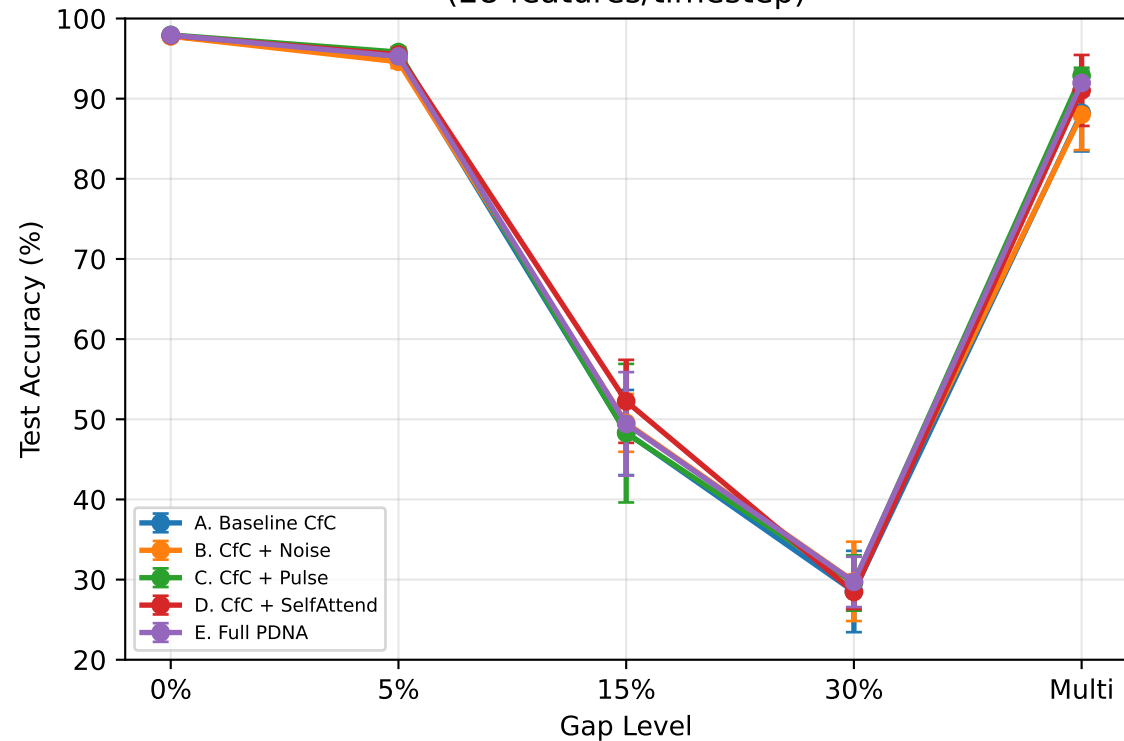(a) sMNIST: High Information Density (28 features/timestep)

(b) psMNIST: Low Information Density (1 feature/timestep)

Legend (a):
- A. Baseline CfC
- B. CfC + Noise
- C. CfC + Pulse
- D. CfC + SelfAttend
- E. Full PDNA

Legend (b):
- A. Baseline CfC (n=5)
- B. CfC + Noise (n=2)

Near-zero degradation (gap-invariant)