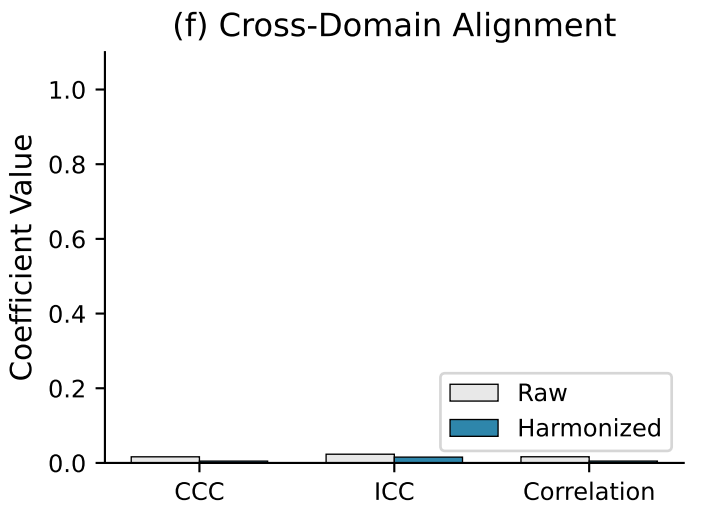
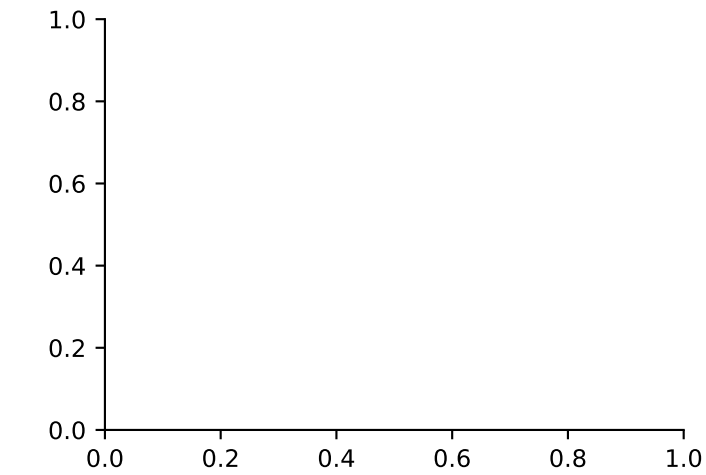
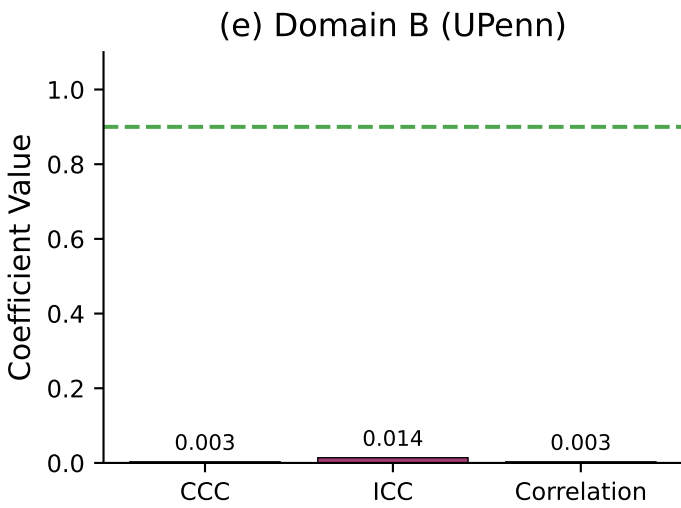
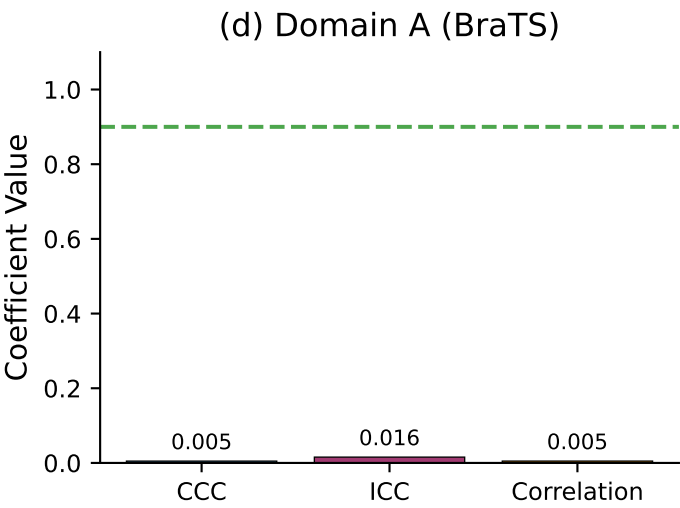
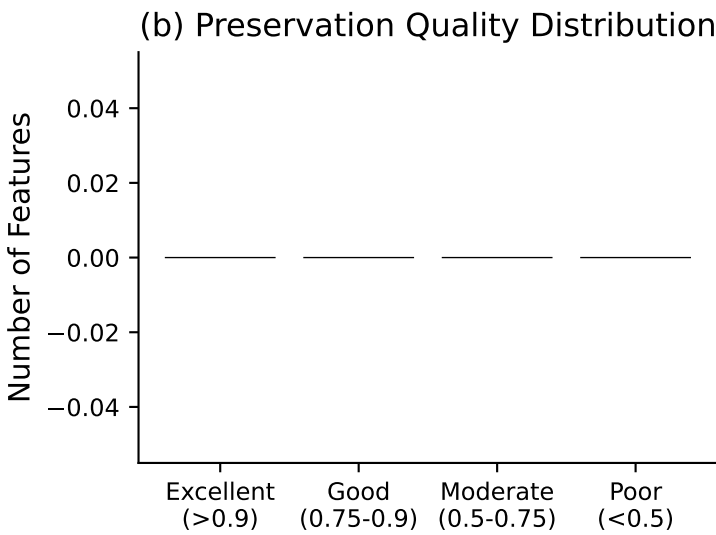
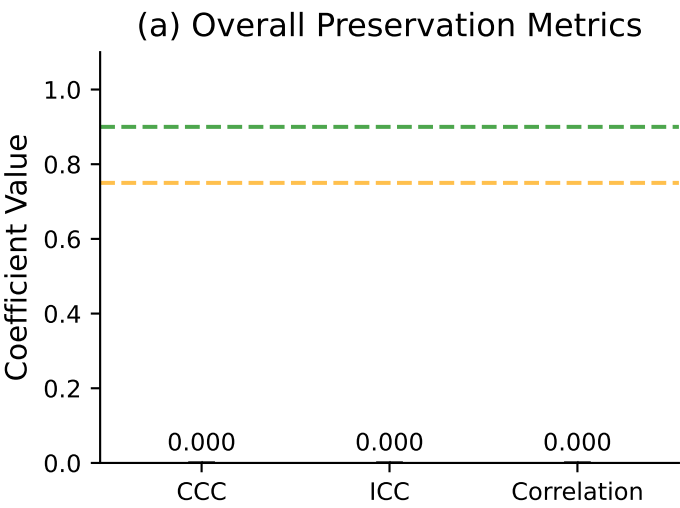


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Key Findings - SA-CycleGAN-2.5D Radiomics Preservation:

1. Overall Preservation: Mean CCC = 0.000, Mean ICC = 0.000
- 0 features with excellent preservation (CCC > 0.9)
- 0 features with good preservation (0.75 < CCC < 0.9)

2. Domain-Specific Results:
- Domain A: CCC = 0.005
- Domain B: CCC = 0.003

3. Cross-Domain Alignment:
- Raw alignment CCC: 0.016
- Harmonized alignment CCC: 0.005
- Improvement: -0.012

Interpretation: Harmonization effectively preserves radiomics features within each domain while improving cross-domain feature alignment, supporting clinical validity.