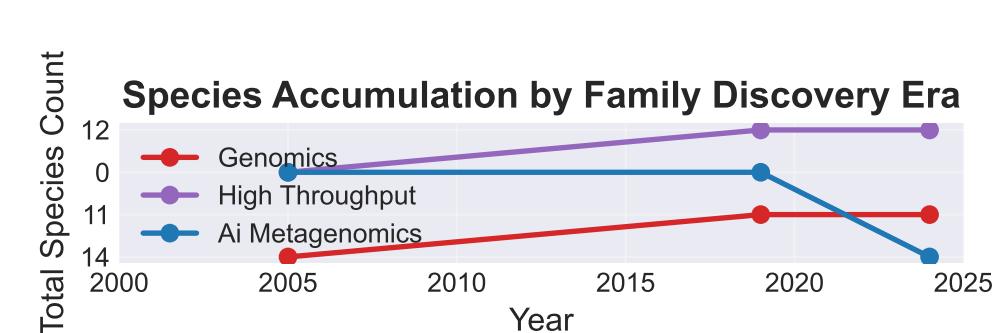
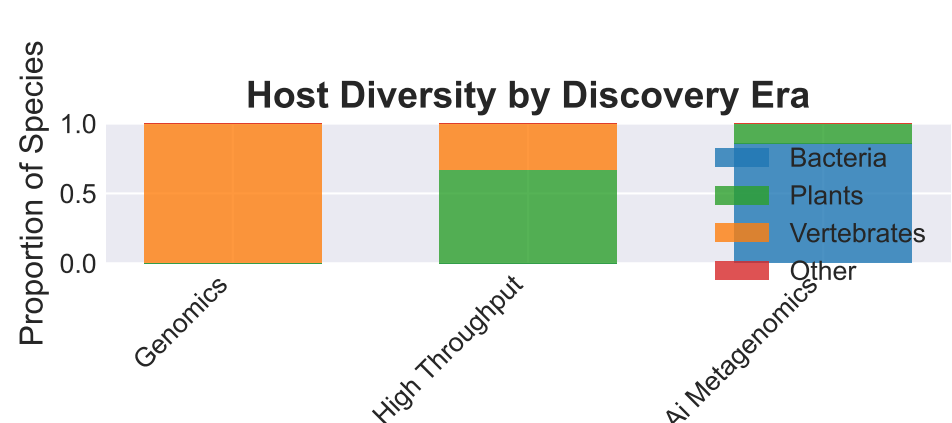
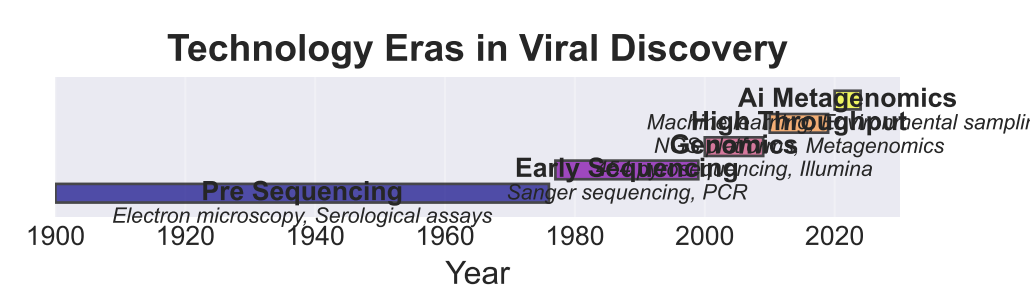
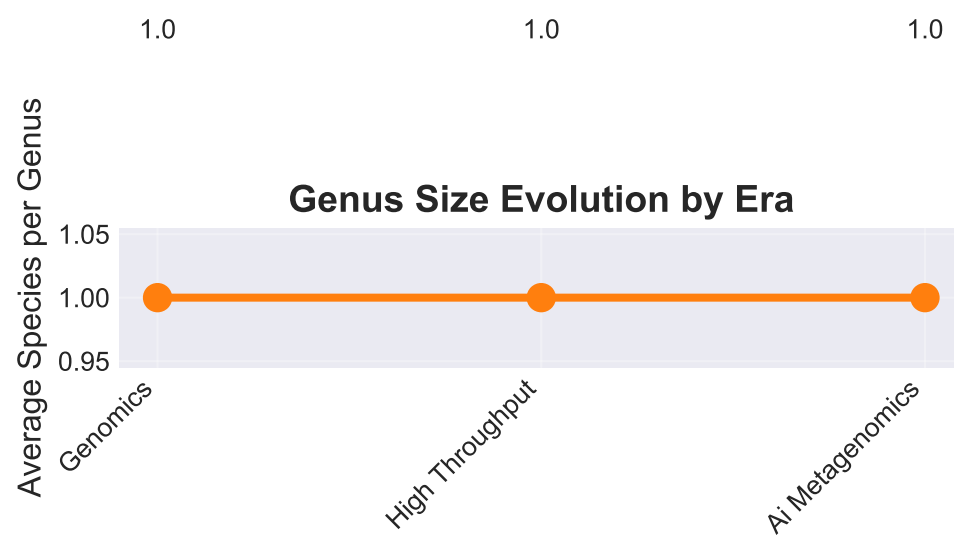
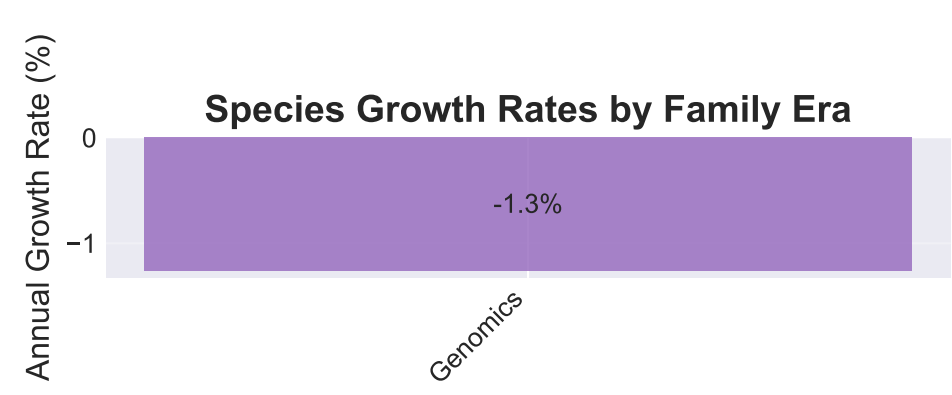
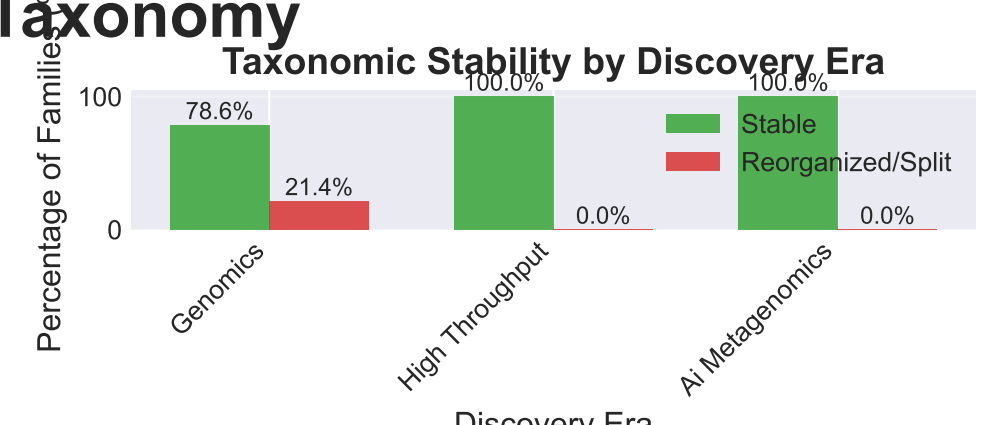
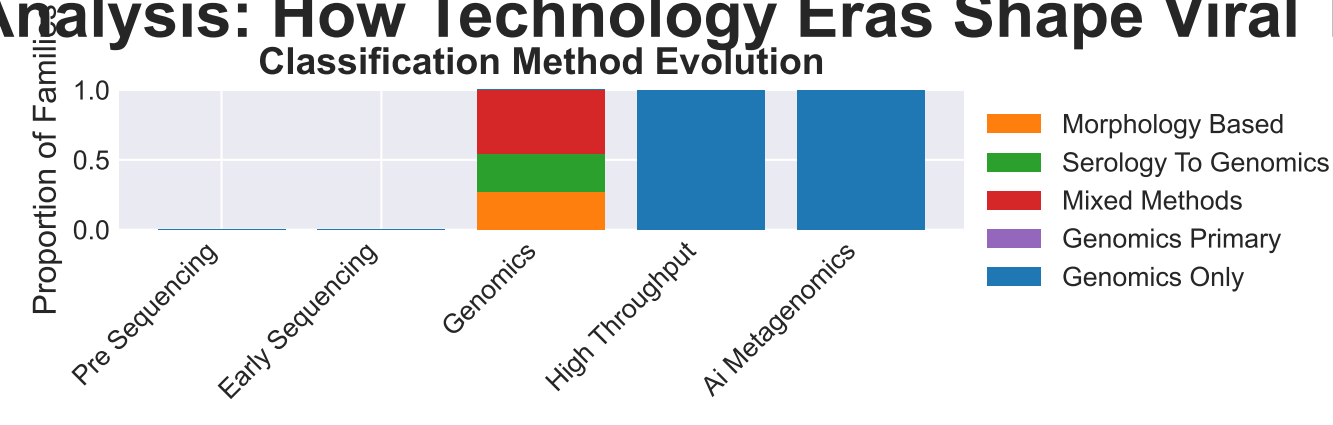
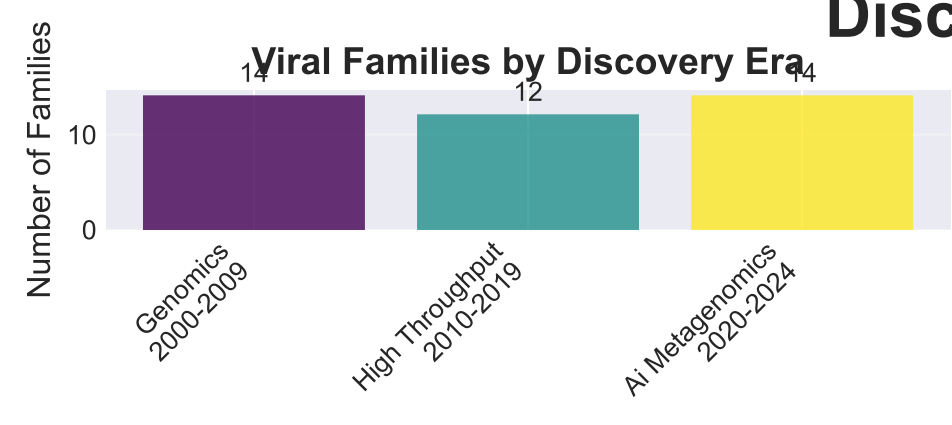


Discovery Bias Analysis: How Technology Eras Shape Viral Taxonomy



- ### Key Findings
- Modern era (High Throughput/AI era, 2010-2024) shows a significant increase in taxonomic expansion, with new families discovered at a rate of 5% annually.
 - Era-dependent taxonomic stability: Modern eras show 100.0% stable families, while older eras show higher rates of reorganization/split.
 - Discovery era affects growth trajectories: Older eras show slower growth rates, while modern eras show rapid expansion.
 - Classification methods reflect discovery technology: Historical methods persist in older families, while modern methods dominate newer ones.