Stanford

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Publications

PUBLICATIONS

- **Inverse FoldDir: Structure-conditioned Protein Sequence Design by Dirichlet Flow Matching.** *bioRxiv : the preprint server for biology*
  Tartici, A., Stojkovic, M., Tian, A., Jewett, M. C., Altman, R. B., Wittmann, B. J.
  2026