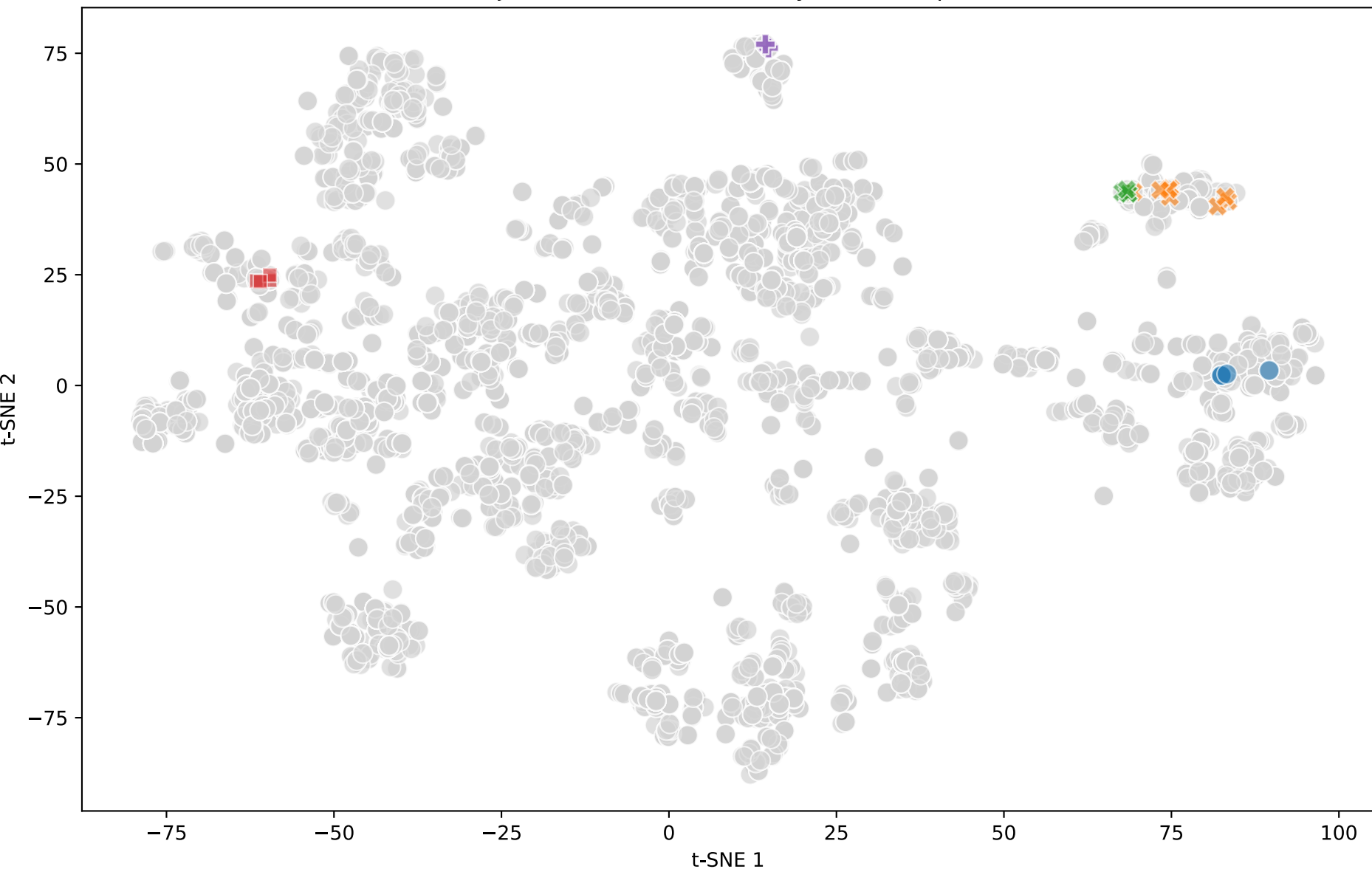


t-SNE Projection of Most Chemically Similar Superfamilies



Superfamily

- GI Alpha 1, domain 2-like
- × Lyase 2-enoyl-coa Hydratase, Chain A, domain 2
- × Single alpha-helices involved in coiled-coils or other helix-helix interfaces
- Fatty acid synthase; domain 2
- + Nitrogenase Molybdenum-iron Protein, subunit B; domain 4

Cluster

- 3
- × 11
- 2
- + 8