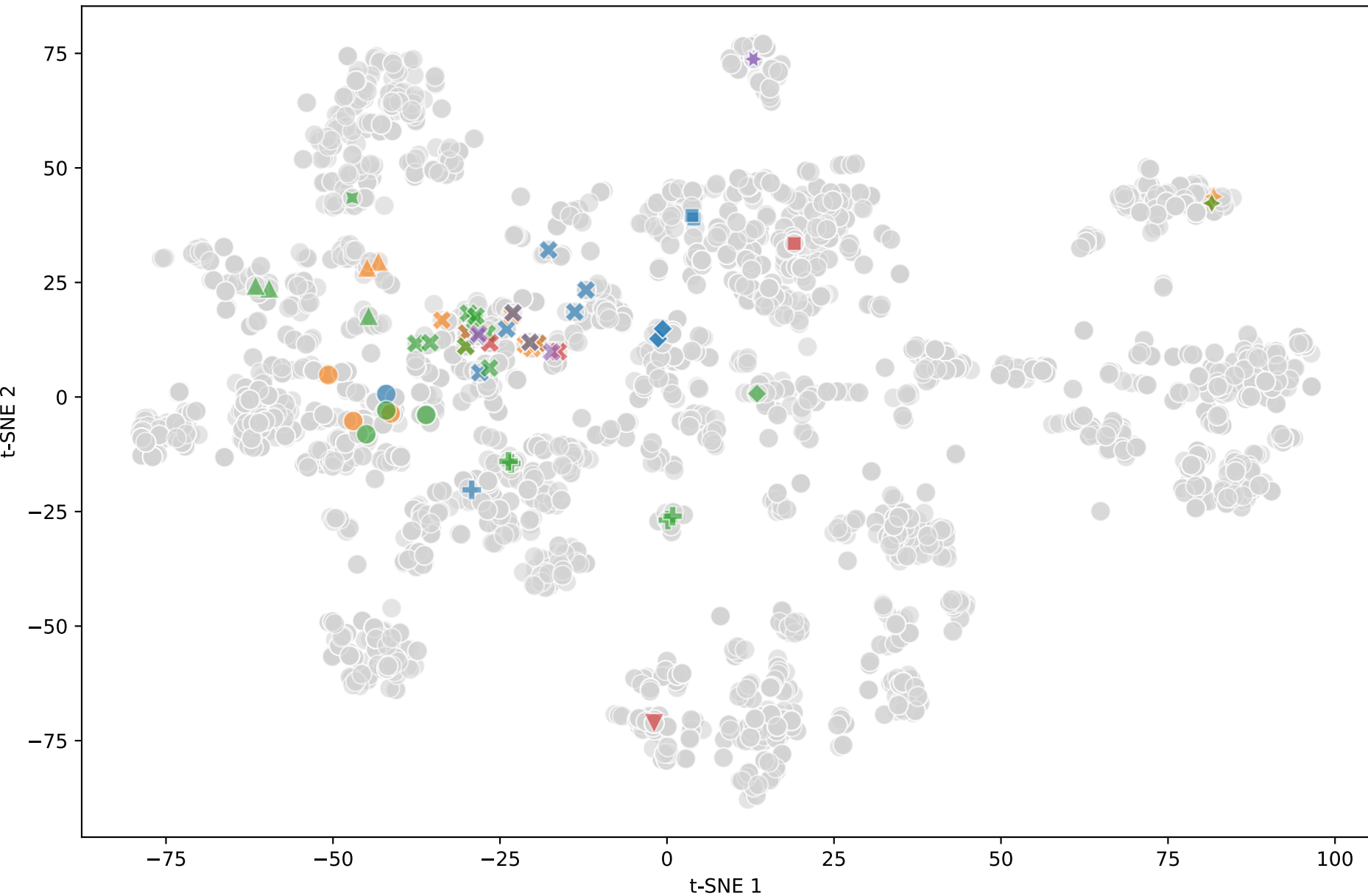


t-SNE Projection of Most Chemically Dissimilar Superfamilies



Superfamily

- | | |
|--|--|
| ● Copper amine oxidase, catalytic domain | ● main proteinase (3clpro) structure, domain 3 |
| ● Ferritin | ● Cytochrome c oxidase-like, subunit I domain |
| ● Ribonucleotide Reductase, subunit A | |

Cluster

- | | | |
|------|------|-----|
| ● 5 | ◆ 12 | ✧ 9 |
| ✕ 13 | ◆ 11 | ▼ 4 |
| ■ 1 | ▲ 2 | ★ 8 |
| ✚ 0 | | |