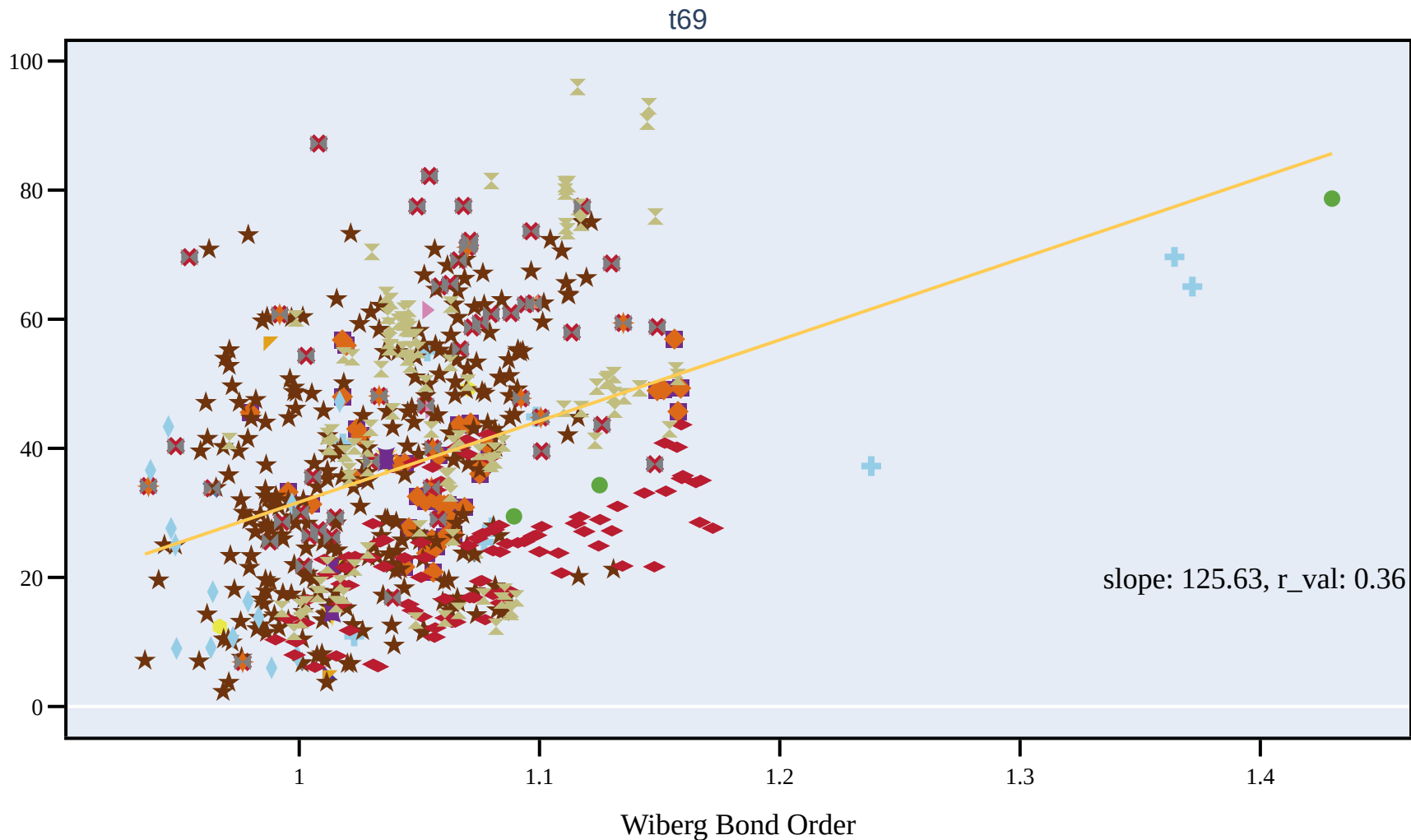


Torsion barrier in KJ/mol



- | | |
|---|---|
| ● Fragment Stability Benchmark.oeb | ■ Fragment Stability Benchmark.oeb |
| ◆ Fragment Stability Benchmark.oeb | + OpenFF Fragmenter Validation 1.0.oeb |
| ✕ OpenFF Full TorsionDrive Benchmark 1.oeb | ▲ OpenFF Gen 2 Torsion Set 1 Roche.oeb |
| ▼ OpenFF Gen 2 Torsion Set 1 Roche 2.oeb | ▶ OpenFF Gen 2 Torsion Set 2 Coverage 2.oeb |
| ▵ OpenFF Gen 2 Torsion Set 3 Pfizer Discrepancy.oeb | ▴ OpenFF Gen 2 Torsion Set 3 Pfizer Discrepancy 2.oeb |
| ▾ OpenFF Gen 2 Torsion Set 4 eMolecules Discrepancy.oeb | ▹ OpenFF Gen 2 Torsion Set 4 eMolecules Discrepancy 2.oeb |
| ● OpenFF Gen 2 Torsion Set 5 Bayer 2.oeb | ★ OpenFF Group1 Torsions.oeb |
| ▼ OpenFF Primary Benchmark 1 Torsion Set.oeb | ✧ OpenFF Primary Benchmark 1 Torsion Set.oeb |
| ◆ OpenFF Primary TorsionDrive Benchmark 1.oeb | ◇ OpenFF Rowley Biaryl v1.0.oeb |
| ◆ OpenFF Substituted Phenyl Set 1.oeb | ⋈ OpenFF-benchmark-ligand-fragments-v1.0.oeb |
| ⋈ Pfizer Discrepancy Torsion Dataset 1.oeb | ● SMIRNOFF Coverage Torsion Set 1.oeb |