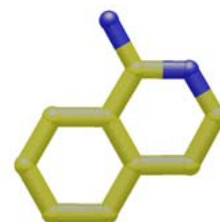
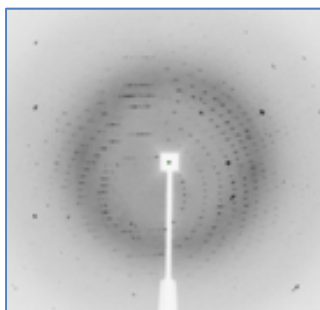
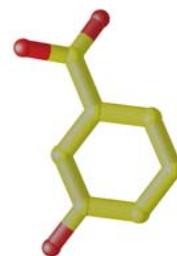


Fragment Libraries

Use the library specially designed for efficient screening by protein crystallography

- ✓ single core compounds
- ✓ compounds that explicitly combine shape and electrostatic diversity
- ✓ fragment classes cross-checked with known fragment binders and substructures from lead-like molecules
- ✓ physical-chemical attributes optimized for detection in protein crystals
- ✓ available compounds
- ✓ flexibly designed to match your experimental needs

Complete your initial screen in a single synchrotron shift



Also available Focused scaffold libraries designed for high concentration chemical screening

Build your leads with fragments

