**Table S2.** Data collection and refinement statistics for caspase‑9 and NleF co-crystallisation.

|  |  |
| --- | --- |
|  | Caspase-9-NleF  |
| **Data collection and Processing**No. of crystals used Wavelength [Å] Space group Unit cell parameters a, b, c [Å] Matthews coefficient [Å³/Da] Solvent content [%]Diffraction dataResolution [Å] Unique reflections R(I)sym [%] Completeness [%] Redundancy I/(I)**Refinement**Resolution range [Å] Reflections used in refinement (work/free) Final R values for all reflections (work/free) [%]Protein atomsRMSDsBonds [Å] Angles [°]Ramachandran plot Residues in most favoured regions [%] Residues in additional allowed regions [%] Residues in generously allowed regions [%] Residues in disallowed regions [%]**Mean B factor [Å2]** Protein | 11.0015P212121100.7, 209.91, 317.2390.0, 90.0, 90.03.362.73.49 (3.77-3.49)84,797 (17,178)12.6 (58.1)98.8 (98.3)3.3 (3.4)8.3 (2.0)174.08 – 3.4983 041 / 1 68922.5 / 26.025,5000.0091.0686.512.60.20.786.4 |