

Table 1. Data collection and refinement statistics of Fgfr1 minimal kinase with PLX052

Data collection	
Resolution, Å	50.0-2.4
Observations, n	247,572
Completeness, %*	88.05 (98.07)
Redundancy	3.5
R_{sym} , %*	8.1 (31.7)

Refinement	
Resolution, Å	15.0-3.3
Reflections, n	8,377
R value, %†	22.5 (27.7)
Bonds, Å	0.008
Angles, °	1.6

*Value in parentheses is for the highest-resolution shell.

† Value in parentheses is the free R value determined from 5% of the data.

