	M_2 receptor:Nb9-8 complex
Data collection[a]	
Number of crystals	17
Space group	$P2_12_12_1$
Unit cell dimensions	
a, b, c (Å)	62.9, 78.1, 163.5
α, β, γ (°)	90, 90, 90
Resolution (Å)	33 – 3.5 (3.6 – 3.5)
R_{merge} (%)	18.8 (74.4)
<I/σI>	5.8 (1.4)
$CC_{1/2}$ (%)	99.1 (35.0)
Completeness (%)	95.9 (83.1)
Redundancy	4.8
Refinement	
Number of reflections	10237
R_{work}/R_{free} (%)	24.9 / 29.8
No. atoms	
Protein	3020
Ligand(s)	14
Average B factors (Å^2)	
Receptor	109.7
Nb689	137.4
Iperoxo	105.5
LY2119620	-
RMS deviation from ideality	
Bond length (Å)	0.004
Bond angles (°)	0.86
Ramachandran statistics[b]	
Favored (%)	97.0
Allowed (%)	3.0
Outliers (%)	0

FIG. 6