

# Molecular Origami of HOBr

given information

ElementNames	[ (O) (Br) (H) ]	
dotted	F	
distance	95.327	O <sup>1</sup> -H <sup>1</sup>
distance	184.999	O <sup>1</sup> -Br <sup>1</sup>
angle	109.998	H <sup>1</sup> -O <sup>1</sup> -Br <sup>1</sup>
	235.3	H <sup>1</sup> -Br <sup>1</sup>
dopage	T	
AutoAlign	F	
showboth	F	

structure type: XAB

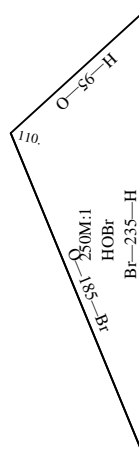
# Molecular Origami of HOBr

!O1  
Br1  
H1  
HOBr

bent

scale 250,000,000 : 1  
units: pm  
offsetx 1.1 offsety 1.01

View -1



Current: (centerx 5.40) (centery 6.01) (scale 250)

%%BoundingBox: 329 339 399 526 actual: 339 349 389 516

center: 364 433

actual size: 50 167

Better: (centerx 4.60) (centery 5.50) (scale 250)

%%BoundingBox: 350 375 420 562

actual: 360 385 410 552

center: 385 469

actual size: 50 167