

Electronic supporting material for

## The Solid-State Structures and Ligand Cavity Evaluation of Lanthanide(III) Complexes of DOTA Analogue with a (Dibenzylamino)methylphosphinate Pendant Arm

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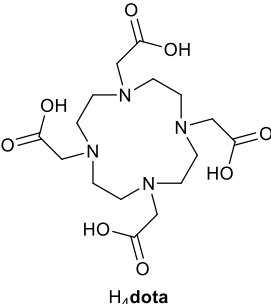
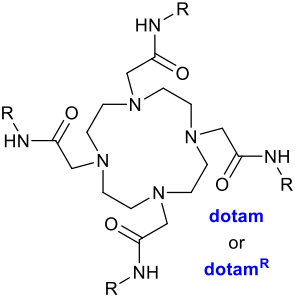
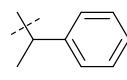
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Table S3. Selected geometric parameters found in the crystal structures of lanthanide(III)– $\text{H}_4\text{do3ap}^{\text{DBAm}}$  and related complexes.

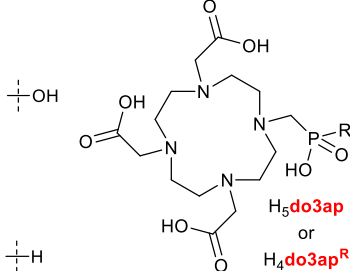
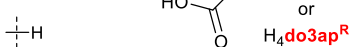
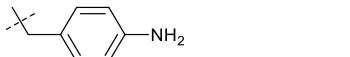
Crystal structures refinement details.

Table S4. Experimental crystallographic data for structures of lanthanide(III)– $\text{H}_4\text{do3ap}^{\text{DBAm}}$  complexes.

Table S1. Crystal structures from CCDC taken for comparison with present complexes.

Ligand type	Ligand structure	Ln	Isomer	CCDC code	Reference	
H <sub>4</sub> dota	 <p style="text-align: center;">H<sub>4</sub>dota</p>	La	2x TSA <sup>a</sup>	nonjuv	1	
H <sub>4</sub> dota		Ce	TSA	luqbii	2	
H <sub>4</sub> dota		Gd	TSA'	ogizou	3	
H <sub>4</sub> dota		Tm	TSA'	luqcef	2	
H <sub>4</sub> dota		Sc	2x TSA' <sup>a</sup>	jogzem	4	
H <sub>4</sub> dota		Pr	SA	luqboo	2	
H <sub>4</sub> dota		Nd	SA	luqbuu	2	
H <sub>4</sub> dota		Sm	SA	ogizag	3	
H <sub>4</sub> dota		Eu	SA	cexkul	5	
H <sub>4</sub> dota		Eu	SA	cexkul01	6	
H <sub>4</sub> dota		Eu	SA	coknas	7	
H <sub>4</sub> dota		Eu	SA	ogizek	3	
H <sub>4</sub> dota		Gd	SA	jopjih	8	
H <sub>4</sub> dota		Gd	SA	jopjih01	9	
H <sub>4</sub> dota		Gd	SA	cokmul	7	
H <sub>4</sub> dota		Gd	SA	ogizio	3	
H <sub>4</sub> dota		Dy	SA	luqcab	2	
H <sub>4</sub> dota		Dy	SA	unefid	10	
H <sub>4</sub> dota		Ho	SA	godkoz	6	
H <sub>4</sub> dota		Lu	SA	nojyiu	11	
H <sub>4</sub> dota		Y	SA	latkog	9	
H <sub>4</sub> dota		Y	SA	latkog01	12	
H <sub>4</sub> dota		Sc	3x SA' <sup>b</sup>	luqcij	2	
dotam		 <p style="text-align: center;">dotam or dotam<sup>R</sup></p>	Pr	TSA	kejbeh	13
dotam			Nd	SA	veckew	22
dotam			Eu	TSA	zacxac <sup>c</sup>	17
dotam	Eu		SA	veckia	22	
dotam	Eu		SA	veckia01	22	
dotam	Eu		SA	veckia02	22	
dotam	Gd		SA	ehovay	30	
dotam	Gd		SA	kehzut	13	
dotam	Gd		2x SA <sup>a</sup>	kejbad	13	
dotam	Gd		SA	sircay	33	
dotam	Gd		SA	sircec	33	
dotam	Gd		SA	sirdon	33	
dotam	Tb		SA	bosjea	36	
dotam	Tb		SA	tusput	37	
dotam	Yb		SA	xohvev	40	
dotam <sup>R</sup>			Ce	2x SA <sup>a</sup>	ufiruw	19
dotam <sup>R</sup>			Pr	2x SA <sup>a</sup>	ufisad	19
dotam <sup>R</sup>		Pr	SA	eqozal	20	
dotam <sup>R</sup>		Nd	SA	eqozep	20	
dotam <sup>R</sup>		Sm	SA	eqozit	20	
dotam <sup>R</sup>		Eu	SA	tuqtoo	24	
dotam <sup>R</sup>		Eu	SA	tuqtoo01	25	
dotam <sup>R</sup>		Eu	SA	tuqtii	24	
dotam <sup>R</sup>		Eu	SA	tuqtii01	25	
dotam <sup>R</sup>		Eu	SA	eqozoz	20	
dotam <sup>R</sup>		Gd	2x SA <sup>a</sup>	ufirac	19	
dotam <sup>R</sup>		Gd	2x SA <sup>a</sup>	ufiseh	19	
dotam <sup>R</sup>		Gd	SA	eqozuf	20	
dotam <sup>R</sup>		Gd	SA	sircig	33	
dotam <sup>R</sup>		Gd	SA	sircom	33	
dotam <sup>R</sup>		Gd	SA	sircus	33	
dotam <sup>R</sup>		Tb	SA	ufireg	19	
dotam <sup>R</sup>		Dy	SA	equbat	20	
dotam <sup>R</sup>		Dy	SA	tuqtuu	24	
dotam <sup>R</sup>		Dy	SA	tuqtuu01	25	
dotam <sup>R</sup>		Er	2x SA <sup>a</sup>	ufisil	19	
dotam <sup>R</sup>		Er	SA	ufirik	19	
dotam <sup>R</sup>		Yb	SA	fibtit	41	

<b>dotam<sup>R</sup></b>		Yb	3x SA <sup>b</sup>	sirdaz	33
<b>dotam<sup>R</sup></b>		Yb	2x SA <sup>a</sup>	sirded	33
<b>dotam<sup>R</sup></b>		Yb	SA	sirdih	33
<b>dotam<sup>R</sup></b>		Lu	SA	ufiroq	19
<b>dotam<sup>R</sup></b>		Eu	SA	zacxac <sup>c</sup>	17
<b>dotam<sup>R</sup></b>		Nd	SA	banxiz	21
<b>dotam<sup>R</sup></b>		Gd	SA	ekezaw	31
<b>dotam<sup>R</sup></b>		Tb	SA	banxof	21
<b>dotam<sup>R</sup></b>		Dy	SA	banxev	21
<b>dotam<sup>R</sup></b>		Yb	SA	banxul	21
<b>dotam<sup>R</sup> (Et, H)</b>		Eu	SA	xiwpic	23
<b>dotam<sup>R</sup> (Et, Me)</b>		Eu	SA	fibfeb	27
<b>dotam<sup>R</sup> (Me, H)</b>		Eu	SA	mubgar	26
<b>dotam<sup>R</sup> (H, H)</b>		Eu	SA	zewsay	28
<b>dotam<sup>R</sup> (Bn, Me)</b>		Tb	SA	ruznuv	34
<b>dotam<sup>R</sup></b>		Eu	TSA	abofom	14
<b>dotam<sup>R</sup></b>		Lu	2x SA <sup>a</sup>	abofus	14
<b>dotam<sup>R</sup></b>		Eu	2x TSA <sup>a</sup>	pekdob	16
<b>dotam<sup>R</sup></b>		Eu	TSA	walvel	15
<b>dotam<sup>R</sup></b>		La	SA	kawmay	18
<b>dotam<sup>R</sup></b>		Eu	SA	abofig	14
<b>dotam<sup>R</sup></b>		Gd	SA	gegcie	29
<b>dotam<sup>R</sup></b>		Dy	SA	goybol	38
<b>dotam<sup>R</sup></b>		Gd	SA	pehfug	32
<b>dotam<sup>R</sup></b>		Gd	SA	pehgan	32
<b>dotam<sup>R</sup></b>		Tb	SA	efumat	35
<b>dotam<sup>R</sup></b>		Tm	SA	tavsiv	39
<b>dotam<sup>R</sup></b>		Lu	SA	iwuqex	42
<b>H<sub>8</sub>dotp</b>		Eu	2x TSA' <sup>a</sup>	axamap	45
<b>H<sub>8</sub>dotp</b>		Eu	2x TSA' <sup>a</sup>	axamet	45
<b>H<sub>8</sub>dotp</b>		Gd	2x TSA' <sup>a</sup>	eqiyim	46
<b>H<sub>4</sub>dotp<sup>R</sup></b>		La	2x TSA' <sup>a</sup>	ciqyoq	43
<b>H<sub>4</sub>dotp<sup>R</sup></b>		Ce	2x TSA' <sup>a</sup>	ciqyuw	43
<b>H<sub>4</sub>dotp<sup>R</sup></b>		Eu	TSA'	ruhmem	44
<b>H<sub>4</sub>dotp<sup>R</sup></b>		La	TSA	ruhmiq	44
<b>H<sub>4</sub>dotp<sup>R</sup></b>		Dy	TSA'	yeqded	48
<b>H<sub>4</sub>dotp<sup>R</sup></b>		Dy	TSA'	yeqded01	48
<b>H<sub>4</sub>dotp<sup>R</sup></b>		Yb	2x TSA' <sup>a</sup>	ruhmov	44
<b>H<sub>4</sub>dotp<sup>R</sup></b>		Y	TSA'	lijfal	49
<b>H<sub>4</sub>dotp<sup>R</sup></b>		Gd	2x TSA' <sup>a</sup>	sonduv	47
<b>H<sub>4</sub>dotp<sup>R</sup></b>			Gd	TSA'	sonfad
<b>trans-H<sub>6</sub>do2a2p</b>		Ce	TSA	babfob	50
<b>trans-H<sub>6</sub>do2a2p</b>		Nd	TSA	babcog	50
<b>trans-H<sub>6</sub>do2a2p</b>		Sm	2x TSA <sup>a</sup>	babhap	50
<b>trans-H<sub>6</sub>do2a2p</b>		Sm	TSA'	babgui	50
<b>trans-H<sub>6</sub>do2a2p</b>		Eu	2x TSA' <sup>a</sup>	babgiw	50
<b>trans-H<sub>6</sub>do2a2p</b>		Tb	TSA'	babhet	50
<b>trans-H<sub>6</sub>do2a2p</b>		Dy	TSA'	babgao	50
<b>trans-H<sub>6</sub>do2a2p</b>		Er	TSA'	babges	50
<b>trans-H<sub>6</sub>do2a2p</b>		Yb	TSA'	babhix	50

H <sub>5</sub> do3ap		Nd	2x TSA <sup>a</sup>	lanqat	51
H <sub>5</sub> do3ap		Tb	TSA	larlia	53
H <sub>5</sub> do3ap		Dy	TSA'	larkof	53
H <sub>5</sub> do3ap		Er	TSA'	larkul	53
H <sub>5</sub> do3ap		Lu	TSA'	larlas	53
H <sub>5</sub> do3ap		Lu	TSA'	larlew	53
H <sub>5</sub> do3ap		Y	TSA'	larlog	53
H <sub>4</sub> do3ap <sup>R</sup>		Eu	TSA	ofayec	53
H <sub>4</sub> do3ap <sup>R</sup>		Y	TSA + SA + TSA'	kejroh <sup>d</sup>	54

<sup>a</sup>Two complex molecules present in the structurally independent part. <sup>b</sup>Three complex molecules present in the structurally independent part. <sup>c</sup>Disorder in macrocyclic part was found but positions of the corresponding carbon atoms only were refined. <sup>d</sup>Two complex molecules were present in the structurally independent part, one of them with disordered macrocyclic part. Positions of corresponding carbon and nitrogen atoms were refined.

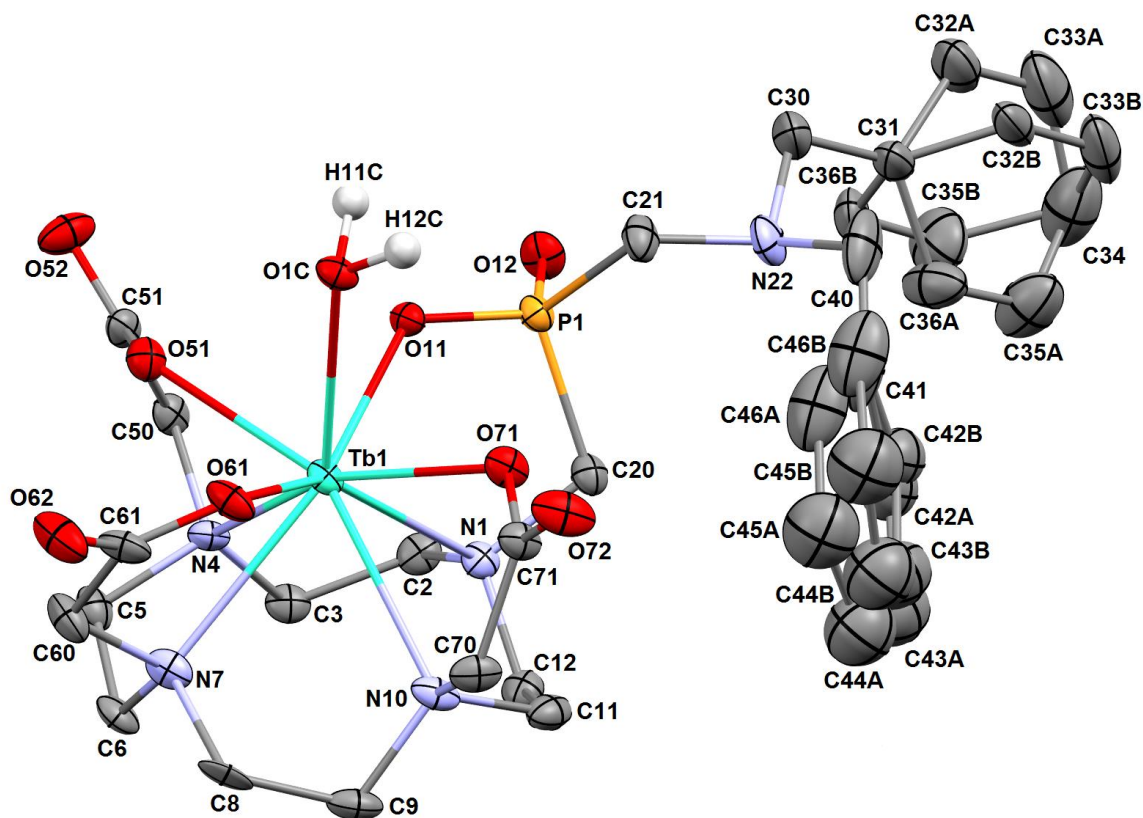


Figure S1. Molecular structure of the  $[\text{Tb}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})]^-$  anion found in the crystal structure of  $(\text{NH}_4)[\text{Tb}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$ . Disordered phenyl rings are shown. All carbon-bound hydrogen atoms are omitted for clarity.

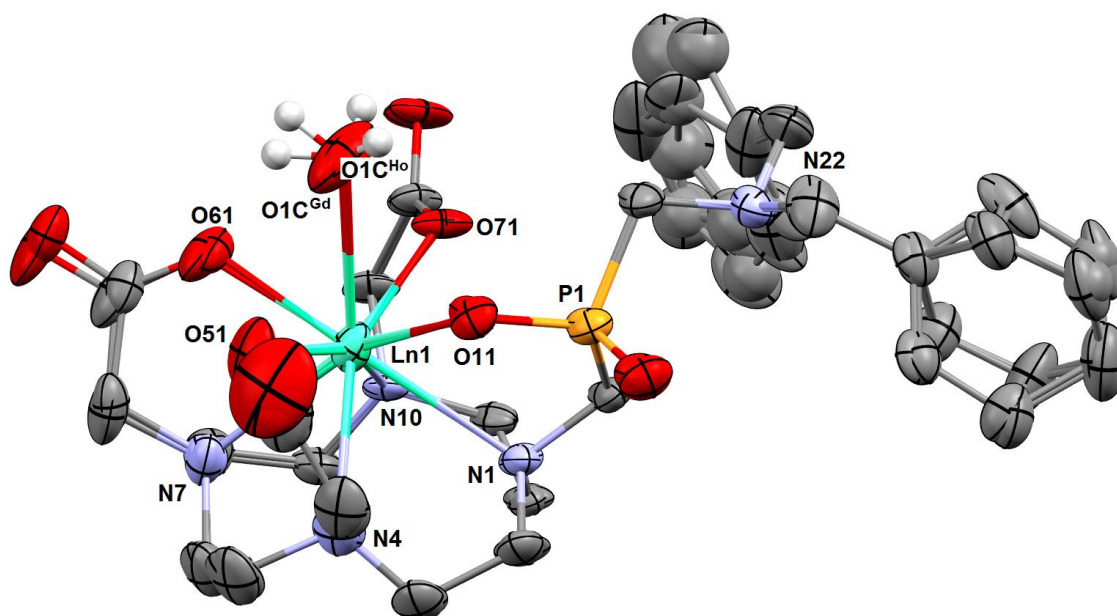


Figure S2. An overlay of the molecular structures of  $[\text{Gd}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})]^-$  and  $[\text{Ho}“(\text{H}_2\text{O})_{0.5}”(\text{do3ap}^{\text{DBAm}})]^-$  anions found in the crystal structures of  $\text{Ca}_{0.5}[\text{Gd}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})] \cdot 6\text{H}_2\text{O}$  and  $\text{Ca}_{0.5}[\text{Ho}(\text{do3ap}^{\text{DBAm}})] \cdot 6\text{H}_2\text{O}$ , showing the half-occupied water molecule apically pseudo-coordinated to Ho(III). For calculation of the overlay, lanthanide Ln1, phosphorus P1, and nitrogen N7 atom pairs from both structures were merged. Disordered phenyl rings are shown. Carbon-bound hydrogen atoms are omitted for clarity.

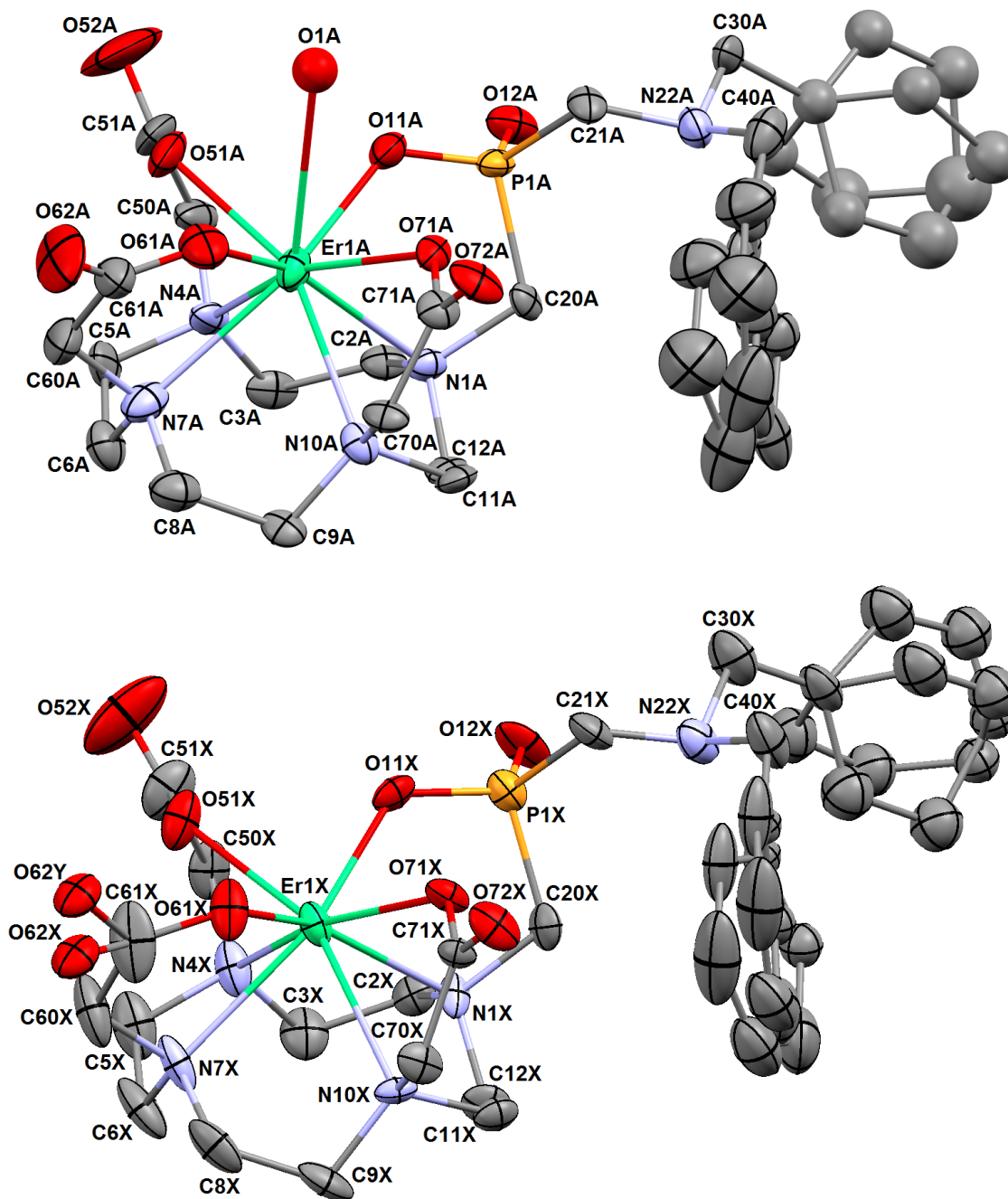


Figure S3. Molecular structure of the independent  $[\text{Er}(\text{H}_2\text{O})_{0.5}(\text{do3ap}^{\text{DBAm}})]^-$  (top) and  $[\text{Er}(\text{do3ap}^{\text{DBAm}})]^-$  (bottom) units found in the crystal structure of  $\text{Na}[\text{Er}(\text{do3ap}^{\text{DBAm}})] \cdot 4.25\text{H}_2\text{O}$ . Disordered phenyl rings are shown. Hydrogen atoms and labels of disordered phenyl rings are omitted for clarity.

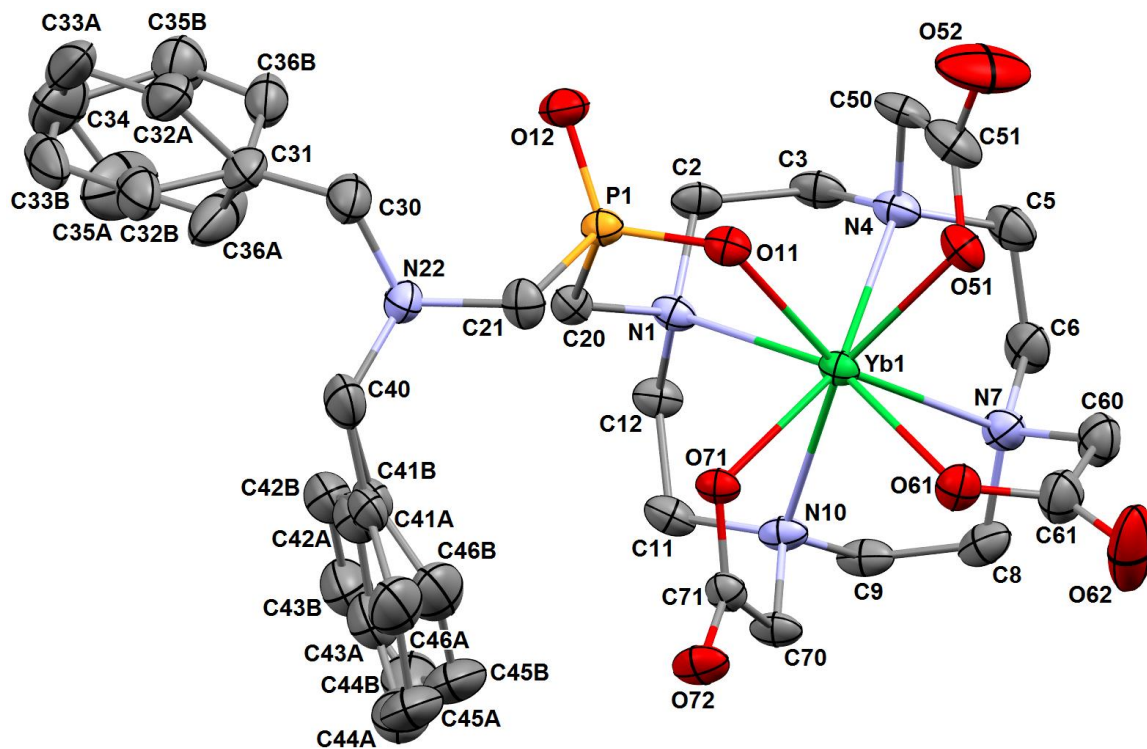


Figure S4. Molecular structure of the  $[\text{Yb}(\text{do3ap}^{\text{DBAm}})]^-$  anion found in the crystal structure of  $\text{Na}[\text{Yb}(\text{do3ap}^{\text{DBAm}})] \cdot 6\text{H}_2\text{O}$ . Disordered phenyl rings are shown. Hydrogen atoms are omitted for clarity.

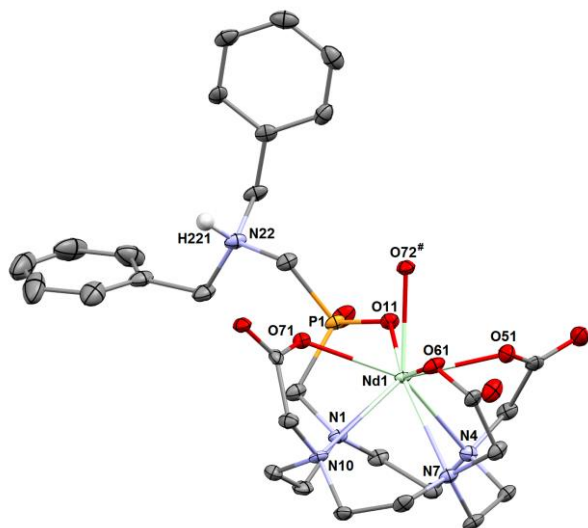
Table S2. Selected geometric parameters found in the crystal structures of lanthanide(III)–H<sub>4</sub>do3ap<sup>DBAm</sup> (Ln–H<sub>4</sub>L1) complexes.

Compound	[Ce(HL1)] ·0.5NH <sub>4</sub> Cl ·6.5H <sub>2</sub> O	[Pr(HL1)] ·0.5NH <sub>4</sub> Cl ·6.5H <sub>2</sub> O	[Nd(HL1)] ·3H <sub>2</sub> O	(NH <sub>4</sub> )[Nd(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O	(NH <sub>4</sub> )[Sm(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O	(NH <sub>4</sub> )[Eu(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O	(NH <sub>4</sub> )[Gd(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O (ref. <sup>[55]</sup> )	Ca <sub>0.5</sub> [Gd(H <sub>2</sub> O) (L1)]·6H <sub>2</sub> O	(NH <sub>4</sub> )[Tb(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O
	Distances / Å								
	Nitrogen plane								
N1···N4	2.961(3)	2.959(3)	2.952(5)	2.964(14)	2.942(8)	2.971(7)	2.966(6)	2.946(7)	2.953(7)
N4···N7	2.968(3)	2.956(3)	2.952(5)	2.975(16)	2.953(8)	2.944(7)	2.932(6)	2.953(8)	2.927(7)
N7···N10	2.953(3)	2.947(3)	2.938(5)	2.945(15)	2.897(9)	2.922(8)	2.918(7)	2.916(7)	2.899(8)
N10···N1	2.984(3)	2.984(3)	2.970(5)	3.014(14)	2.983(8)	2.968(7)	2.960(6)	2.954(7)	2.964(7)
N1···N7	4.185(3)	4.174(3)	4.166(5)	4.223(14)	4.207(9)	4.198(7)	4.189(6)	4.180(7)	4.176(7)
N4···N10	4.205(3)	4.202(3)	4.186(5)	4.190(15)	4.119(8)	4.148(7)	4.137(6)	4.141(7)	4.128(7)
	Oxygen plane								
O11···O51	3.289(3)	3.242(3)	3.223(4)	3.053(13)	2.997(7)	2.992(6)	2.983(5)	2.956(7)	2.959(6)
O51···O61	3.197(3)	3.187(3)	3.143(4)	3.279(15)	3.225(8)	3.213(7)	3.197(6)	3.115(8)	3.170(6)
O61···O71	3.655(2)	3.614(3)	3.559(4)	3.248(14)	3.185(8)	3.175(7)	3.159(6)	3.170(7)	3.120(6)
O71···O11	3.213(2)	3.201(3)	3.210(4)	3.285(12)	3.235(7)	3.227(6)	3.207(5)	3.132(6)	3.182(6)
O11···O61	4.723(2)	4.693(3)	4.653(4)	4.526(13)	4.476(7)	4.433(6)	4.414(5)	4.362(7)	4.377(6)
O51···O71	4.715(2)	4.666(3)	4.629(4)	4.568(14)	4.462(7)	4.480(6)	4.456(5)	4.385(7)	4.413(6)
	Angles / °								
	Nitrogen plane								
N10···N1···N4	90.05(8)	89.99(8)	90.0(1)	89.0(4)	88.1(2)	88.6(2)	88.5(2)	89.2(2)	88.5(2)
N1···N4···N7	89.79(8)	89.76(8)	89.8(1)	90.6(4)	91.1(2)	90.5(2)	90.5(2)	90.3(2)	90.5(2)
N4···N7···N10	90.51(8)	90.76(8)	90.6(1)	90.1(4)	89.5(2)	90.0(2)	90.0(2)	89.8(2)	90.2(2)
N7···N10···N1	89.63(8)	89.46(8)	89.7(1)	90.3(4)	91.3(2)	90.9(2)	90.9(2)	90.8(2)	90.8(2)
	Oxygen plane								
O71···O11···O51	92.95(6)	92.80(6)	92.0(1)	92.2(3)	91.4(2)	92.1(2)	92.0(1)	92.1(2)	91.8(2)
O11···O51···O61	93.46(6)	93.76(7)	93.9(1)	91.2(4)	91.9(2)	91.1(2)	91.1(1)	91.8(2)	91.1(2)
O51···O61···O71	86.67(6)	86.39(6)	87.1(1)	88.9(3)	88.2(2)	89.1(2)	89.0(1)	88.5(2)	89.1(2)
O61···O71···O11	86.63(6)	86.82(6)	86.7(1)	87.7(3)	88.4(2)	87.7(2)	87.8(1)	87.6(2)	88.0(2)
N <sub>4</sub> -plane vs. O <sub>4</sub> -plane	2.00(9)	2.3(1)	1.5(1)	1.0(4)	1.2(3)	1.0(2)	0.9(2)	1.1(3)	0.9(2)

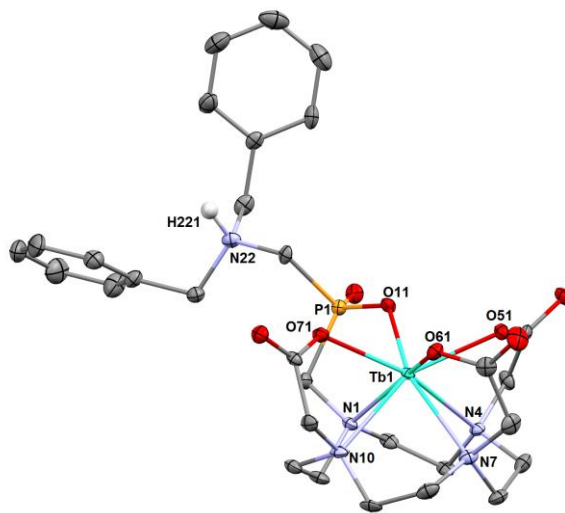


Table S2. Selected geometric parameters found in the crystal structures of lanthanide(III)–H<sub>4</sub>do3ap<sup>DBAm</sup> (Ln–H<sub>4</sub>L1) complexes – continuation.

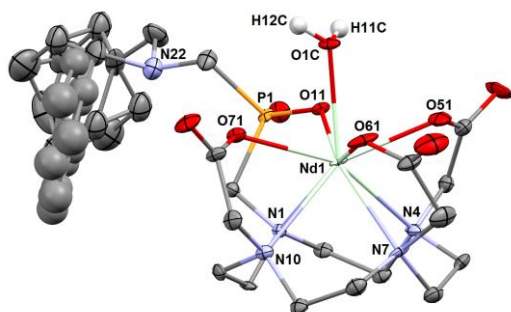
Compound	[Tb(HL1)] ·NH <sub>4</sub> Cl·H <sub>2</sub> O	Na[Dy(H <sub>2</sub> O)(L1)] ·4H <sub>2</sub> O (ref. <sup>[55]</sup> )	Ca <sub>0.5</sub> [Ho(L1)] ·6H <sub>2</sub> O	Na[Er(L1)] ·4.25H <sub>2</sub> O	(NH <sub>4</sub> )[Er(L1)] ·4H <sub>2</sub> O	Na[Yb(L1)] ·6H <sub>2</sub> O	(NH <sub>4</sub> )[Sc(L1)] ·4H <sub>2</sub> O		
Distances / Å									
Nitrogen plane									
N1···N4	2.913(5)	2.932(4)	2.935(4)	2.926(15)	2.934(18)	2.935(19)	2.957(4)	2.920(6)	2.898(2)
N4···N7	2.924(6)	2.926(4)	2.925(4)	2.946(16)	2.885(23)	2.947(19)	2.920(4)	2.913(7)	2.862(2)
N7···N10	2.939(7)	2.909(4)	2.911(4)	2.891(15)	2.884(21)	2.903(19)	2.928(4)	2.896(6)	2.869(2)
N10···N1	2.983(7)	2.968(4)	2.951(4)	2.933(14)	2.902(19)	2.964(18)	2.950(4)	2.925(6)	2.893(2)
N1···N7	4.206(6)	4.172(4)	4.168(4)	4.175(16)	4.151(22)	4.174(18)	4.174(4)	4.157(6)	4.096(2)
N4···N10	4.107(7)	4.124(4)	4.121(4)	4.096(14)	4.053(19)	4.134(19)	4.137(4)	4.083(6)	4.050(2)
Oxygen plane									
O11···O51	2.969(5)	2.940(3)	2.955(3)	2.862(14)	2.897(18)	2.798(17)	2.829(3)	2.794(5)	2.664(2)
O51···O61	2.890(5)	3.061(4)	3.099(3)	2.998(15)	2.910(23)	2.899(17)	2.847(3)	2.842(5)	2.670(2)
O61···O71	2.941(4)	3.213(4)	3.109(3)	2.977(14)	3.013(17)	2.869(16)	2.855(3)	2.853(5)	2.656(2)
O71···O11	2.969(5)	3.094(3)	3.084(3)	2.992(13)	2.991(16)	2.897(14)	2.891(3)	2.854(5)	2.683(2)
O11···O61	4.066(5)	4.349(3)	4.316(3)	4.151(14)	4.160(19)	3.997(16)	3.988(3)	3.961(5)	3.719(2)
O51···O71	4.248(5)	4.347(3)	4.341(3)	4.209(13)	4.189(17)	4.103(15)	4.086(3)	4.053(5)	3.825(2)
Angles / °									
Nitrogen plane									
N10···N1···N4	88.3(2)	88.7(1)	88.9(1)	88.7(4)	88.0(7)	89.0(5)	88.9(1)	88.6(2)	88.76(5)
N1···N4···N7	92.2(2)	90.8(1)	90.7(1)	90.6(4)	91.0(5)	90.4(6)	90.5(1)	90.9(2)	90.66(5)
N4···N7···N10	89.0(2)	90.0(1)	89.8(1)	89.1(4)	89.3(6)	89.9(5)	90.1(1)	89.3(2)	89.93(5)
N7···N10···N1	90.5(2)	90.5(1)	90.6(1)	91.6(4)	91.7(6)	90.7(5)	90.5(1)	91.2(2)	90.62(5)
Oxygen plane									
O71···O11···O51	91.4(1)	92.15(9)	91.88(9)	91.9(4)	90.7(5)	92.2(5)	91.17(8)	91.7(2)	91.35(5)
O11···O51···O61	87.9(2)	92.88(9)	90.91(9)	90.2(4)	91.5(5)	89.1(4)	89.28(8)	89.3(2)	88.43(5)
O51···O61···O71	93.5(1)	87.69(9)	88.75(9)	89.6(4)	90.0(5)	90.7(5)	91.55(8)	90.8(2)	91.83(5)
O61···O71···O11	87.0(1)	87.17(9)	88.37(9)	88.1(4)	87.7(5)	87.8(4)	87.92(8)	87.9(1)	88.32(5)
N <sub>4</sub> -plane···O <sub>4</sub> -plane	2.9(2)	0.9(1)	1.0(1)	1.0(5)	1.0(6)	1.5(7)	1.8(1)	1.2(2)	1.5(7)



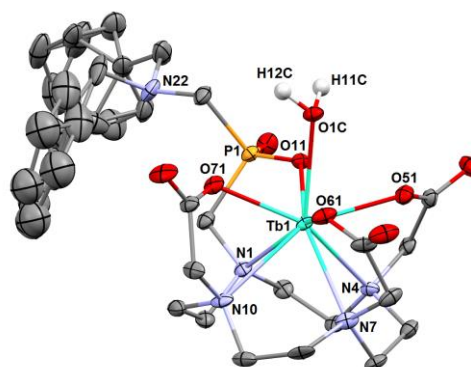
$[\text{Nd}(\text{O}^\#)(\text{Hdo3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$



$[\text{Tb}(\text{Hdo3ap}^{\text{DBAm}})] \cdot \text{NH}_4\text{Cl} \cdot \text{H}_2\text{O}$

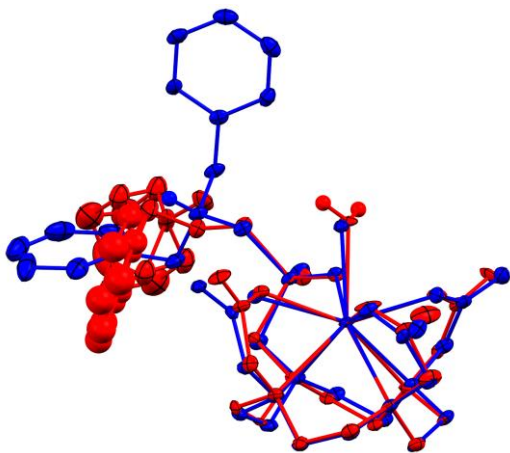


$(\text{NH}_4)[\text{Nd}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$

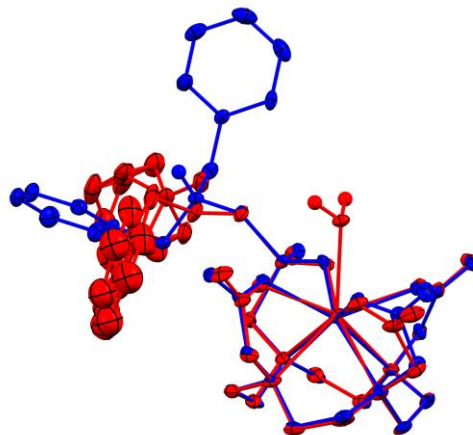


$(\text{NH}_4)[\text{Tb}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$

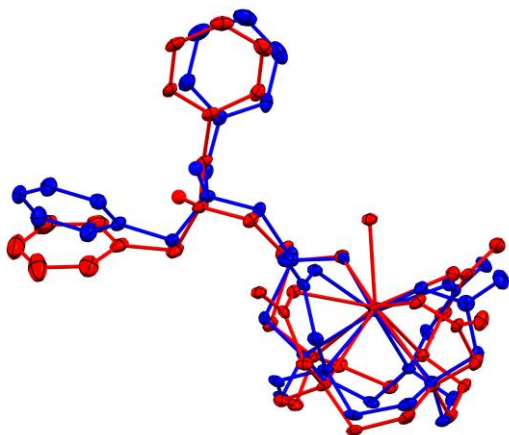
Figure S5. Molecular structures of Nd(III) and Tb(III) complex species in different protonation states found in the crystal structures of  $[\text{Nd}(\text{O}^\#)(\text{Hdo3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$  ( $\text{O}^\#$  = apically coordinated oxygen atom belonging to the neighbouring unit in the dimeric species),  $[\text{Tb}(\text{Hdo3ap}^{\text{DBAm}})] \cdot \text{NH}_4\text{Cl} \cdot \text{H}_2\text{O}$  and  $(\text{NH}_4)[\text{Ln}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$  ( $\text{Ln} = \text{Nd}, \text{Tb}$ ). Carbon-bound hydrogen atoms are omitted and only selected atoms are labelled for clarity. Only a half of the dimer present in  $[\text{Nd}(\text{O}^\#)(\text{Hdo3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$  with the apically coordinated oxygen atom ( $\text{O72}^\#$ ) from the neighbouring unit is shown. In structures of the deprotonated species, the disorder of the benzyl groups is shown.



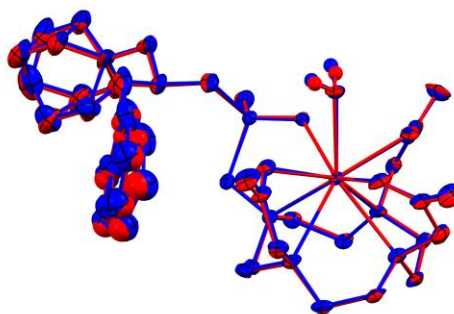
Overlay of  $[\text{Nd}(\text{O}^\#)(\text{Hdo3ap}^{\text{DBAm}})]$  (blue) and  $[\text{Nd}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})]^-$  (red).



Overlay of  $[\text{Tb}(\text{Hdo3ap}^{\text{DBAm}})]$  (blue) and  $[\text{Tb}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})]^-$  (red).



Overlay of  $[\text{Nd}(\text{Hdo3ap}^{\text{DBAm}})]$  (red) and  $[\text{Tb}(\text{Hdo3ap}^{\text{DBAm}})]$  (blue).



Overlay of  $[\text{Nd}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})]^-$  (red) and  $[\text{Tb}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})]^-$  (blue).

Figure S6. Overlays of molecular structures of the Nd(III) and Tb(III) complex species in the different protonation states as found in the crystal structures of  $[\text{Nd}(\text{O}^\#)(\text{Hdo3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$  ( $\text{O}^\#$  = apically coordinated oxygen atom belonging to the neighbouring unit in the dimeric species),  $[\text{Tb}(\text{Hdo3ap}^{\text{DBAm}})] \cdot \text{NH}_4\text{Cl} \cdot \text{H}_2\text{O}$  and  $(\text{NH}_4)[\text{Ln}(\text{H}_2\text{O})(\text{do3ap}^{\text{DBAm}})] \cdot 3\text{H}_2\text{O}$  ( $\text{Ln} = \text{Nd}, \text{Tb}$ ). Carbon-bound hydrogen atoms and atom labels are omitted for clarity. To overlay the molecules, distances between central Ln(III) ions, phosphorus atoms P1, and nitrogen atoms N7 were minimalized.

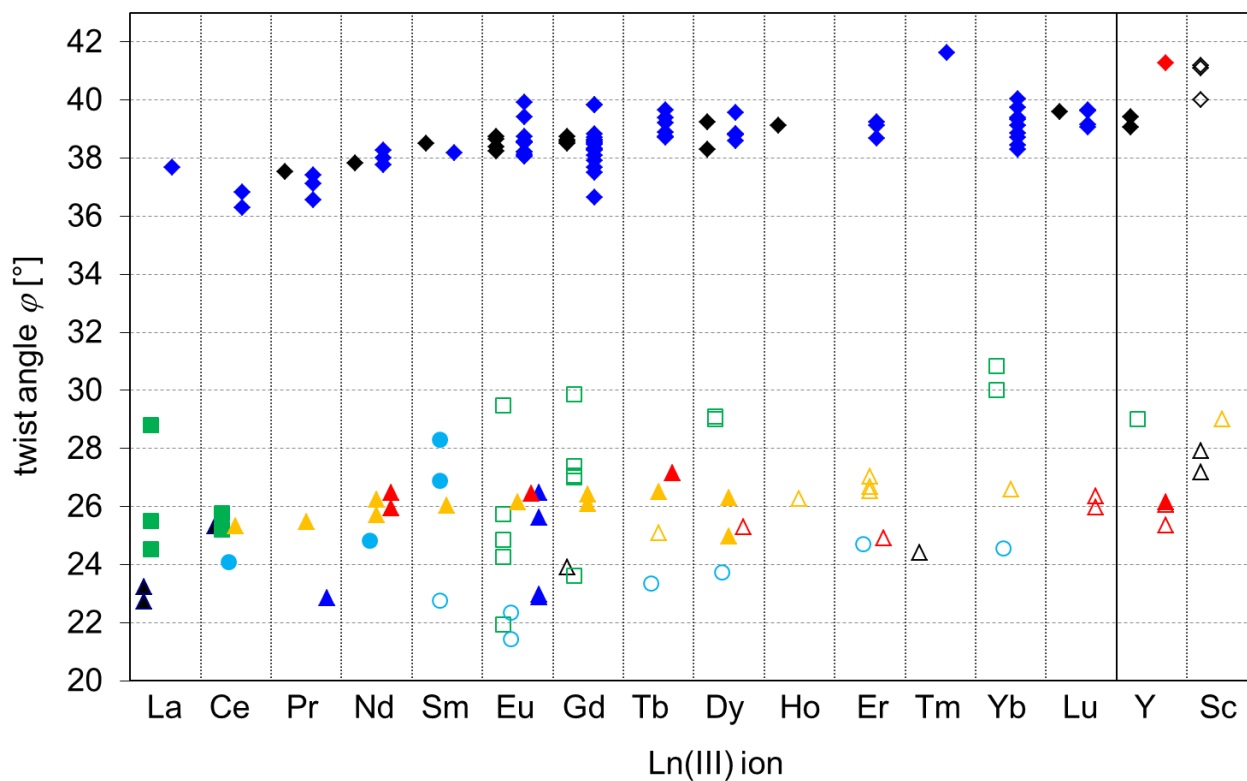


Figure S7. Dependence of mean twist angle  $\varphi$  of the pendant arms in Ln(III)/Y(III)/Sc(III) complexes of selected ligands on metal ion. Ligand colour codes:  $\text{H}_4\text{dota}$ ,  $\text{dotam}^{(R)}$ ,  $\text{H}_8\text{dotp}/\text{H}_4\text{dotp}^R$ ,  $\text{trans-H}_6\text{do2a2p}$ ,  $\text{H}_5\text{do3ap}/\text{H}_4\text{do3ap}^R$ ,  $\text{H}_4\text{do3ap}^{\text{DBAm}}$ . Isomer coding: diamonds are used for the SA/SA’ isomers ( $\text{H}_4\text{dota}$ ,  $\text{dotam}^{(R)}$  and  $\text{H}_4\text{do3ap}^R$ ), and triangles ( $\text{H}_4\text{dota}$ ,  $\text{dotam}^{(R)}$ ,  $\text{H}_5\text{do3ap}/\text{H}_4\text{do3ap}^R$  and  $\text{H}_4\text{do3ap}^{\text{DBAm}}$ ), squares ( $\text{H}_8\text{dotp}/\text{H}_4\text{dotp}^R$ ) and circles ( $\text{trans-H}_6\text{do2a2p}$ ) for the TSA/TSA’ species. Full symbols mark nonacoordinated “hydrated” species (*i.e.* TSA/SA), and open symbols stay for octacoordinated “anhydrous” species (*i.e.* TSA’/SA’).

Table S3. Selected geometric parameters found in the crystal structures of lanthanide(III)–H<sub>4</sub>do3ap<sup>DBAm</sup> and related complexes.

CCDC code	ligand type	ion	configuration	distances / Å				opening angles / °		twist angles / °			
				M–O <sub>w</sub>	QO <sub>4</sub> –QN <sub>4</sub>	M–QO <sub>4</sub>	M–QN <sub>4</sub>	O1–Ln–O3	O2–Ln–O4	N1–Q–O1 <sup>a</sup>	N2–Q–O2 <sup>a</sup>	N3–Q–O3 <sup>a</sup>	N4–Q–O4 <sup>a</sup>
nonjuv1 <sup>b</sup>	H <sub>4</sub> dota	La	TSA	2.569 <sup>c</sup>	2.537	0.702	1.835	147.9	146.6	24.7	20.1	24.6	21.5
nonjuv2 <sup>b</sup>	H <sub>4</sub> dota	La	TSA	2.538 <sup>c</sup>	2.539	0.731	1.810	146.2	145.2	24.3	21.5	22.9	24.2
luqbii	H <sub>4</sub> dota	Ce	TSA	2.598	2.520	0.755	1.765	145.0	143.5	24.9	25.9	25.7	24.9
ogizou	H <sub>4</sub> dota	Gd	TSA'	- <sup>c</sup>	2.546	1.070	1.476	126.6	125.5	25.0	22.9	25.0	22.9
luqcef	H <sub>4</sub> dota	Tm	TSA'	- <sup>c</sup>	2.530	1.064	1.466	124.0	124.7	24.7	24.6	24.0	24.3
jogzem1 <sup>b</sup>	H <sub>4</sub> dota	Sc	TSA'	- <sup>c</sup>	2.473	1.172	1.301	116.4	116.4	27.9	27.9	27.9	27.9
jogzem2 <sup>b</sup>	H <sub>4</sub> dota	Sc	TSA'	- <sup>c</sup>	2.525	1.209	1.316	122.3	122.3	27.2	27.2	27.2	27.2
luqboo	H <sub>4</sub> dota	Pr	SA	2.530	2.366	0.671	1.695	149.1	146.8	38.5	36.6	39.0	36.1
luqbuu	H <sub>4</sub> dota	Nd	SA	2.508	2.360	0.684	1.676	148.2	146.0	38.8	36.8	39.3	36.5
ogizag	H <sub>4</sub> dota	Sm	SA	2.404	2.325	0.642	1.684	147.8	149.5	38.4	40.0	35.4	40.2
cexkul	H <sub>4</sub> dota	Eu	SA	2.480	2.361	0.711	1.650	147.5	143.4	34.7	42.4	35.5	42.0
cexkul01	H <sub>4</sub> dota	Eu	SA	2.484	2.355	0.710	1.645	146.3	144.1	39.1	37.5	39.8	37.1
coknas	H <sub>4</sub> dota	Eu	SA	2.475	2.347	0.724	1.623	143.9	145.4	37.4	38.8	37.7	39.1
ogizek	H <sub>4</sub> dota	Eu	SA	2.389	2.320	0.646	1.675	147.7	149.1	38.5	40.5	35.6	40.5
jopjih	H <sub>4</sub> dota	Gd	SA	2.458	2.353	0.720	1.634	143.6	145.4	37.8	39.2	37.5	39.9
jopjih01	H <sub>4</sub> dota	Gd	SA	2.463	2.347	0.715	1.632	145.9	143.7	39.1	37.2	40.1	37.6
cokmul	H <sub>4</sub> dota	Gd	SA	2.455	2.345	0.729	1.616	143.6	144.9	38.0	39.0	37.9	39.4
ogizio	H <sub>4</sub> dota	Gd	SA	2.437	2.318	0.731	1.587	143.0	145.1	38.8	40.0	37.3	38.9
luqcab	H <sub>4</sub> dota	Dy	SA	2.479	2.340	0.755	1.585	141.6	143.1	38.1	38.5	38.4	38.2
unefid	H <sub>4</sub> dota	Dy	SA	2.466	2.340	0.721	1.649	144.8	143.1	39.9	38.4	40.7	38.0
godkoz	H <sub>4</sub> dota	Ho	SA	2.443	2.335	0.728	1.608	144.6	142.6	39.9	38.2	40.5	37.9
nojyiu	H <sub>4</sub> dota	Lu	SA	2.417	2.318	0.732	1.586	143.3	141.7	40.4	38.5	41.1	38.4
latkog01	H <sub>4</sub> dota	Y	SA	2.425	2.332	0.717	1.615	145.1	143.0	40.3	38.4	41.0	38.1
latkog	H <sub>4</sub> dota	Y	SA	2.436	2.334	0.718	1.617	144.8	143.1	40.0	37.8	40.4	38.1
luqci1 <sup>e</sup>	H <sub>4</sub> dota	Sc	SA'	- <sup>d</sup>	2.334	1.007	1.328	124.1	124.1	41.1	41.1	41.1	41.1
luqci2 <sup>e</sup>	H <sub>4</sub> dota	Sc	SA'	- <sup>d</sup>	2.325	0.989	1.336	125.0	125.0	41.2	41.2	41.2	41.2
luqci3 <sup>e</sup>	H <sub>4</sub> dota	Sc	SA'	- <sup>d</sup>	2.349	1.019	1.330	123.7	123.7	40.0	40.0	40.0	40.0
kejbeh	dotam	Pr	TSA	2.516	2.529	0.793	1.736	143.3	140.9	24.7	21.1	23.8	22.0
abofom	dotam <sup>R</sup> (R: <i>p</i> -NO <sub>2</sub> -Ph)	Eu	TSA	2.450 <sup>f</sup>	2.455	0.831	1.624	141.1	138.3	25.3	25.9	25.0	26.2
walvel	dotam <sup>R</sup> (R: <i>m</i> -pyridyl)	Eu	TSA	2.370	2.450	0.811	1.639	141.8	139.6	28.6	25.8	26.3	25.2
pekdob1 <sup>b</sup>	dotam <sup>R</sup> (R: <i>p</i> -CF <sub>3</sub> -benzyl)	Eu	TSA	2.532	2.544	0.881	1.663	135.3	138.0	21.6	24.1	21.6	24.1
pekdob2 <sup>b</sup>	dotam <sup>R</sup> (R: <i>p</i> -CF <sub>3</sub> -benzyl)	Eu	TSA	2.532	2.503	0.840	1.663	140.3	136.5	23.7	22.2	23.7	22.2
zaxcac	dotam	Eu	TSA	2.442	- <sup>g</sup>	0.793	- <sup>g</sup>	142.4	140.1	- <sup>g</sup>	- <sup>g</sup>	- <sup>g</sup>	- <sup>g</sup>
kawmay	dotam <sup>R</sup>	La	SA	- <sup>h</sup>	2.345	0.538	1.807	155.2	155.2	37.7	37.7	37.7	37.7

	(R: HO(CH <sub>2</sub> ) <sub>2</sub> O(CH <sub>2</sub> ) <sub>2</sub> -)												
ufiruw1 <sup>b</sup>	dotam <sup>R</sup> (R: α-phenethyl)	Ce	SA	2.491	2.407	0.643	1.763	148.3	150.7	35.5	37.1	36.0	36.6
ufiruw2 <sup>b</sup>	dotam <sup>R</sup> (R: α-phenethyl)	Ce	SA	2.488	2.384	0.646	1.738	148.6	149.7	36.2	37.5	36.4	37.2
ufisad1 <sup>b</sup>	dotam <sup>R</sup> (R: α-phenethyl)	Pr	SA	2.463	2.390	0.658	1.732	147.6	149.6	35.7	36.9	36.7	37.0
ufisad2 <sup>b</sup>	dotam <sup>R</sup> (R: α-phenethyl)	Pr	SA	2.479	2.361	0.652	1.709	148.0	149.1	36.5	38.2	36.7	37.0
eqozal	dotam <sup>R</sup> (R: α-phenethyl)	Pr	SA	2.516	2.363	0.665	1.698	147.1	149.1	38.2	37.2	37.9	36.5
eqozep	dotam <sup>R</sup> (R: α-phenethyl)	Nd	SA	2.501	2.357	0.682	1.676	146.0	148.2	38.4	37.6	38.1	37.0
banxiz	dotam <sup>R</sup> (R: prop-2-yn-1-yl)	Nd	SA	2.472	2.354	0.697	1.657	147.7	146.0	38.7	37.6	39.4	37.4
veckew	dotam (deuterated amides)	Nd	SA	2.435	2.347	0.695	1.653	146.8	146.5	37.2	37.8	37.8	39.2
eqozit	dotam <sup>R</sup> (R: α-phenethyl)	Sm	SA	2.495	2.348	0.707	1.641	144.4	146.1	38.8	38.3	38.5	37.1
xiwpic	dotam <sup>R</sup> (R: EtO <sub>2</sub> CCH <sub>2</sub> -)	Eu	SA	2.414	2.343	0.758	1.585	141.1	143.2	37.8	39.8	36.5	40.0
tuqtoo	dotam <sup>R</sup> (R: α-phenethyl)	Eu	SA	2.425	2.349	0.676	1.673	144.8	148.8	37.0	40.1	37.0	40.1
tuqtoo01	dotam <sup>R</sup> (R: α-phenethyl)	Eu	SA	2.425	2.349	0.676	1.673	144.8	148.8	37.0	40.1	37.0	40.1
abofig	dotam <sup>R</sup> (R: Ph)	Eu	SA	2.431	2.383	0.709	1.673	145.7	144.9	38.9	36.5	39.4	37.7
mubgar	dotam <sup>R</sup> (R: MeO <sub>2</sub> CCH <sub>2</sub> -)	Eu	SA	2.444 <sup>i</sup>	2.287	0.671	1.616	147.1	147.1	39.9	39.9	39.9	39.9
fibfeb	dotam <sup>R</sup> (R: EtO <sub>2</sub> C(CH <sub>3</sub> )CH-)	Eu	SA	2.414	2.330	0.733	1.597	144.4	144.6	39.9	40.7	38.8	38.4
tuqtii	dotam <sup>R</sup> (R: α-phenethyl)	Eu	SA	2.438	2.366	0.717	1.649	145.2	144.6	38.4	39.0	37.6	37.2
tuqtii01	dotam <sup>R</sup> (R: α-phenethyl)	Eu	SA	2.438	2.366	0.717	1.649	145.2	144.6	38.4	39.0	37.6	37.2
eqozoz	dotam <sup>R</sup> (R: α-phenethyl)	Eu	SA	2.482	2.352	0.716	1.637	143.7	145.9	38.8	38.1	38.6	37.4
veckia	dotam (deuterated amide)	Eu	SA	2.409	2.332	0.713	1.620	144.9	145.9	39.5	38.1	38.4	38.3
veckia01	dotam	Eu	SA	2.406	2.340	0.712	1.629	145.6	145.0	37.7	37.8	38.3	39.1
veckia02	dotam	Eu	SA	2.402	2.336	0.715	1.622	145.4	144.9	37.9	38.3	38.4	39.5
zaxcac	dotam	Eu	SA	2.442	- <sup>g</sup>	0.793	- <sup>g</sup>	142.4	140.1	- <sup>g</sup>	- <sup>g</sup>	- <sup>g</sup>	- <sup>g</sup>
zewsay	dotam <sup>R</sup> (R: HO <sub>2</sub> CCH <sub>2</sub> -)	Eu	SA	2.365 <sup>c</sup>	2.320	0.667	1.653	147.8	147.1	38.6	39.1	37.8	39.3
ufirac1 <sup>b</sup>	dotam <sup>R</sup> (R: α-phenethyl)	Gd	SA	2.351	2.336	0.724	1.611	144.1	144.3	38.9	36.4	38.9	36.4
ufirac2 <sup>b</sup>	dotam <sup>R</sup> (R: α-phenethyl)	Gd	SA	2.383	2.340	0.708	1.632	146.1	143.9	39.5	37.2	39.5	37.2
ufiseh1 <sup>b</sup>	dotam <sup>R</sup> (R: α-phenethyl)	Gd	SA	2.408	2.364	0.701	1.664	144.1	147.1	36.9	38.4	37.9	38.5
ufiseh2 <sup>b</sup>	dotam <sup>R</sup> (R: α-phenethyl)	Gd	SA	2.420	2.334	0.699	1.636	145.3	145.3	38.3	39.5	38.3	38.4
eqozuf	dotam <sup>R</sup> (R: α-phenethyl)	Gd	SA	2.460	2.350	0.723	1.627	143.2	145.4	39.0	38.4	38.6	37.8
gegcie	dotam <sup>R</sup> (R: Me)	Gd	SA	2.461	2.332	0.742	1.591	145.1	142.3	40.3	38.3	38.1	36.6
ehovay	dotam	Gd	SA	2.466	2.340	0.739	1.601	142.7	144.7	37.2	38.5	38.3	40.3
ekezaw	dotam <sup>R</sup> (R: prop-2-yn-1-yl)	Gd	SA	2.437	2.349	0.733	1.616	143.3	145.3	37.6	39.3	37.8	40.1
kehzut	dotam	Gd	SA	2.395	2.330	0.717	1.614	144.7	145.1	37.7	37.6	38.9	38.3
kejbad1 <sup>b</sup>	dotam	Gd	SA	2.395	2.328	0.732	1.597	142.7	145.3	38.2	39.7	37.3	39.2
kejbad2 <sup>b</sup>	dotam	Gd	SA	2.474	2.341	0.741	1.601	144.3	1423.0	38.4	38.9	38.8	37.1
pehfug	dotam <sup>R</sup> (R: H <sub>2</sub> O <sub>3</sub> PCH <sub>2</sub> -)	Gd	SA	2.425	2.306	0.710	1.597	145.2	144.4	39.5	38.8	40.2	40.7
pehgan	dotam <sup>R</sup> (R: H <sub>2</sub> O <sub>3</sub> PCH <sub>2</sub> -)	Gd	SA	2.395	2.299	0.671	1.628	147.8	146.7	41.2	38.4	41.2	38.4
sircay	dotam	Gd	SA	2.467	2.346	0.731	1.616	144.8	143.2	38.2	37.6	38.8	39.6

sircec	dotam	Gd	SA	2.461	2.360	0.737	1.624	144.4	143.2	38.1	37.3	38.3	39.5
sircig	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Gd	SA	2.435	2.351	0.690	1.661	146.2	146.0	38.3	38.0	39.3	38.2
sircom	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Gd	SA	2.380	2.372	0.655	1.717	148.9	146.5	37.5	35.9	37.5	35.9
sircus	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Gd	SA	2.381	2.364	0.672	1.691	147.5	146.0	38.0	37.0	38.0	37.0
sirdon	dotam	Gd	SA	2.384	2.332	0.715	1.617	146.0	143.9	39.2	39.1	39.0	38.1
ufireg	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Tb	SA	2.461	2.345	0.727	1.618	142.7	144.9	39.3	38.7	38.9	38.0
ruznuv	dotam <sup>R</sup> (R: BnO <sub>2</sub> C(CH <sub>3</sub> )CH-)	Tb	SA	2.380	2.312	0.714	1.599	144.6	145.2	39.2	40.8	37.7	40.9
efumat	dotam <sup>R</sup> (R: PhC(O)CH <sub>2</sub> -)	Tb	SA	2.437	2.319	0.746	1.574	142.6	143.9	39.2	39.5	38.6	39.7
banxof	dotam <sup>R</sup> (R: prop-2-yn-1-yl)	Tb	SA	2.437	2.337	0.741	1.596	144.4	142.9	40.0	39.3	39.8	38.5
bosjea	dotam	Tb	SA	2.450	2.330	0.727	1.603	143.5	144.8	38.2	39.1	37.4	40.2
tusput	dotam	Tb	SA	2.382	2.330	0.732	1.599	144.7	143.0	39.8	38.8	38.8	38.2
goybol	dotam <sup>R</sup> (R: Me)	Dy	SA	2.427	2.353	0.754	1.599	144.7	140.4	39.6	38.1	40.4	36.3
equbat	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Dy	SA	2.454	2.340	0.730	1.610	142.3	144.6	39.1	39.1	38.8	38.3
tuqtuu	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Dy	SA	2.422	2.342	0.725	1.617	144.1	143.6	39.2	38.1	38.3	39.7
tuqtuu01	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Dy	SA	2.422	2.342	0.725	1.617	144.1	143.6	39.2	38.1	38.3	39.7
banxev	dotam <sup>R</sup> (R: prop-2-yn-1-yl)	Dy	SA	2.427	2.335	0.747	1.588	142.4	144.0	39.4	40.1	38.6	40.2
ufisil1	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Er	SA	2.374	2.356	0.716	1.640	142.9	145.6	38.5	38.8	38.6	38.8
ufisil2	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Er	SA	2.380	2.336	0.719	1.617	143.5	144.0	38.7	40.4	38.6	39.3
ufirik	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Er	SA	2.432	2.335	0.743	1.593	141.4	143.5	39.4	39.4	39.0	38.8
tavsiv	dotam <sup>R</sup> (R: CF <sub>3</sub> CH <sub>2</sub> -)	Tm	SA	2.347	2.281	0.727	1.554	143.1	143.3	41.5	41.8	41.5	41.8
xohvev	dotam	Yb	SA	2.335	2.304	0.739	1.566	143.2	141.9	39.8	39.1	40.5	39.6
fibtit	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Yb	SA	2.440	2.328	0.740	1.588	141.1	143.2	39.7	39.7	39.1	39.1
banxul	dotam <sup>R</sup> (R: prop-2-yn-1-yl)	Yb	SA	2.417	2.322	0.758	1.564	141.0	142.7	38.9	40.7	40.0	40.5
sirdaz1 <sup>e</sup>	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Yb	SA	2.408	2.317	0.735	1.583	142.5	142.0	41.4	38.2	39.1	38.6
sirdaz2 <sup>e</sup>	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Yb	SA	2.388	2.326	0.747	1.580	141.5	142.0	36.9	39.9	38.6	39.5
sirdaz3 <sup>e</sup>	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Yb	SA	2.448	2.335	0.757	1.578	141.7	140.8	40.0	39.0	39.3	38.3
sirded1 <sup>b</sup>	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Yb	SA	2.445	2.344	0.766	1.578	140.9	140.3	38.4	39.4	37.2	38.8
sirded2 <sup>b</sup>	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Yb	SA	2.439	2.347	0.770	1.576	140.9	140.2	37.8	38.0	38.1	39.4
sirdih	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Yb	SA	2.397	2.349	0.747	1.602	140.7	141.9	37.5	39.3	37.4	41.3
abofus1 <sup>b</sup>	dotam <sup>R</sup> (R: <i>p</i> -NO <sub>2</sub> -Ph)	Lu	SA	2.346	2.271	0.760	1.511	142.0	141.5	39.6	39.5	39.5	39.9
abofus2 <sup>b</sup>	dotam <sup>R</sup> (R: <i>p</i> -NO <sub>2</sub> -Ph)	Lu	SA	2.369	2.290	0.790	1.500	140.0	139.9	39.2	39.0	38.4	39.7
ufiroq	dotam <sup>R</sup> (R: $\alpha$ -phenethyl)	Lu	SA	2.426	2.329	0.747	1.583	140.8	142.7	39.8	40.1	39.2	39.4
iwuqex	dotam <sup>R</sup> (R: <i>m</i> -pyridylmethyl)	Lu	SA	2.381	2.364	0.741	1.623	143.9	141.5	39.2	39.9	38.6	38.9
ciqyoq1 <sup>b</sup>	H <sub>4</sub> dotp <sup>R</sup> (R: Ph)	La	TSA	2.656	2.727	0.793	1.935	140.8	143.7	24.5	26.3	22.5	24.8
ciqyoq2 <sup>b</sup>	H <sub>4</sub> dotp <sup>R</sup> (R: Ph)	La	TSA	2.628	2.692	0.789	1.903	143.3	141.4	23.7	25.8	25.4	27.2
ruhmiq	H <sub>4</sub> dotp <sup>R</sup> (R: Bn)	La	TSA	2.664	2.661	0.812	1.850	142.3	140.4	29.6	28.1	29.3	28.3
ciquw1 <sup>b</sup>	H <sub>4</sub> dotp <sup>R</sup> (R: Ph)	Ce	TSA	2.652	2.714	0.806	1.909	139.5	142.8	25.2	26.7	23.8	25.2

ciqyuw2 <sup>b</sup>	H <sub>4</sub> dotp <sup>R</sup> (R: Ph)	Ce	TSA	2.583	2.689	0.792	1.897	140.7	142.7	23.9	26.0	25.8	27.4
ruhmem	H <sub>4</sub> dotp <sup>R</sup> (R: Bn)	Eu	TSA'	- <sup>d</sup>	2.646	0.970	1.676	130.0	131.4	28.8	29.9	29.1	30.2
axamap1 <sup>b</sup>	H <sub>8</sub> dotp	Eu	TSA'	- <sup>d</sup>	2.727	1.032	1.695	126.5	128.4	24.9	26.1	26.1	25.9
axamap2 <sup>b</sup>	H <sub>8</sub> dotp	Eu	TSA'	- <sup>d</sup>	2.749	1.058	1.692	122.9	128.9	23.8	24.8	23.3	25.2
axamet1 <sup>b</sup>	H <sub>8</sub> dotp	Eu	TSA'	- <sup>d</sup>	2.795	1.067	1.728	125.9	125.9	24.9	24.9	24.9	24.9
axamet2 <sup>b</sup>	H <sub>8</sub> dotp	Eu	TSA'	- <sup>d</sup>	2.792	1.067	1.724	125.4	125.4	21.9	21.9	21.9	21.9
eqiyim1 <sup>b</sup>	H <sub>8</sub> dotp	Gd	TSA'	- <sup>d</sup>	2.718	1.057	1.661	125.6	125.6	27.0	27.0	27.0	27.0
eqiyim2 <sup>b</sup>	H <sub>8</sub> dotp	Gd	TSA'	- <sup>d</sup>	2.685	1.042	1.643	126.4	126.4	23.6	23.6	23.6	23.6
sonfad	H <sub>4</sub> dotp <sup>R</sup> (R: H)	Gd	TSA'	- <sup>d</sup>	2.656	1.050	1.605	124.6	127.4	29.2	30.5	29.2	30.5
sonduv1 <sup>b</sup>	H <sub>4</sub> dotp <sup>R</sup> (R: OEt)	Gd	TSA'	- <sup>d</sup>	2.688	1.050	1.639	124.7	127.6	26.7	27.4	27.0	28.6
sonduv2 <sup>b</sup>	H <sub>4</sub> dotp <sup>R</sup> (R: OEt)	Gd	TSA'	- <sup>d</sup>	2.701	1.049	1.652	127.2	124.9	27.7	26.1	28.2	26.3
yeqded	H <sub>4</sub> dotp <sup>R</sup> (R: Bn)	Dy	TSA'	- <sup>d</sup>	2.625	0.978	1.647	130.3	128.3	29.8	28.6	29.2	28.9
yeqded01	H <sub>4</sub> dotp <sup>R</sup> (R: Bn)	Dy	TSA'	- <sup>d</sup>	2.626	0.984	1.642	127.5	129.2	28.7	29.6	27.9	29.9
ruhmow1 <sup>b</sup>	H <sub>4</sub> dotp <sup>R</sup> (R: Bn)	Yb	TSA'	- <sup>d</sup>	2.606	1.002	1.604	126.3	127.7	31.9	30.9	31.2	29.4
ruhmow2 <sup>b</sup>	H <sub>4</sub> dotp <sup>R</sup> (R: Bn)	Yb	TSA'	- <sup>d</sup>	2.626	1.015	1.612	124.8	127.8	29.3	29.4	32.9	28.5
lijfal	H <sub>4</sub> dotp <sup>R</sup> (R: Bn)	Y	TSA'	- <sup>d</sup>	2.612	0.973	1.639	129.0	128.9	30.4	27.9	29.3	28.3
babfob	<i>trans</i> -H <sub>6</sub> do2a2p	Ce	TSA	2.545	2.596	0.822	1.774	136.9	144.4	26.4	21.9	26.1	21.9
babcog	<i>trans</i> -H <sub>6</sub> do2a2p	Nd	TSA	2.542	2.584	0.835	1.749	136.1	143.2	26.9	23.1	26.5	22.8
babhap1 <sup>b</sup>	<i>trans</i> -H <sub>6</sub> do2a2p	Sm	TSA	2.555	2.516	0.801	1.715	137.4	143.8	30.0	27.1	29.1	27.0
babhap2 <sup>b</sup>	<i>trans</i> -H <sub>6</sub> do2a2p	Sm	TSA	2.602	2.547	0.819	1.729	135.3	143.8	27.7	25.9	28.2	25.8
babgui	<i>trans</i> -H <sub>6</sub> do2a2p	Sm	TSA'	- <sup>d</sup>	2.647	1.063	1.584	116.3	135.2	20.8	24.7	20.8	24.7
babgiw1 <sup>b</sup>	<i>trans</i> -H <sub>6</sub> do2a2p	Eu	TSA'	- <sup>d</sup>	2.716	1.101	1.616	115.5	132.9	19.6	25.1	19.6	25.1
babgiw2 <sup>b</sup>	<i>trans</i> -H <sub>6</sub> do2a2p	Eu	TSA'	- <sup>d</sup>	2.700	1.086	1.615	115.1	134.6	19.7	23.2	19.7	23.2
babhet	<i>trans</i> -H <sub>6</sub> do2a2p	Tb	TSA'	- <sup>d</sup>	2.631	1.079	1.552	114.7	133.6	21.5	25.2	21.5	25.2
babgao	<i>trans</i> -H <sub>6</sub> do2a2p	Dy	TSA'	- <sup>d</sup>	2.610	1.077	1.532	114.4	132.6	22.2	25.3	22.2	25.3
babges	<i>trans</i> -H <sub>6</sub> do2a2p	Er	TSA'	- <sup>d</sup>	2.610	1.086	1.524	114.5	131.4	23.4	26.0	23.4	26.0
babhix	<i>trans</i> -H <sub>6</sub> do2a2p	Yb	TSA'	- <sup>d</sup>	2.601	1.094	1.507	113.0	130.8	23.4	25.7	23.4	25.7
lanqat1 <sup>b</sup>	H <sub>5</sub> do3ap	Nd	TSA	2.519	2.499	0.783	1.738	142.2	143.0	30.0	23.3	28.6	24.0
lanqat2 <sup>b</sup>	H <sub>5</sub> do3ap	Nd	TSA	2.536	2.591	0.792	1.746	140.3	143.4	28.9	23.3	28.1	23.5
ofayec	H <sub>4</sub> do3ap <sup>R</sup> (R: H)	Eu	TSA	2.555	2.524	0.871	1.653	134.6	139.5	25.8	25.6	26.6	27.8
larlia	H <sub>5</sub> do3ap	Tb	TSA	2.678	2.514	0.853	1.663	135.8	138.6	26.9	27.2	27.4	27.1
kejroh1 <sup>j</sup>	H <sub>4</sub> do3ap <sup>R</sup> (R: <i>p</i> -NH <sub>2</sub> -Bn)	Y	TSA	2.486	2.504	0.763	1.742	140.3	142.6	27.1	24.5	25.7	27.2
larkof	H <sub>5</sub> do3ap	Dy	TSA'	- <sup>d</sup>	2.564	1.035	1.531	123.5	129.5	25.5	24.7	25.7	25.2
larkul	H <sub>5</sub> do3ap	Er	TSA'	- <sup>d</sup>	2.525	1.054	1.472	122.2	128.2	26.4	24.2	24.0	25.0
larlas	H <sub>5</sub> do3ap	Lu	TSA'	- <sup>d</sup>	2.547	1.065	1.483	120.8	127.1	26.7	26.0	26.4	26.3
larlew	H <sub>5</sub> do3ap	Lu	TSA'	- <sup>d,c</sup>	2.489	1.068	1.468	120.5	126.9	26.6	25.9	25.9	25.5
larlog	H <sub>5</sub> do3ap	Y	TSA'	- <sup>d</sup>	2.566	1.043	1.525	123.0	129.1	25.8	25.0	25.4	25.3
kejroh3 <sup>j</sup>	H <sub>4</sub> do3ap <sup>R</sup> (R: <i>p</i> -NH <sub>2</sub> -Bn)	Y	TSA'	- <sup>d</sup>	2.562	1.030	1.533	122.4	130.4	25.1	27.7	24.5	27.0



kejroh2 <sup>j</sup>	H <sub>4</sub> do3ap <sup>R</sup> (R: <i>p</i> -NH <sub>2</sub> -Bn)	Y	SA	2.486	2.348	0.763	1.586	140.3	142.6	42.2	40.0	40.8	42.2
1959625	H <sub>4</sub> do3ap <sup>DBAm</sup>	Ce	TSA	2.529	2.531	0.716	1.817	143.4	148.6	24.3	26.3	21.4	29.5
1959634	H <sub>4</sub> do3ap <sup>DBAm</sup>	Pr	TSA	2.537	2.529	0.730	1.801	142.9	147.3	24.7	26.0	21.8	29.4
1959633	H <sub>4</sub> do3ap <sup>DBAm</sup>	Nd	TSA	2.623	2.541	0.796	1.746	139.7	143.0	27.3	23.8	25.7	26.1
1959632	H <sub>4</sub> do3ap <sup>DBAm</sup>	Nd	TSA	2.476	2.510	0.718	1.794	142.9	148.0	25.8	27.0	22.5	29.6
1959636	H <sub>4</sub> do3ap <sup>DBAm</sup>	Sm	TSA	2.612	2.533	0.802	1.732	139.0	142.0	27.8	24.1	25.9	26.5
1959629	H <sub>4</sub> do3ap <sup>DBAm</sup>	Eu	TSA	2.594	2.521	0.804	1.718	138.7	141.9	28.3	23.9	26.3	26.2
diydat <sup>k</sup>	H <sub>4</sub> do3ap <sup>DBAm</sup>	Gd	TSA	2.583	2.517	0.807	1.710	138.5	141.5	28.5	24.1	26.6	26.4
1959630	H <sub>4</sub> do3ap <sup>DBAm</sup>	Gd	TSA	2.655	2.533	0.864	1.671	135.3	138.3	27.6	24.7	25.2	26.8
1959637	H <sub>4</sub> do3ap <sup>DBAm</sup>	Tb	TSA	2.591	2.505	0.809	1.697	138.2	141.0	28.4	24.3	26.7	26.6
diycum <sup>b,k</sup>	H <sub>4</sub> do3ap <sup>DBAm</sup>	Dy	TSA	2.623	2.542	0.856	1.687	135.2	138.6	26.0	23.8	23.5	26.7
diycum <sup>b,k</sup>	H <sub>4</sub> do3ap <sup>DBAm</sup>	Dy	TSA	2.747	2.509	0.834	1.675	136.2	139.4	27.3	24.9	26.1	26.9
1959638	H <sub>4</sub> do3ap <sup>DBAm</sup>	Tb	TSA'	- <sup>d</sup>	2.568	1.010	1.560	124.8	131.4	24.5	25.9	24.7	25.2
1959631	H <sub>4</sub> do3ap <sup>DBAm</sup>	Ho	TSA'	2.753 <sup>l</sup>	2.533	0.950	1.584	129.0	133.2	27.4	25.2	26.0	26.6
1959626	H <sub>4</sub> do3ap <sup>DBAm</sup>	Er	TSA'	- <sup>d</sup>	2.533	1.052	1.481	123.2	126.7	27.7	26.5	27.0	26.9
1959628 <sup>b</sup>	H <sub>4</sub> do3ap <sup>DBAm</sup>	Er	TSA'	2.803 <sup>l</sup>	2.545	0.947	1.599	129.7	132.7	27.9	26.1	25.7	27.1
1959628 <sup>b</sup>	H <sub>4</sub> do3ap <sup>DBAm</sup>	Er	TSA'	- <sup>d</sup>	2.502	1.007	1.496	124.4	129.9	27.2	25.7	26.4	26.9
1959627	H <sub>4</sub> do3ap <sup>DBAm</sup>	Yb	TSA'	- <sup>d</sup>	2.534	1.050	1.485	121.7	127.7	27.1	26.1	26.3	26.9
1959635	H <sub>4</sub> do3ap <sup>DBAm</sup>	Sc	TSA'	- <sup>d</sup>	2.457	1.040	1.417	120.6	123.9	29.2	28.6	29.2	29.0

<sup>a</sup>Q represents pair of centroids, QN<sub>4</sub>-QO<sub>4</sub>. <sup>b</sup>Two structurally independent molecules present in the crystal structure. <sup>c</sup>Carboxylate oxygen atom from neighbouring unit coordinated in the apical position. <sup>d</sup>No apically coordinated water/oxygen atom. <sup>e</sup>Three structurally independent molecules present in the crystal structure. <sup>f</sup>Methanol oxygen atom coordinated in the apical position. <sup>g</sup>Not applicable due to disorder. <sup>h</sup>Apical coordination of chloride. <sup>i</sup>Trifluoromethylsulfonate oxygen atom coordinated in the apical position. <sup>j</sup>Two structurally independent molecules present in the crystal structure, one of them disordered in TSA/SA arrangement. <sup>k</sup>ref.<sup>[55]</sup>. <sup>l</sup>“Semi-coordinated” water molecule.

### Crystal structures refinement details

In general, all non-hydrogen atoms were refined anisotropically. Only some disordered groups/atoms (typically carbon atoms of benzyl groups and oxygen atoms of water molecules of crystallization) were treated isotropically. Almost all hydrogen atoms were localized in difference density map; however, those bound to the carbon atoms were placed in theoretical positions and hydrogen atoms bound to heteroatoms (O, N) were usually kept in original positions using  $U_{eq}(H) = 1.2 U_{eq}(X)$  to keep the number of refinement parameters low. Alterations from these general procedures are described below.

In the crystal structures of  $[Ln(HL1)] \cdot 0.5NH_4Cl \cdot 6.5H_2O$  ( $Ln = Ce, Pr$ ), one formula unit forms the structurally independent unit. Chloride atom was best refined with half occupancy to get reliable thermal factor, and its charge is compensated by half-occupied ammonium ion, of which nitrogen atom shares position with oxygen atom of half-occupied water molecule. In these fragments, four tetrahedrally-located hydrogen atoms of ammonium cation were found, two of them were defined as shared with water molecule, and EXYZ/EADP commands for corresponding N/O atoms were used. Two water molecules of crystallization were best refined with oxygen atoms disordered over two close positions and sharing corresponding hydrogen atoms.

In the crystal structure of  $[Nd(HL1)] \cdot 3H_2O$ , one formula unit forms the structurally independent unit. Two water molecules of crystallization were best refined with oxygen atoms disordered over two and three close positions, respectively. Positions of corresponding hydrogen atoms cannot be reliably determined.

One formula unit forms the structurally independent unit also in the crystal structures of  $(NH_4)[Ln(H_2O)(do3ap^{DBAm})] \cdot 3H_2O$  ( $Ln = Nd, Sm, Eu, Tb$ ). Similarly to isostructural Gd(III)-complex reported previously,[55] the same kind of disorder of benzylic side groups was found. One of phenyl rings was best refined to be staggered in two positions with pivot and “para” carbon atoms common for both possibilities. The other phenyl ring was best refined as regular hexagon (AFIX 66) disordered slit over two or three positions, which are nearly parallel. However, in Sm(III)-complex, probably more complicated disorder is present, but it cannot be reliably modelled. As the result of such complication, large maxima in difference density map are present.

However, all hydrogen atoms belonging to coordinated water molecule, ammonium cation as well as to water molecules of crystallization were found in the difference density map, making rich system of hydrogen bonds.

In the structure of  $Ca_{0.5}[Ho(L1)] \cdot 6H_2O$ , one formula unit forms the structurally independent unit. Occupancy of Ca(II) ion was set to 50% as the ion is close to its symmetrically related position; such disorder is reflected also in Ca(II)-coordinated water molecules, which were refined also with occupancy 50%. However, more complicated disorder is present, which results in large electronic density map maxima/minima close to disordered water molecules. Occupancy of water molecule present in apical position of the Ho(III) complex was set to 50% to keep its thermal factor reliable. In benzylic side groups, similar disorder as described above was found (one phenyl staggered in two positions, other phenyl slid in two positions).

In the structure of  $Ca_{0.5}[Gd(H_2O)(L1)] \cdot 6H_2O$ , one formula unit forms the structurally independent unit. The Ca(II) occupies special position with 50% occupancy factor, and is coordinated with water molecules forming distorted octahedron. Two of such water molecules are placed in special position, two other molecules are disordered over two close positions (but large difference density map maxima close to these positions point to more complicated disorder), corresponding hydrogen atoms cannot be found. Hydrogen atoms of coordinated water molecule and of other three water molecules of crystallization were located in the difference density map and form hydrogen bond

network. One side-group phenyl was refined staggered in two positions and other phenyl was slid in two positions in similar disorder as described above.

In the crystal structures of  $[\text{Tb}(\text{HL1})]\cdot\text{NH}_4\text{Cl}\cdot\text{H}_2\text{O}$ ,  $(\text{NH}_4)[\text{Er}(\text{L1})]\cdot 4\text{H}_2\text{O}$  and  $(\text{NH}_4)[\text{Sc}(\text{L1})]\cdot 4\text{H}_2\text{O}$ , formula unit corresponds to the independent part and all hydrogen atoms were located in the difference density map. No disorder was found.

In the crystal structure of  $\text{Na}[\text{Er}(\text{L1})]\cdot 4.25\text{H}_2\text{O}$ , two formula units form the independent part of the unit cell. In both independent complex molecules, disorder of phenyl rings was found (one phenyl staggered, other phenyl slid). Sodium aquaions are disordered as well. The disorder was best refined by splitting Na(I) into two close positions with half-occupancy. Remaining large maxima in difference density map were attributed to water molecules of crystallization, however, some of them lie very close to each other, and needed to be refined with half-occupancy. Hydrogen atoms could not be located.

In the crystal structure of  $\text{Na}[\text{Yb}(\text{L1})]\cdot 6\text{H}_2\text{O}$ , one formula unit forms the structurally independent unit. Disorder of phenyl rings was found similar to previous cases (one phenyl staggered, other phenyl slid). Some of water molecules coordinated to Na(I) counter ion were best refined disordered in two close positions; corresponding hydrogen atoms could not be found.

All the data for the structures reported have been deposited with the Cambridge Crystallographic Data Centre. Experimental crystal data are outlined in Table S4.

Table S4. Experimental crystallographic data for structures of lanthanide(III)–H<sub>4</sub>do3ap<sup>DBAm</sup> (Ln–H<sub>4</sub>L1) complexes.

Compound	[Ce(HL1)]·0.5NH <sub>4</sub> Cl ·6.5H <sub>2</sub> O	[Pr(HL1)]·0.5NH <sub>4</sub> Cl ·6.5H <sub>2</sub> O	[Nd(HL1)] ·3H <sub>2</sub> O	(NH <sub>4</sub> )[Nd(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O	(NH <sub>4</sub> )[Sm(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O	(NH <sub>4</sub> )[Eu(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O	Ca <sub>0.5</sub> [Gd(H <sub>2</sub> O) (L1)]·6H <sub>2</sub> O
Formula	C <sub>30</sub> H <sub>56</sub> CeCl <sub>0.5</sub> N <sub>5.5</sub> O <sub>14.5</sub> P	C <sub>30</sub> H <sub>56</sub> Cl <sub>0.5</sub> N <sub>5.5</sub> O <sub>14.5</sub> PPr	C <sub>30</sub> H <sub>47</sub> N <sub>5</sub> NdO <sub>11</sub> P	C <sub>30</sub> H <sub>52</sub> N <sub>6</sub> NdO <sub>12</sub> P	C <sub>30</sub> H <sub>52</sub> N <sub>6</sub> O <sub>12</sub> P <sub>Sm</sub>	C <sub>30</sub> H <sub>52</sub> EuN <sub>6</sub> O <sub>12</sub> P	C <sub>30</sub> H <sub>54</sub> Ca <sub>0.5</sub> GdN <sub>5</sub> O <sub>15</sub> P
<i>M<sub>r</sub></i>	914.61	915.40	828.93	863.98	870.09	871.70	933.04
Shape	bar	prism	bar	plate	prism	prism	plate
Colour	colourless	slightly green	colourless	colourless	colourless	colourless	colourless
Crystal system	triclinic	triclinic	triclinic	monoclinic	monoclinic	monoclinic	monoclinic
Space group	<i>P</i> –1	<i>P</i> –1	<i>P</i> –1	<i>C</i> 2/ <i>c</i>	<i>C</i> 2/ <i>c</i>	<i>C</i> 2/ <i>c</i>	<i>P</i> 2/ <i>c</i>
<i>a</i> / Å	11.7568(8)	11.7803(8)	10.5893(5)	47.142(2)	47.041(8)	47.136(3)	24.2222(7)
<i>b</i> / Å	12.4231(9)	12.4274(9)	12.4277(6)	10.3053(4)	10.2877(14)	10.3082(5)	10.1613(2)
<i>c</i> / Å	14.5497(10)	14.5862(11)	14.6008(8)	15.3151(7)	15.205(2)	15.2253(8)	15.4093(4)
<i>α</i> / °	68.256(2)	68.147(2)	112.071(2)	90	90	90	90
<i>β</i> / °	79.612(2)	79.566(2)	92.429(2)	90.722(2)	90.6500(10)	90.685(2)	95.4550(10)
<i>γ</i> / °	78.900(2)	78.962(2)	109.292(2)	90	90	90	90
<i>U</i> / Å <sup>3</sup>	1923.0(2)	1931.1(2)	1649.94(15)	7439.7(5)	7357.9(19)	7397.3(7)	3775.50(16)
<i>Z</i>	2	2	2	8	8	8	4
<i>D</i> <sub>calc</sub> / g cm <sup>–3</sup>	1.580	1.574	1.669	1.543	1.571	1.565	1.641
<i>μ</i> / mm <sup>–1</sup>	1.332	1.409	1.690	1.505	1.706	1.806	1.941
Unique refl.	8828	8886	7581	7322	8668	8492	8663
Obsd. refl.	8293	8518	6521	6678	8435	8117	7451
<i>R</i> <sub>1</sub> ( <i>I</i> > 2σ( <i>I</i> ))	0.0271	0.0286	0.0433	0.1067	0.0626	0.0567	0.0616
<i>R</i> <sub>2</sub> (all data)	0.0301	0.0300	0.0613	0.1147	0.0637	0.0590	0.0714
<i>wR</i> <sub>2</sub> ( <i>I</i> > 2σ( <i>I</i> ))	0.0718	0.0776	0.0742	0.2324	0.1671	0.1358	0.1484
<i>wR</i> <sub>2</sub> (all data)	0.0732	0.0785	0.0798	0.2357	0.1674	0.1369	0.1548
CCDC-number	1959625	1959634	1959632	1959633	1959636	1959629	1959630

Compound	(NH <sub>4</sub> )[Tb(H <sub>2</sub> O) (L1)]·3H <sub>2</sub> O	[Tb(HL1)] ·NH <sub>4</sub> Cl·H <sub>2</sub> O	Ca <sub>0.5</sub> [Ho(L1)] ·6H <sub>2</sub> O	Na[Er(L1)] ·4.25H <sub>2</sub> O	(NH <sub>4</sub> )[Er(L1)] ·4H <sub>2</sub> O	Na[Yb(L1)] ·6H <sub>2</sub> O	(NH <sub>4</sub> )[Sc(L1)] ·4H <sub>2</sub> O
Formula	C <sub>30</sub> H <sub>52</sub> N <sub>6</sub> O <sub>12</sub> PTb	C <sub>30</sub> H <sub>47</sub> ClN <sub>6</sub> O <sub>9</sub> PTb	C <sub>30</sub> H <sub>52</sub> Ca <sub>0.5</sub> HoN <sub>5</sub> O <sub>14</sub> P	C <sub>30</sub> H <sub>48.5</sub> ErN <sub>5</sub> NaO <sub>12.25</sub> P	C <sub>30</sub> H <sub>52</sub> ErN <sub>6</sub> O <sub>12</sub> P	C <sub>30</sub> H <sub>52</sub> N <sub>5</sub> NaO <sub>14</sub> PYb	C <sub>30</sub> H <sub>52</sub> N <sub>6</sub> O <sub>12</sub> PSc
<i>M<sub>r</sub></i>	878.66	861.07	922.70	896.46	887.00	933.76	764.70
Shape	prism	plate	plate	plate	plate	plate	prism
Colour	colourless	colourless	colourless	slightly pink	slightly pink	colourless	colourless
Crystal system	monoclinic	monoclinic	monoclinic	triclinic	monoclinic	monoclinic	monoclinic
Space group	<i>C2/c</i>	<i>Pn</i>	<i>P2/c</i>	<i>P</i> -1	<i>P2<sub>1</sub>/c</i>	<i>C2/c</i>	<i>P2<sub>1</sub>/c</i>
<i>a</i> / Å	47.048(2)	9.9701(2)	24.1798(13)	10.2212(10)	24.3204(4)	49.3984(12)	24.0303(16)
<i>b</i> / Å	10.2943(4)	15.3519(3)	10.1265(5)	15.3230(13)	10.42140(10)	10.1801(2)	10.3449(6)
<i>c</i> / Å	15.2057(7)	11.4210(3)	15.2799(8)	25.162(2)	15.1910(2)	15.3464(4)	15.1430(11)
<i>α</i> / °	90	90	90	77.957(3)	90	90	90
<i>β</i> / °	90.717(2)	94.9903(8)	95.775(2)	78.577(3)	107.6020(10)	102.544(2)	107.497(3)
<i>γ</i> / °	90	90	90	89.761(3)	90	90	90
<i>U</i> / Å <sup>3</sup>	7363.9(6)	1741.47(7)	3722.4(3)	3774.9(6)	3669.93(9)	7533.2(3)	3590.2(4)
<i>Z</i>	8	2	4	4	4	8	4
<i>D</i> <sub>calc</sub> / g cm <sup>-3</sup>	1.585	1.642	1.646	1.577	1.605	1.647	1.415
<i>μ</i> / mm <sup>-1</sup>	2.031	2.213	2.309	2.341	2.397	2.608	0.318
Unique refl.	7234	6788	8553	14762	8434	7410	8266
Obsd. refl.	6599	6464	7679	13327	6985	6343	7311
<i>R</i> <sub>1</sub> ( <i>I</i> > 2σ( <i>I</i> ))	0.0522	0.0228	0.1022	0.0915	0.0300	0.0402	0.0389
<i>R</i> <sub>2</sub> (all data)	0.0566	0.0252	0.1103	0.1010	0.0404	0.0495	0.0455
<i>wR</i> <sub>2</sub> ( <i>I</i> > 2σ( <i>I</i> ))	0.1142	0.0484	0.2400	0.2019	0.0653	0.0990	0.0983
<i>wR</i> <sub>2</sub> (all data)	0.1159	0.0495	0.2429	0.2082	0.0702	0.1039	0.1028
CCDC-number	1959637	1959638	1959631	1959628	1959626	1959627	1959635

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