

Supplementary Information

Elucidating the role of metal ions in carbonic anhydrase catalysis

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This PDF file includes:

Supplementary Figure 1 - HCO₃⁻ binding geometry in Zn-, Co-, Ni-CA II.

Supplementary Figure 2 - Substrate binding in Zn-CA II at pH 7.8 and 11.0.

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Supplementary Figure 4 - Absorbance of Apo- & Zn-CA II in esterase kinetic assay at pH 7.8

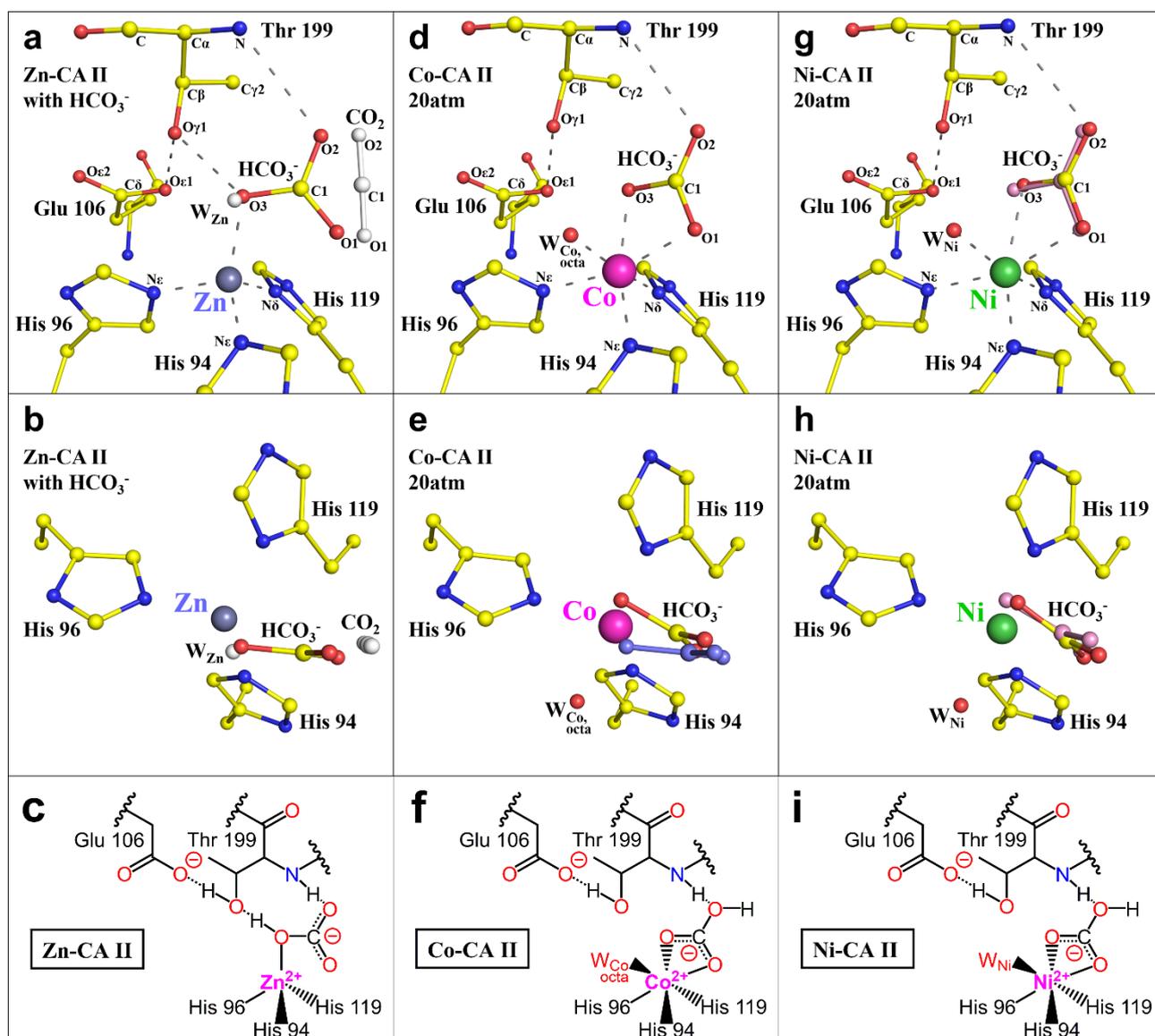
Supplementary Figure 5 - Partial occupancy determination of His64 in CA II structures.

Supplementary Table 1 - Data collection and refinement statistics for the CA II structures.

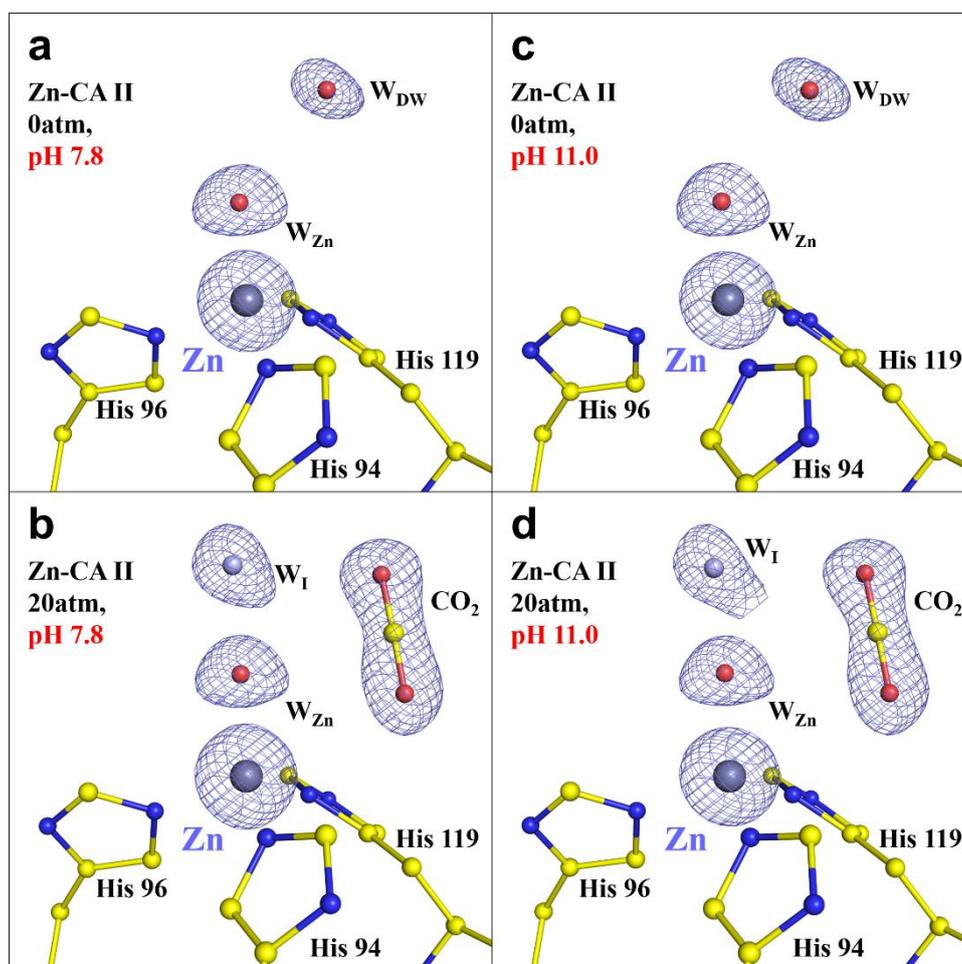
Supplementary Table 2 - List of key bound water molecules in the CA II structures.

Supplementary Table 3 - Distance geometry in the CA II structures.

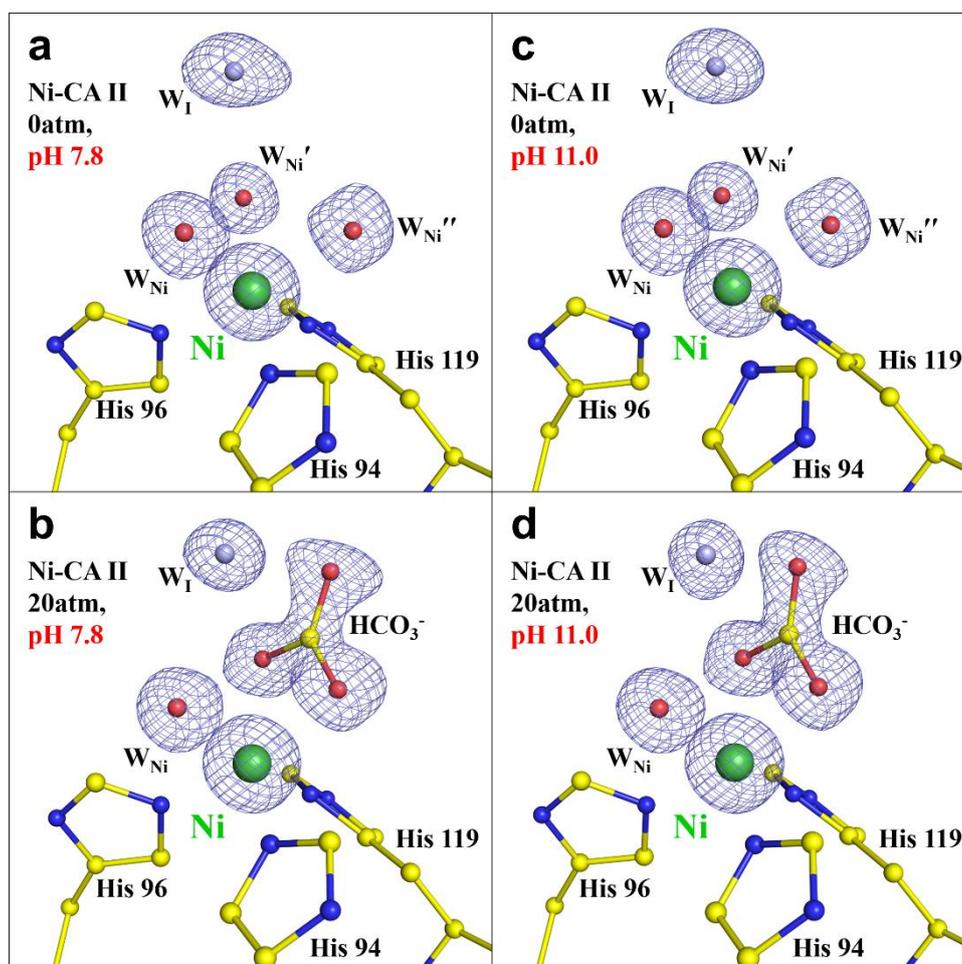
Supplementary References



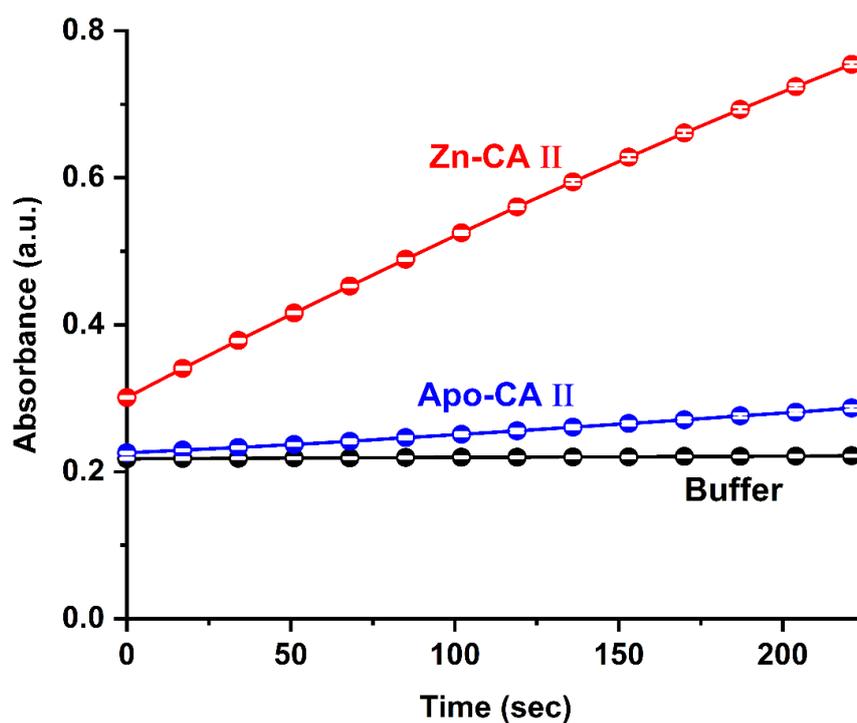
Supplementary Fig. 1. HCO_3^- binding geometry in Zn-, Co-, Ni-CA II. **a-c)** Monodentate binding of HCO_3^- in Zn-CA II (PDB code: 2vvb)¹. The CO_2 molecule and W_{Zn} (white) from Zn-CA II 20atm are superimposed for comparison. Note that HCO_3^- lies on the plane made by the CO_2 molecule and W_{Zn} . **d-f)** Bidentate binding of HCO_3^- in Co-CA II 20atm pH 7.8. The HCO_3^- molecule from Zn-CA II (light purple, PDB 2vvb) is superimposed for comparison. Note that the HCO_3^- molecule in Co-CA II is tilted by $\sim 31^\circ$ to the HCO_3^- molecule in Zn-CA II. **g-i)** Bidentate binding of HCO_3^- in Ni-CA II 20atm. The HCO_3^- molecule in Co-CA II (pink) is superimposed for comparison.



Supplementary Fig. 2. Substrate binding in Zn-CA II at pH 7.8 (a-b) and 11.0 (c-d). The metal coordination maintains tetrahedral geometry upon CO₂ binding regardless of pH values. The electron density ($2F_o - F_c$, blue) is contoured at 2.2σ . The intermediate water (W_I) is colored in steel blue for clarity.

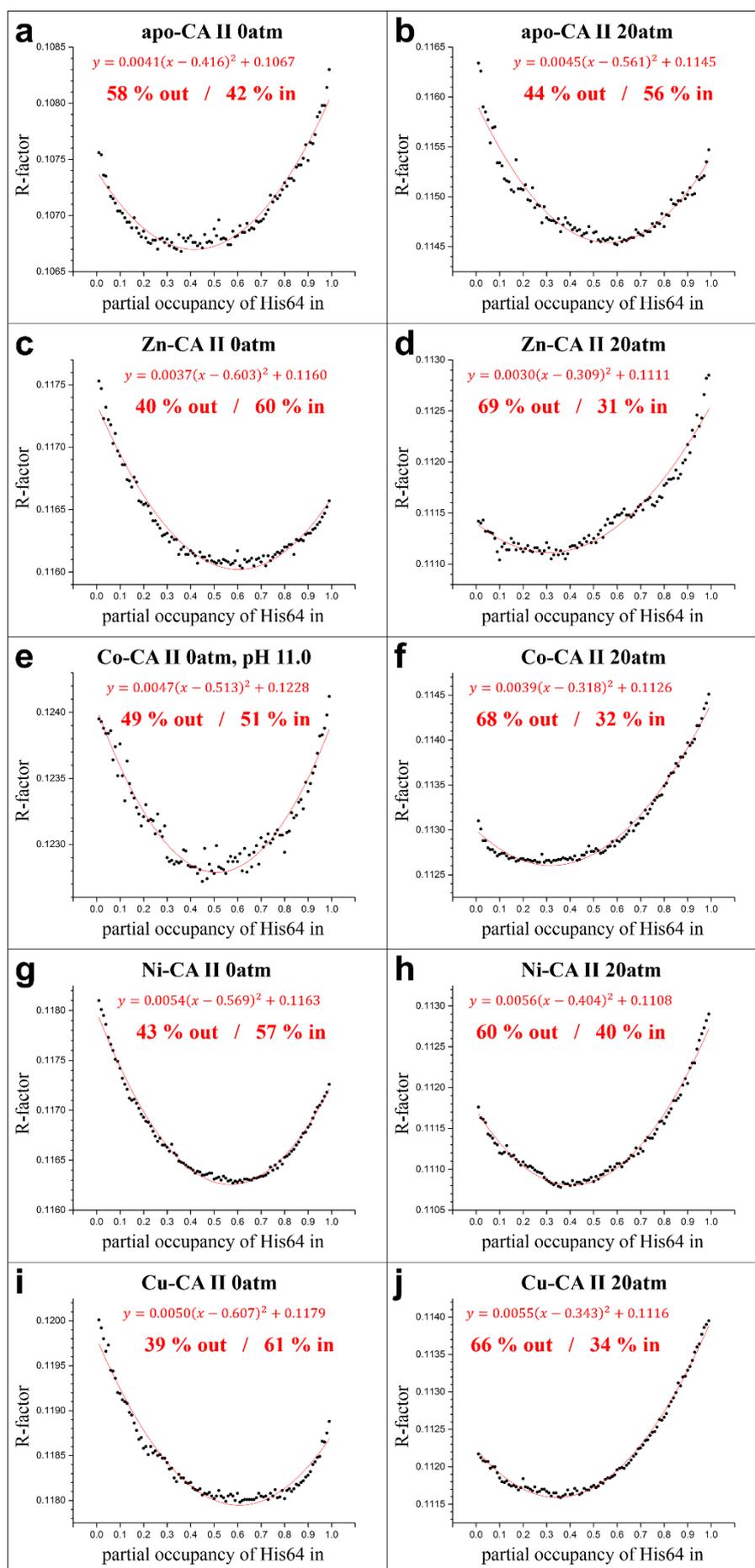


Supplementary Fig. 3. Product binding in Ni-CA II at pH 7.8 (a-b) and 11.0 (c-d). The metal coordination maintains octahedral geometry upon HCO_3^- binding regardless of pH values. The intermediate water (W_I) is colored in steel blue for clarity. The electron density ($2F_o-F_c$, blue) is contoured at 2.2σ .



Supplementary Fig. 4. Absorbance of Apo- and Zn-CA II in esterase kinetic assay at pH 7.8.

CA II esterase activity was measured spectroscopically at 348nm, indicative of substrate 4-nitrophenyl acetate hydrolysis. Compared to Zn-CA II, Apo-CA II and buffer show little to no esterase activity. The standard deviation errors (white) are presented in the data points and are ranging from 0.2 % ~ 1.1 %. Source data are provided as a Source Data file.



Supplementary Fig. 5. See next page for figure caption.

Supplementary Fig. 5. Partial occupancy determination of His64. The His64 out/in conformations are determined for the selected structures of apo-CA II (**a, b**) Zn-CA II (**c, d**), Co-CA II (**e, f**), Ni-CA II (**g, h**) Cu-CA II (**i, j**). For each data set, systematic refinements were carried out on 99 structures with manually adjusted His64 in/out occupancies. The obtained data points were then fitted to quadratic functions, showing the minimum points in the overall R factors. The His64 out/in conformations for the other structures can be found in Supplementary Table 1. Source data are provided as a Source Data file.

Supplementary Table 1. Data collection and refinement statistics for the CA II structures.

	apo-CA II 0atm 6LUU	apo-CA II 20atm 6LUV	Zn-CA II 0atm 6LUW	Zn-CA II 20atm 6LUX	Zn-CA II 0atm pH11.0 6LUY	Zn-CA II 20atm pH11.0 6LUZ	Co-CA II 0atm 6LV1	Co-CA II 20atm 6LV2
Data collection								
Space group	$P2_1$	$P2_1$	$P2_1$	$P2_1$	$P2_1$	$P2_1$	$P2_1$	$P2_1$
Cell dimensions								
a, b, c (Å)	42.13, 41.30, 72.21	42.26, 41.38, 72.00	42.21, 41.28, 72.15	42.37, 41.44, 72.13	42.28, 41.26, 72.07	42.39, 41.47, 72.15	42.31, 41.22, 72.05	42.32, 41.32, 72.22
β (°)	104.27	104.18	104.18	104.05	104.19	104.11	104.16	104.03
Resolution (Å)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)
R_{sym} (%)	5.8 (36.8)	7.1 (48.1)	6.5 (37.6)	6.8 (66.9)	8.5 (17.7)	5.6 (20.3)	9.2 (60.7)	8.2 (65.6)
$I/\sigma(I)$	28.8 (6.8)	21.2 (5.1)	29.7 (6.3)	29.8 (3.6)	25.2 (12.7)	29.9 (11.8)	24.3 (3.2)	20.1 (3.8)
Completeness (%)	95.5 (92.6)	94.3 (91.2)	98.8 (97.5)	96.2 (92.9)	98.0 (96.0)	94.6 (91.9)	96.0 (93.3)	98.4 (96.4)
Redundancy	7.5 (7.3)	7.5 (7.4)	7.3 (7.1)	7.4 (7.3)	7.4 (7.4)	7.6 (7.5)	7.6 (7.6)	7.4 (7.2)
Refinement								
Resolution (Å)	1.20	1.20	1.20	1.20	1.20	1.20	1.20	1.20
No. reflections	72,011	71,305	74,561	73,183	73,898	72,112	72,478	74,640
$R_{\text{work}} / R_{\text{free}}$ (%)	11.0 / 13.8	11.7 / 14.8	11.7 / 14.1	11.3 / 14.2	11.1 / 13.1	10.3 / 12.7	12.0 / 15.0	11.4 / 13.9
No. atoms								
Protein	4,285	4,256	4,253	4,245	4,253	4,245	4,259	4,240
Ligand/ion	1 glycerol	2 CO ₂ , 1 glycerol	1 glycerol	2 CO ₂ , 1 glycerol	1 glycerol	2 CO ₂ , 1 glycerol	1 HCO ₃ ⁻ , 1 glycerol	1 HCO ₃ ⁻ , 1 CO ₂ , 1 glycerol
Water	371	373	263	367	264	365	283	333
B-factors								
Protein (main / side chain)	9.78 / 12.75	11.48 / 14.38	10.09 / 13.22	10.80 / 13.62	7.90 / 10.99	8.87 / 11.51	10.84 / 14.07	10.31 / 13.31
Ligand/ion	20.76 (glycerol)	13.90 (first CO ₂), 26.78 (second CO ₂), 18.39 (glycerol)	18.32 (glycerol)	10.63 (first CO ₂), 23.50 (second CO ₂), 17.17 (glycerol)	14.79 (glycerol)	8.80 (first CO ₂), 21.13 (second CO ₂), 14.78 (glycerol)	11.23 (HCO ₃ ⁻), 24.46 (glycerol)	9.25 (HCO ₃ ⁻), 23.17 (second CO ₂), 21.27 (glycerol)
Water	30.25	30.67	26.12	30.58	24.33	27.75	27.41	29.71
R.m.s. deviations								
Bond lengths (Å)	0.031	0.030	0.029	0.028	0.029	0.028	0.031	0.028
Bond angles (°)	2.515	2.435	2.437	2.423	2.506	2.368	2.644	2.284
Partial occupancy (%)								
His64 (out/in) conformation (%)	58 / 42	44 / 56	40 / 60	69 / 31	32 / 68	63 / 37	49 / 51	68 / 32
	Co-CA II 0atm pH11.0 6LV3	Co-CA II 20atm pH11.0 6LV4	Ni-CA II 0atm 6LV5	Ni-CA II 20atm 6LV6	Ni-CA II 0atm pH11.0 6LV7	Ni-CA II 20atm pH11.0 6LV8	Cu-CA II 0atm 6LV9	Cu-CA II 20atm 6LVA
Data collection								
Space group	$P2_1$	$P2_1$	$P2_1$	$P2_1$	$P2_1$	$P2_1$	$P2_1$	$P2_1$
Cell dimensions								
a, b, c (Å)	42.33, 41.26, 72.08	42.36, 41.46, 72.31	42.40, 41.29, 71.92	42.40, 41.37, 72.13	42.52, 41.23, 71.87	42.42, 41.40, 72.19	42.33, 41.23, 72.09	42.36, 41.40, 72.31
β (°)	104.13	104.02	104.03	104.01	104.06	104.04	104.17	103.97
Resolution (Å)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)	30-1.20 (1.22-1.20)
R_{sym} (%)	8.5 (46.4)	5.8 (42.4)	8.6 (33.1)	7.2 (37.7)	8.5 (29.7)	4.7 (22.7)	9.9 (42.8)	7.8 (44.8)
$I/\sigma(I)$	23.0 (4.3)	31.8 (5.6)	20.7 (7.6)	23.1 (7.2)	23.1 (8.4)	40.6 (10.1)	19.5 (6.0)	25.3 (5.5)
Completeness (%)	95.0 (91.7)	94.6 (91.2)	96.6 (93.8)	95.4 (92.1)	97.5 (94.7)	96.1 (93.0)	95.1 (92.3)	96.9 (94.3)
Redundancy	7.6 (7.7)	7.6 (7.6)	7.5 (7.4)	7.6 (7.5)	7.4 (7.3)	7.5 (7.3)	7.7 (7.6)	7.5 (7.5)

Refinement								
Resolution (Å)	1.20	1.20	1.20	1.20	1.20	1.20	1.20	1.20
No. reflections	71,822	72,158	73,007	72,507	73,791	73,217	71,843	73,841
$R_{\text{work}} / R_{\text{free}}$ (%)	12.3 / 14.9	11.3 / 13.9	11.6 / 14.1	11.2 / 13.8	11.7 / 13.9	11.2 / 13.6	11.7 / 14.4	11.3 / 14.2
No. atoms								
Protein	4,252	4,240	4,259	4,240	4,259	4,240	4,259	4,240
Ligand/ion	1 glycerol	1 HCO ₃ ⁻ , 2 CO ₂ , 1 glycerol	1 glycerol	1 HCO ₃ ⁻ , 1 CO ₂ , 1 glycerol	1 glycerol	1 HCO ₃ ⁻ , 1 CO ₂ , 1 glycerol	1 glycerol	1 CO ₂ , 1 glycerol
Water	265	333	279	347	279	346	286	360
<i>B</i> -factors								
Protein (main / side chain)	10.84 / 14.06	10.39 / 13.16	8.73 / 11.74	7.93 / 10.50	9.15 / 12.17	8.38 / 10.83	9.71 / 12.55	8.23 / 10.89
Ligand/ion	21.51 (glycerol)	15.78 (HCO ₃ ⁻), 14.19 (first CO ₂), 21.78 (second CO ₂), 19.39 (glycerol)	18.86 (glycerol)	9.15 (HCO ₃ ⁻), 21.92 (second CO ₂), 14.87 (glycerol)	19.13 (glycerol)	7.40 (HCO ₃ ⁻), 21.95 (second CO ₂), 13.93 (glycerol)	17.19 (glycerol)	19.67 (second CO ₂), 14.81 (glycerol)
Water	26.98	28.72	24.56	26.58	26.51	26.38	26.62	29.36
R.m.s. deviations								
Bond lengths (Å)	0.031	0.028	0.032	0.029	0.030	0.028	0.031	0.029
Bond angles (°)	2.484	2.358	2.600	2.382	2.552	0.352	2.550	2.460
Partial occupancy (%)								
His64 (out/in) conformation (%)	49 / 51	69 / 31	43 / 57	60 / 40	46 / 54	52 / 48	39 / 61	66 / 34

*Values in parentheses are for the highest-resolution shell.

Supplementary Table 2. List of key bound water molecules in the CA II structures.

	apo-CA II 0atm 6LUU	apo-CA II 20atm 6LUV	Zn-CA II 0atm 6LUW	Zn-CA II 20atm 6LUX	Co-CA II 0atm pH11.0 6LV3	Co-CA II 20atm 6LV2	Ni-CA II 0atm 6LV5	Ni-CA II 20atm 6LV6	Cu-CA II 0atm 6LV9	Cu-CA II 20atm 6LVA
CO ₂	–	A 302		A 302	–		–		–	–
HCO ₃ [–]	–	–			–	A 304	–	A 305	–	–
W _{Zn}	–	–	A 481	A 426	–		–	–	–	–
W _{Co, tetra}	–	–	–	–	A 490		–	–	–	–
W _{Co, octa}	–	–	–	–	–	A 485	–	–	–	–
W _{Ni}	–	–	–	–	–	–	A 573	A 583	–	–
W _{Ni'}	–	–	–	–	–	–	A 421	–	–	–
W _{Ni''}	–	–	–	–	–	–	A 602	–	–	–
W _{Cu}	–	–	–	–	–	–	–	–	A 451	A 421
W _{Cu'}	–	–	–	–	–	–	–	–	A 580	–
W _{apo}	A 486	A 475		–	–	–	–	–	–	–
W _{DW}	A 618	–	A 581	–	A 569	–	–	–	A 587	A 496
W _{DW'}	–	–	–	–	–	–	–	–	–	A 695
W _{DW''}	–	–	–	–	–	–	–	–	–	A 675
W _I	–	A 591	–	A 483	–	A 448	A 611	A 495	–	A 427
W _{I'}	–	–	–	A 643	–	–	–	–	–	A 663
W _I	A 591	–	A 487		A 546	–	–	–	A 563	
W ₂	A 641	A 666	A 615	A 678	A 584	A 667	A 627	A 662	A 636	A 681
W _{2'}	–	–	–	A 401	–	A 401	–	–	–	A 401
W _{3a}	A 541	A 556	A 518	A 543	A 529	A 537	A 532	A 507	A 540	A 528
W _{3b}	A 462	A 430	A 451	A 451	A 454	A 432	A 453	A 479	A 442	A 452
W _{3b'}	–	–	–	A 695	–	–	–	–	–	A 684
W _{3bapo'}	–	A 606	–	–	–	–	–	–	–	–
W _{3bapo'}	–	A 703	–	–	–	–	–	–	–	–
W _{3bCo}	–	–	–	–	–	A 648	–	–	–	–
W _{EC1}	A 680	–	A 614	A 665	A 611	A 630	–	–	A 619	A 648
W _{EC1'}	–	–	–	A 702	–	–	–	–	–	A 699
W _{EC1''}	–	–	–	–	–	–	A 598	A 626	–	–
W _{EC2}	A 668	A 700	A 608	–	A 609	–	–	–	A 603	A 641
W _{EC2'}	–	–	–	A 697	–	A 661	A 649	A 669	–	A 697
W _{EC2''}	–	–	–	–	–	–	A 613	A 642	–	–
W _{EC2'''}	–	–	–	–	–	A 673	–	–	–	–
W _{EC3}	A 707	A 712	A 634	A 720	A 639	A 706	A 658	A 704	A 649	A 703
W _{EC3'}	–	A 714	–	–	–	–	–	–	–	–
W _{EC3''}	–	A 710	–	–	–	–	–	–	–	–
W _{EC4}	A 761	A 760	A 655	A 755	A 660	A 720	A 672	A 734	A 676	
W _{EC5}	A 447	A 421	A 438	A 417	A 437	A 430	A 442	A 428	A 420	A 502

Supplementary Table 3. Distance geometry (Å) of CO₂, HCO₃⁻ and key bound water molecules in the CA II structures.

	apo- CA II 0atm 6LUU	apo- CA II 20atm 6LUV	Zn-CA II 0atm 6LUW	Zn-CA II 20atm 6LUX	Co-CA II 0atm pH11.0 6LV3	Co-CA II 20atm 6LV2	Ni-CA II 0atm 6LV5	Ni-CA II 20atm 6LV6	Cu-CA II 0atm 6LV9	Cu-CA II 20atm 6LVA
Zn – W _{Zn}	–	–	1.88	1.92	–	–	–	–	–	–
Zn – CO ₂ (O1)	–	–	–	3.31	–	–	–	–	–	–
Co – W _{Co, tetra}	–	–	–	–	1.72	–	–	–	–	–
Co – W _{Co, octa}	–	–	–	–	–	2.03	–	–	–	–
Co – CO ₂ (O1)	–	–	–	–	–	–	–	–	–	–
Co – HCO ₃ ⁻ (O1)	–	–	–	–	–	2.07	–	–	–	–
Co – HCO ₃ ⁻ (O3)	–	–	–	–	–	2.02	–	–	–	–
Ni – W _{Ni}	–	–	–	–	–	–	2.09	2.13	–	–
Ni – W _{Ni'}	–	–	–	–	–	–	2.14	–	–	–
Ni – W _{Ni''}	–	–	–	–	–	–	2.22	–	–	–
Ni – HCO ₃ ⁻ (O1)	–	–	–	–	–	–	–	2.16	–	–
Ni – HCO ₃ ⁻ (O3)	–	–	–	–	–	–	–	2.18	–	–
Cu – W _{Cu}	–	–	–	–	–	–	–	–	2.17	2.26
Cu – W _{Cu'}	–	–	–	–	–	–	–	–	2.36	–
Cu – W _{DW''}	–	–	–	–	–	–	–	–	–	3.14
CO ₂ (C1) – W _{Zn}	–	–	–	2.87	–	–	–	–	–	–
CO ₂ (C1) – W _{Co, tetra}	–	–	–	–	–	–	–	–	–	–
HCO ₃ ⁻ (O1) – W _{Co, octa}	–	–	–	–	–	2.85	–	–	–	–
HCO ₃ ⁻ (O3) – W _{Co, octa}	–	–	–	–	–	2.72	–	–	–	–
HCO ₃ ⁻ (O1) – W _{Ni}	–	–	–	–	–	–	–	2.92	–	–
HCO ₃ ⁻ (O3) – W _{Ni}	–	–	–	–	–	–	–	2.85	–	–
CO ₂ (C1) – W _{apo}	–	2.94	–	–	–	–	–	–	–	–
W _{Zn} – W _{DW}	–	–	2.58	–	–	–	–	–	–	–
W _{Zn} – W1	–	–	2.63	–	–	–	–	–	–	–
W _{Zn} – W _I	–	–	–	2.75	–	–	–	–	–	–
W _{Zn} – W2	–	–	4.41	4.65	–	–	–	–	–	–
W _{Co, tetra} – W _{DW}	–	–	–	–	2.61	–	–	–	–	–
W _{Co, tetra} – W1	–	–	–	–	2.73	–	–	–	–	–
W _{Co, tetra} – W _I	–	–	–	–	–	–	–	–	–	–
W _{Co, tetra} – W2	–	–	–	–	4.55	–	–	–	–	–
W _{Co, octa} – W1	–	–	–	–	–	2.75	–	–	–	–
W _{Co, octa} – W2	–	–	–	–	–	2.71	–	–	–	–
W _{Co, octa} – W2'	–	–	–	–	–	4.90	–	–	–	–
W _{Ni} – W _{Ni'}	–	–	–	–	–	–	2.82	–	–	–
W _{Ni} – W _{Ni''}	–	–	–	–	–	–	3.08	–	–	–
W _{Ni'} – W _{Ni''}	–	–	–	–	–	–	2.78	–	–	–
W _{Ni} – W _I	–	–	–	–	–	–	2.95	2.88	–	–
W _{Ni} – W2	–	–	–	–	–	–	2.73	2.68	–	–
W _{Cu} – W1	–	–	–	–	–	–	–	–	2.54	–
W _{Cu} – W _{Cu'}	–	–	–	–	–	–	–	–	3.06	–

W _{Cu} – W _{DW}	–	–	–	–	–	–	–	–	3.23	2.84
W _{Cu'} – W _{DW}	–	–	–	–	–	–	–	–	2.84	–
W _{DW} – W _{DW'}	–	–	–	–	–	–	–	–	–	1.52
W _{DW'} – W _{DW''}	–	–	–	–	–	–	–	–	–	1.60
W _{Cu} – W _I	–	–	–	–	–	–	–	–	–	2.84
W _{Cu} – W ₂	–	–	–	–	–	–	–	–	3.44	3.90
W _{apo} – W _{DW}	2.71	–	–	–	–	–	–	–	–	–
W _{apo} – W ₁	2.82	–	–	–	–	–	–	–	–	–
W _{apo} – W _I	–	2.88	–	–	–	–	–	–	–	–
W _{apo} – W ₂	4.90	3.96	–	–	–	–	–	–	–	–
W ₁ – W ₂	2.73	–	2.74	–	2.70	–	–	–	2.78	–
W ₁ – W _{I'}	–	–	–	2.02	–	–	–	–	–	2.09
W ₂ – W _{2'}	–	–	–	1.43	–	2.79	–	–	–	1.51
W ₂ – W _{3a}	2.75	2.80	2.77	3.12	2.77	3.40	2.81	2.72	2.81	3.14
W ₂ – W _{3b}	2.72	2.99	2.73	2.49	2.68	2.41	2.95	3.08	2.79	2.57
W ₂ – W _{3b'}	–	–	–	1.68	–	2.69	–	–	–	1.65
W _{3b} – W _{3b'}	–	–	–	1.37	–	1.86	–	–	–	1.39
Thr199(N) – HCO ₃ ⁻ (O2)	–	–	–	–	–	2.93	–	3.05	–	–
Thr199(N) – CO ₂ (O2)	–	3.20	–	3.62	–	–	–	–	–	–
Thr199(N) – W _{DW}	2.93	–	2.92	–	2.90	–	–	–	2.90	2.74
Thr199(N) – W _I	–	3.34	–	3.52	–	3.71	3.04	3.52	–	3.68
Thr199(Oγ1) – HCO ₃ ⁻ (O3)	–	–	–	–	–	2.44	–	2.46	–	–
Thr199(Oγ1) – W _{Zn}	–	–	2.73	2.62	–	–	–	–	–	–
Thr199(Oγ1) – W _{Co, tetra}	–	–	–	–	2.75	–	–	–	–	–
Thr199(Oγ1) – W _{Co, octa}	–	–	–	–	–	3.75	–	–	–	–
Thr199(Oγ1) – W _{Ni}	–	–	–	–	–	–	3.68	3.65	–	–
Thr199(Oγ1) – W _{Ni'}	–	–	–	–	–	–	2.62	–	–	–
Thr199(Oγ1) – W _{Ni''}	–	–	–	–	–	–	5.04	–	–	–
Thr199(Oγ1) – W _{Cu}	–	–	–	–	–	–	–	–	2.68	2.61
Thr199(Oγ1) – W _{Cu'}	–	–	–	–	–	–	–	–	4.37	–
Thr199(Oγ1) – W _{apo}	2.72	2.69	–	–	–	–	–	–	–	–
Thr199(Oγ1) – Glu106(Oε1)	2.60	2.59	2.61	2.56	2.62	2.53	2.91	2.55	2.61	2.59
Glu106(Oε1) – W _{Zn}	–	–	4.15	4.01	–	–	–	–	–	–
Glu106(Oε1) – W _{Co, tetra}	–	–	–	–	4.08	–	–	–	–	–
Glu106(Oε1) – W _{Co, octa}	–	–	–	–	–	4.85	–	–	–	–
Glu106(Oε1) – W _{Ni}	–	–	–	–	–	–	4.75	4.84	–	–
Glu106(Oε1) – W _{Ni'}	–	–	–	–	–	–	2.78	–	–	–
Glu106(Oε1) – W _{Cu}	–	–	–	–	–	–	–	–	4.20	4.19

Glu106(Oε1) – W _{apo}	4.01	4.09	–	–	–	–	–	–	–	–
His64 _{in} (Nδ1) – W2	3.02	3.16	3.22	3.28	2.94	3.88	3.29	3.22	3.43	3.48
His64 _{in} (Nδ1) – W2'	–	–	–	1.86	–	1.26	–	–	–	1.99

Supplementary References

1 Sjoebloom, B., Polentarutti, M. & Djinovic-Carugo, K. Structural study of X-ray induced activation of carbonic anhydrase. *Proc Natl Acad Sci U S A* **106**, 10609-10613, doi:10.1073/pnas.0904184106 (2009).