

Supplementary Data

From Cluster Halides to Catalysts: Nanostructured Molybdenum Carbides for Efficient Hydrogen Evolution Reaction

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Fig. S13. Comparison between the XRD patterns of the AHM precursor (in blue) and that of the product (AHM + sucrose (S)) after treatment at 100°C (in orange).

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Table S1. Details of the Rietveld refinements and crystallites size for Mo₂C-AHM, Mo₂C-HMC and Mo₂C-TMC obtained with the FullProf Suite Software.

	Mo ₂ C-AHM	Mo ₂ C-HMC	Mo ₂ C-TMC
Lattice			
<i>a</i> (Å)	4.747(1)	4.726(1)	4.741(1)
<i>b</i> (Å)	6.004(1)	6.015(3)	6.015(2)
<i>c</i> (Å)	5.196(1)	5.198(1)	5.205(1)
<i>V</i> (Å ³)	148.1(1)	147.8(1)	148.1(1)
Space group	<i>Pbcn</i>	<i>Pbcn</i>	<i>Pbcn</i>
Refinement			
N0. Diffraction peaks (K α /K β)	110	110	110
N0. Background points	28	31	35
N0. Refined parameters	11	40	42
N0. atoms	0	0	0
R _p (%)	34.7	32.1	44.3
R _{wp} (%)	24.3	27.8	27.6
R _{exp} (%)	12.6	11.2	11.5
χ^2	3.74	6.19	5.77
Microstructure			
Crystallites size (nm)	14.3(1)	30.3(1)	26.9(1)

Table S2: Atomic ratio and stoichiometry of Mo carbides species calculated from XPS data.

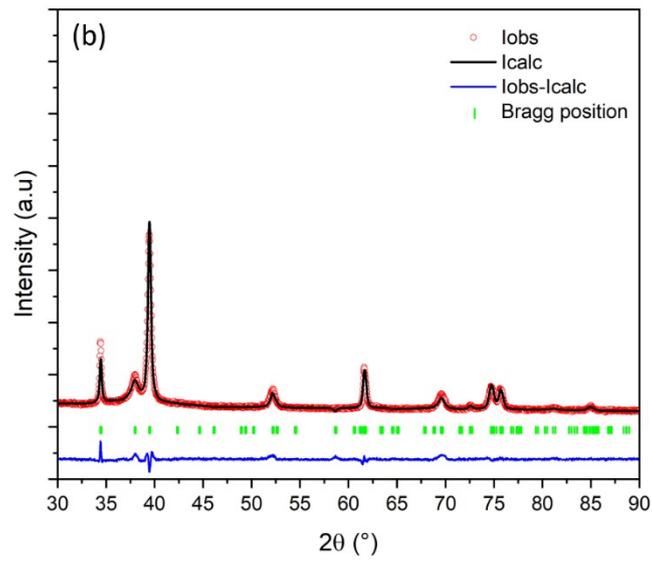
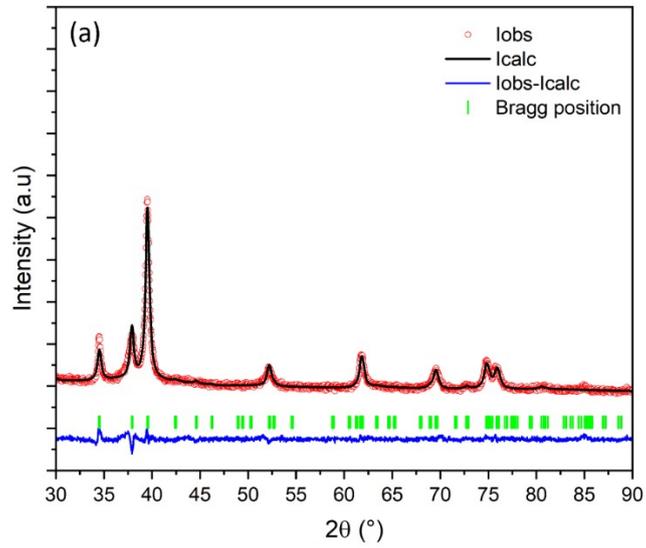
	atomic %C sp ² ^a	Mo/C ^b	Average stoichiometry (x) of Mo _x C species ^c	Mo ²⁺ /Mo ³⁺ ^d
Mo ₂ C-TMC	72.6	1.79	1.76	3.1
Mo ₂ C-HMC	79.5	1.66	1.58	1.4
Mo ₂ C-AHM	51.3	1.62	1.63	1.7

^aversus total carbon at the surface. ^bratio calculated by taking account the Mo carbides species only from C1s and Mo3d signal. ^ccalculated from XPS atomic ratio from decomposition of Mo3d signal according to (%at.Mo²⁺x2 + %at.Mo³⁺) / (%at.Mo²⁺+ %at.Mo³⁺). ^dratio calculated form atomic% of Mo²⁺ and Mo³⁺

Table S3. Comparison of the electrocatalytic performance of Mo₂C-TMC with other HER catalysts reported in the literature in alkaline solutions.

Catalyst	Precursor	SSA (m ² .g ⁻¹)	η_{10} (mV)	Tafel Slope (mV.dec ⁻¹)	Electrolyte	Charge loading (mg.cm ⁻²)	Ref
Mo₂C-TMC	(TBA) ₂ Mo ₆ Cl ₁₄ sucrose	103 ± 1	212	95	KOH 1 M	0.15	This work
Mo₂C@NCS	Ammonium molybdate, glucose, melamine, organic acid	101	147	77	KOH 1 M	0.25	1
β-Mo₂C nanotubes	(NH ₄) ₆ [Mo ₇ O ₂₄].4H ₂ O, hydrochloric dopamide	127	112	55	KOH 0.1 M	0.75	2
Mo₂C@NPC	PMo ₁₂ , melamine, silicate spheres	47	121	73.5	KOH 1 M	0.57	3
α-Mo₂C	MoCl ₅ , urea	9	176	58	KOH 0.1 M	0.102	4
γ-Mo₂N	MoCl ₅ , urea	10	353	108	KOH 0.1 M		4
NiMo	(NH ₄) ₆ [Mo ₇ O ₂₄].4H ₂ O and nickel nitrate	N/D	100	ND	KOH 2 M	0.1	5
Mo₂C	Glucose, NaCl and (NH ₄) ₆ Mo ₇ O ₂₄ .4H ₂ O	N/D	430	143	KOH 1 M	ND	6
Mo₂C-C	(NH ₄) ₆ Mo ₇ O ₂₄ .6H ₂ O, glucose and ethylene glycol	N/D	149	66	KOH 1 M	0.84	7
Mo/Mo₂C/N-CNFs	MoO ₂ (acac), cellulose acetate, polyacrylonitrile	N/D	162	47.9	KOH 1 M	1.68	8
Mo₂C-GNR	(NH ₄) ₆ [Mo ₇ O ₂₄].4H ₂ O, glucose, MWNTs	N/D	217	64	NaOH 1 M	0.28	9
Mo₂C@CS	(NH ₄) ₆ [Mo ₇ O ₂₄].4H ₂ O, glucose	816	178	50	KOH 1 M	0.4	10
Mo₂(CN)	(NH ₄) ₆ [Mo ₇ O ₂₄].4H ₂ O,	148	127	50	KOH 1 M	0.2	11

	melamine						
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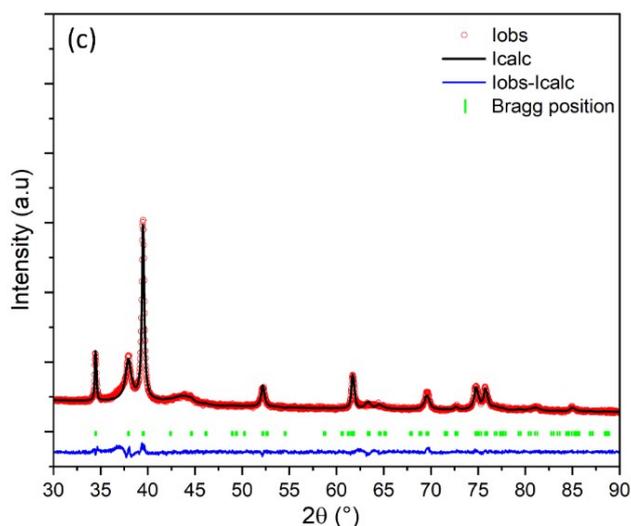


Fig. S1. LeBail refinement patterns for Mo₂C-AHM (a), Mo₂C-HMC (b) and Mo₂C-TMC (c): observed (red dotted line), calculated (black line) and difference X-ray powder diffraction profiles from the pattern matching plot obtained with the FullProf Suite Software. The vertical green markers correspond to the position of the Bragg reflections.

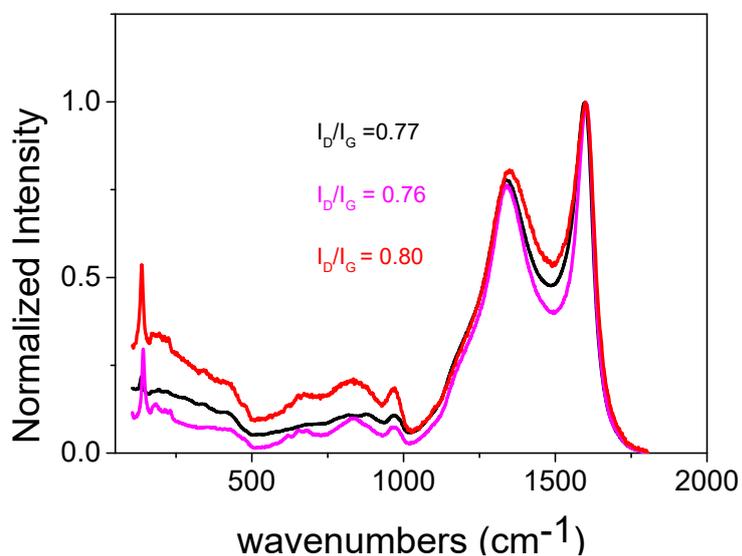


Fig. S2. Normalized Raman spectra for Mo₂C-TMC (black line); Mo₂C-HMC (magenta line) and Mo₂C-AHM (red line). The G and D bands due to carbon are clearly observed at 1597 and 1340-1348 cm⁻¹, respectively with a small downshift of the latter for the clusters- based compounds.

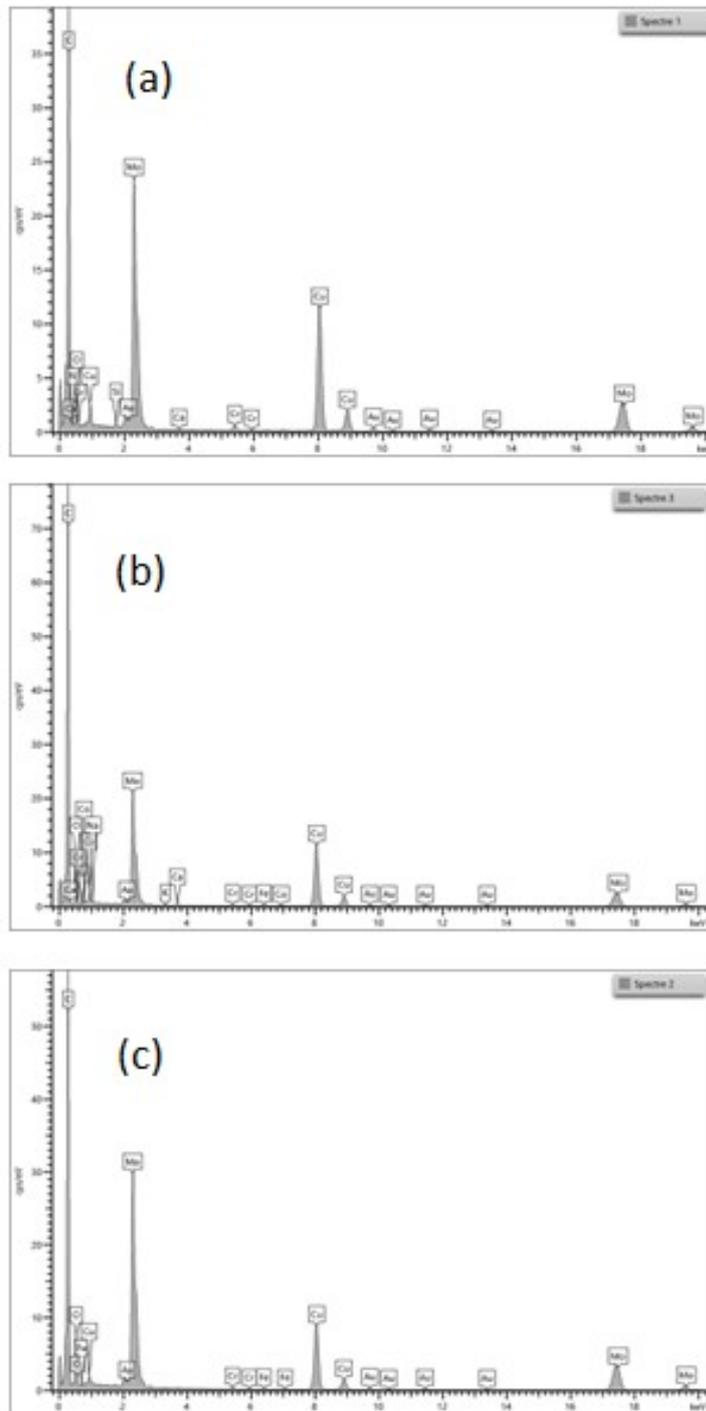


Fig. S3. EDS-TEM for the Mo₂C-AHM (a), Mo₂C-HMC (b) and Mo₂C-TMC (c). Cu, Fe, Cr and Au elements correspond to the material used for TEM analyses (Cu grid and specimen holder).

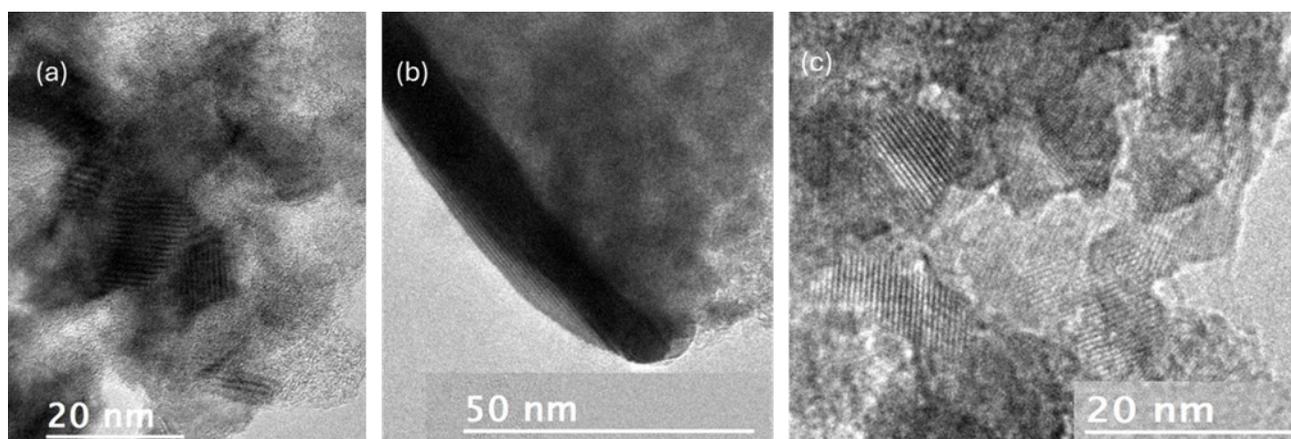


Fig. S4. HRTEM images of Mo₂C-AHM (a), Mo₂C-HMC (b) and Mo₂C-TMC (c) showing the lattice fringes of nanocrystals.

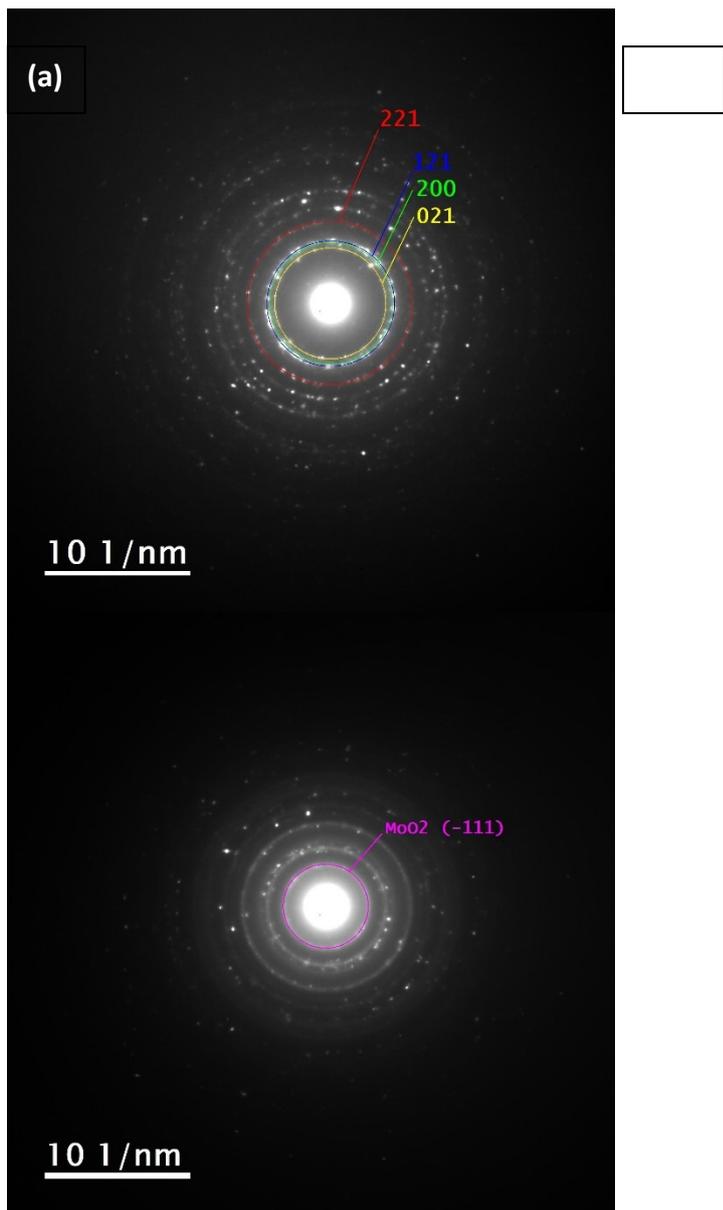


Fig. S5. Indexed SAED patterns displaying the (021), (200), (121) and (221) reflexions for Mo₂C-AHM (a), and (-111) reflexion of MoO₂ for Mo₂C-HMC (b).

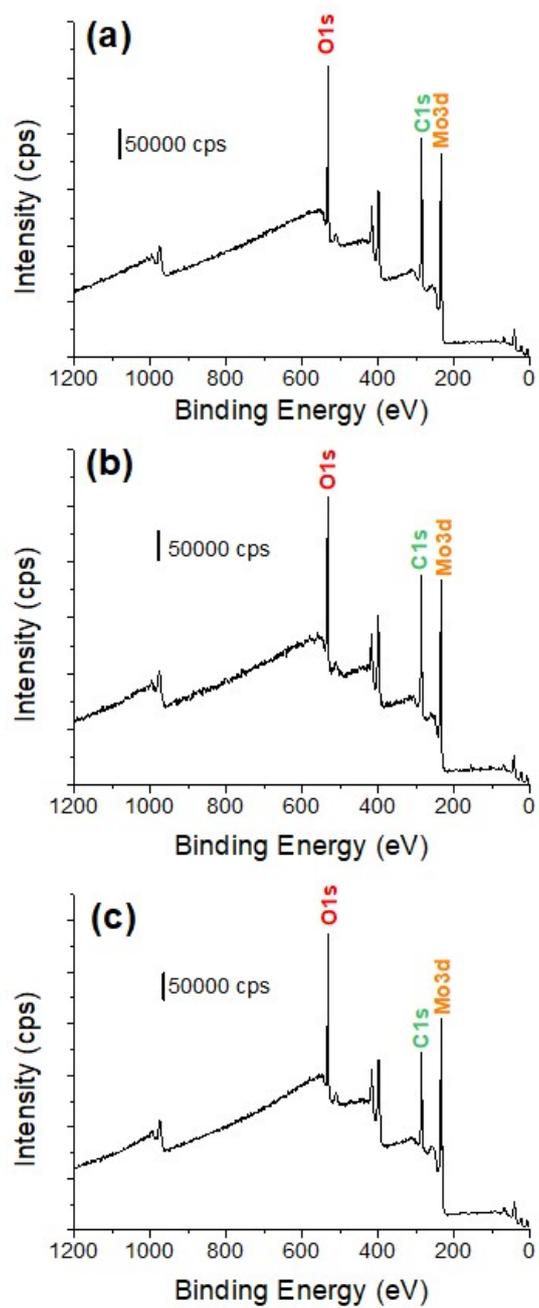


Fig. S6. XPS survey spectra for Mo₂C-TMC (a), Mo₂C-HMC (b) and Mo₂C-AHM (c)

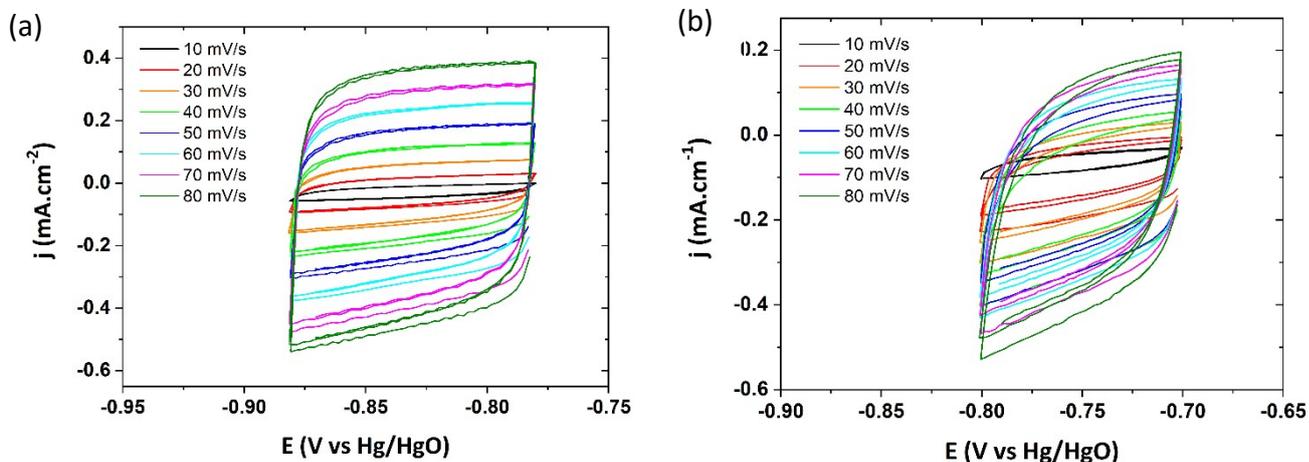


Fig. S7. Cyclic voltammeteries in a non-faradaic region of the Mo₂C-AHM (a) before and (b) after 18 -chronopotentiometry at different scan rates (10, 20, 30, 40, 50, 60, 70 and 80 mV/s) in aqueous 1 M KOH.

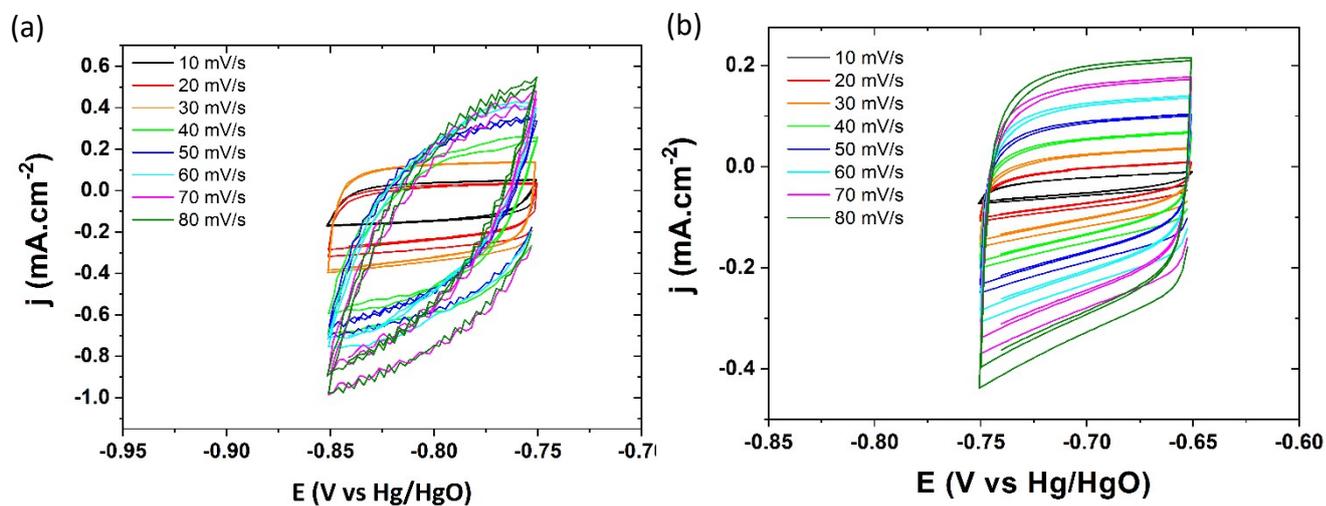


Fig. S8. Cyclic voltammeteries in a non-faradaic region of the Mo₂C-HMC (a) before and (b) after 18h-chronopotentiometry at different scan rates (10, 20, 30, 40, 50, 60, 70 and 80 mV/s) in aqueous 1 M KOH.

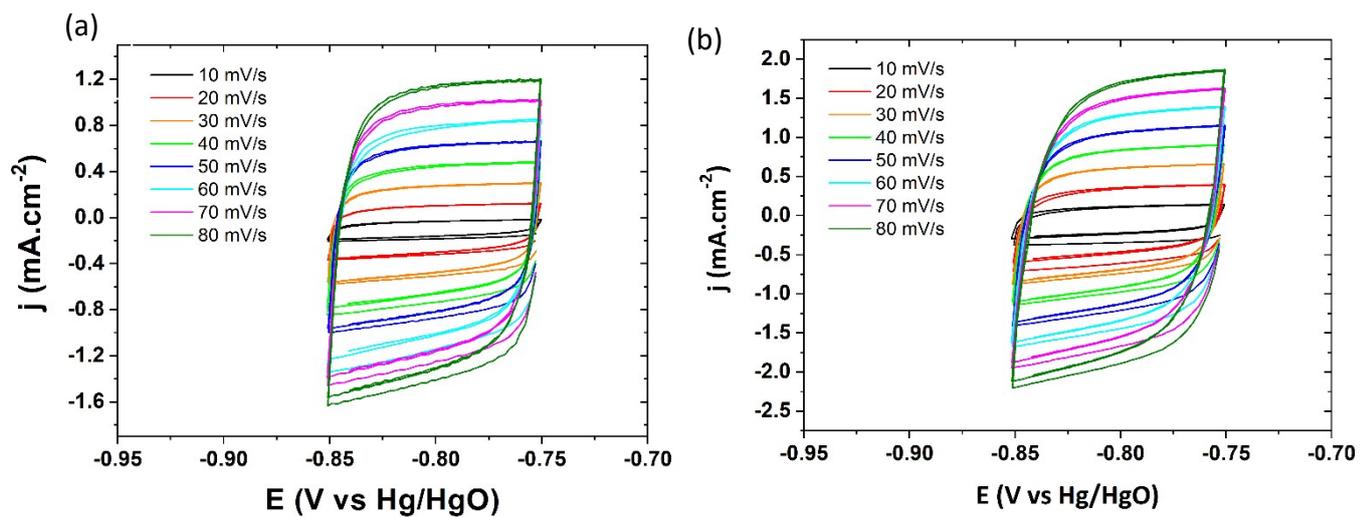


Fig. S9. Cyclic voltammeteries in a non-faradaic region of the $\text{Mo}_2\text{C-TMC}$ (a) before and (b) after 18h-chronopotentiometry at different scan rates (10, 20, 30, 40, 50, 60, 70 and 80 mV/s) in aqueous 1 M KOH.

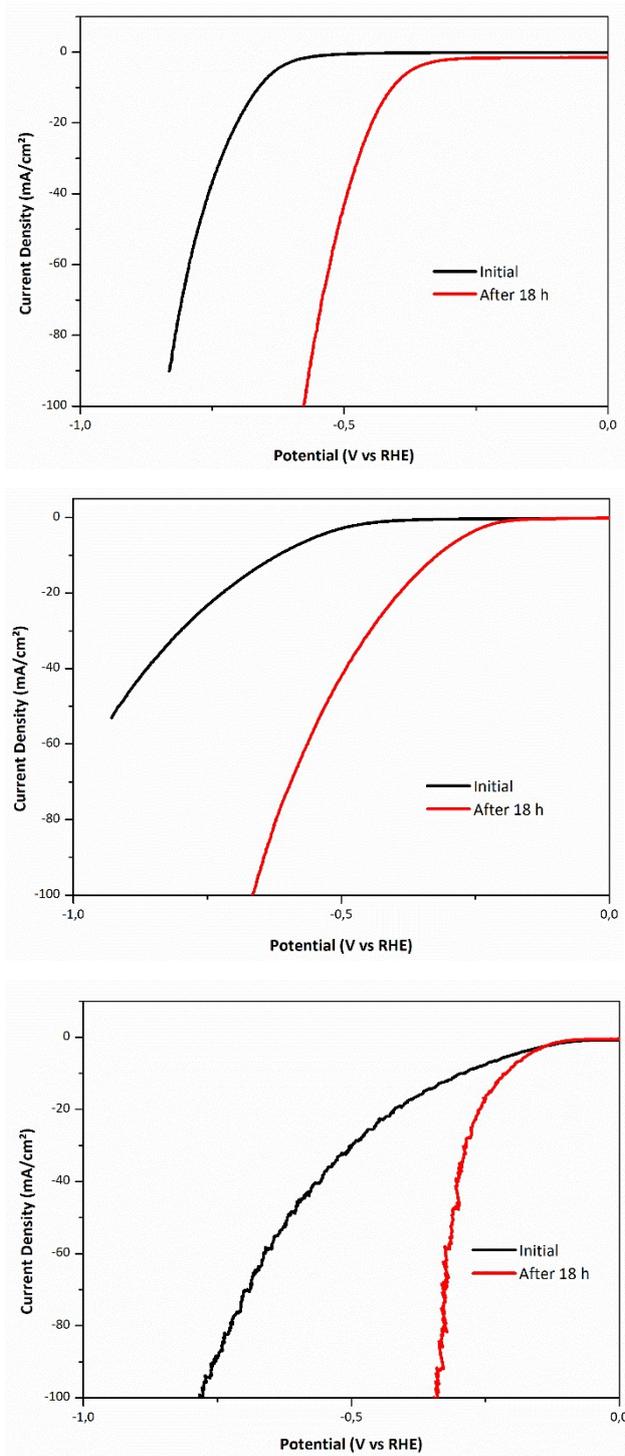


Fig. S10. iR-corrected HER polarization curves in aqueous 1 M KOH at 10 mV/s for Mo₂C-TMC (bottom), Mo₂C-HMC (middle), Mo₂C-AHM (top) before and after 18h-chronopotentiometry.

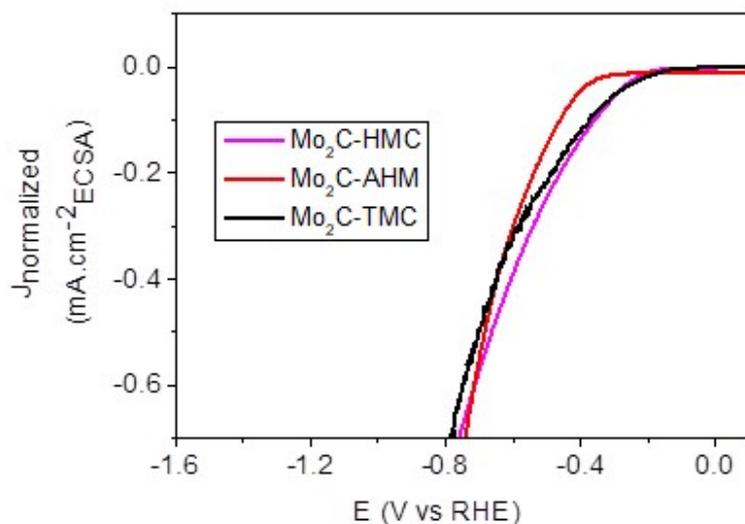


Fig. S11. iR-corrected HER polarization curves normalized by ECSA in aqueous 1 M KOH at 10 mV/s for Mo₂C-TMC, Mo₂C-HMC, Mo₂C-AHM after 18h-chronopotentiometry.

Note that ECSA value is roughly estimated here from C_{dl}/C_s where C_s is the specific capacitance. This value is taken equal to 0.040 mF/cm² following Jaramillo et al.¹² this value is considered as a typical standard value (measured for metallic planar electrode) because we do not know the true C_s value. The electrochemical capacitance is estimated through CVs in non-Faradaic regions.

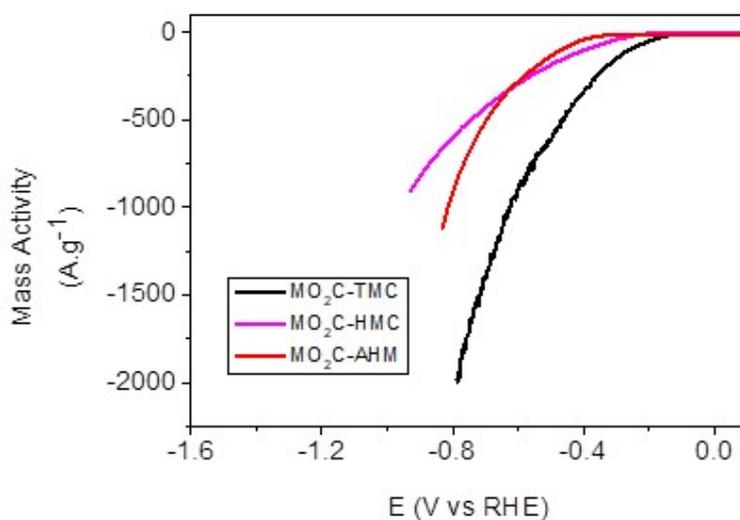


Fig. S12. iR-corrected HER polarization curves normalized by mass of the catalyst in aqueous 1 M KOH at 10 mV/s for Mo₂C-TMC, Mo₂C-HMC, Mo₂C-AHM after 18h-chronopotentiometry – Mass Activity.

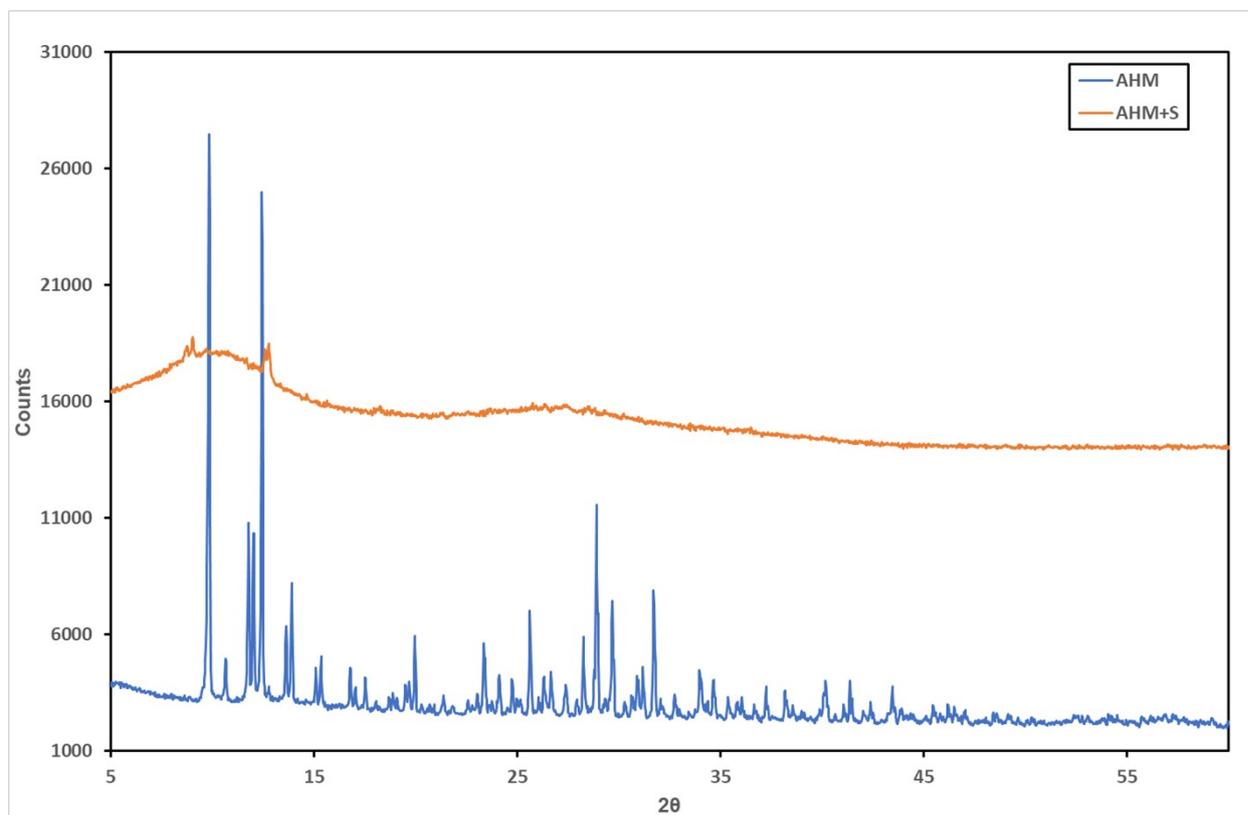


Fig. S13. Comparison between the XRD patterns of the AHM precursor (in blue) and that of the product (AHM + sucrose (S)) after treatment at 100°C (in orange).

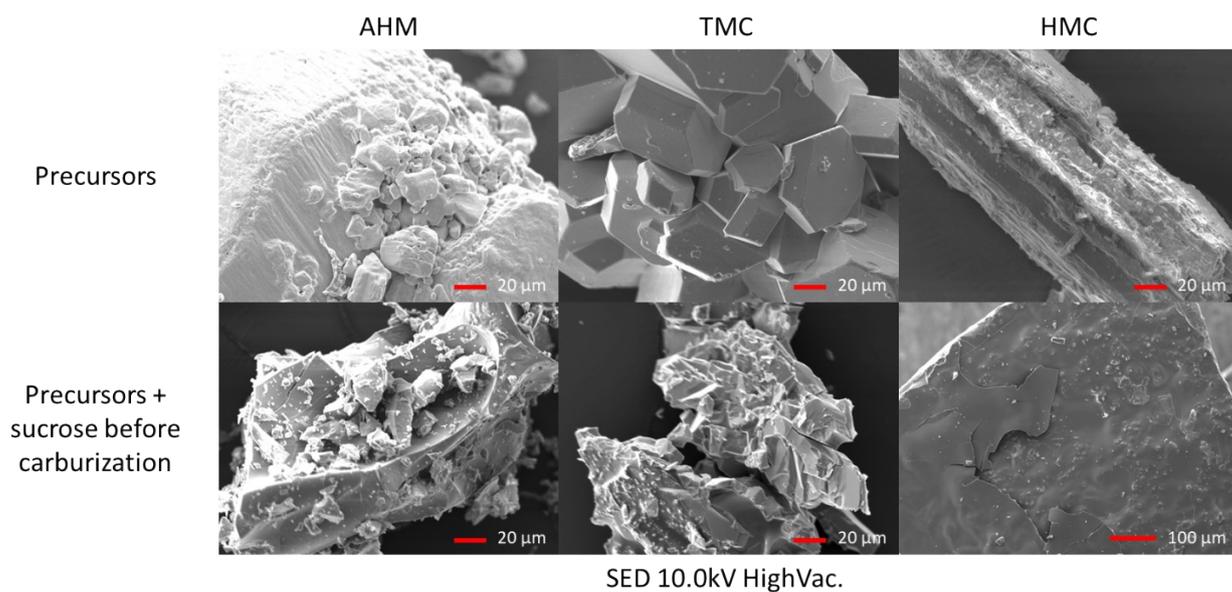


Fig. S14. SEM of the precursors (AHM, TMC and HMC) before (first line) and after (second line) sucrose addition

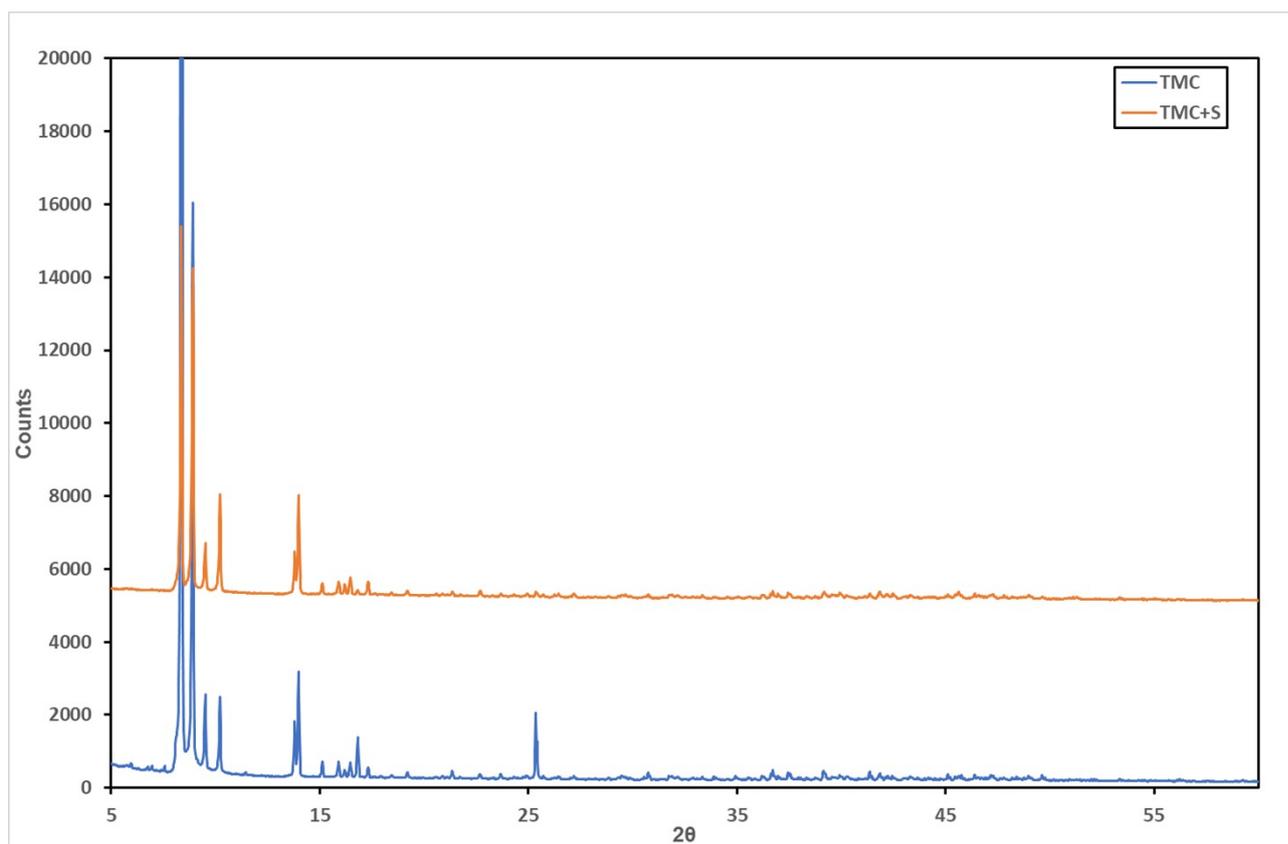


Fig. S15. Comparison between the XRD patterns of the TMC precursor (in blue) and that of the product (TMC + sucrose (S)) after treatment at 100°C (in orange).

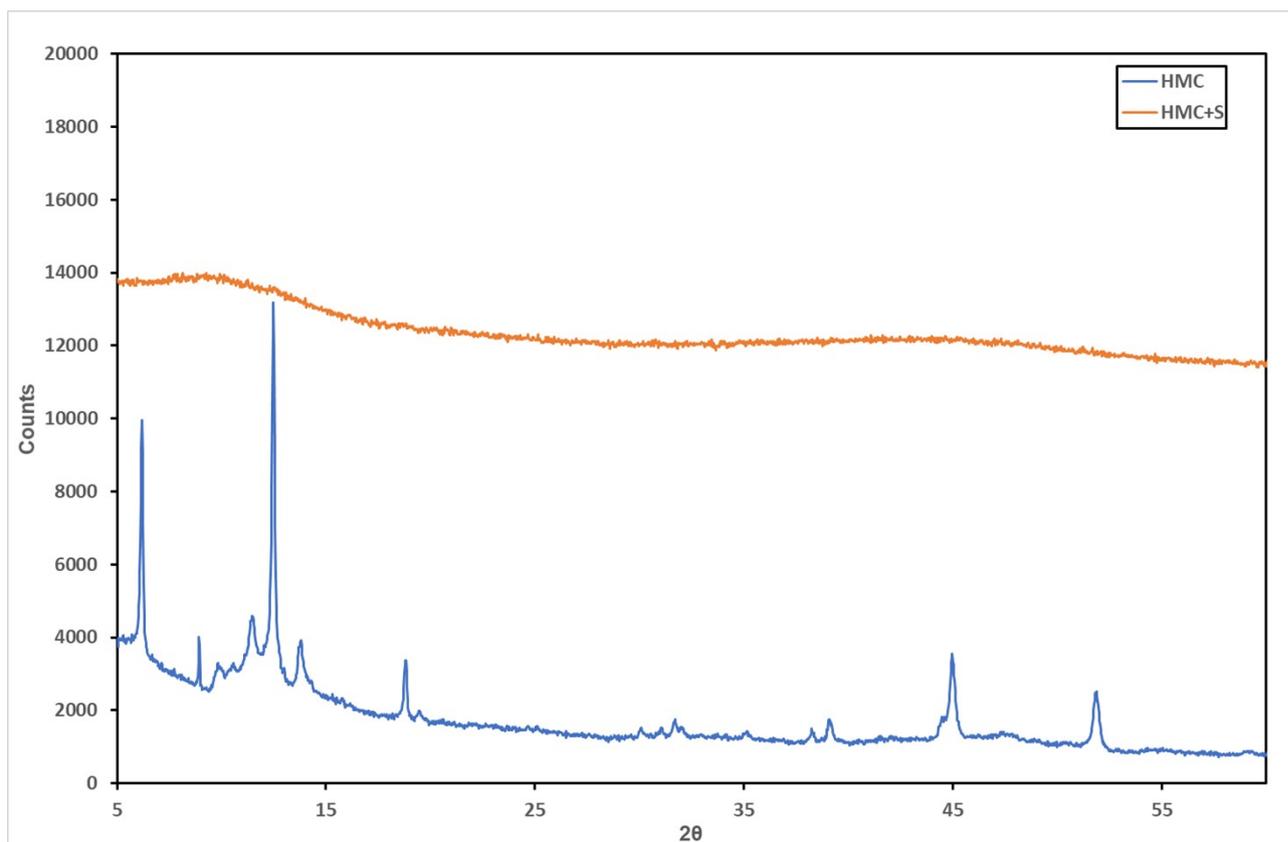


Fig. S16. Comparison between the XRD patterns of the HMC precursor (in blue) and that of the product (HMC + sucrose (S)) after treatment at 100°C (in orange).

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