Nitrogen-fixation catalyst based on graphene: every part counts

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Table S1. The bond distance, bond angle of the MoN_3 part (the label of N atoms is shown in Figure S1) in $C_{33}H_{15}MoN_3$, $C_{69}H_{21}MoN_3$ and $C_{117}H_{27}MoN_3$.

bond length	Mo-N1	Mo-N2	Mo-N3	Mo-N1N2	Mo-N2N3	Mo-N1N3	NNN-Mo
C ₃₃ H ₁₅ MoN ₃	1.96	1.96	1.96	91.3	91.3	91.3	53.8
$C_{69}H_{21}MoN_3$	1.96	1.96	1.96	91.3	91.3	91.3	53.8
C ₁₁₇ H ₂₇ MoN ₃	1.95	1.95	1.95	91.8	91.8	91.8	53.4

Table S2. The bond distances of the MoN₃ part (the label of N atoms is shown in Figure S1) of $C_{33}H_{15}MoN_3/N_2$ and $C_{69}H_{21}MoN_3/N_2$ before and after the addition of the first H⁺ in the enzymatic catalytic reaction path.

Bond	N	$N_2(ad)$	Н	1	Н	1R
length	C ₃₃ H ₁₅ Mol	$N_3 C_{69} H_{21} Mo N_3$	C ₃₃ H ₁₅ MoN ₃	C ₆₉ H ₂₁ MoN ₃	C ₃₃ H ₁₅ MoN ₃	C ₆₉ H ₂₁ MoN ₃
Mo-N1	2.11	2.10	2.13	2.12	2.07	2.07
Mo-N2	1.99	1.98	2.02	2.02	2.05	2.05
Mo-N3	1.99	1.98	2.02	2.00	2.05	2.05
Να-Νβ	1.21	1.21	1.29	1.28	1.30	1.30
Μο-Να	2.02	2.02	1.90	1.97	1.91	1.91
Μο-Νβ	2.04	2.03	2.01	1.98	2.04	2.04
Νβ-Η			1.02	1.02	1.02	1.02

Table S3. Mulliken charge of the three groups (the division into three groups is illustrated in Figures S1 and S2) in $C_{33}H_{15}MoN_3$, $C_{69}H_{21}MoN_3$ and $C_{117}H_{27}MoN_3$.

net charge	grp3(t)	grp3(a)	grp2(t)	grp2(a)	Mo	N1	N2	N3
C ₃₃ H ₁₅ MoN ₃	1.24	0.082	0.00	0.000	0.86	-0.70	-0.70	-0.70
C ₆₉ H ₂₁ MoN ₃	1.74	0.083	-0.43	-0.006	0.90	-0.73	-0.73	-0.73
C ₁₁₇ H ₂₇ MoN ₃	2.27	0.084	-0.94	-0.008	0.89	-0.74	-0.74	-0.74

Table S4. Variation of Mulliken charge of different groups (grp) in Mo/N graphenes

before and after loss of an electron.

charge variation	$C_{33}H_{15}MoN_3$	$C_{69}H_{21}MoN_3$	C117H27MoN3
grp1	0.05	0.03	-0.02
grp2	0.37	0.43	0.51
grp3	0.58	0.54	0.51

Table S5. Reaction energies(kJ/mol), N-N bond length (Å) and bond order in the
Schrock(Schrk) and enzymatic(Enzym) reaction as shown in Figure 3. Hn is the addition
of the nth proton and HnR is the reduction after the nth protonation (as shown in Figure
6). The number (1-15) in parenthesis of reaction step is the order of reaction process in

Figure 2. S in parenthesis stands for Schrock and E in parenthesis stands for enzymatic.

Reaction s	tepReaction	energy	bond let	ngth/N-N	bond or	der/N-Nt	otal bond	order/Mo
	Schrk	Enzym	Schrk	Enzym	Schrk	Enzym	Schrk	Enzym
N_2 +Mo(1)	0	0					4.17	4.17
$N_2(ad)(2)$	-104.1	-143.9	1.13	1.21	2.58	2.03	4.53	4.94
H1(3)	-131.1	-156.0	1.23	1.29	1.75	1.53	4.81	5.28
H1R(4)	-181.4	-211.3	1.24	1.30	1.74	1.47	4.48	4.65
H2(5)	-323.1	-294.7	1.32	1.38	1.25	1.21	4.59	4.51
H2R(6)	-331.6	-326.4	1.33	1.39	1.20	1.18	4.79	4.93
H3(7)	-228.2	-420.4	1.44	1.42	0.97	1.05	4.73	4.59
H3R(8)	-318.6	-484.9	1.43	1.43	0.99	1.02	4.43	4.15
H4(9)	-678.3	-516.8	2.91	1.46	0.03	1.01	4.54	3.89
H4R(10)	-682.2	-497.0	3.01	1.46	0.02	1.00	4.73	4.33
NH(S)/H5(E)(1	1) -661.3	-789.8		3.02				
H5(S)/H5R(E)(12) -758.5	-858.7		3.09				
$H5R(S)/NH_2(E)$	(13) -839.0	-839.0						
H6(14)	-898.8	-898.8						
H6R(15)	-890.3	-890.3						

Table S6. Variation of Mulliken charge of the three groups (as shown in Figure 5) in the catalyst during the enzymatic protonation of N_2 (the charge difference of the present step from that of the previous step). The number in parenthesis is the average charge variation per atom. More effective numbers of digit is kept for some values for comparison.

charge variation	1 st	2 nd	3 rd	4 th	5 th	6 th	average
grp1	0.12	0.09	0.07	-0.12	0.14	-0.16	0.022
(active center)	(0.03)	(0.02)	(0.02)	(-0.03)	(0.03)	(-0.04)	(0.006)
grp2	0.05	0.02	-0.048	-0.06	0.23	-0.04	0.025
(transmitter)	(0.002)	(0.001)	(-0.002)	(-0.002)	(0.01)	(-0.002)	(0.001)
grp3	0.75	0.79	0.65	0.66	1.00	0.66	0.75
(reservoir)	(0.031)	(0.03)	(0.027)	(0.03)	(0.04)	(0.03)	(0.031)
total	0.92	0.90	0.67	0.48	1.37	0.46	0.80

Table S7. Variation of Mulliken charge (the charge difference of the present step from that of the previous step) of the three groups (shown in Figure 5) in the catalyst during the enzymatic reduction.

charge variation	1 st	2 nd	3 rd	4 th	5 th	6 th	average
grp1	-0.04	-0.07	-0.09	-0.05	-0.04	-0.06	-0.06
(active center)	(-0.01)	(-0.02)	(-0.02)	(-0.01)	(-0.01)	(-0.02)	(-0.02)
grp2	-0.02	-0.08	0.03	-0.08	-0.07	-0.09	-0.05
(transmitter)	(-0.001)	(-0.003)	(0.001)	(-0.003)	(-0.003)	(-0.004)	(-0.002)
grp3	-0.82	-0.79	-0.79	-0.82	-0.84	-0.83	-0.81
(reservoir)	(-0.03)	(-0.03)	(-0.03)	(-0.03)	(-0.04)	(-0.04)	(-0.03)
total	-0.88	-0.94	-0.85	-0.95	-0.95	-0.98	-0.92

Table S8. Variation of natural charge of the three groups (as shown in Figure 5) in the catalyst during the enzymatic protonation of N_2 (the charge difference of the present step from that of the previous step) in $C_{33}H_{15}MoN_3$.

charge variation	1 st	2 nd	3 rd	4 th	5 th	6 th
grp1 (active center)	-0.13	0.04	0.03	-0.01	0.26	-0.12
grp2 (transmitter)	0.26	-0.03	0.00	-0.04	0.12	-0.15
grp3 (reservoir)	1.04	0.78	0.62	0.52	1.22	0.74
total	1.17	0.79	0.65	0.47	1.6	0.47

Table S9. Variation of natural charge (the charge difference of the present step from that of the previous step) of the three groups (shown in Figure 5) in the catalyst during the enzymatic reduction.

charge variation	1 st	2 nd	3 rd	4 th	5 th	6 th
grp1 (active center)	0.01	-0.02	-0.04	-0.17	0.04	-0.06
grp2 (transmitter)	-0.03	-0.02	-0.13	-0.12	-0.004	0.00
grp3 (reservoir)	-0.85	-0.86	-0.76	-0.69	-1.00	-0.92
total	-0.87	-0.90	-0.93	-0.95	-0.95	-0.98

Reaction step	grp1		Į	grp2		rp3	N ₂ +H	total
	total	average	total	average	total	average	total average	
N ₂ +Mo(1)	-1.24	-0.309	1.58	0.067	-0.34	-0.014		0.0
N ₂ (ad)(2)	-1.13	-0.284	1.62	0.067	-0.23	-0.010	-0.25 -0.125	0.0
H1(3)	-0.89	-0.223	1.75	0.073	0.54	0.023	-0.40 -0.132	1.0
H1R(4)	-0.95	-0.237	1.71	0.071	-0.29	-0.012	-0.48 -0.159	0.0
H2(5)	-0.90	-0.226	1.76	0.073	0.52	0.022	-0.38 -0.095	1.0
H2R(6)	-0.97	-0.243	1.69	0.070	-0.62	-0.011	-0.45 -0.112	0.0
H3(7)	-0.91	-0.226	1.64	0.068	0.39	0.016	-0.12 -0.023	1.0
H3R(8)	-0.99	-0.248	1.67	0.070	-0.40	-0.017	-0.28 -0.056	0.0
H4(9)	-1.11	-0.278	1.61	0.067	0.26	0.011	0.24 0.040	1.0
H4R(10)	-1.17	-0.292	1.53	0.064	-0.56	-0.023	0.20 0.032	0.0
NH(S)/H5(E)(11)	-1.03	-0.257	1.76	0.073	0.44	0.018	-0.17 -0.024	1.0
H5(S)/H5R(E)(12)	-1.07	-0.267	1.69	0.070	-0.40	-0.017	-0.22 -0.031	0.0
$H5R(S)/NH_2(E)(13)$	-1.04	-0.259	1.70	0.071	-0.34	-0.014	-0.33 -0.110	0.0
H6(14)	-1.18	-0.294	1.65	0.069	0.32	0.013	0.21 0.052	1.0
H6R(15)	-1.24	-0.309	1.56	0.065	-0.51	-0.021	0.19 0.047	0.0

Table S10. Mulliken charge of the three groups (as shown in Figure 5) in the catalystduring the reaction steps.



Figure S1. Three models for the Mo/N-doped graphene catalyst and the lowest unoccupied molecular orbital of Mo/N-doped graphenes ($C_{69}H_{21}MoN_3$ and $C_{117}H_{27}MoN_3$).



Figure S2. The structures of the intermediates in the Schrock reaction path. 2-N₂

adsorption, 3/4-1st hydrogenation (protonation and reduction), 5/6-2nd hydrogenation, 7/8-3rd hydrogenation, 9/10-4th hydrogenation, 11-ammonia desorption, 12/13-5th hydrogenation, 14/15-6th hydrogenation. The number of intermediate is the same to that in Fig. 2. (N in blue, C in gray, H in white, Mo in light blue).



Figure S3. The structures of the intermediates in the enzymatic reaction path. 2-N₂ adsorption, 3/4-1st hydrogenation (protonation and reduction), 5/6-2nd hydrogenation, 7/8-3rd hydrogenation, 9/10-4th hydrogenation,11/12-5th hydrogenation, 13-ammonia desorption, 14/15-6th hydrogenation. The number of intermediate is the same to that in Fig. 2. (N in blue, C in gray, H in white, Mo in light blue).



Figure S4. The variation of natural charge (the charge difference of the present step from that of the previous step) of different groups (grp) in Mo/N graphene ($C_{33}H_{15}MoN_3$) during the enzymatic catalytic N₂ fixation. The numbering of the reaction steps is shown in Figure 2.



Figure S5. The N-N bond order and the total bond order of Mo during the N₂ fixation in the Schrock and enzymatic reaction paths. 1-catalyst, 2-N₂ adsorption, 3/4-1st hydrogenation (protonation and reduction), 5/6-2nd hydrogenation, 7/8-3rd hydrogenation, 9/10-4th hydrogenation.