Supporting Information

Nitrogen-Doped Graphene Quantum Dots Combined Sodium 10-Amino-2-Methoxyundecanoate: Studies of Pro-Inflammatory Gene Expression and Live Cell Imaging

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Figure S1 HPLC and FT-IR analysis of the compoundS-3									
Figure SAM	S2	1H	NMR	and	13C	NMR	of	the	purified 5-3 & S-4
Figure S3	2D-NM	R analys	is of the S A	M			•••••		S-4
Figure S4	ESI MS	analysis	of the SAN	A				•••••	S-5
Figure S5	Structur	al analys	is of <i>Sodiu</i>	m 10-ami	no-2-me	thoxyundec	canoate	(SAM)	S-5
Figure S6	Absorpt	ion spect	ra and emi	ssion spe	ctrum				S-6
Figure macropha	S7 ge	Cell	survival	assay	of	SAM	on	RAW	264.7 S-7
Figure S8 Binding constant plot for fluorescence quenching of N-GQDs in the presence of SAM									
Table S1 Structural validation of SAM by NMR analysis									
Table S2 List of primers used for RT-PCR analysis									



Figure S1. HPLC and FT-IR spectrum of SAM fraction from Lyngbya sp.,





Figure S2. 1H NMR & 13C NMR of SAM from Lyngbya sp.



Figure S3. 2D-NMR of SAM from *Lyngbya sp.*



Figure S4. ESI-MS of SAM from Lyngbya sp. major peak observed is of mass 253.09.



Figure S5. Structural of Sodium 10-amino-2-methoxyundecanoate (SAM).



Figure S6. Spectral overlap of absorption spectrum of (a) SAM with emission spectrum of N-GQDs (b).



Figure S7. Effect of compound SAM on survival percentage of RAW 264.7 cells.



Figure S8. Binding constant plot for fluorescence quenching of N-GQDs in the presence of SAM.

δ (ppm)	No., of H	Attachment/position of H in		
o (ppm)	corresponding	the molecule		
0.85, 0.87	3	CH ₃ (C ₁₁)		
1.12	2	CH ₂ (C ₃)		
1.25	10	$CH_2 (C_4 - C_8)$		
1.59	2	CH ₂ (C ₉)		
1.99	1	CH (C ₂)		
2.31	1	CH (C ₁₀)		
3.47	3	OCH ₃ (C ₂)		
4.13	2	NH ₂		

 Table S1. Structural validation of SAM by NMR.

Table S2. List of primers used for RT-PCR analysis for the expression of the inflammatory genes:

Gene symb ol	Gene Name	Forward Primer	Reverse Primer
Act	Actin	CTGACAGACTACCTCATGAG	CTCGAAGTCTAGAGCAACAT
		ATCC	AGCAC
IL-1α	Interleukin-	ΔΤΓΔGΓΔCCTTΔCΔCCTΔCC	GCTGAGATAGTGTTTGTCCA
1 alpha	1 alpha		CATC
		AGAGI	
IL-1β	Interleukin-	GTTCCCATTAGACAACTGCA	CTTGGTTCTCCTTGTACAAA
	1 beta	CTAC	GCTC
IL-4	Interleukin-	CTTCTTTCTCGAATGTACCA	GAGCTCACTCTCTGTGGTGT
	4	GGAG	ТСТ
IL-6	Interleukin-	CCAGAGTCCTTCAGAGAGAT	CTGTGACTCCAGCTTATCTG

	6	ACAGA	TTAGG
COX-	Cycloxygen	GACCGAGTTTACTGAGAAAG	CTCCTGGAAGGTACCAAAG
2	ase-2	AGGAG	ATAGAG
TNF-	Tumor	GATEGETTETACCTTETCTA	GAGGTTGACTTTCTCCTGGT
α	necrosis	CTCC	
	factor		AIGAG
NF-	Nuclear	ACTACCACCTATCATCCCAC	CTCATACACACCTCAAAC
κβ	factor-	AGTACCACCTATGATGGGAC	
	kappa beta	TACAC	TTCTCC
iNOS	Inducible		
	nitric oxide	CCTCCTCCACCCTACCAAGT	CACCCAAAGTGCTTCAGTCA
	synthase		