

Supplementary file

Table S1: Isotherm Equations.

Isotherm	Equation	Parameters
Langmuir	$\frac{C_e}{Q_e} = \left(\frac{1}{Q_m}\right)C_e + \left(\frac{1}{K_L Q_m}\right)$	Q_m = maximum adsorption capacity (mg/g) K_L = Langmuir constant (L/mg).
Freundlich	$\log Q_e = \log K_f + \frac{1}{n} \log C_e$	K_f is the Freundlich constant (represents performance) $1/n$ is the heterogeneity factor (represents effectiveness).
Temkin	$Q_e = B \ln K_T + B \ln C_e$	$B = RT/b$, T = temperature (K), R = Gas constant ($8.314 \text{ JK}^{-1} \text{ mol}^{-1}$), b = Temkin constant related to the heat of adsorption (J mol^{-1}), and K_T = Equilibrium binding constant (L mg^{-1})

Table S2: Physiochemical Characteristics of Seawater

S.No.	Physiochemical Characteristics	Quantity (Units)
1.	Temperature	27±0.3°C
2.	pH	8.1±0.07
3.	Salinity	35±0.1 ppt
4.	Dissolved Oxygen	7.17 ±0.3 mg/L
5.	Conductivity	14.65±0.54 mS/cm
6.	Total Hardness	100±2.3 mg/L
7.	Calcium	400±1.2 mg/L
8.	Magnesium	1300±15.2 mg/L
9.	Potassium	380±0.8 mg/L
10.	Sulphate	3708±23.5 mg/L

Table S3: Zeta potential values of various PSNPs dispersed in Milli-Q and seawater

Particles	Zeta potential (mV)	
	In Milli-Q water	In Seawater
MH	0.1±0.09	-0.98±0.04
PS-100 nm	-47.3±1.2	-58.2±1.5
PS-COOH	-64.5±1.0	-75.6±0.8
PS-NH ₂	-53.1±0.5	-66.3±1.4
PS-500 nm	-47.3±0.9	-55.5±0.6

Table S4: Hydrodynamic sizes of various PSNPs and PS-MH complexes dispersed in Milli-Q and seawater

Particles	Hydrodynamic Size (nm)	
	Milli-Q water	Sea water
PS-100 nm	100±4	409±2
PS-100 nm + MH	114±2	570±5
PS-COOH	102±5	430±6
PS-COOH + MH	120±5	620±3
PS-NH ₂	100±7	415±7
PS-NH ₂ + MH	115±3	585±6
PS-500 nm	501±9	895±2
PS-500 nm + MH	560±8	954±3

Table. S5 Independent action model for the PSNPs+MH complex

S.NO	Test samples	Concentration of Complex pollutants ($\mu\text{g/ml}$)	RI value	Mode of action
1.		2.5	1.10	Synergistic
2.	PS-100 + MH	5	1.14	Synergistic
3.		10	1.20	Synergistic
4.		2.5	1.06	Synergistic
5.	PS-COOH+MH	5	1.14	Synergistic
6.		10	1.20	Synergistic
7.		2.5	1.06	Synergistic
8.	PS-NH ₂ +MH	5	1.12	Synergistic
9.		10	1.17	Synergistic
10.		2.5	1.00	Additive
11.	PS-500 +MH	5	1.68	Synergistic
12.		10	1.69	Synergistic

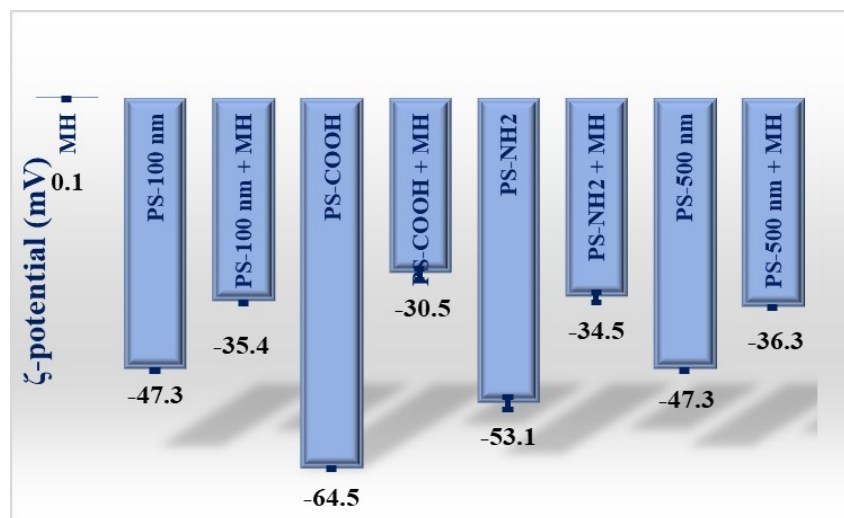


Fig. S1. Zeta potential of NPs, MH, and their corresponding complexes. [NPs], [MH], [NPs, MH – 1:0.5] = 10mg/L, pH=7. Data were the mean of triplicate experiments with SD.

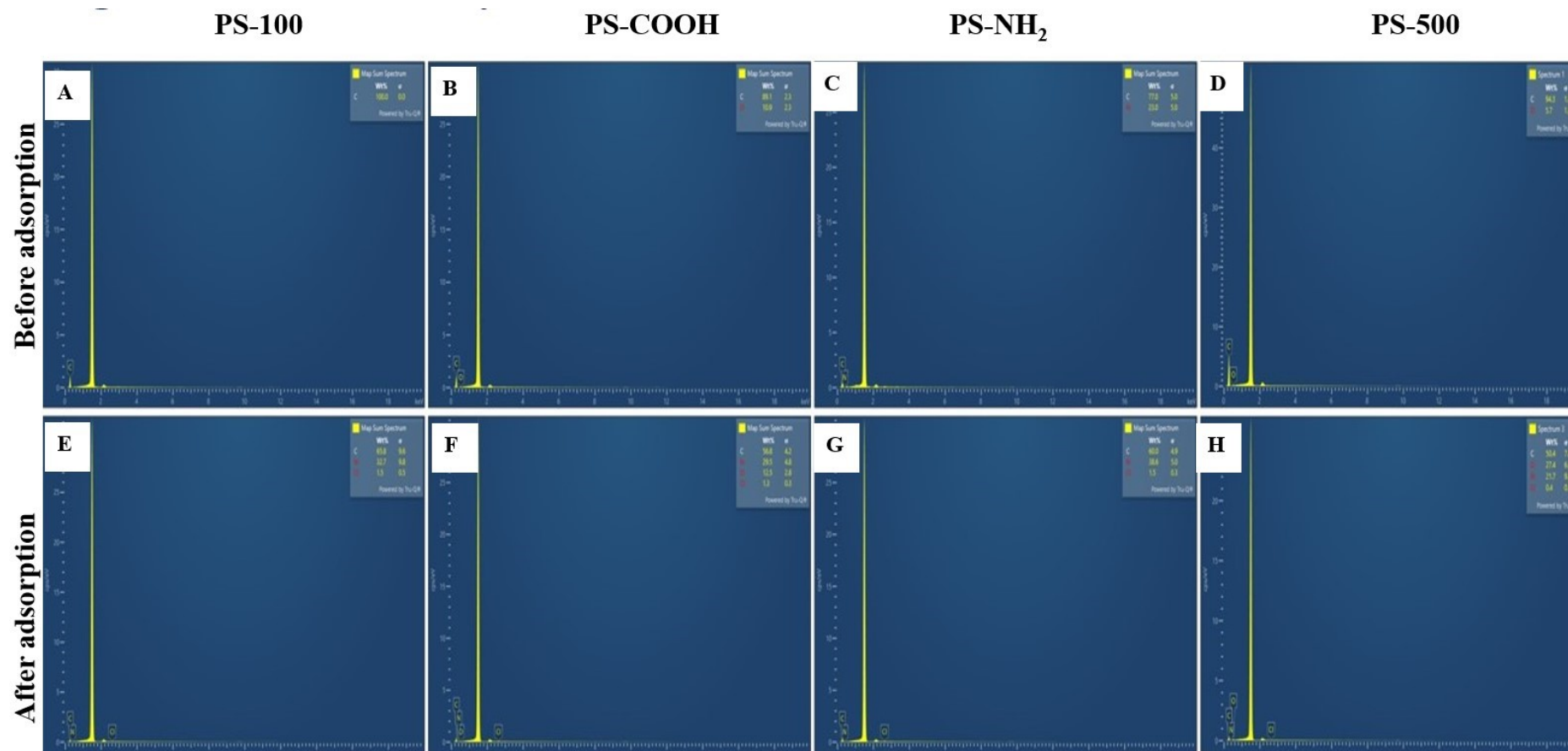


Fig. S2. A-D: EDX analysis of NPs before the adsorption process. (A) PS-100, (B) PS-COOH, (C) PS-NH₂, (D) PS-500. E-H: EDX analysis of NPs after adsorption of MH. (E) PS-100+MH, (F) PS-COOH+MH, (G) PS-NH₂+MH, (H) PS-500+MH.

Decapsulation	Instar 1	Instar 2	Instar 3	Observation
				PS-100nm Greater intake of PS compared to PS-200, 500 and PS-COOH, with slight difference in the body structure.
				PS-COOH-100 nm Intake of PS-COOH is also visible, however it is least significant than PS and PS-NH ₂ .
				PS-NH₂-100 nm Intake of particles is observed greater with developmental abnormality in the Instar 1 and 2 stages.
				PS-500 nm Morphology was not affected but the accumulation of the particles is found in the gut region.
				PS-200 nm (FI) The fluorescent particles are seen attached to the cyst and up taken by larvae.
				PS-100+MH When adsorbed with MH, PS-100 nm delayed the hatching at 16 h followed by intake, developmental deformity at the 1st instar stage and accumulation in the gut region.
				PS-COOH-100+MH Similar to PS-100 nm with a slightly reduced toxicity
				PS-NH₂-100+MH Highly significant among all others, where the cyst is observed to completely spherical showing delayed hatching. Delay in developmental growth as seen in 2 nd instar stages and 3 rd instar stages.
				PS-500 +MH Accumulation of particles are visible.
				PS-200 (FI)+MH Accumulation of complex were observed in the gut region
				MH No toxic effects observed But little accumulation
				CONTROL No developmental abnormalities

Fig. S3: Developmental changes of *A. salina* on exposure to different NPs and MH complexes.

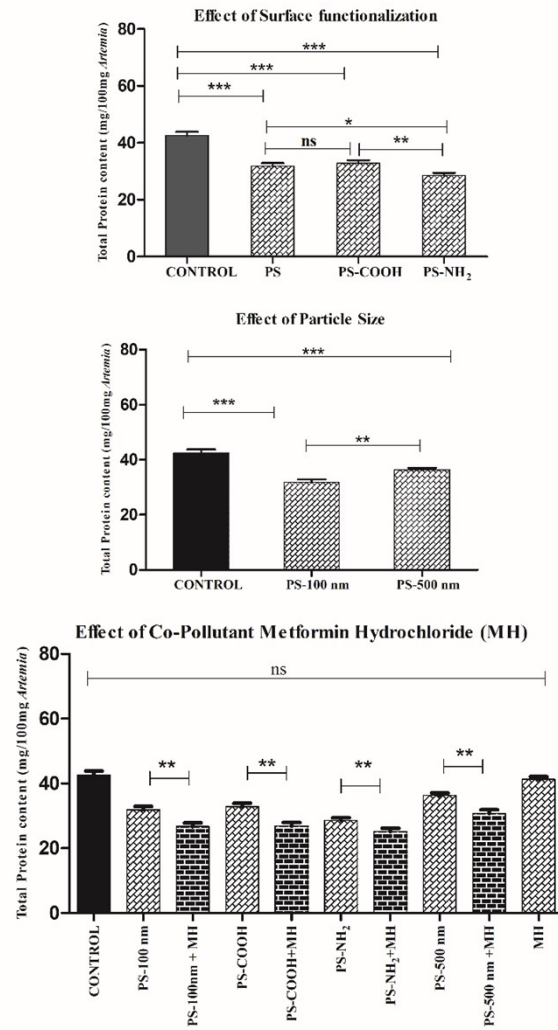


Fig. S4: Total protein content of *A. salina* on interaction with NPs and MH complexes (n=3). All values expressed as mean±SD. (***)P<0.0001)