

DNA damage and systemic toxicity induced by silver and copper oxide nanoparticles, and their mixture in *Clarias gariepinus* (Burchell, 1822)

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Introduction

- Nanotechnology industry is a **rapidly growing industry**, with numerous benefits (Nelson *et al.*, 2016)
- Silver (Ag) and Copper oxide (CuO) nanoparticles (NPs) have unique **antimicrobial properties** (Makwana *et al.*, 2015).
- AgNPs constitute 24% of NPs containing products (Vance *et al.*, 2015)
- 63 – 91% of ~ 200 metric tons of Cu or CuONPs in 2010 was estimated to have end up in landfill (Keller *et al.*, 2014)



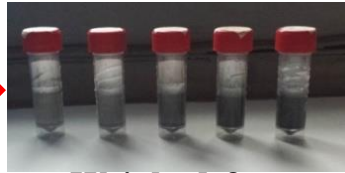
Justification and Hypothesis

- Rapid increase in manufacture and use of NPs but insufficient studies on potential adverse health effect (Koederith *et al.*, 2014).
- Existing genotoxicity data are largely on the *in vitro* system with limited *in vivo* studies (Pattan and Kaul, 2012).
- Non - existence of information on the *in vivo* impacts of the co - exposure of the selected NPs on aquatic biota
- As part of our ongoing studies on genotoxicity of metal and metal oxide nanoparticle, **We hypothesize that AgNPs and CuONPs and their 1:1 mixture might have DNA damaging and systemic toxicity effect in aquatic model *C. gariepinus*.**

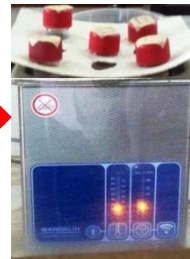
Study Design



AgNPs (<100 nm)
CuONPs (< 50 nm)
1:1 AgNPs + CuONPs



Weighed &
dispersed
in distilled water



Sonicated for
10 min



Vortexed for
5 min



C. gariepinus were exposed to 5
Sub-lethal concentrations (6.25 -
100 mg/L) for 28 days



Juveniles of *C. gariepinus* (25.56
± 4.79g and length 13.85±1.06
cm ; n = 270)

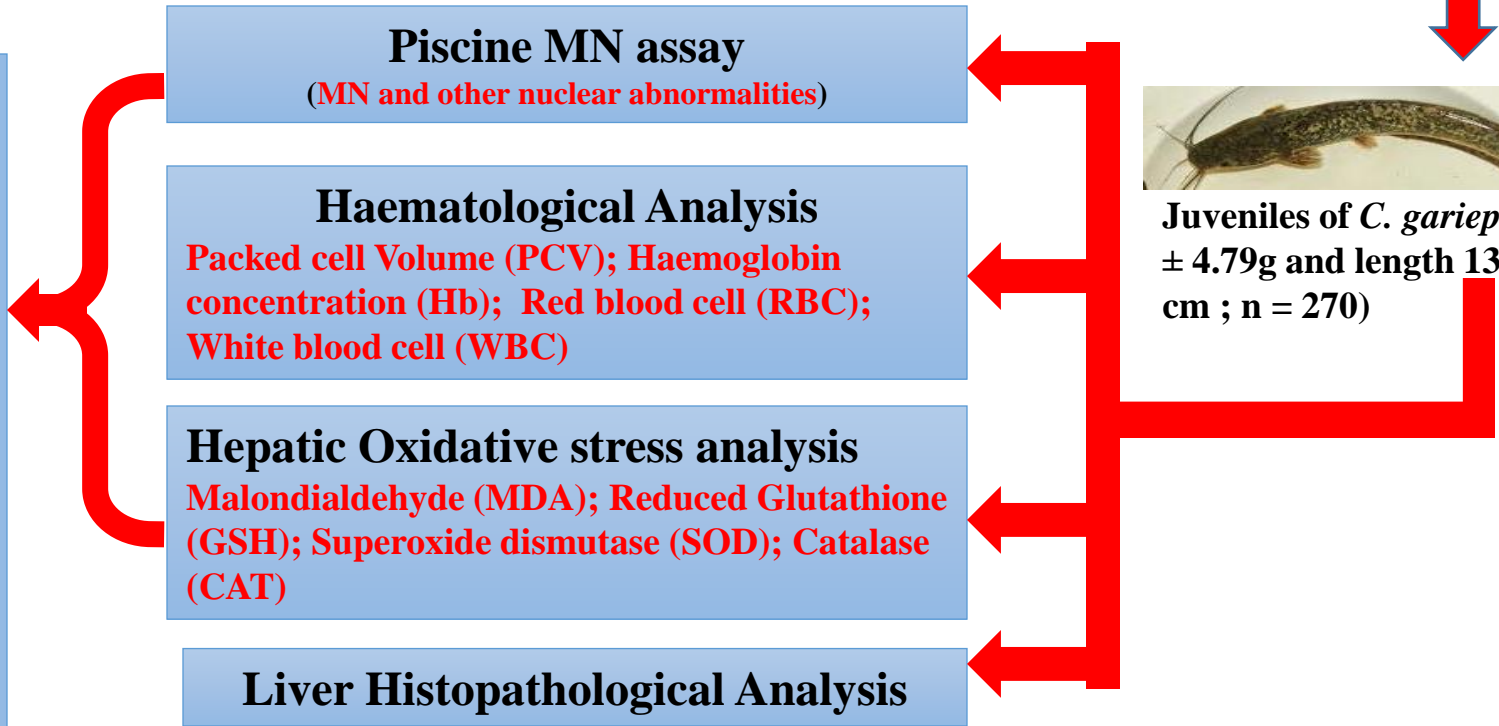
One way analysis of variance
and Interaction factor analysis

Piscine MN assay
(MN and other nuclear abnormalities)

Haematological Analysis
Packed cell Volume (PCV); Haemoglobin
concentration (Hb); Red blood cell (RBC);
White blood cell (WBC)

Hepatic Oxidative stress analysis
Malondialdehyde (MDA); Reduced Glutathione
(GSH); Superoxide dismutase (SOD); Catalase
(CAT)

Liver Histopathological Analysis



Results

- Significant induction of MN which indicates DNA damage
- PCV, Hb and RBC ↓ while WBC ↑
- Presence of histopathological lesions such as diffuse vacuolation of hepatocytes, vacuolar degeneration, atrophy and necrosis livers of exposed fish
- Levels of MDA, GSH and SOD increased while CAT decreased in in the liver
- Interaction analysis of data indicates:
 - antagonistic DNA damage, oxidative damage (MDA), GSH and SOD activities in mixture group while
 - CAT activity was synergistic

RESULTS

Table 1: Effects of AgNPs, CuONPs and the interaction of their mixture on MN frequencies and total nuclear abnormalities (TNA) in peripheral erythrocytes of juvenile *Clarias gariepinus* following 28 days exposure period

Concentrations (mg/L)	MN Frequency per 1000 erythrocyte (Mean ± SE)			Total Nuclear Abnormalities (TNA) abnormalities per 1000 erythrocytes (Mean ± SE)		
	AgNPs	CuONPs	Mixture (Ag + CuO NPs)	AgNPs	CuONPs	Mixture (Ag + CuO NPs)
NC	4.07± 0.55	4.07± 0.55	4.07± 0.55	0.00± 0.00	0.00 ± 0.00	0.00 ± 0.00
6.25 mg/L	4.40± 0.34	5.50± 0.44	6.47± 0.54	0.53± 0.53	0.07 ± 0.00	3.47 ± 1.42
12.5 mg/L	5.27± 0.16	7.27± 0.64*	8.27± 0.71***	1.67 ± 1.29	2.20 ± 0.90	4.20 ± 0.71
25 mg/L	6.33± 0.21*	8.80± 0.90***	10.67± 0.65***	0.00 ± 0.00	0.13 ± 0.13	0.00 ± 0.00
50 mg/L	7.73± 0.86***	7.20± 0.50*	10.20± 0.68***	0.00 ± 0.00	3.53 ± 0.58	1.13 ± 1.13
100 mg/L	7.00± 0.11**	8.58±1.16**	7.83± 0.32***	0.00 ± 0.00	4.08 ± 2.22	0.00 ± 0.00
PC	13.00 ± 0.99***	13.00 ± 0.99***	13.00 ± 0.99***	6.00 ± 3.38*	6.00 ± 3.38*	6.00 ± 3.38*

NC: Negative control (dechlorinated tap water), PC: Positive control 0.05mL/L Benzene *, **, *** values significantly different from the control group at p<0.05, 0.01 and 0.001 respectively following one way ANOVA

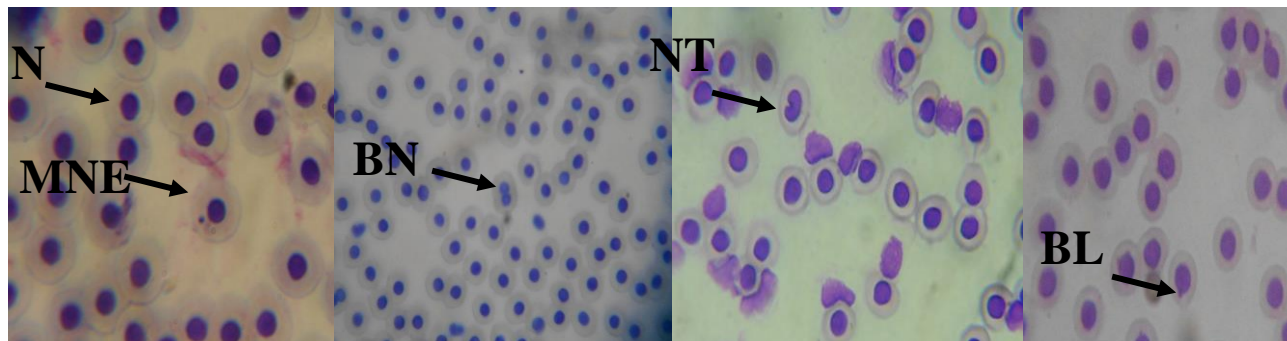


Fig. 2: Normal erythrocyte (N), micronucleated erythrocytes (MNE), binucleated cells (BN), notched (NT) and blebbed nuclei (BL) in peripheral blood of *C. gariepinus* exposed to AgNPs, CuONPs and their mixtures.

RESULTS

TABLE 2: Effect of AgNPs, CuONPs and mixture on haematological indices and levels of oxidative stress biomarkers in *C. gariepinus* following 28 days exposure.

Concentration (mg/L)		Mean \pm SD					
		RBC (x 10 ⁶ /μl)	WBC (x10 ⁶ /μl)	MDA (unit/mg protein)	GSH (unit/mg protein)	SOD(unit/mg protein)	CAT(μmol H ₂ O ₂ consumed/min/mg protein)
Ag NPs	NC	3.77 \pm 0.10	24190 \pm 2966	15.91 \pm 2.50	774.3 \pm 150.4	0.1658 \pm 0.01	120.1 \pm 43.74
	6.25	3.92 \pm 0.12	21440 \pm 1479	7.276 \pm 2.39	240.7 \pm 26.05*	0.0906 \pm 0.01	39.67 \pm 10.93*
	12.5	3.38 \pm 0.27*	21670 \pm 1337	28.35 \pm 17.1	481.2 \pm 144.6	0.07609 \pm 0.03	119.2 \pm 4.10
	25	2.03 \pm 0.40*	12530 \pm 1267*	18.22 \pm 1.97	912 \pm 224.4	0.3689 \pm 0.08***	141 \pm 29.4
	50	2.15 \pm 0.39	1585 \pm 1234*	24.26 \pm 2.37	1127 \pm 166.7	0.3014 \pm 0.04**	80.28 \pm 0.60
	100	3.20 \pm 0.17	13840 \pm 1156*	15.65 \pm 1.79	778.3 \pm 141.4	0.1526 \pm 0.03	87.06 \pm 18.09
CuO NPs	NC	3.77 \pm 0.10	24190 \pm 2966	15.91 \pm 2.50	774.3 \pm 150.4	0.1658 \pm 0.01	120.1 \pm 43.74
	6.25	2.40 \pm 0.47	15910 \pm 2044	20.12 \pm 2.7	1257 \pm 136.3**	0.3493 \pm 0.03***	41.11 \pm 4.84
	12.5	2.19 \pm 0.28	16800 \pm 2310	20.42 \pm 3.18	1143 \pm 130.4*	0.1922 \pm 0.04	12.06 \pm 2.21
	25	2.78 \pm 0.52	17910 \pm 1162*	37.86 \pm 2.5***	845.6 \pm 174.8	0.1789 \pm 0.02	17.25 \pm 0.00
	50	1.36 \pm 0.04	19430 \pm 3346	16.23 \pm 2.2	840 \pm 224.3	0.2747 \pm 0.02**	3.908 \pm 2.22
	100	2.78 \pm 0.52	17910 \pm 1162*	29.06 \pm 1.25***	1244 \pm 63.1**	0.2523 \pm 0.05*	3.137 \pm 2.56
Mixture	NC	3.77 \pm 0.10	24190 \pm 2966	15.91 \pm 2.50	774.3 \pm 150.4	0.1658 \pm 0.01	120.1 \pm 43.74
	6.25	2.24 \pm 0.33	16430 \pm 2015	13.39 \pm 2.9	629.5 \pm 77.83	0.1052 \pm 0.07	88 \pm 8.93
	12.5	1.63 \pm 0.05	18900 \pm 1778	13.83 \pm 3.2	784.3 \pm 178.3	0.08154 \pm 0.02*	80.42 \pm 11.39
	25	3.34 \pm 0.25*	16540 \pm 2482*	5.682 \pm 0.18*	285.6 \pm 79.84*	0.07482 \pm 0.03**	53.13 \pm 0.00
	50	2.25 \pm 0.43*	17160 \pm 1204*	14.85 \pm 3.97	622.6 \pm 162.4	0.1349 \pm 0.04	50.77 \pm 6.42*
	100	1.94 \pm 0.27	14963 \pm 1740*	7.326 \pm 1.65*	696.9 \pm 240.7	0.08106 \pm 0.02*	85.01 \pm 20.95

Data are presented as mean \pm SD. Values with asterisks (*) indicate significant difference from control at *p < 0.05, ** p < 0.01,

***p < 0.001

RESULTS

TABLE 3: Interaction factor analysis of AgNPs and CuONPs mixture (1:1) on MN frequencies in peripheral erythrocytes and hepatic oxidative stress biomarkers in juvenile *Clarias gariepinus* after 28 days exposure

Conc. (mg/L)	**Interaction factor (IF± SE_{IF})				
	MN	MDA	GSH	SOD	CAT
NC	-	-	-	-	-
6.25	-2.54 ± 0.61	1.904± 2.84	-93.9 ± 88.9	-0.17 ± 0.05	128.22 ± 9.35
12.50	-1.79 ± 1.31	-19.03± 12.37	-65.6 ± 152.4	-0.02 ± 0.03	70.16 ± 7.29
25.00	-7.72 ± 2.09	-34.49± 1.84	-697.7 ± 149.52	-0.31 ± 0.05	15.88± 20.79
50.00	-4.13 ± 2.17	-9.73± 2.95	-570.1 ± 186.63	-0.28 ± 0.03	87.58 ± 4.73
100.0	-4.06 ± 1.57	-21.47± 1.64	-551.1± 186.99	-0.16 ± 0.03	115.81 ± 19.62

**** NB: Interaction factor IF = (Mixture - Control) - [(AgNPs - NC) + (CuONPs- NC)] = (Mixture - AgNPs - CuONPs + NC)**
- VE IF = antagonistic interaction, +VE IF = synergistic interaction

RESULTS

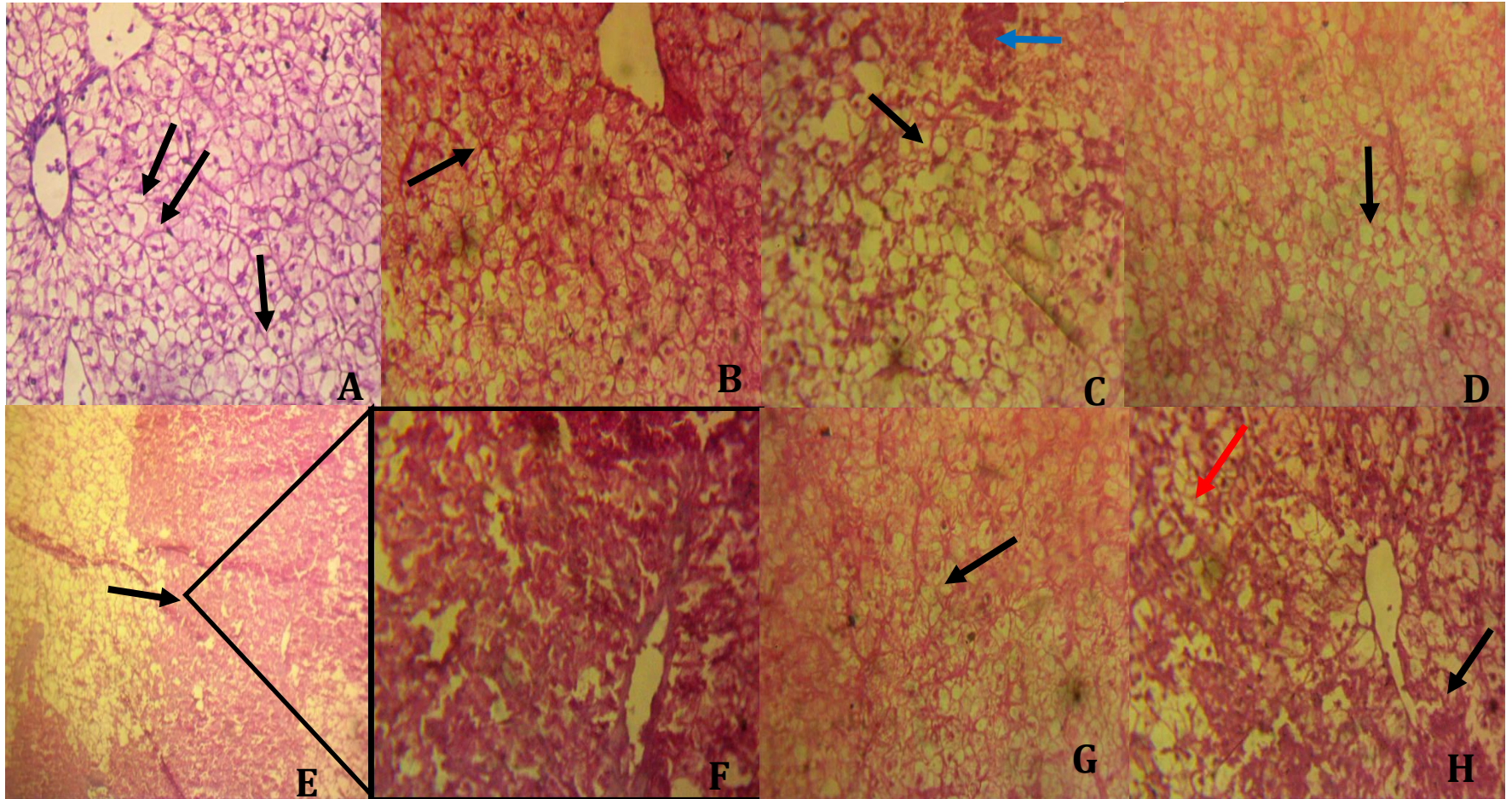


Figure 3: Haematoxylin & Eosin stained sections showing hepatic histological changes in juvenile *C. gariepinus* exposed to AgNPs, CuONPs, 1:1 mixture (B – H) and control

(A) Negative control: the hepatic plates are closely-packed with the hepatocytes (arrows) having large clear cytoplasmic appearance with the nucleus centrally placed; **(B) Diffuse vacuolation of hepatocytes (arrows; fat accumulation, normal)** **(C) Centrilobular vacuolar degeneration (black arrow) of hepatocytes and atrophy of adjacent hepatocytes (blue arrow)** **(D) Diffuse hepatocellular degeneration and atrophy (arrow)** **(E-F) Vacuolar degeneration and necrosis of hepatocytes (arrow)** **(G) Diffuse vacuolar degeneration (red arrow) of hepatocytes and foci of hepatocellular necrosis (black arrow)** **(H). moderate diffuse swelling of hepatocytes (arrow)**
Magnification $\times 400$.