



Occurrence, Toxicodynamics, and Mechanistic Insights for Atrazine Degradation in the Environment

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Abstract Atrazine, a herbicide used for controlling broadleaf weeds, has been one of the predominant pollutants constituting 80–90% of detection frequency in the samples collected from rivers, estuaries, oceans, sediments, agricultural lands, and crops. The fate of atrazine is highly unpredictable depending on the physio-chemical, physiological and geographical conditions. Range of metabolites such as deethylatrazine (DEA), deisopropyl atrazine (DIA), and didealkylatrazine (DDA) are formed as a result of biotic as well as abiotic degradation process in the environment following cyanuric acid, ammelide, CO₂ and NH₃ are formed as final products. Atrazine degraded products has shown more hazardous nature than the parent compound, atrazine. Atrazine is banned in Italy, India, Germany and European union but widely used in China, Australian, Canadian and US agriculture. To date, reviews evaluating the assimilation of synergistic treatment technologies and comparative degradation mechanism have not been highlighted. This work focuses on (1) the spatiotemporal distribution of atrazine and its metabolites globally and the factors governing it (2) provides an in-depth discussion about the various studies showing the toxicity of atrazine in microbes,

cattle, human, terrestrial and aquatic organisms; (3) discusses the contaminants of emerging concern which are continuously replacing atrazine like terbutylazine and their intermediate compounds posing more risk to wildlife and humans; (4) summarises the different treatment technologies which have been predominantly applied for the removal of atrazine in water and soil systems and also discusses the synergistic or mutualistic aspects of treatment methods in degrading atrazine.

Keywords Atrazine · Occurrence · Biodegradation · Metabolites · Treatment · Synergistic

1 Introduction

Developed in 1958, atrazine is a herbicide used prominently for controlling broadleaf weeds and grasses in crops like corn, maize, millets, and sugarcane. Being a herbicide, it works by inhibiting the photosynthetic electron transport chain in plants (Rodríguez-González et al., 2017). The molecular formula of atrazine (2-chloro-4-ethylamino-6-isopropylamino-1,3,5-triazine) is C₈H₁₄ClN₅, its melting point varies between 173 °C–175 °C. It has low biodegradability and solubility ranges between 30–33 mg L⁻¹ at 20 °C (Hong et al., 2022). Atrazine (ATZ) such as simazine, and terbutylazine (TBA) belong to the chlorotriazine class of herbicides. Soil adsorption coefficient (K_{OC}) and half-life

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(DT₅₀) are fundamentals in assessing pesticide persistence and dissipation risk in different environment matrices (Góngora-Echeverría et al., 2019). Adsorption of atrazine to soil constituents is moderate with K_d values ranging in between 0.4–3.1 L kg⁻¹ (Salazar-Ledesma et al., 2018). The half-life of atrazine with no previous application on soil was 14.5 days while with previous history application the half-life reduced to 2.3 days suggesting widespread application over time in soil leads the development of atrazine degrading bacteria (Barrios et al., 2019). It is a weak base, its pK_a is 1.7. If the pH of soil solution comes around pK_a leading to migration of atrazine into groundwater or surface water (Salazar-Ledesma et al., 2018). In aquatic ecosystems half-life ranges from a few months to several years (Tulcan et al., 2021).

The ubiquitous nature of herbicides and their metabolites in different environments is posing a risk to human health and other living organism. The annual usage of atrazine is approximately 70,000–90,000 tons worldwide (Zhang et al., 2018). Canada, China and USA are the major consumers of atrazine. However, owing to toxicity issues, atrazine was banned in Italy and Germany in 1991 (Scherr et al., 2017). Even after 20 years of ban of atrazine by European union, atrazine and its by-product, deethylatrazine presence in groundwater was detected above the threshold value of 0.1 µg L⁻¹ in Germany (Vonberg et al., 2014). Several reports are available across countries like USA, China, and Spain where atrazine was frequently detected crossing the regulatory levels, exceeding the toxicological end points for organisms indicating the prevalence of atrazine at high concentrations (de Araújo et al., 2022).

The permissible limits of atrazine and its metabolites (diaminochlorotriazine, desethylatrazine, hydroxyatrazine, deisopropylatrazine, and atrazine mercapturate) are 5 µg L⁻¹ as per WHO guidelines (Dehghani et al., 2022) and 3 µg L⁻¹ in the United states while in Europe it is 0.1 µg L⁻¹. Several reports are available in literature where the atrazine and its metabolite concentrations are above the permissible limits. The lipophilic nature of atrazine leads to bioaccumulation in the exposed organisms and the food chain. The Lipophilicity of atrazine, Log D at pH 7.4 is 2.20 representing more water soluble nature of this compound whereas for

atrazine degraded products Log D values are even more smaller indicating the presence of more polar groups in the chemical structure (Furtado et al., 2019). Atrazine has been reported to be mutagenic, genotoxic, and morphotoxic in different exposed organisms (de Oliveira et al., 2020). Several epidemiological studies have shown that atrazine, even at lower than the maximum contaminant level (MCL), causes alteration in the functioning and formation of different parts of the brain, reproductive system, neuronal cell development, and neuroendocrine developmental process (Abass et al., 2021; Sadeghnia et al., 2021; Shan et al., 2021; Stradtman & Freeman, 2021). Some metabolites of atrazine such as desethylatrazine and deisopropylatrazine have been reported to have similar toxicity and structure to that atrazine, while hydroxyatrazine shows different toxicological properties from atrazine (Geng et al., 2013). Terbutylazine (TBA), a triazine family herbicide have been classified in carcinogen category 3 by European Food Safety Authority (EFSA) (Bottoni et al., 2013). It has already replaced atrazine, causing frequent detection in most of the EU countries, including Italy, Spain and Portugal (Álvarez et al., 2016; Bottoni et al., 2013). It has been detected in pristine marine ecosystem of North Sea, Mediterranean basin achieving the status of chemical of emerging concern because of its persistence and hazardous nature even at lower concentrations (Brumovský et al., 2017; Mai et al., 2013; Navarro et al., 2004). At cellular and animal organism level, TBA induces toxicological risk as it causes low-level DNA instability, DNA cross-links and cytotoxic effects (Želježić et al., 2018).

Several processes for the mineralization of Atrazine in aqueous and soil system have been reported (Arar et al., 2023; Liu et al., 2023; Tuğaç et al., 2023; Wang et al., 2022; Zhang et al., 2023). The treatment technologies include bioremediation, biological adsorption, oxidation, reduction, immobilization, nanofiltration, and adsorption (Gao et al., 2018; Khandarkhaeva et al., 2017; Moeini et al., 2019). Also, conventional drinking water treatment methods such as chlorination have limited degradation capacity for atrazine removal.

Although several reviews on atrazine treatment have been published (Mili et al., 2022; Rostami et al., 2021; Singh et al., 2018) few have discussed the occurrence, toxic effects, and advanced oxidation

processes (AOPs). Hybrid approaches to the remediation of atrazine have also not been discussed. This review discusses the spatio-temporal distribution of atrazine in several countries, and also an overview of the toxicity in different organisms at different stages of their growth. In addition, this review discusses the various treatment processes, new technologies, and their efficiency in remediating atrazine from the water and soil environment.

2 Occurrence

Due to greater application, atrazine presence is detected in biotic and abiotic components like water, sediments, different organisms, and food (Yin et al., 2020). Due to its symmetrical molecular structure, hydrophobic nature, and low solubility, it retains in the aqueous environment for a longer time (de Souza et al., 2020).

2.1 Occurrence in the Water Resources

Atrazine has been applied for more than 60 years in several countries worldwide, with an estimated use of 60,000–70,000 tons annually (Y. Xue et al., 2021a, b). The frequent detection of atrazine in coastal bay areas worldwide implies that the continuous application on agricultural lands also leads to transportation and leaching of atrazine in open seas due to various geographical, environmental, and climatic factors (Nowell et al., 2018). The half-life period of atrazine ranges between 41 and 237 days in freshwater, with an average of 159 days, but in saltwater, it varies between 3–190 days (Almasi et al., 2020). In a study conducted in the seawater of Xiangshan Harbor, the concentration of atrazine was between 3.99 and 73.0 ng L⁻¹, which was higher than the maximum permissible limit for coastal areas in the European union (Y. Xue et al., 2021a, b). In the southern Baltic Sea area, the mean value of atrazine was 2.2 ng L⁻¹ (Fisch et al., 2021). Atrazine was also detected in Bohai (31.2–112.20 ng L⁻¹ range), yellow sea, the Liaodong Peninsula (23.3 ng L⁻¹ mean concentration) and Laizhou Bay (6.8–83.0 ng L⁻¹ range) (Xie et al., 2019). On the Portuguese coast, the concentration of atrazine was detected around 35.3 ng L⁻¹, which was related to medium risk in the sea (Sousa et al., 2020). In the Jiaozhou

Bay of eastern China, the spatial distribution of atrazine and its degraded primary and secondary metabolite products were compared in water, suspended particulate sediment and surface sediment in an estuary-to-bay system. Deisopropylhydroxy-atrazine (DIHA), Hydroxy-atrazine (HyA), and Deethylhydroxy-atrazine (DEHA) in seawater were estimated with concentrations around 80.77 ng L⁻¹, 12.54 ng L⁻¹, 15.11 ng L⁻¹, and 4.95 ng L⁻¹, respectively. Higher atrazine concentrations was detected in the sea water near the coast then in the estuary and the bay showing the settlement of contaminants in sediments during transportation with seawater. (Ouyang et al., 2021). In the Jiaozhou Bay, atrazine and its metabolites was observed, reaching maximum in August, suggesting that difference in temperature and other climatic factors affects the transportation as well as degradation of atrazine and metabolites (Z. Wang et al., 2021a, b). In the ctalmochita River basin, atrazine detection was higher in the spring season as monitored for the year (2011–2015), stating that season plays a role in fluctuating the atrazine level (Bachetti et al., 2021) In the St. Lawrence Estuary and Gulf aquatic environments, predominantly atrazine was found at a high range of 34 ng L⁻¹. (Picard et al., 2021).

The continuous application of pesticides on agricultural lands makes them enter into surface water through continuous leaching and surface runoff. Proactive precautionary measures already taken as European union banned the use of atrazine as it exceeded permissible drinking water concentrations limits of 0.1 µg/L (van Rensburg et al., 2022b). In the drinking water treatment plants in Brazil, atrazine as well as its metabolite HA, DIA and DEA had a detection frequency of 100% from 2 to 2744 ng L⁻¹ (Vizioli et al., 2023). In rural areas of Nigeria, the atrazine levels ranged from 10 to 80 µg L⁻¹ when assessed in drinking water (Loureiro et al., 2023). Atrazine was detected at the highest concentration in more than 99% of samples taken from water bodies near agricultural land in the Mid-Atlantic area with concentrations up to 1.9 µg L⁻¹, indicating atrazine was a major pollutant (Zhu et al., 2021). In Paranoa lake, during the rainy season, surface water samples detect the concentration of atrazine at a range of 1.2 ± 0.9 to 5.5 ± 0.4 ng L⁻¹ (Bachetti et al., 2021). Atrazine and its metabolites were also equally detected in Danube River, Hungary River, Brittany

River (France), the Arno River, Klodnica River (Poland), Spanish Duero River, Miño River, Ebro River, Scheldt River (Netherlands), Tagus River (Portugal), the concentration ranges from 18.1 to 105.5 ng L⁻¹ with a mean value of 54.4 ng L⁻¹ (Triassi et al., 2022).

Atrazine presence was evaluated by the U.S. EPA National Lake assessment, the 2021 U.S. Department of agriculture cropscape and the global HydroLAB HydroLAKES databases where atrazine was detected in 32% of the U.S. waterbodies with a mean concentration of a 0.17 µg L⁻¹ (Beaulieu et al., 2020). In the lower Gangetic River basins (WBB) of West Bengal, India, Atrazine (0.95 – 3.93 µg L⁻¹) was detected at a very high concentration, surpassing the maximum level by 46 times (Duttagupta et al., 2020; Organization, 2004). Currently, TBA, an alternative to atrazine is the most detected herbicide in the natural waters of Spain due to its continuous and increased usage (Herrero-Hernández et al., 2017). Table 1, 2, 3, 4, 5, 6, 7, and 8.

2.2 Occurrence in Agricultural Land

Atrazine is a persistent organic pollutant whose half-life can range from 2 to 57 weeks in environment, depending upon soil characteristics, application history and various other factors whereas its metabolites can reside up to 4 months post-application (Dehghani et al., 2022) (Y. Zhang et al., 2021a, b). The persistence and soil interaction of atrazine is governed by the concentrations of dissolved organic matter (DOM) and soluble soil organic matter (SOM). Atrazine is predominantly sorbed by SOM whereas presence of humic acids, clay minerals play a huge role in migration, degradation, availability, sorption and accumulation of atrazine in soils. Clay minerals and smectites soil particularly have higher potential of atrazine sorption (Salazar-Ledesma et al., 2018). In a study done on agricultural land of Brazil on red and yellow soil for 70 days, atrazine shows a half-life of 10 days in sunlight while in shadow the half life is 19 days (de Paula et al.,

Table 1 This table depicts atrazine occurrence in the aquatic regions and agricultural lands of several countries

Rivers/Land	Countries	Concentration ATR (Mean) ng/L	Detection rate	References
Biana river	China	70–1120	100%	(Sun et al., 2019)
Volturno River	South Italy	3.42	90.1 ± 6.8%	(Triassi et al., 2022)
Red river	Canada	500	65%	(Challis et al., 2018)
Sea water	Portugal	35.3	-	(Sousa et al., 2020)
St. Lawrence River	Canada	0.67	82%	(Montiel-León et al., 2019)
Alqueva reservoir	South Portugal	> 4–666	100%	(Palma et al., 2014)
Great lake basin Lebo drain	Canada	>500	100%	(Metcalf et al., 2019)
Barratta Creek River	Australia	13,000	9%	(Wood et al., 2017)
Crocodile river	South Africa	483–940	(52–55) %	(Rimayi et al., 2018)
Ohre river	Czech Republic	3	–	(Fikarová et al., 2018)
Stream system	Germany	21	–	(Weber et al., 2018)
Red River	Canada	500	65%	(Challis et al., 2018)
River Nile	Egypt	84	23%	(Eissa et al., 2021)
Niger River valley	West Africa	217	40%	(Illatou et al., 2023)
Mojui River	North central Brazil	1.41	–	(Illatou et al., 2023)
Subsoils and soil (0–10 cm)	Northern Greece	50000	–	(Vryzas et al., 2012)
Soil (0–15 cm)	Central Mexico	700,000 ± 710,000	–	(Salazar-Ledesma et al., 2018)
Arable soils	Czech Republic	28,000	89%	(Hvězdová et al., 2018)
Shiraz Farmland	Iran	15,0000–55,0000	–	(Dehghani et al., 2022)
Venado Tuerto City	Argentina	28,000	–	(Singh et al., 2023)

Table 2 The table displays the toxicity of atrazine in different organisms affecting various subsets of physiology, morphology, and developmental process

Sample/ species	Atrazine exposure	Toxic level	Effects	References
Microalgae	4–8 days	(0.1–0.2) mg L ⁻¹	Microalgae growth was suppressed entirely	(Sun et al., 2020)
Diatom <i>Phaeodactylum tricornutum</i> pt-1	01–07 days	(15–50) nmol L ⁻¹	The cell structure was destroyed, leading to the decomposition of chloroplast	(Yang et al., 2019)
Zebrafish	7 days dpf (Days post-fertilization)	(2,2.5.5) mg L ⁻¹	Morphological defects, cardiac edema, tail reduction, head malformation, decreased head length	(Zaluski et al., 2022)
Fish		(0.30 ± 0.03) ng g ⁻¹	Adverse effects on genotoxicity, DNA damage, and hematology, different fish organs are found to be affected	(Destro et al., 2021)
Soybean plants		(5–15) mg Kg ⁻¹	Inhibition % of plant height increased from (17.22–41.75) %, and inhibition % of root elongation increased from (10.19–31.42) %	(Sánchez et al., 2020)
Human kidney cell lines (hek 293t)	24 h	(3–30) ppb	Drop in meCPG (23–32) %, altering DNA methylation levels, aberrant alterations in h3k9me observed in cancerous cells	(Sánchez et al., 2020)
Male Wistar rats	14 days	100, 200 or 400 mg Kg ⁻¹	Neuronal apoptosis and mitochondrial autophagy of the neurons at all concentrations, severity increases with increasing dose	(Li et al., 2020)
<i>Caenorhabditis elegans</i>	Dose-dependent	> 0.04 mg L ⁻¹	Decreased brood size, locomotory disability, and aberration in metabolic dynamics	(Yu et al., 2020)
Freshwater shrimp <i>Caridina africana</i>		106.8 and 5557 µg L ⁻¹	Biomarker response affects the levels of glutathione, catalase, and SOD (antioxidant enzyme), alleviating oxidative damage of cells	(van Rensburg et al., 2022a)
<i>Anaxyrus americanus</i>	96 h post-hatching	(0.2–20) mg L ⁻¹	Tail flexures, facial edema, axial shortening, blistering	(Slaby et al., 2019)
Marine blue mussel (<i>Mytilus galloprovincialis</i>)	7 days	(1 µm, 10 µm)	Mussel lysosomal stability decreases	(Slaby et al., 2019)
Strain <i>Azotobacter vinelandii</i> az6	10 days	900 µg mL ⁻¹	Inhibitory effects on IAA production (33%), PGP traits affected	(Shahid et al., 2019)

Table 2 (continued)

Sample/ species	Atrazine exposure	Toxic level	Effects	References
<i>Drosophila melanogaster</i> (germ-free flies)	72 h	2.0 mm	In females, no. of deregulated genes (92) and downregulation of reproductive genes in both males and females cause disturbed vitelline membrane formation	(Brown et al., 2021)
Rats	2 weeks	100 mg Kg ⁻¹	Acute exposure causes decreased cell Purkinje fibers. Locomotion activity disturbed	(MOSELHY et al., 2016)

Table 3 Different physiochemical properties of atrazine and its metabolites

Metabolites	Molecular formula	Molecular weight	Density	Melting point	Boiling point
Atrazine	C ₈ H ₁₄ ClN ₅	215.68	1.23 gcm ⁻³	(173–175) °C	200 °C
de-ethylatrazine[DEA]	C ₆ H ₁₀ ClN ₅	188	1.90 ^d	135–137	308 °C
de-isopropylatrazine[DIA]	C ₅ H ₈ ClN ₅	174	1.19 ^d	177–179	237 °C
de-ethylhydroxyatrazine[DEHA]					
Hydroxyatrazine[HA]	C ₈ H ₁₄ ClN ₅ O	197.24	1.30 gcm ⁻³	>320 °C	409.52 °C
di-dealkylatrazine[DDA]		215.68	–	–	–
de-ethyldeisopropyl atrazine[DEDIHA]	C ₅ H ₈ ClN ₅	173.60	–	–	–
de-isopropylhydroxyatrazine[DIHA]	C ₅ H ₉ N ₅ O	155.16	–	–	–
di-aminochlorotriazine[DACT]	C ₃ H ₄ ClN ₅	145.55	1.700 gcm ⁻³	>320 °C	237.97 °C

2016). It has been reported that atrazine undergoes many transformations depending on geographical, agricultural, temperature conditions and climatic factors (Srivastava et al., 2017) (Yu et al., 2020). Atrazine was detected at deeper layers of soil, where the corn root was unable to reach, showing the ease of migration and adsorption in the soil system (de Oliveira et al., 2019). The distribution of atrazine in major aquatic bodies and soils around the world is depicted in Table 1. Another study reported that even after 20 years of application in agricultural soil of Germany and Belgium atrazine was detected at 19.5 g ha⁻¹ (Jablonski et al., 2011). In a study done for the adsorption–desorption characteristic of atrazine in three soils, atrazine sorption capacity ranked as paddy soil > alluvial soil > laterite stating sorption is governed by the presence of sorption sites on the soil surfaces irrespective of atrazine concentration (Yue et al., 2017).

In the cultivated corn field in Croatia, when atrazine was applied at a concentration of 1 kg ha⁻¹ in soil, it remained detected until 5 months of corn harvesting (Stipičević et al., 2015). We summarized and described how the contaminant atrazine enters the food web, hampering the exposed organism's life cycle at different stages in Fig. 1.

3 Toxicity of Atrazine on Non-Target Organisms

The causal effect of Atrazine and its metabolite on the living system depends upon the time the organism has been exposed, the residual concentration of the metabolite in the living system, and physiochemical, physiological, geographical, as well as biological factors. Atrazine has shown genotoxic and mutagenic effects on yeasts and plants, while chromosomal aberrations and DNA damage have been reported in mammalian cell lines.

Table 4 This table describes the various bacterial strains degrading atrazine under different conditions

Microbial strains	Phase/Medium	Microbial source	Biodegradation efficiency	References
<i>Nocardioides</i> sp. DN36	Liquid	Paddy field	100%, 6 days	(Satsuma, 2010)
<i>Enterobacter cloacae</i> , <i>Burkholderia cepacia</i>	Liquid	Sugarcane fields	82% 54 days	(Ngigi et al., 2012)
<i>Chelatobacter heintzii</i> SalB	Liquid	Arable soil	100%, 18 days	(Cheyins et al., 2012)
<i>Arthrobacter</i> sp. DNS10 DNS9, <i>Bacillus subtilis</i> DNS4, <i>Variovorax</i> sp. DNS12	Liquid	Corn planted soil	40 h	(Zhang et al., 2012)
<i>Arthrobacter</i> sp.	Solid	Manfredi soils, Argentina	100%, 30-36 h	(Fernández et al., 2013)
<i>Frankia alni</i> , ACN14a	Liquid		57%, 6th day	(Rehan et al., 2014)
<i>Shewanella</i> sp.	Liquid	Corn field	100%, 36 h	(Ye et al., 2016)
<i>Ensifer</i> sp.	Liquid		100%, 30 h	(Ma et al., 2017)
<i>Leucobacter triazinivorans</i> <i>JW-1</i>	Liquid	Prometryn contaminated sludge	43%, about 50mgL ⁻¹ , 2 days	(Liu et al., 2017)
<i>Citricoccus</i> sp. TT3	Liquid	Wastewater outfall	50% in 66 h	(Yang et al., 2018)
<i>Bacillus badius</i> ABP6 and <i>Bacillus encimenis</i> ABP8 on α -Fe ₂ O ₃ mag- netic nanoparticle	Liquid		(90.56 ± 1.69)%, 20 days	(Khatoon & Rai, 2020)
<i>Klebsiella variicola</i>			81.5%, 11th day	(Zhang et al., 2019)
<i>Arthrobacter</i> sp. DNS10 and <i>Enterobacter</i> sp. P1	Liquid		99.18 ± 1.00%	(Jiang et al., 2019)
<i>Klebsiella variicola</i> FH- 1	Liquid	Agricultural University, China	72.5% in Zn ²⁺	(J. Zhang et al., 2021a, 2021b)
<i>Arthrobacter</i> sp. DNS4, DNS9 <i>Bacillus subtilis</i> DNS10, <i>Variovorax</i> sp. DNS12	Liquid	Corn planted soil	40 h, >90%	(Zhang et al., 2012)
<i>Stenotrophomonas</i> <i>maltophilia</i> , <i>Arthrobac-</i> <i>ter</i> sp.	Liquid	Isolated from a microbial consortium	41.2 ± 2.4%	(Galíndez-Nájera et al., 2011)
<i>Bacillus badius</i> A BP6	Contaminated field soil, tarai agro- eco- system	89.7%, at PH 7.04, temp 30.4 °C		(Khatoon & Rai, 2020)

Table 5 This table describes the different photocatalysts in the degradation of atrazine under different conditions

Photocatalysts	Degradation efficiency	References
N doped TiO ₂ (Ndt) + syndiotactic polystyrene (sps) matrix	47% and 25% UV or Vis light, 180 min	(Navarra et al., 2022)
Heterojunction photocatalyst MIL-53(Fe/Co)/CeO ₂ + PMS	99%, 60 min of visible light	(Roy et al., 2022)
La-doped ZnO nanofibers	98%, 1 h, light	(Krishnasamy et al., 2022)
Bulk carbonitride + phosphoric acid (PCN-H)	(18.4 to 45.7) %, 1 h, visible light	(L. Xue et al., 2021a, 2021b)
BaTiO ₃ @MWCNT Photocatalyst	40 min, Visible light	(Sobahi & Amin, 2021)
HPEI-0.5NaCl-BiVO ₄ - Bi ₂ O ₃	95%, after 40 min, mercury visible light	(Mahlalela et al., 2020)

Table 6 This table depicts the ozonation process for removing atrazine under different conditions

Ozonation catalyst	Removal efficiency	References
MnCe-ceramic membranes	99.99%, 40 min, pH 7	(He et al., 2022)
Synthetic 4A zeolite	87.5%, 6 min	(Su et al., 2022)
Mn-loaded biochar, Fe-loaded biochar	83%, 100% 30 min pH7	(Tian et al., 2021)
0.5NaCl-BiVO ₄ -Bi ₂ O ₃ heterojunction	>95%, 40 min, pH 7.7	(Mahlalela et al., 2020)
Microbubble ozonation	95.3%, 15 min, pH 9	(Liu et al., 2020)
Ozonation of atrazine on raw tourmaline	98%, 10 min, pH 7	(Wang et al., 2018)

Table 7 This table explains the persulfate degradation method for removing atrazine under different conditions

Persulfate degradation	Degradation efficiency	References
Dielectric barrier discharge plasma O ₃ /peroxydisulfate	82.46%, 25 min	(Shen et al., 2022)
nZVI sulfurized	96.8% pH 7	(Jiang et al., 2022)
UV/PS	98.4% pH 7	(Liu et al., 2022)
Cu ²⁺ /LDH-MoS ₄ /persulfate	30%, pH 3.0–9.0	(Li et al., 2022)
Persulfate (1 mm)/Dielectric barrier discharge/microbubbles	89%, 75 min	(Q. Wang et al., 2021a, 2021b)
Natural montmorillonite + Fe (III) + Peroxymonosulfate	94.1, 60 min	(Wang et al., 2020)

Table 8 This table explains the Fenton treatment process for removing atrazine under different conditions

Fenton treatment	Degradation efficiency	References
FecL ₃ /H ₂ O ₂ , FecL ₃ /H ₂ O ₂ + UV	92.40% 120 min 97.02% 120 min	(Fareed et al., 2021)
nZVI/H ₂ O ₂ /UVA	80% pH 3, 60 min of irradiation	(Plaza et al., 2021)
H ₂ O ₂ + ACF (carbon fiber/cobalt ferrite) with/without light	39.0%, 15 h and 97.6%	(Jiang et al., 2021)
Fe/TiO ₂ + visible irradiation + catalyst + H ₂ O ₂	95%, pH 3, 30 min	(Yang et al., 2020)
NHPC (N doped porous carbon) + FeOx NPs	96%, 90 min	(Cao et al., 2020)
Fe ₃ O ₄ /peroxymonosulphate / Hydroxylamine	40%, 5.0–6.8pH	(J. Li et al., 2019a, 2019b)

3.1 Inhibitory Effects on Mammals

Atrazine has been observed to have broad implications on the human body as its continuous presence in drinking water makes it more pernicious to multiple organs affecting the human body's reproductive, neuronal, and immunological cells (Surmeier et al., 2017). Being an endocrine-disrupting chemical, it plays an intertwined role in targeting the reproductive and neuroendocrine system. (Stradtman & Freeman, 2021).

Atrazine exposure is associated with many neurodegenerative problems, including Parkinson's disease, schizophrenia, and attention deficit disorder (P. Li et al., 2021a, 2021b). Substantia nigra pars compacta, and striatal dopaminergic neurons are most prone to atrazine toxicity as it has caused degeneration of dopaminergic neurons in the substantia nigra. (Li et al., 2020) (P. Li et al., 2021a, 2021b). Phosphatidylcholine and CDP-choline are important metabolites associated with neurotransmitter synthesis and transmission, lipid transportation and

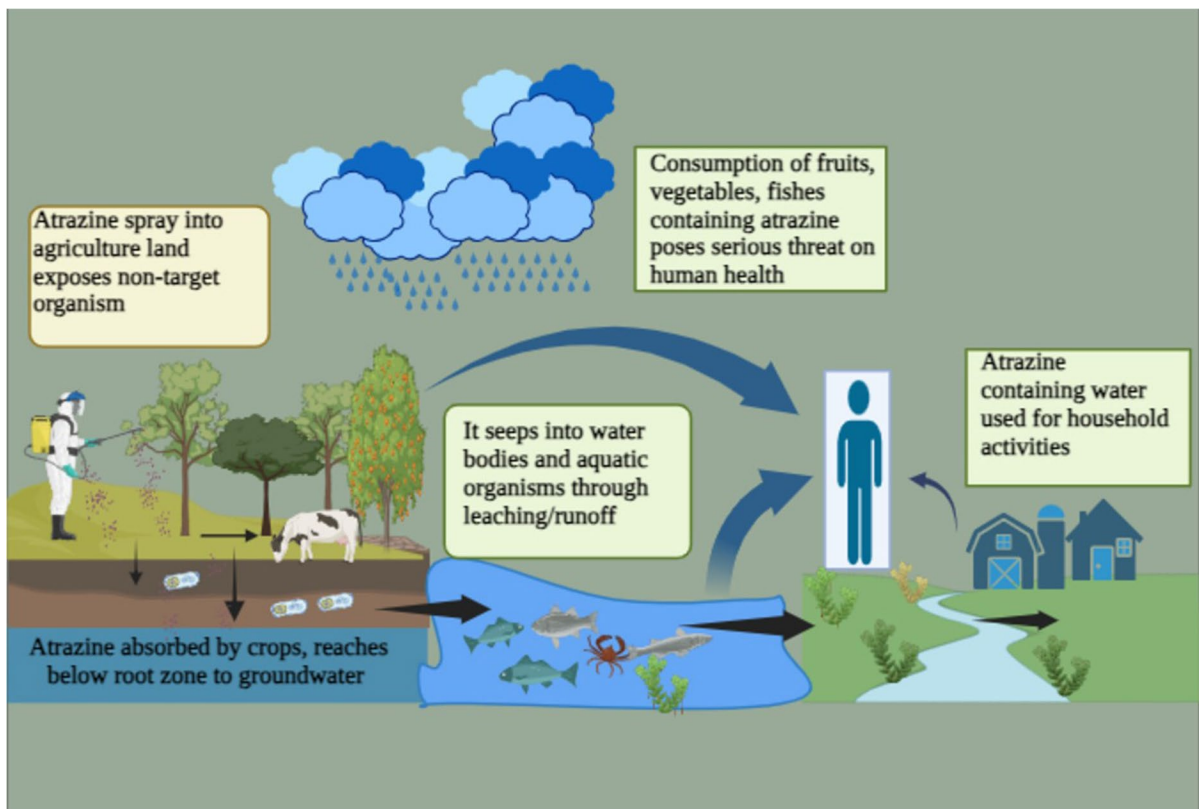


Fig. 1 The dispersion of atrazine from the agricultural crop, soil microbiota, aquatic life, and biomagnification to the human body is depicted

bile acid secretion (Kennelly et al., 2018; McDaniel et al., 2002; Trousil et al., 2014). Atrazine exposure decreased the production of phosphatidylcholine and CDP-choline indicating the altered metabolism of these compounds (Yin et al., 2020). It has been observed in mice that Perinatal exposure, even to a lower dose of atrazine (1.4 mg kg^{-1}), has been associated with abnormal behavior patterns linked with brain dopamine and serotonin disruption (Lin et al., 2014). Exposing rats to atrazine has caused alteration in the expression levels of proteins like transferrin receptor (TFR), divalent metal transporter1 (DMT1), Hephaestion (HEPH), and ferroportin1 (fpn1), which are essential regulators of Fe transporter maintaining homeostasis of the mid brain (B. Li et al., 2021a, b). In the zebrafish, atrazine exposure has shown adverse effects on the development of primary and secondary sexual characters in males and females, affecting LH surge, delay in mammary gland development, delayed

vaginal opening, and GnRH release (Gonadotropin releasing hormone) (Stradtman & Freeman, 2021).

Atrazine has been shown a significant decrease in methylated CPG (meCPG) proteins, histone 3 lysine, and 9 methylated (H3K9me3) proteins, causing altered enzymatic expression levels overall decreasing the cell growth rate in human kidney cell line (HEK293T) (Sánchez et al., 2020). At the genetic level, metabolomics studies indicate that atrazine alters the expression of various anti-apoptotic Bcl-2, Lc3-II, Mtorc1, TNF- α , IL6, BEX 2, and GSH genes, causing a change in the expression of several proteins (Li et al., 2020). To understand the toxicity and underlying mechanism involved, various toxicology tests have been performed on human SH-SY5Y cell lines showing decreased viable neuroblastoma cells and increased MCF7 cell proliferation (Sogos et al., 2021); (Lu et al., 2022). Figure 2a represents the adverse effects of atrazine exposure on different organs of the human body, disturbing regular

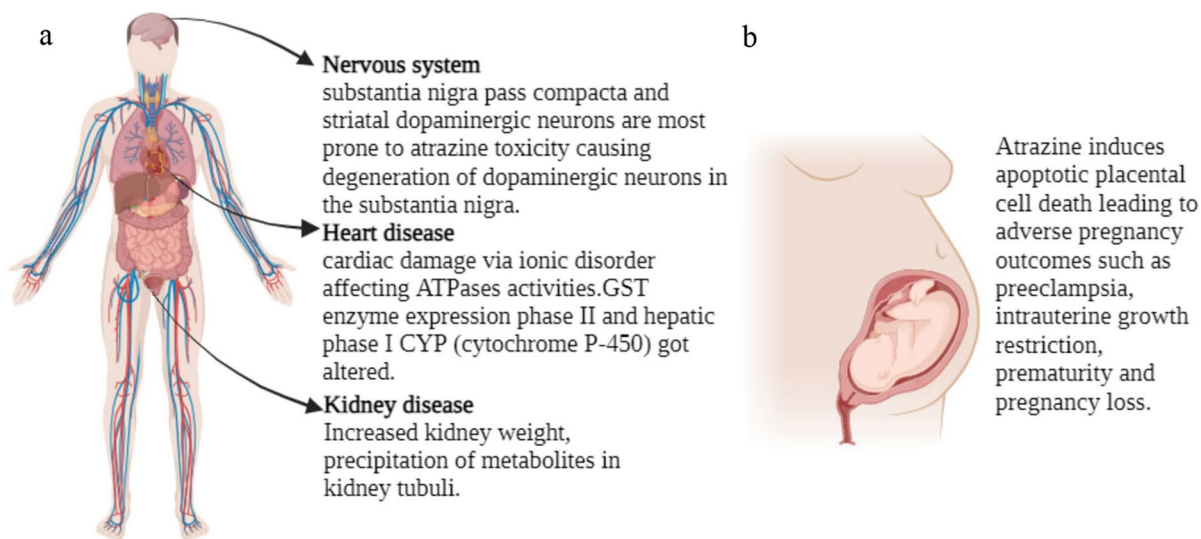


Fig. 2 (a) The inhibitory effects of atrazine on different human organs affecting normal human physiology and (b) the adverse outcomes during pregnancy

physiological activity, while 2b depicts the adverse outcome on the health of pregnant women when exposed to atrazine, causing many abnormalities during and post partum.

Atrazine exposure showed metabolic alterations in folate biosynthesis, affecting purine and pyrimidine synthesis and suppressing various physiological processes like cell division (Lu et al., 2022). In metabolomics study determining the alteration in metabolic pathways, the metabolites synthesis is getting affected has been linked to oxidative stress, downregulation of gene expression with an increase in gluconeogenesis and reduced oxidative phosphorylation and ATP synthesis mechanism such as glycolysis and citric acid cycle. (Lin et al., 2014); (Yin et al., 2020). Intoxication occurred in cattle when ten of 40 cows died because of, multi-organ mitochondrial dysfunction and oxidative stress indicating acute toxicity (Props et al., 2021). Atrazine mercapturate (metabolite) presence was detected in urine samples of corn farmers', but no major association has been detected between the atrazine metabolite and the three markers of oxidative stress that are malondialdehyde (MDA), 8-hydroxy-2'-deoxyguanosine (8-OHdG), and 8-isoprostaglandin- $F_{2\alpha}$ (8-isoPGF). MDA is created when Reactive oxygen species reacts with polyunsaturated fatty acids, 8-hydroxy-2'-deoxyguanosine (8-OHdG) is

a marker of oxidative injury causing lesion in DNA (Muniz et al., 2008) while (8-isoPGF) is a prostaglandin compound produced during non-enzymatic lipoprotein peroxidation (Lee et al., 2006). However when atrazine mercapturate measured above limit of detection (LOD) an association with 8-OHdG was observed (Lerro et al., 2017). The no observed adverse effect level (NOAEL) and lowest observed adverse effect level (LOAEL) for infants and children are considered at 6.25 mg/kg/day and 12.5 mg/kg/day for short-term oral dietary exposure. Meanwhile for females aged (13–50 years), US EPA NOAEL is 10 mg/kg/day whereas the LOAEL is 70 mg/kg/day (Stradtman & Freeman, 2021).

3.2 Inhibitory Effects on Aquatic Organisms

Atrazine has been shown to induce histopathological, hematological, reproductive, carcinogenic, and genotoxic changes in aquatic organisms. In the aquatic system, the half-life of atrazine is about 168 days (Stradtman & Freeman, 2021). we summarized the inhibitory aspects of atrazine on various aquatic and non-aquatic organisms exposed at a specific concentration and periods in Table 2. Atrazine has been shown to cause oxidative damage in microalgae, studied by calculating antioxidant response and oxidative stress in algae (Castro et al., 2021).

Atrazine and its metabolites desethyl atrazine and de isopropyl atrazine are phytotoxic, directly affecting various phytoplankton (Yang & Zhang, 2020). Office of environmental health hazard management (OEHHA) officially declared atrazine had endocrine disruptor activity mainly in reproduction toxicity (Zheng et al., 2017). Reproductive toxicity has been observed to cause population declines in various aquatic organisms, including amphibians, gastropods, shellfish, fishes, and crustaceans (Lopes-Lima et al., 2017).

At a concentration range of (0.02–20 mg L⁻¹), abnormalities ranging from axial shortening, tail flexure, and facial edema have been noticed in *Anaxyrus americanus*, *Lithobates pipiens*, and *Lithobates sylvaticus* frogs (Rohr & McCoy, 2010). The continuous leaching of pesticides in aquatic ecosystems shows the highest bioaccumulation in fishes in the food chain (Yang et al., 2021). Meanwhile, the gonadal intersex abnormality is observed due to atrazine toxicity in fishes, wherein an egg yolk precursor protein, the vitellogenin gets abnormal. This vitellogenin protein is responsible for improper sex development, abnormal steroid levels, fertility, and reproductive problems in aquatic organisms (Rohr & McCoy, 2010). Atrazine reduces the expression of vitellogenin protein in the female crayfish ovaries and hepatopancreas forming smaller oocytes (Silveyra et al., 2018). In estuarine crabs, it has been seen that atrazine exposure has decreased the growth of ovaries (Silveyra et al., 2017). In fathead minnows, an atrazine concentration range of (0.5–50 g L⁻¹) downregulated vitellogenin protein, decreasing oocyte maturation (Ali et al., 2018). At a concentration range of 2.5 g L⁻¹ atrazine, feminine characteristics were observed in male *Xenopus laevis* and increased production of testicular oocytes was observed on consecutive exposure of 7 days (Ali et al., 2018).

The embryotoxicity of atrazine and its degraded products (DE isopropyl, atrazine, and desethyl atrazine) was studied in zebrafish, observing mortality, hatching, and edema at 24, 48, 72 and 96 h post fertilization (hpf) showing retardation in hatching at 96 hpf and pericardial edema just at 48 hpf at varying concentrations (Zheng et al., 2017). Atrazine exposure of 30–300 µg L⁻¹ disturbs the swimming pattern of larval zebrafish (Tai et al., 2021). In common carp, mortality was found at an atrazine concentration of 0.3–300 g L⁻¹ when exposed

for 33 days (Blahova et al., 2020). Increased axial malformations were reported when tadpoles were exposed to atrazine (Hanson et al., 2019). Over six days, exposure to 0.003 mg L⁻¹ atrazine in goldfish has been shown to cause severe stress on circadian rhythm (Ren et al., 2019). Atrazine exposure has been shown to cause detrimental effects on oxidative stress, protein carbonyl level, cell injury, and lysosomal stability by generating free radicals when exposed for 7 days in marine blue mussels (Shaw et al., 2019). However, in green mussels, exposure of atrazine did not affect immunity and differentiation in the male and female sexes (Juhel et al., 2017). In oysters, high concentrations of atrazine exposure induce GST (glutathione s-transferase), further activating antioxidant enzymes in different organs (Geret et al., 2013).

3.3 Inhibitory Effects on Microbiota

Various studies have shown the effects of atrazine on regulating parameters like growth activity (phytohormones), biochemical functions, cell physiology, cell morphology and molecular activity, and microbial communities (Shahid et al., 2019). It has been reported that atrazine causes decreased plant growth-promoting traits in *Azotobacter vinelandii* strain AZ6, decreasing IAA (Indole 3 acetic acid) production, phosphate solubilization, and phenolate siderophore production, causing oxidative damage and membrane destruction (Shahid et al., 2019).

Gut microbiota is vital in maintaining gut and overall health in humans and animals. The presence of atrazine in food and crops has significantly disturbed gut microbes (Luo et al., 2021). It has also been reported that atrazine has altered the microbiome composition, decreasing *Acetobacter acetii* and *Rhodospirillales* compared to *Lactobacillus acidophilus* and other genera impacting the gut microbiome of *Drosophila melanogaster* (Brown et al., 2021). Atrazine either increases or has no significant effect on soil microbial biomass, as it has been reported that the respiration rate increases, boosting the soil microbial metabolic rate (Bonfleur et al., 2015). In a study on the effects of atrazine on the bench microbial nutrient assessment by analyzing nutrient absorption and remineralization of phosphate, ammonium, and nitrate, atrazine has been shown not to have a significant impact on nutrient cycling

(Elias & Bernot, 2014). When atrazine concentration (2–10 mg kg⁻¹) was applied, microbial diversity decreased from 2.59 to 2.23, showing *Microvirga* species., *Haplosporidium* species., and *Sphingopyxis* species., absence compared to the control test (Chen et al., 2015). Applying atrazine has increased bacterial strains of *Betaproteobacteria* belonging to *Methylophilicea* and *Nitrosomonadacea* family (Briceño et al., 2010). At the genetic level, various changes have been detected when exposed to atrazine, causing the downregulation of some genes and affecting cell morphology and physiology (Brown et al., 2021). In mice, abundant *Rodentibacter pneumotropicus* has been reported when exposed to atrazine, which might cause ailments like conjunctivitis and autophagy signaling disturbance in the liver, causing abnormal enzyme cascades (Liu et al., 2021).

3.4 Inhibitory Effects on Non Target Aquatic and Crop Plants

The various physiochemical properties of soil, such as its binding capacity, phytotoxicity, and biotoxicity, determine the bioavailability, migration, and half-life of herbicides in the soil ecosystem and subsequently in the plants also (Y. Zhang et al., 2021a, b). *Wolffia brasiliensis*, an aquatic plant, showed a mortality rate of 16% at 11.2, 36.5, and 118.0 mg L⁻¹ concentrations of atrazine, indicating the herbicide risk in the aquatic environment (Pereira et al., 2019). In *Arabidopsis thaliana*, plant atrazine interactions have shown several energy changes, causing mitochondrial dysfunction, increased ROS, and PSII inhibition (Alberto et al., 2017). When in carrots, cucumber, lettuce, onion, perennial ryegrass, and tomato, rate-response trends were observed, causing photosynthesis starvation in plants due to the generation of reactive oxygen species capable of degrading cell membranes (Brain & Hoberg, 2016). When exposed to more tolerant plants or to lower herbicide concentrations, this deleterious effect caused by reactive oxygen species reduces the photosynthetic rate of plants, reducing the accumulation of dry matter and altering the normal development of plant growth. In seagrass, *Halophila ovalis* exposure of herbicides has reduced the photosynthetic ability to block the electron transport chain by binding to protein in the thylakoid membrane and displacing the plastoquinone affecting the synthesis of ATP and NADPH (Wilkinson et al., 2015).

4 Atrazine and the Transformed Compounds

Atrazine belongs to the chloro s triazine class of compounds constituting atrazine, simazine, propazine, and chlorinated by-products. The metabolites formed as a result of the natural degradation of atrazine in the environment are de-ethyl ATZ (DEA), de-ethyldeisopropyl hydroxy ATZ (DEDIHA), de-isopropyl ATZ (DIA), di-dealkyl ATZ (DDA), hydroxy ATZ (HA), atrazine mercapturate (AM), desethylterbutylazine (DET), de-ethylhydroxy ATZ (DEHA) and de-isopropylhydroxy ATZ (DIHA). The primary degraded products formed as a result of Atrazine degradation are deethylatrazine (DEA), deisopropylatrazine (DIA), and didalkylatrazine (DDA), show equal or more toxicity than atrazine (Bhatti et al., 2022). The occurrence, distribution, and degradation of atrazine and its metabolites depend on various parameters like seasonal variation, solar radiation, temperature, precipitation, dissolved oxygen, salinity, wind, sea currents, pH, the mode of transportation, and geographical area (Y. Xue et al., 2021a, b) (Bhatti et al., 2022). After the mineralization of atrazine and the intermediate metabolites, the final products consist of ammelide, cyanuric acid, CO₂, and NH₃. Table 3 depicts the different physio-chemical natures of atrazine and its metabolites formed due to the biodegradation of atrazine.

The potential of desethylterbutylazine (DET) as a continuous leachate is notable (Bozzo et al., 2013) as its leaching potential (Gustafson, 1989) or Groundwater Ubiquity score (GUS) was estimated at 3.07 (Hertfordshire, 2017), showing higher leaching potential in soil. (Jian et al., 2021), whereas in seawater of Xiangshan Harbor atrazine and its metabolites detection fraction ranges ATZ (89.7%), DIA (4.6%), DEA (0%) and DDA (5.7%) (Y. Xue et al., 2021a, b). In soils, DEA (2700–3200 mgL⁻¹) and DIA (670–980 mgL⁻¹) showed more mobility as well as leachate fractions than the parent compound, ATZ (30–33 mgL⁻¹), explaining the presence of less alkyl groups on the s-triazine rings increasing the polarity (Bhatti et al., 2022). In a study, it was corroborated that the parent compound (ATZ) degrades faster than its metabolites (DEA, DIA) as the presence of more alkyl subunits in atrazine enhanced the hydrolytic degradation suggesting metabolites may be more recalcitrant than the parent compound (Bhatti et al., 2022).

A study showed that the acute, carcinogenic, and mutagenic toxicity of atrazine and its metabolites

pose a continuous threat to organism's lives (Li et al., 2014). In other significant findings, it was found that not only ATZ but its metabolites, DACT, DIA, and DEA have the same or even more inhibitory effects on developmental and locomotory stages of zebrafish constituting even at $300 \mu\text{g L}^{-1}$ exposure of ATZ causes no inhibitory effects but $100 \mu\text{g L}^{-1}$ and $300 \mu\text{g L}^{-1}$ of DIA and DEA induces developmental toxicity (Liu et al., 2016). Another study showed that the concentration of atrazine and the degraded by-products in different aquatic organisms in Xiangshon harbor seawater shows that benthic organisms Ditrema and Black sea bream have increased bio-concentration of ATZ, posing low risk to fish and invertebrate (Y. Xue et al., 2021a, b). For soil species, springtail and earthworm, metabolites DEA and DIA pose unacceptable hazards, while for avian and mammalian species, it poses moderate to mild threats (Bhatti et al., 2022). Among the 15 metabolites formed due to atrazine degradation, the significant metabolites formed are DEA and DIA by the N dealkylation of the parent molecule's ethyl or isopropyl side chains (Lin et al., 2008). Figure 3 depicts the primary and secondary metabolites formed through different metabolic pathways.

5 Atrazine Remediation

Scopus database was used to search for the literature on the degradation of atrazine using the keywords (ATRAZINE AND DEGRADATION) OR (REMOVAL) OR (TREATMENT) OR (CONTROL). A total of 7,317 documents from the year 2012 to the current year (2023) were obtained out of which 3,585 documents were found to be relevant. VOS VIEWER network visualization tool was used to better understand the various treatment technologies applied and also the upcoming technologies. The threshold value selected was 5 showing the number of times the keyword is reoccurring. Figure 4 a shows the co-occurrence map. The different classes of Clusters show the topic area studied for the degradation of atrazine. A total of 15 clusters were observed out of which 6 clusters were found to be distinct and each is represented by a different colour. Figure 4 corroborates the research focus of the article. The map shows the diversity and uniqueness of degradation methods for the atrazine pollutant and their inter relatedness. Cluster 1 (dark green) Fig. 4 (a) focusses on advanced oxidation processes (AOPs) covering ozonation, UV, photocatalysis, hydroxide radical, UV/H₂O₂. Cluster 2 (dark

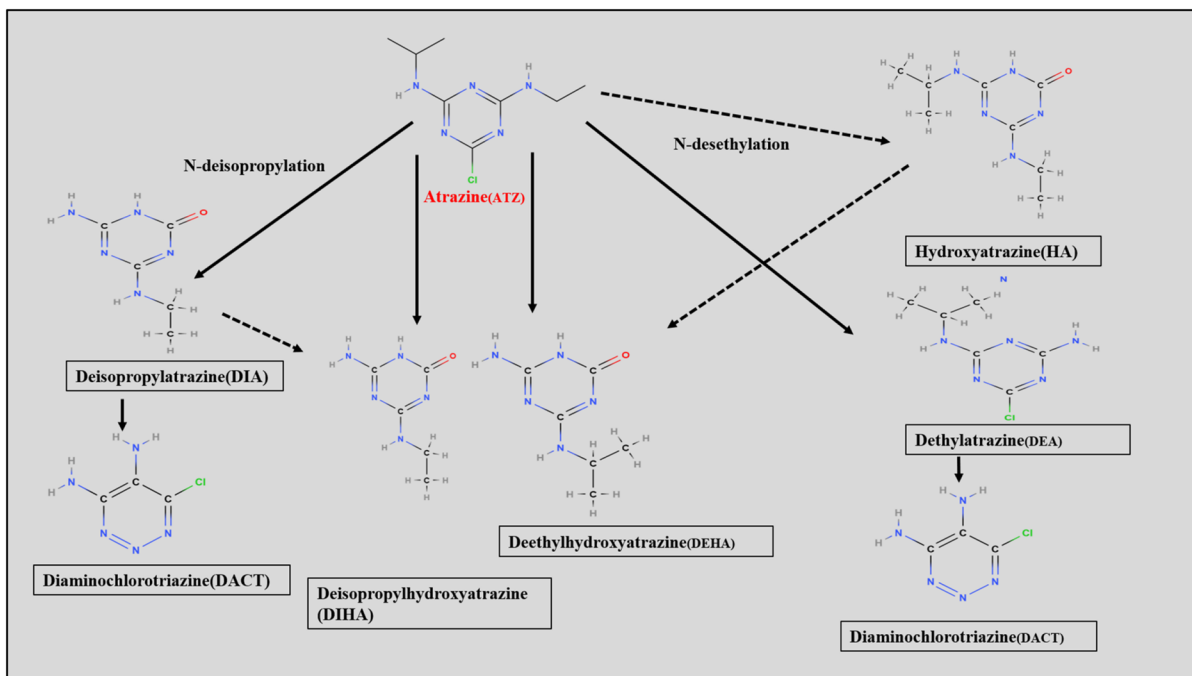


Fig. 3 Degradation pathways of atrazine following the formed transformed metabolites through different metabolic pathways

brown) Fig. 4 (b) focusses on sorption, desorption and adsorption based on biochar, activated carbon as well as the related kinetic and mechanistic studies. Cluster 3 (dark orange) Fig. 4 (c) highlights biodegradation aspect of atrazine removal including mineralization, encapsulation, bioaugmentation involving different bacterial communities. Cluster 4 (light blue) highlights the monitoring of pollutant and its transformed products in surface water, drinking water using gc–ms, ic–ms/ms analysis. Cluster 5 (yellow) highlights the histopathological and immunotoxicity of atrazine in different organisms. Cluster 6 (dark red) covers the efficacy of atrazine in controlling the growth as well its effect on different crops. According to this cluster analysis, biodegradation of atrazine, AOPs, and toxicity have been the major areas of research focus. VOS VIEWER overlay visualization also points out to the recent application of synergistic removal techniques

like photoelectrocatalysis, photo – fenton etc. and also the new classes of catalyst applied for AOPs.

5.1 Microbial Degradation of Atrazine

Microbial remediation has been extensively used for atrazine removal from contaminated soils as it has a low application cost and is less toxic to the environment (Rehan et al., 2014). The efficiency and mechanism of atrazine degradation depend on the soil microbial community and their physiology. Dichlorination, dealkylation, and deamination are the main degradation pathways utilized by bacteria for atrazine degradation. Some bacteria break atrazine through dichlorination, replacing the OH group with a chlorine atom. Hydroxy-atrazine formed was further degraded into N- isopropylammelide or N-ethylammelide proceeding through hydrolytic deamination reactions, producing cyanuric

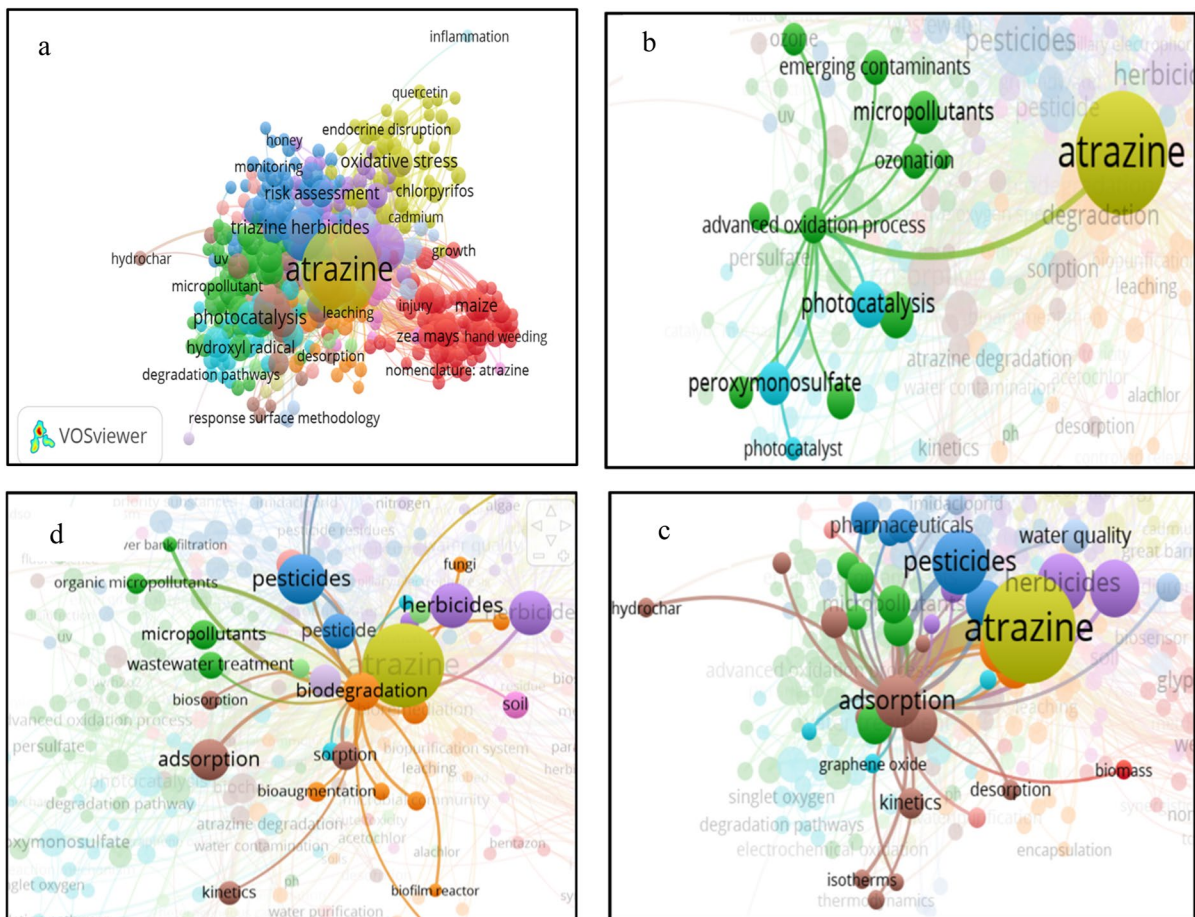


Fig. 4 The figure displays the keyword Atrazine Degradation network analysis using VOS viewer software. a) Shows the co-occurrence map b) Highlights the advanced oxidation processes c) focuses adsorption methods and d) specifies Biodegradation technique

acid's final degraded product. (Rostami et al., 2021). Gram-negative and Gram-positive bacteria follows different degradation pathways. Gram-positive bacteria degrade atrazine following a hydrolysis reaction catalyzed by enzymes such as TrzN, (triazine hydrolase) while gram-negative bacteria catalyze the reaction by the enzyme AtzA (atrazine chlorohydrolase) playing an essential role in the breakdown process. (Huang et al., 2017). *Paenarthrobacter ureafaciens* ZF1 a Gram+ bacteria has the potential of degrading atrazine (99.3%) (100 mg kg^{-1}) from soil while in liquid it could completely remove atrazine within 2 h (Zhang et al., 2022). *Bacillus pumilus* and *Bacillus subtilis*, which are gram+ bacteria could, successfully degrade 95% and 98% of atrazine respectively (Zhu et al., 2022). Figure 5, shows the degradation pathway utilized by gram negative bacteria involving atrazine chlorohydrolase (AtzA) enzyme, hydroxydechloroatrazine ethylaminohydrolase (atzB), and N-isopropylammelide isopropylaminohydrolase (atzC).

New techniques, like microbial encapsulation, have been developed for treating soil contaminated with atrazine. (Rostami et al., 2021). It has also been observed that the microbial strain *Bacillus velezensis* *MHNK1* producing surfactin lipopeptide resulted in $100 \pm 1.20\%$ degradation within 4 days. Presence of atrazine degrading genes and surfactin are potential source in removing atrazine (Jakinala et al., 2019). It has been shown that bioaugmentation has enhanced the atrazine-degrading efficiency of constructed wetland. *Pseudomonas* and *Arthrobacter* sp. were dominant among the atrazine degrading microbial

community because of high adaptability and atrazine degrading capability in the constructed wetland (Zhao et al., 2019). Co-culture of *Arthrobacter* sp. DNS10 and *Enterobacter* sp. P1 degraded $99.18 \pm 1.00\%$ of atrazine as compared to $38.57 \pm 7.39\%$ by the single microbial strain DNS10. The expression of the atrazine degradation-associated genes *trzN*, *atzB*, and *atzC* was also more as compared to single microbial strain treatment (Jiang et al., 2019). We summarized the different microbial strains used for treatment of atrazine and the biodegradation efficiency in Table 4.

5.2 Fungal Remediation

Fungi, an essential constituent of the soil ecosystem, play a significant role in degrading atrazine after bacteria. Fungi degrade or transform recalcitrant compounds into biotransformed products which are further broken down by other soil microorganisms (Maqbool et al., 2016). The plasmidial genes in bacteria *atzA*, *atzB*, and *atzC* encode the various enzymes breaking the compound through several metabolic pathways. Gene *atzA* follows dichlorination and s-triazine ring cleavage, gene *atzB* (hydroxyatrazine N-ethylamino hydrolase enzyme) proceeds the reaction by hydrolytic conversion of hydroxyatrazine to N-isopropylammelide whereas *atzC* (N-isopropylammelide isopropylaminohydrolase) catalyses the hydrolysis of N-substituted amino dihydroxy-s-triazines as well as N-isopropylammelide to cyanuric acid and isopropylamine (Fan & Song, 2014b). Further, genes encoding *atzD* metabolises cyanuric acid whereas *atzE* and *atzF* hydrolase

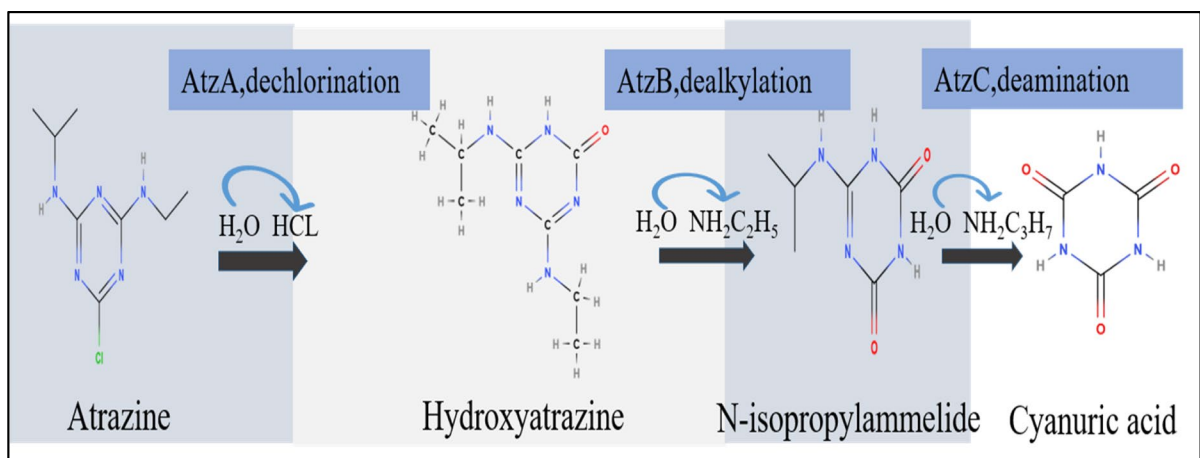


Fig. 5 Metabolic pathway of atrazine degradation through bacterial communities

cyanuric acid further yielding carbon dioxide and ammonia each of three moles (Fan & Song, 2014b). So far, these proteins have not been characterized in fungi. Enzyme atzA has been reported to catalyse the chlorohydrolysis of atrazine, deisopropylatrazine (DIA) and desethylatrazine (DEA), but not desethyldeisopropylatrazine (DEDIA) although this enzyme has been described only in bacteria, some researchers identified the chlorohydrolysis products from atrazine breakdown by fungus (Lopes et al., 2020). Conversely, Fungi have a complex set of hydrolytic and oxidative enzymes with N-dealkylation, deamination steps, or both forming DEA, DIA following diverse metabolic pathway (Esparza-Naranjo et al., 2021). Studies have shown that mycorrhizal and nonmycorrhizal fungi were associated with degrading atrazine. However, ericoid mycorrhizal fungi showed the best mineralization capacity, highlighting that the degradation depends upon herbicide and fungus, irrespective of fungal ecotype (mycorrhizal or free-living) (Donnelly et al., 1993).

Arbuscular mycorrhizal fungi (AMF), which forms a symbiotic association with the plants, has been observed to play a great role in removing atrazine. In studies it has been reported that atrazine removal

efficiency was upto 74.65% in *Medicago sativa* mycorrhizal stating atrazine degradation rate is higher in mycorrhizal treatments than those in non-mycorrhizal treatments (Song et al., 2016). During the contaminants degradation process by AMF, the exudates secretion by fungi may change the dynamics of rhizosphere soil.

microbial activity affecting the rate of atrazine dissipation in the soil (Fan et al., 2020). Lignolytic enzymes produced by white rot fungi have been classified into three types of peroxidases-manganese, lignin peroxidases,

and laccases. Ligninolytic enzymes applied under the controlled or symbiotic association of deuteromycetes with soil bacteria have shown complete mineralization of atrazine (Chan-Cupul et al., 2016; Jin et al., 2016). Fungi follow the intracellular and extracellular enzymatic degradation pathways wherein Basidiomycetes and ascomycetes follow the extracellular enzymatic ligninolytic complex degradation pathway leading to high degradation efficiency of atrazine (Deshmukh et al., 2016; Fan & Song, 2014a). Figure depicts the degradation of atrazine favours N-dealkylation of ethylamine and/or isopropylation Figure 6.

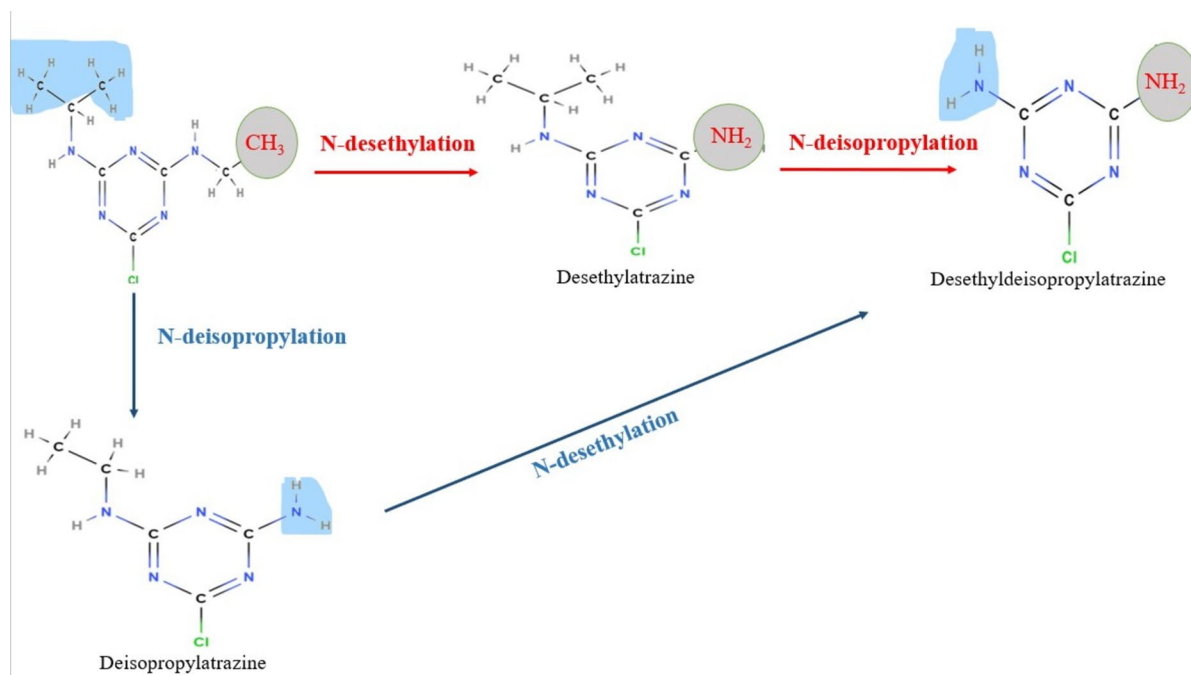


Fig. 6 The mechanism N-desethylation and N-deisopropylation showing breakdown pathway of atrazine and subsequent metabolites in *pleurotus ostreatus* fungus adapted from (Lopes et al., 2020)

5.3 Phytoremediation

Phytoremediation is a widely used technique in which plants remove pollutants through degradation, uptake, evaporation, and by increasing the rhizobacterial communities. Plants with dense root systems help to reduce soil contaminants by decreasing the transport rate of herbicides in the soil, thereby acting as a natural filter in the environment (Chellaiah & Yule, 2018). Phytoremediation capacity is mainly shown by vascular plants, macrophytes, and some gramineous plants. Potentiality of some plants in remediating atrazine is well depicted because of its fast growth even in adverse conditions, large biomass exhibiting large specific surface area with ample macromolecules, and fibrous dense root system. The mechanism is generally associated with absorption, accumulation, and detoxification through enzymatic activity of glutathione-S-transferase (GST) as well as bacterial assisted rhizodegradation (Loureiro et al., 2023; McKnight et al., 2022; Zhang et al., 2023). The biosorption approach includes the absorption either by submerged roots and by the leaves. Plants secretion of root exudates provides energy source and favourable niche for rhizospheric bacterial communities increasing the activity of the microbial population by which the breakdown of herbicide is enhanced (Zhang et al., 2023). In a study on *Cyperus alternifolius* plant exposed to atrazine concentration at 20mgL^{-1} , the phytoremediation efficiency was $91.28 \pm 6.35\%$ (Ameri Siahouei et al., 2020). Other findings showed that an increase of three times of *Typha latifolia* rhizomes causes an increase in the degradation rate of atrazine at twice the rate showing the result that terbuthylazine and its metabolite get accumulated in plant tissues (Papadopoulos & Zalidis, 2019). Macrophytes *potamogeton crispus* and *Myriophyllum spicatum* absorbed atrazine and DEDIA from the sediment, remediating atrazine from the sediment and water (H. Li et al., 2019a, b). Plants like eucalyptus, *Hymenaea coubaril*, and *Cecropia hololeuca* showed phytoremediation efficiency in the Quartzarenic Neosol soil (Heemann et al., 2018). In a study, it was found that the two treatments BR (Bean Rhizobium) and BRT (bean Rhizobium—Trichoderma) significantly caused the removal of 20 mg of atrazine from 50 g of soil (Madariaga-Navarrete et al., 2017). The plant species *Iris versicolor* has the potential to

phytoremediate atrazine by 58.7% after 112 days of treatment, with no plant death reported. However, stunted growth or leaf injury was reported suggesting plants could produce more above-ground biomass to recover from herbicide contamination. (McKnight et al., 2022). The U.S environmental protection agency (2006) suggested less than $1\mu\text{gL}^{-1}$ of chronic atrazine for non-vascular plant species. However, metabolization of atrazine by crop plants are yet to be explored.

Prairie grasses have shown promising phytoremediation efficiency in removing atrazine which gets degraded into its metabolite, which later is accumulated in leaves at approximately (60 – 80)% (Madariaga-Navarrete et al., 2017). Another study showed that the Maize plant had more potential to accumulate atrazine in its tissues. (Sánchez et al., 2020). With the inoculation of *Cannabis indica* with *Funneliformis mosseae*, removal percentage rate of atrazine increased from 68.064% to 95.670%, indicating a viable phytoremediation approach for in situ remediation (Dong et al., 2016). A study found that maize plants planted with *Penisetum clandestinum* degraded 45% of atrazine in about 80 days, showing *Zea mays* have the potential for phytoremediation of soils contaminated with atrazine (Ibrahim et al., 2013).

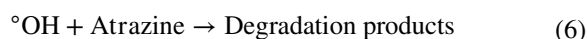
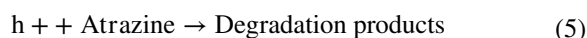
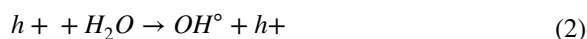
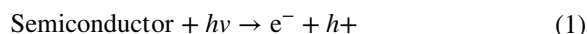
5.4 Advanced Oxidation Process

Advanced oxidation processes are chemical processes that include a set of chemical treatments, like ozone, hydroxide radicals, and UV irradiation. AOPs treatments are based on releasing active radicals like $\text{O}_2^{\bullet-}$, $\cdot\text{OH}$, and $\text{SO}_4^{\bullet-}$ breaking the contaminants into small inorganic molecules. Oxidants like ozone, oxygen, hydrogen peroxide, sulfite ion, and catalysts like titanium oxide, metal oxides, and various compounds in the synergistic effect produce the radicals.

5.4.1 Photocatalytic Degradation

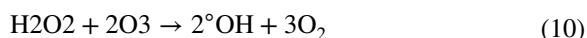
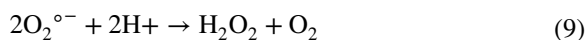
Photocatalysis is one of the most powerful technology used to treat organic contaminants due to its powerful mineralizing and oxidizing capacity (Xu et al., 2013). Photocatalytic degradation is associated with the formation of $e^- - h^+$ pair (Eq. 1) when irradiated with light, having a wavelength equal to or more

than the bandgap energy on the semiconductor photocatalyst (Poonia et al., 2022). Further, the generated electrons and holes react with water or oxygen to form superoxide, hydrogen peroxide, hydroxyl radicals (Eq. 3,4), and hydroperoxyl radicals. Reactive oxygen species (ROS) generated plays a ubiquitous role in degrading atrazine (Eq. 5,6 and 7) using an array of photocatalysts such as metal oxides (TiO₂, ZnO) and sulfides (ZnS, CuS) (Poonia et al., 2022) and nano or mesoporous compounds (BaTiO₃, Bi₂MoO₆) (Sobahi & Amin, 2021) (Sharma et al., 2019). A study showed that when an aqueous solution of atrazine was photolyzed ($\lambda=254$ nm) under the optimum condition of low pressure (LP/UV/H₂O₂), about 90% of ATZ got degraded in one hour (Li et al., 2012). Various studies have proved that Titania is an effective photocatalyst in the degradation of atrazine, where titania film is applied at the surface of quartz crystal, making degradation two times higher than noncoated titania films (Zhang et al., 2013). It has also been observed that the doped TiO₂ with the lowest nitrogen loading degraded atrazine at higher rate (Samsudin et al., 2015). Another study evaluated the photocatalytic activity of N- TiO₂/ZSP (ZnS-based phosphors microparticle) to remove atrazine under UVA light radiation (Sacco et al., 2015). It has also been observed that AC/g-C₃N₄ composites with peroxymonosulphate enhance the atrazine's photodegradation efficiency by 57.90%, mainly due to the efficient charge carrier distinguishing ability of the composite and greater absorption capacity (Dikdim et al., 2019). Table 5 describes the degradation of atrazine by different photocatalysts.



5.4.2 Ozonation Process

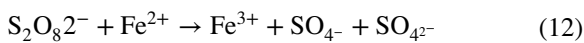
Ozone is a powerful oxidizing agent that proved to be a powerful alternative yielding a higher degradation rate than conventional oxidation (Glaze et al., 1987). O₃/UV removes the contaminants by producing hydroxy radicals (Eq. 8 and 10) as the formed H₂O₂ (Eq. 9) acts as a promoter in ozonation process maintaining the redox cycle reaction. Table 6 summarizes the present scenario; along with the ozonation process, catalysts are added, enhancing the degradation rate of contaminants. MnCe-CM oxides showed the best efficiency with 99.99% atrazine degradation in 40 min, showing that the novel MnCe-CM has dual functions of filtration and catalytic ozonation (He et al., 2022) Plasmon-enhanced catalytic ozonation with silver doped spinel ferrite has shown excellent degradation efficiency compared with ozonation and catalytic ozonation processes (Yang & Wu, 2022). A study demonstrated that synthetic 4A zeolite showed promising results in removing atrazine at a rate of 87.5% in 6 min (Su et al., 2022). In MnOx/biochar and FeOx/biochar, the ozonation efficiency of ATZ increased to 83% and 100%, respectively, reducing the acute toxicity of atrazine from 38.3% to 6.3% (Tian et al., 2021).



5.4.3 Persulfate Oxidation

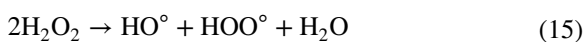
Persulfate-based remediation technology is one of the most reliable techniques. Light, (Eq. 11) heat, metal ions, and carbon compounds can easily activate persulfate to form SO₄^{•-} radicals (Diao et al., 2021). These compounds enhance the oxidative degradation (Eq. 12) of atrazine in different environments. In this process, the Sulfate and hydroxy radicals react faster with atrazine, forming DIA and DEA (Lutze et al., 2015). In the persulfate advanced oxidation process, catalysts enhance the degradation capability shown in Table 7. A study found that the Co₃O₄ catalyst at high peroxymonosulphate (PMS) concentration removed about 20 μM ATZ with 2.0 mM PMS and 0.4 g/L Co₃O₄ at pH 6.0 (Fan et al., 2017). The Boron-doped diamond (BDD) anode

used to activate persulfate (PS) demonstrated that degradation of ATZ got increased by 78.2% with the rise in the current density and quantity of PS (not more than 1.0 mM). However, the mechanism underlying these processes needs to be studied more (Bu et al., 2018). It was observed that persulfate activation with the mass ratio of 5:1 for nanoscale zero-valent iron to graphene showed the highest potential for atrazine catalytic degradation, removing 92.1% of atrazine within 21 min, showing degradation efficiency increased with the rise in persulfate concentration (Wu et al., 2018).



5.4.4 Fenton Treatment

Fenton treatment is an advanced oxidation process of ferrous salt (Fe^{2+}) and H_2O_2 called Fenton's reagent degrading organic contaminants. In the Fenton oxidation–reduction reaction, the free radicals are generated (Eq. 13 and 14) upon the reaction with iron ions with H_2O_2 . Hydrogen peroxide acts as an oxidizer and catalyst, accelerating the redox cycle ($\text{Fe}^{2+}/\text{Fe}^{3+}$) and increasing toxicants' degradation efficiency rate. Fe/TiO_2 heterogenous Fenton and visible light photocatalytic activity showed atrazine degradation of 10 mg \cdot L $^{-1}$ at pH 3 (Yang et al., 2020). In a study, the ability of the photo Fenton process on the atrazine degradation was demonstrated, showing a mineralization rate of 62.5% in about 60 min (Benzaquén et al., 2012). Atrazine gets degraded at the rate of 20%, 60%, and 70% respectively under fenton, UV-A photo fenton and UV-C photo fenton treatments (De Luca et al., 2013). In the visible TaON (Tantalum) Fenton-like system, the degradation rate was higher, but dissolved oxygen reversed the degradation efficiency as the residue of atrazine left was 10% and 15% at 60 min in the absence and presence of DO, respectively (Zhang et al., 2014).



6 Challenges and Future Perspectives

In most countries, atrazine is the maximum percent of detected herbicide in different sampling sites across rivers, estuaries, bays, and agricultural lands. The physiochemical properties of this compound make it less degradable, enhancing the retention time in the environment. The herbicide is banned in some countries, but its presence crossing the minimum residual level is continuously detected in rivers, soils, groundwater, tap water, and surface water, posing a severe threat to aquatic and non-aquatic organisms, microbes, and mammals. It is an area of concern when multiple exposures occur. There is always a potential for multigenerational effects in an organism's different developmental stages, making it more hazardous to living forms.

Some of the critical future research perspectives are given below:

- Bioremediation is currently the most reliable technique for the degradation of atrazine. However, the removal of atrazine from soils require a consortia of microbes or fungi for better remediation. Bio-augmented mixtures have already proved to be of more excellent value in remediating herbicides.
- Recently, bioorganic fertilizers have been proven to reduce the phytotoxicity of atrazine, and future research should lay more importance on the behavior of atrazine and its metabolites in different environments.
- Selecting green compounds/ biomaterials that are budget-friendly and cause no harm to the environment should be the focus of further research. Chemical methods come with many challenges risking soil and groundwater ecosystems. Thus, sustainable and cost-friendly biomaterials are ideal for atrazine degradation.
- Most phytoremediation studies are done under controlled conditions. Few studies have been based on natural circumstances like field based, so to understand the effects of treatment, field-based mitigation studies should be done extensively.
- The combination effects of the two treatment technologies should be the idea of future research implementing sustainability and low capital. AOPs have shown potential for the degradation of Atrazine-contaminated waters and soils. However, many reports have not observed the complete deg-

radation of Atrazine to cyanuric acid. Enhancing the methods available for the complete degradation of Atrazine is necessary.

- Most toxicity studies are done on aquatic organisms. Studies on the toxicity of atrazine on terrestrial organisms are not vigorous, although it has been directly exposed to soil microbes/organisms first, so researchers can target more on studying the toxicological inhibitory aspects of atrazine and its degraded products on terrestrial organisms.

7 Conclusion

The long shelf life of atrazine in different environments has put an enormous threat to every life form it has been exposed to, but more concerning is the intermediate metabolites being produced over time in the environment. Further research should be made to assess the ecotoxicological risk of atrazine metabolites and the substituted compounds (TBA) and its metabolites, due to the accumulation of atrazine in different plants, animals, water bodies through different routes. In most European countries, including Spain, Italy, and Portugal, atrazine has been replaced by more calcitrant compounds like terbuthylazine(TBA), which have more natural retention capacity and induce more hazards to living organisms even at lower doses. This implies that the atrazine ban is not foreseeable in countries like China, India, the USA, and some EU countries. Fungal/microbial culture intervention have broad application prospects in soil as well as water. More understanding and knowledge of the interaction of adsorbents and microbial or fungal strains will facilitate more sorption and mineralization of atrazine. Some widely applied atrazine-contaminated soil/ aqueous treatment technologies were discussed, including bioremediation, phytoremediation, and AOPs. AOPs application time and degradation rate are comparatively faster, and potential associated risks need further assessment. Among the various strategies incorporated, some challenges still need to be focused on for pilot scale or in-situ application of AOPs.

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Data Availability The articles analyzed during the current study are available in the literature and listed in the references.

Declarations

Ethics Approval and Consent to Participate Not applicable.

Consent to Participate Not applicable.

Consent for Publication Not applicable.

Competing Interests The authors have no competing interests to declare that are relevant to content of this article.

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