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# Assessment of the cytotoxic and mutagenic potential of the Jialu River and adjacent groundwater using human-hamster

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## <sup>3</sup> hybrid cells

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### ABSTRACT

The Jialu River in China has been seriously polluted by the direct discharge of industrial and 21 domestic wastewater. The predominant contaminants of the Jialu River and its adjacent 22 groundwater were recently investigated. However, the potential genotoxic impact of 23 polluted water on human health remains to be clarified. Here, we used human-hamster 24 hybrid (AL) cells, which are sensitive for detecting environmental mutagens. We found 25 that the cytotoxicity and mutagenicity of the groundwater in the Jialu River basin were 26 influenced by the infiltration of the Jialu River. Hydrological periods significantly affected 27 the cytotoxicity, but not the mutagenic potential, of surface and groundwater. Further, the 28 mutagenic potential of groundwater samples located < 1 km from the Jialu River (S<sub>M-2</sub> water 29 samples) was detected earlier than that of groundwater samples located approximately 30 20 km from the Jialu River (S $_{
m N}$  water samples). Because of high cytotoxicity, the mutagenic  $\,$  31 potential of water samples from the Jialu River ( $S_{M-1}$  water samples) was not significantly 32 enhanced compared with that of untreated controls. To further assess the mutagenic 33 dispersion potential, an artificial neural network model was adopted. The results showed 34 that the highest mutagenic potential of groundwater was observed approximately 10 km 35 from the Jialu River. Although further investigation of mutagenic spatial dispersion is 36 required, our data are significant for advancing our understanding of the origin, dispersion, 37 and biological effects of water samples from polluted areas. 38

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### 53 Introduction

54 Accidental pollution of surface water occurs frequently 55 worldwide, and rivers act as the most important medium for the transportation of environmental pollutants and transfor- 56 mation (White and Rasmussen, 1998; Ohe et al., 2003, 2004). 57 The United States Environmental Protection Agency (USEPA)'s 58 Toxic Release Inventory reported that approximately 0.1 billion 59

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kilograms of chemical pollutants were directly released 60 into surface water in 2014 (USEPA, 2016). Moreover, airborne 61 62 emission and surface runoff are largely responsible for the transport of chemical pollutants to aquatic systems 63 (Motelay-Massei et al., 2006). In contrast, chemical pollutants 64 in surface water are readily soluble and move slowly through 65 66 layers of soil, sand, and rocks and may be subject to transport into other media such as groundwater (Mehler et al., 2010; Chen 67 68 et al., 2012; Ma et al., 2013). The latter is important for 69 agricultural, recreational, and domestic activities, particularly 70 as a source of drinking water in most parts of the world (Wang 71 et al., 2011). Numerous studies assessed the potential risk of the 72 adverse effects of polluted surface and groundwater on human health (Fries and Puttmann, 2004; Nakada et al., 2008; Ma et al., 73 74 2012b). In addition, disinfection by-products (DBPs), formed by 75 the reaction of disinfectants with natural organic matter in surface and groundwater, were also considered to be cytotoxic 76 and genotoxic (Richardson and Postigo, 2015; Wagner and 77 78 Plewa, 2017). Evidence indicates that the pollution of drinking water is closely related to the mortality rate of patients with 79 80 esophageal (Zhang et al., 2003), liver (Li et al., 1994), and 81 urothelial cancers (Markovic, 1993). Moreover, the risk to health of polluted drinking water is not solely the sum of the effects of 82 83 individual compounds. Therefore, there is growing concern for 84 the additive, synergistic, and antagonistic effects of such 85 complex mixtures of pollutants (Wilkinson et al., 2000), partic-86 ularly the genotoxicity of organic residues in polluted 87 groundwater.

The Jialu River, which is located in Fugou County, Zhoukou 88 89 City, Henan Province, is 256 km long, and its basin is 5896 km<sup>2</sup>. 90 The Jialu River is an important tributary of the Huaihe River, which is seriously polluted by the direct discharge of contam-91 92 inants associated with economic growth and urbanization, 93 such as industrial and domestic wastewater (Ma et al., 2012a), industrial wastes, and untreated or lightly treated sewage 94 95 (Zhang et al., 2009). Large numbers of treated and untreated sewage from alongshore cities and villages were estimated to 96 be  $25,124 \times 10^4$  tons per year from 1996 to 1999, 81% of which 97 98 were discharged from Zhengzhou city (Xiao et al., 1999). As one of the six most important industrial cities by "the development 99 of central zones" stratagem of the Chinese Government, 100 101 Zhengzhou City has a long history of textile and metallurgy 102 industries and is therefore regarded to exert a strong impact 103 on surface water quality of the Jialu River basin. In rural areas, the main source of drinking water for many residents living 104 105 along the river is shallow, untreated groundwater. In this area, the incidence of cancers of the digestive system, such 106 as carcinoma of the esophagus, stomach, and liver during 107 2004–2006, was  $6.1 \times 10^{-4}$ , which is 1.71-fold higher than that 108 during 1973–1975 (Yang, 2010). Epidemiological studies of 109 110 verbal accounts of autopsies have shown that mortality rates 111 of patients with digestive cancers are three to four times 112 higher along the Huai River basin compared with those in the control areas (Wan et al., 2011). Nitrosamines and secondary 113 114 amines, which are mutagenic and carcinogenic (Bogovski and Bogovski, 1981; Eichholzer and Gutzwiller, 1998), are 115 116 the predominant toxic compounds in the Jialu River and its adjacent groundwater (Ma et al., 2012a). Comparative 117 genotoxicity of nitrosamine in drinking water DBPs showed 118 119 that genotoxic potencies of five nitrosamine DBPs in Salmonella

typhimurium and CHO cells showed identical descending rank Q8 order for genotoxicity and were highly correlated (Wagner 121 et al., 2012). In addition, Wagner et al. (2014) also found that the 122 genotoxicity of analogous N-nitrosamines and N-nitramines 123 relevant to  $CO_2$  capture present a potential risk for contami- 124 nating airsheds and drinking water supplies. Flumequine 125 and nitroarenes are the main contributors to the genotoxicity 126 of adjacent groundwater around the Jialu River because of 127 their high soil permeability and lateral seepage (Ma et al., 128 2012b). We reasoned therefore that assessing the overall 129 effects of these compounds will be facilitated by investigating 130 the cytotoxicity and genotoxicity of the river water and its 131 adjacent groundwater.

Numerous in vivo tests on the genotoxicity of water samples 133 were conducted, particularly using indigenous aquatic organ- 134 isms such as fish (Bahari et al., 1994; Alsabti and Metcalfe, 1995; 135 Minissi et al., 1996), sea urchins (Hose et al., 1983), mussels 136 (Ericson et al., 2002; Klobucar et al., 2003), oysters (Burgeot et al., 137 1995), newts (Jaylet et al., 1986; Fernandez et al., 1993), and 138 marine worms (Jha et al., 1995). Nevertheless, because of the 139 insensitivity of aquatic organisms to genotoxic compounds 140 and complexities involved in manipulating these organisms 141 in the laboratory, the application of in vitro genotoxic tests of 142 water samples is more helpful for investigating the presence 143 and distribution of genotoxins (Ohe et al., 1993; Eckl, 1995; 144 Schnurstein and Braunbeck, 2001). For example, the Ames 145 test was used to demonstrate a dose-dependent increase of 146 five orders of magnitude in the number of TA98 revertants 147 associated with industrial effluent extracts, as well as a 148 lower but significant increase of three orders of magnitude 149 associated with river water extracts 6 km downstream (Cerna 150 et al., 1996). Blue rayon extracts collected downstream of 151 wastewater treatment plants from the Katsura, Nishitakase, 152 and Kamo rivers induce a higher frequency of sister chromatid 153 exchange in cultured Chinese hamster lung cells than those 154 collected upstream, with and without metabolic activation 155 (Ohe et al., 1993). 156

Although numerous in vitro studies demonstrate the 157 genotoxicity of polluted waters, mutations that play impor- 158 tant roles in tumor development are not well documented. 159 Human-hamster hybrid (AL) cells stably express a single 160 copy of the human chromosome 11, which encodes the CD59 161 cell surface antigen, rendering AL cells sensitive to killing 162 by specific monoclonal antibodies in the presence of rabbit 163 serum complement (Hei et al., 1998). These hybrids were used 164 to detect the mutagenic effects of ionizing radiation (Wu et al., 165 1999; Hong et al., 2010) and chemicals (Bao et al., 2009; Zhao 166 et al., 2011). For example, a 50-fold increase in mutations 167 at the CD59 locus occurs compared with the HPRT locus in 168 crocidolite-treated  $A_L$  cells (Hei et al., 1992). Other studies 169 using this system demonstrated that arsenic, which was 170 considered a serious contaminant of drinking water and 171 a nongenotoxic carcinogen for decades, is actually a strong 172 mutagen (Jacobson-Kram and Montalbano, 1985; Lee et al., 173 1985; Hei et al., 1998; Kessel et al., 2002). Similarly, the 174 mutagenicity of cadmium (Filipic and Hei, 2004) and asbestos 175 fibers (Xu et al., 1999, 2002) in contaminated environmental 176 compartments (sediment, water, and air) was demonstrated 177 using the  $A_L$  mutagen detection system. Meanwhile, using 178 the Ames test, raw water and drinking water samples from 179

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