

1 **A comprehensive assessment of environmental emissions, fate and**
2 **risks of veterinary antibiotics in China: an environmental fate**
3 **modelling approach**

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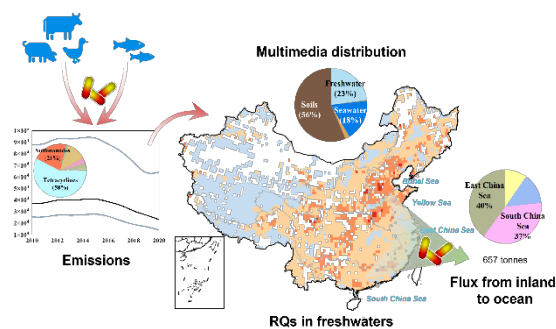
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17 **ABSTRACT**

18 China is one of the major global consumers of veterinary antibiotics. Insufficient
19 recognition of emissions and environmental contamination hampers global efforts to
20 prevent antibiotic resistance development. This pioneering study combined empirical
21 data and modelling approaches to predict total 2010–2020 emissions of 80 veterinary
22 antibiotics ranging from 23,110 to 40,850 tonnes/year, after 36–50% antibiotic removal
23 by manure treatment. Following an initial increase of 10% from 2010 to 2015,
24 emissions declined thereafter by 43%. While 85% of emissions discharged into soils,
25 approximately 56%, 23% and 18% of environmental residue were ultimately
26 distributed in soils, freshwaters and seawaters under steady state conditions. In 2020,
27 657 (319–1470) tonnes entered ocean from inland freshwaters. Median Σ antibiotics
28 concentrations were estimated at 4.7×10^3 ng/L in freshwaters and 2.9 ng/g in soils, with
29 tetracyclines and sulfonamides the predominant components. We identified 44
30 veterinary antibiotics potentially posing high risks of resistance development in
31 freshwaters, with seven exhibiting high risks in >10% of Chinese freshwater areas.
32 Tetracyclines were the category with the most antibiotics exhibiting elevated risks,
33 however sulfamethylthiazole demonstrated the highest individual compound risk. The
34 Haihe River Basin displayed the highest susceptibility overall. The findings offer
35 valuable support for control of veterinary antibiotic contamination in China.



36

37 **Keywords:** veterinary antibiotics, antibiotic emission, antibiotic multimedia
38 distribution, antibiotic discharge to ocean, risk of resistance development

39 **Synopsis:** Veterinary antibiotic emission and contamination probably decreased after
40 2015, with 44 antibiotics exhibiting potential high risks of resistance development in

41 Chinese freshwaters in 2020. Soil is the major sink of most antibiotics.

42 **INTRODUCTION**

43 Antibiotic resistance is now one of the major threats to human health worldwide. It
44 is estimated to have caused 4.95 million deaths globally in 2019.^{1,2} Demand for animal
45 protein drives meat production, which consumes ~75% of antibiotics sold globally for
46 growth promotion and disease prevention of livestock.^{3,4} The subsequent environmental
47 releases and contamination of antibiotics could prompt the development and spread of
48 antibiotic resistance.^{5,6} This increases the likelihood of human infections by pathogens
49 containing resistant genes and compromises the effectiveness of human medicines.^{3,5-7}
50 China is one of the top consumers of veterinary antibiotics, and is one of the largest
51 hotspots of resistance.^{3,4,8,9} Rising incomes in China have resulted in an unparalleled
52 expansion in the demand for animal protein, which may elevate veterinary antibiotic
53 use.¹⁰ Meanwhile, various regulations have been issued in China since 2016 to reduce
54 the use of antibiotics in food animals and to control environmental contamination of
55 antibiotics as high-priority novel contaminants.¹¹⁻¹⁵ It is therefore essential to know the
56 status of emission and contamination of veterinary antibiotics in different media across
57 China, to combat development of resistance both in China and worldwide.

58 Recent studies have reported widely different emissions sourced from food animals
59 in China. One estimated that 45,360 tonnes of 36 widely used antibiotics were emitted
60 by food animals into the Chinese environment in 2013; in contrast, a much lower release
61 from food animals was recently reported - only 4,131 tonnes in China in 2014.^{16,17}
62 Existing studies on antibiotic emission, as well as environmental transport and
63 distribution in China have limitations, such as: on spatial resolution and substance
64 coverage, or on considerations of antibiotic removal during livestock excretion
65 treatment, inter-media chemical transport processes and ionizable features of antibiotics
66 in the modelling approach.¹⁷⁻²² Global studies including China have mainly investigated
67 the use of veterinary antibiotics, but rarely assessed environmental release and
68 contamination.^{3,23,24} Furthermore, previous estimates normally assumed identical
69 application of antibiotics to individual animals within the same livestock species in a
70 country. However, notable geographical disparities were found in residues of some
71 veterinary antibiotics in manures, implying potential differences in regional antibiotic

72 application.²⁵ Such limitations have introduced large uncertainties in our knowledge of
73 contamination and fate of antibiotics sourced from food animals in China, which limits
74 our understanding of antibiotic environmental exposure.

75 This study - for the first time - comprehensively assesses environmental emissions,
76 spatial and multimedia concentrations, and the risks of a relatively complete range of
77 veterinary antibiotics used in livestock farming and aquaculture in China. Temporal
78 patterns of antibiotic emissions and concentrations are analyzed for 2010–2020, during
79 which veterinary antibiotic use policies have been considered. We conducted a literature
80 survey of measurements in livestock faeces and aquaculture water to compile the
81 emission inventories of veterinary antibiotics, taking account of spatially varied
82 application preferences. A previously well-constructed and validated national scale
83 multimedia model - the SESAMe v3.4 model (with a spatial resolution at 0.5°) - was
84 utilized to predict multimedia concentrations of antibiotics across China, with a full
85 consideration of ionizable features of substances (e.g. triclosan and triclocarban with
86 different pKa values, and thus showing distinct degrees of ionization and behaviors in
87 the environment) and relatively complete inter-media transport processes, compared
88 with many other multimedia environmental fate models.¹⁹ The study identifies: i. the
89 antibiotics with higher risks of resistance development; ii. highly-contaminated regions
90 and media requiring priorities in risk control, and iii. the predominant emission
91 pathways and sinks of antibiotics, with discharges to ocean being estimated. This wide-
92 ranging assessment can support contamination prevention and control of veterinary
93 antibiotics in China. The same framework is applicable worldwide, especially for
94 countries with serious antibiotic emission and contamination issues.

95 **MATERIALS AND METHODS**

96 **Data search and selection strategy for collecting antibiotic residual measurements**

97 To acquire a relatively complete list of veterinary antibiotics being widely used in
98 China, a literature survey was conducted to review measurements of antibiotics in
99 livestock faeces (including swine, cattle, and poultry) and aquaculture water. Papers
100 related to livestock farming and aquaculture were searched from Web of Science and
101 the Chinese Academic Journal Network Database by December 1, 2023. Key search

102 terms including ((livestock OR pig OR swine OR cattle OR cow OR yak OR sheep OR
103 goat OR chicken OR poultry OR feedlots OR farms) AND (antibiotic* OR veterinary
104 antibiotic*)) AND (faeces OR urine OR manure OR animal wastes OR animal dung OR
105 residues) AND (China)) and ((antibiotic* OR antimicrobial OR veterinary) and
106 (aquaculture farm OR mariculture OR cultured freshwater OR aquafarm OR fishpond)
107 and (China OR Chinese)) were searched via Web of Science to find English articles for
108 livestock and aquaculture, respectively. ((farms OR livestock manure OR "pig manure"
109 OR "cattle manure" OR "poultry manure" OR "sheep manure") AND (antibiotic* OR
110 veterinary antibiotic*)) AND ("antibiotic residue")) and ((antibiotic) AND (fishery OR
111 aquaculture OR marine aquaculture OR freshwater aquaculture OR pond)) were
112 searched via the Chinese Academic Journal Network Database to find Chinese articles
113 for livestock and aquaculture, respectively.

114 A total of 3773 publications were found (see [Figure S1](#) in the Supporting Information
115 (SI)). A total of 780 review articles and non-peer-reviewed publications including thesis
116 and conference papers were then excluded. Another 2888 articles with irrelevant
117 contents were excluded after examining the title and abstract of remaining articles. A
118 careful full-text reading was then conducted to exclude 18 studies without specific
119 location information of sampling sites and available concentrations for individual
120 antibiotics. Then 55 (38 English articles and 17 Chinese articles) and 32 (18 English
121 articles and 14 Chinese articles) relevant papers for measured antibiotics in livestock
122 faeces and aquaculture water, respectively, were finally included in this study.
123 Antibiotics widely used in food animals were targeted and their measurements were
124 collected from these papers for estimating emissions. Relevant information is presented
125 in [Table S1–S2](#) in SI and [Appendix A](#).

126 **Emissions**

127 Emissions from both livestock farming and aquaculture were estimated during 2010–
128 2020. Livestock farms were categorized as either intensive farms or family farms in
129 this study. Family farms are operated on a small to moderate scale, and use land
130 contracted by a family for production activities, with family members as primary labour
131 forces.²⁶ Intensive farms are industrial agriculture practices. Three livestock types were

132 considered, namely swine, cattle, and poultry. Veterinary antibiotics can be released to
 133 soils and freshwaters with livestock excrement after being (partially) treated.²⁷ All
 134 excreta of poultry, consisting of a mixture of faeces and urine, was assumed to be
 135 released into soils.²⁸ Regarding swine and cattle, since family farms in China typically
 136 involve both crop cultivation and animal husbandry, all excretion from livestock will
 137 commonly be used as organic fertilizer through land application with or without
 138 treatment.²⁹ Therefore, excreted antibiotics in this sector were assumed to be all
 139 released to soils. Untreated urine in intensive farms was assumed to be directly
 140 discharged into freshwaters, while the treated urine and all treated and untreated faeces
 141 were assumed to be release to soils.³⁰ The calculation of the emission for individual
 142 antibiotics from livestock farming is shown as equations (Eqs.) 1-4.

$$143 \quad Ex_i = \sum_i [Exu_i + Exf_i] = \sum_i [(F_i \times C_i \div f_i \times u_i) + (F_i \times C_i)] \quad (1)$$

$$144 \quad F_i = \sum_i N_i \times p_i \quad (2)$$

$$145 \quad Ems_i = \sum_{i,j,x} [r_{a,i} \times (Exf_i \times P_{a,i,j} \times R_j + Exu_i \times P_{a,i,x} \times R_x) + r_{b,i} \times Ex_i \times P_{b,i,j} \times$$

$$146 \quad R_j] \quad (3)$$

$$147 \quad Emw_i = \sum_i r_{a,i} \times Exu_i \times P_{a,i,l} \times R_l \quad (4)$$

148 Where i represents the type of livestock; Ex_i , Exu_i and Exf_i represent the
 149 antibiotics excreted with all livestock excrement, and those excreted with urine and
 150 faeces, respectively, for livestock i (tonnes/year). F_i and C_i refer to the annual amount
 151 of faeces produced by livestock i (kg/year) and residual concentrations of antibiotics
 152 therein ($\mu\text{g}/\text{kg}$). f_i and u_i represent excretion rates of antibiotics (%) through faeces and
 153 urine, respectively, from livestock i (Table S3). N_i stands for the number of livestock i .
 154 The p_i represents the faeces production volume per unit of livestock i per year
 155 (kg/unit/year) (Table S4), which was calculated by multiplying the faeces production
 156 (kg/unit/day) by growth period (days) of livestock i .³¹ Because limited literature has
 157 reported urinary concentrations for livestock, urinary excretion has been estimated from
 158 an adjustment by fecal excretion rates, i.e. (1) a backward calculation of antibiotic usage
 159 using fecal excretion rates and fecal excretion amount of antibiotics, and then (2)
 160 forward estimation of urinary excretion using usage and urinary excretion rates.

161 The emission through different pathways was calculated as Eqs. 3-4. Ems_i and

162 Emw_i represent the amount of antibiotics (tonnes/year) discharged into soils and
 163 freshwaters, respectively, by livestock i . Where j represents the faecal treatment
 164 process, including agricultural utilization, composting, and anaerobic digestion; and x
 165 represents the urinary treatment process: agricultural utilization and anaerobic
 166 digestion. $r_{a,i}$ and $r_{b,i}$ represent the proportion (%) of intensive livestock farms (a) and
 167 family livestock farms (b) respectively, for livestock i , which are provincially and
 168 temporally varied. Their detailed calculation method is given in [section S1](#) in [SI](#). $P_{a,i,j}$
 169 and $P_{a,i,x}$ represent the proportion (%) of the faecal and urinary treatment process in
 170 intensive livestock farms for livestock i ; $P_{a,i,l}$ represents the proportion of direct
 171 discharges from intensive farms for livestock i (%). The three parameters vary
 172 regionally, with values given in [Appendix A](#).³⁰ $P_{b,i,j}$ represents the proportion of the
 173 faecal treatment process in family livestock farms for livestock i . The multi-year (2015-
 174 2019) average value was taken for $P_{b,i,j}$ at a provincial level ([Appendix A](#)).³²⁻³⁶ R_j and
 175 R_x represents the antibiotic residue rate (%) in the faecal and urinary treatment process,
 176 respectively ([Table S5](#)). R_l represents the antibiotic residue rate of the direct discharge
 177 ($R_l=100\%$).

178 Both freshwater aquaculture and coastal marine aquaculture were considered in this
 179 study. According to different aquaculture practices, freshwater aquaculture comprises
 180 pen culture, cage culture, and industrialized culture, while marine aquaculture includes
 181 pond culture, cage culture, deep-sea cage culture, raft culture, hanging culture,
 182 industrialized culture, and bottom-sowing culture.³⁷ Bottom-sowing culture allows
 183 organisms to grow naturally without using antibiotics (GB/T 20014.18-2013), so it was
 184 not considered in this study, while all other practices were considered. All antibiotic use
 185 in aquaculture is released into freshwaters or seawaters directly without treatment.^{38,39}

186 The emission was calculated as Eqs. 5-7.¹⁶

$$187 \quad Emw_{a,f} = \sum_{f=1}^3 C_f \times A_f \times D_f \times T_f \quad (5)$$

$$188 \quad Emw_{a,m} = \sum_{m=1}^6 C_m \times A_m \times D_m \times T_m \quad (6)$$

$$189 \quad Emw_{aqu} = Emw_{a,f} + Emw_{a,m} \quad (7)$$

190 $Emw_{a,f}$ and $Emw_{a,m}$ represent the amount of antibiotics (tonnes/year) discharged

191 into freshwaters and seawaters, respectively, in aquaculture. C_f and C_m represent the
192 concentrations of antibiotics (ng/L) in freshwater aquaculture and marine aquaculture,
193 respectively (Appendix A). A_f and A_m represent the areas of freshwater and marine
194 aquaculture, respectively (m^2) (Appendix A). D_f and D_m represent the water depth in
195 freshwater and marine aquaculture area (m) (Appendix A). T_f and T_m represent the
196 water exchange times between aquaculture area and surrounding waters in freshwater
197 and marine aquaculture per year. Instantaneous homogenization was assumed for each
198 water exchange event. Ten times water exchange was assumed for pond culture per
199 year, whilst everyday water exchange (365 times/year) was assumed for other
200 aquaculture practice types.⁴⁰

201 Livestock production data of swine, cattle, and poultry (N_i) on a county level, as well
202 as freshwater and marine aquaculture area on a provincial level from 2010 to 2020 were
203 collected from the China Statistical Yearbook or local statistical bureaus.⁴¹⁻⁴³ For
204 counties lacking livestock data in the literature, information was collected by contacting
205 individual local survey teams of the National Bureau of Statistics. Based on empirical
206 data collected in the last section, an average antibiotic concentration was taken for
207 samples collected within the same province to acquire the provincial-level residue in
208 faeces and aquaculture water. For provinces lacking measurements, a national average
209 was taken. The data on livestock production, aquaculture area and measured antibiotic
210 residue in faeces and aquaculture water on a provincial level is given in Appendix A.

211 To thoroughly consider the effect of use policy on veterinary antibiotics in China and
212 uncertainties of adopted values for parameters, a best-guess (BG) emission scenario
213 and two extreme emission scenarios, i.e. low-emission (LE) and high-emission (HE)
214 scenarios, were constructed to demonstrate the most appropriate and reliable total
215 emissions with a potential range in China. The BG scenario generally followed the
216 prohibition of veterinary antibiotic use in China, unless any banned veterinary
217 antibiotics were still detected in livestock faeces or aquaculture water after the
218 prohibition.^{44,45} Meanwhile, average values of the faeces production volume per unit of
219 livestock i per year (p_i), water depth in aquaculture (D_f and D_m) collected from the
220 literature, and national or industrial standards were taken in the BG scenario. Within

221 contrast to the BG scenario, the minimum values were used for p_i , D_f and D_m in the
222 LE scenario. In the HE scenario, it was assumed that all identified veterinary antibiotics
223 were used throughout the 11 years (2010–2020) without prohibition. The maximum
224 values were used for p_i , D_f and D_m , and all excrement was assumed to be directly
225 discharged without treatment (R_j and $R_x=100\%$). The investigation was mainly under
226 the BG scenario.

227 **Environmental fate modelling**

228 The SESAMe v3.4 (Sino Evaluative Simplebox-MAMI model) model was utilized
229 to predict antibiotic concentrations in environments.⁴⁶ The model is able to predict
230 concentrations in multiple environmental compartments (i.e., air, different types of soils,
231 freshwaters and sediments, seawaters and sediments, and vegetations) in the Chinese
232 mainland with a high resolution of 0.5°. It has been applied and well validated on
233 chemicals with varying characteristics, such as semi-volatile organic compounds and
234 ionizable chemicals (e.g. active ingredients of personal care products).^{19,46,47} As
235 indicated in the previous study, due to introduction of the activity approach, the model
236 has advantages in predicting multimedia concentrations of ionizable chemicals, such as
237 antibiotics, as it considers varying partitioning and transport behaviors of ionic and
238 neutral forms of organic chemicals and has incorporated spatially varying pH in
239 freshwaters, sediments and soils, such that the model has the potential to consider a
240 spatially varied degree of ionization and the consequent varying behavior of a specific
241 substance in different media and areas.¹⁹ The model principle and basic physical
242 processes are introduced in [section S2](#) in [SI](#). The elimination half-life of various
243 targeted antibiotics in the Chinese environment was calculated using the model. The
244 results indicated that different antibiotics reached the steady-state concentrations (after
245 ~5 elimination half-lives) within a maximum of 96 days, which is less than one year.⁴⁸
246 This justifies the utilization of the steady-state model for investigating the annual
247 pattern of concentrations.

248 Estimated emissions and physicochemical properties (molecular weight (g/mol),
249 vapor pressure at 25°C (Pa), water solubility at 25°C (mg/L), log-transformed octanol-
250 water partition coefficients ($\log K_{ow}$) (-), pKa acid and pKa basic (-), and half-life (hr)

251 in the air, water, soil and sediment) of the identified target veterinary antibiotics (SI
252 Table S1) were input to the model for predictions. A complete literature review was
253 conducted to collect measurements of animal-use-only (AUO) antibiotics in
254 freshwaters and soils across China for model validation. Information on locations of
255 sampling sites and sampling years collected from 74 peer-reviewed papers are given in
256 Table S6. It was found that only nine AUO antibiotics have sufficient measured data for
257 a comparison with predictions, and these were therefore used for model validation.
258 Monte Carlo simulation was performed to conduct the uncertainty analysis by running
259 the model 10,000 times. Parameter values were randomly taken from the environmental
260 parameter databases of the model and emission databases developed in this study for
261 the simulation.

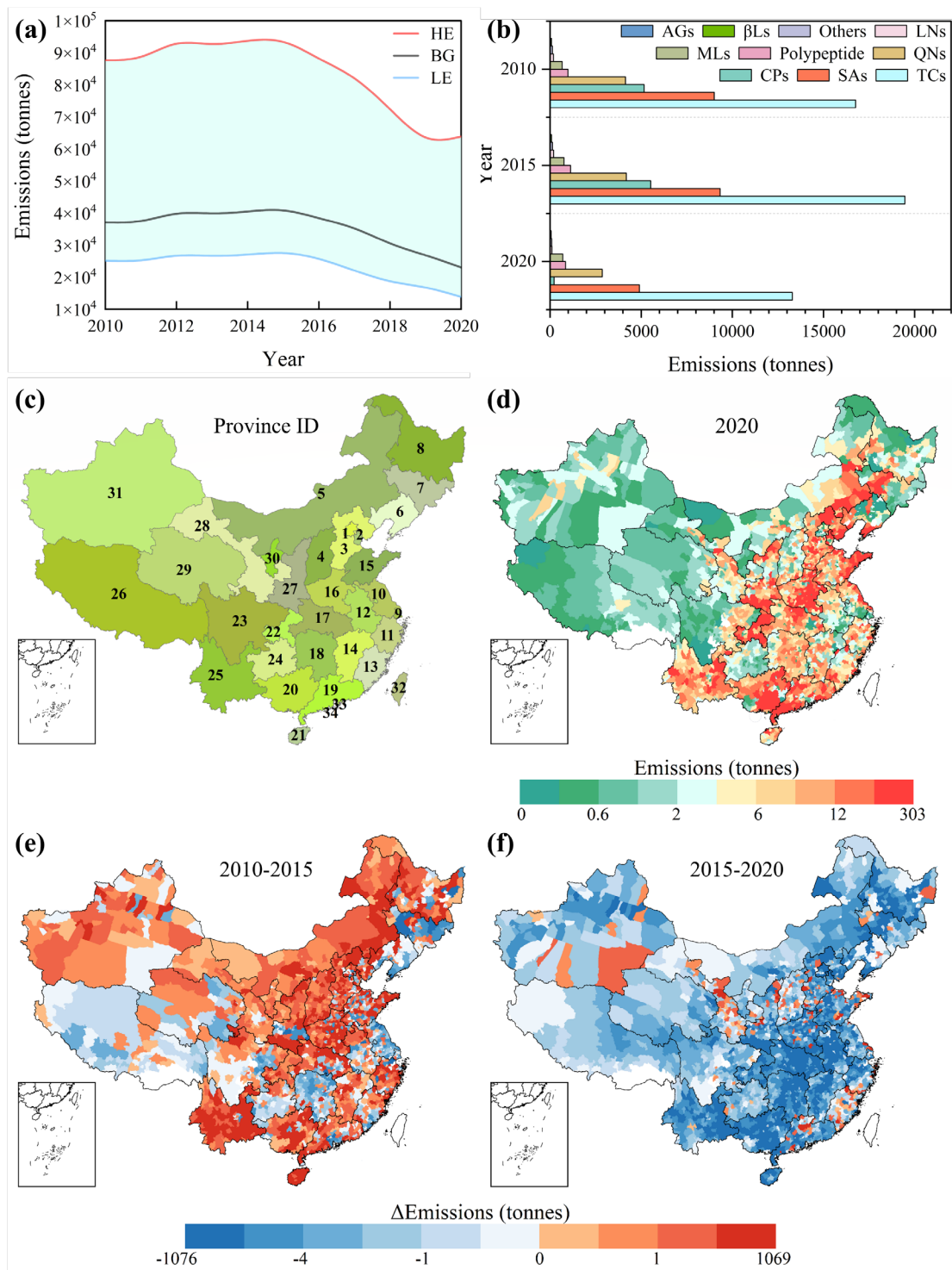
262 Environmental risk assessment

263 As the risk of antibiotic resistance selection is currently one of the major global
264 concerns on public health, this study mainly assessed this type of risk induced by
265 antibiotic contamination in freshwaters. However, to implement more comprehensive
266 assessment and offer a broader perspective, risk for aquatic toxicity has been
267 additionally evaluated. Values of predicted no effect concentrations for resistance
268 development ($PNEC_{res}$) and aquatic toxicity ($PNEC_{tox}$) were collected from the
269 literature (Table S7). The $PNEC_{res}$ value serves as a guide to the maximum residue level
270 of antibiotics in the environment, below which resistance is less likely to develop.⁴⁹
271 They were derived from experiments on resistance to clinically relevant bacteria. The
272 values of $PNEC_{res}$ for most antibiotics were collected from studies by Bengtsson-Palme
273 et al. and Zhang et al.,^{50,51} and 100 ng/L from a study by Le Page et al. was taken for
274 the remaining antibiotics without a chemical-specific value, for a conservative
275 assessment.⁵² In addition, aquatic toxicity data (LC_{50} or EC_{50}) were available for 57
276 antibiotics, mostly obtained from toxicity assays on algae (Table S7). $PNEC_{tox}$ was
277 calculated as LC_{50} or EC_{50} / AF (Assessment factor = 1000).⁵³ The risk quotient (RQ),
278 obtained by dividing predicted environmental concentrations (PECs) by PNECs (i.e.,
279 $RQ = PECs / PNECs$) was employed to predict the environmental risks in freshwaters
280 for individual chemicals across China. A nominal RQ classification of $RQ < 0.1$, $0.1 <$

281 RQ < 1 and RQ > 1 predicts low (or insignificant), medium and high risks in the
 282 environment, respectively.^{54,55}

283 **RESULTS AND DISCUSSION**

284 **Emissions**



285
 286 **Figure 1.** (a) Temporal trends of total antibiotic emissions in low-emission (LE), best-

287 guess (BG), and high-emission (HE) scenarios from 2010 to 2020; (b) emissions of
288 various antibiotic categories in 2010, 2015, and 2020; (c) Province IDs of China (1.
289 Beijing; 2. Tianjin; 3. Hebei; 4. Shanxi; 5. Inner Mongolia; 6. Liaoning; 7. Jilin; 8.
290 Heilongjiang; 9. Shanghai; 10. Jiangsu; 11. Zhejiang; 12. Anhui; 13. Fujian; 14. Jiangxi;
291 15. Shandong; 16. Henan; 17. Hubei; 18. Hunan; 19. Guangdong; 20. Guangxi; 21.
292 Hainan; 22. Chongqing; 23. Sichuan; 24. Guizhou; 25. Yunnan; 26. Tibet; 27. Shaanxi;
293 28. Gansu; 29. Qinghai; 30. Ningxia; 31. Xinjiang; 32. Taiwan; 33. Hong Kong; 34.
294 Macau); (d) spatial distribution of total antibiotic emissions in 2020; (e-f) difference in
295 emissions at the county level between 2010 and 2015, as well as between 2015 and
296 2020.

297 A total of 84 veterinary antibiotics were measured in livestock excrement and
298 aquaculture water based on the comprehensive literature review, of which 80 were
299 detected above the limits of detection (LOD) and investigated in this study. A total of
300 20 are AUO antibiotics (7 quinolones, 5 macrolides, 4 sulfonamides, florfenicol,
301 ceftiofur, monensin, and carbadox ([Table S1](#))), and the other 60 are also used on humans.
302 The 80 antibiotics mainly belong to 9 classes: (1) 23 sulfonamides (SAs); (2) 6
303 tetracyclines (TCs); (3) 17 quinolones (QNs); (4) 12 macrolides (MLs); (5) 5 β -lactams
304 (β Ls); (6) 2 lincomycins (LNs); (7) 3 chloramphenicols (CPs); (8) 2 Aminoglycosides
305 (AGs); (9) 3 Polypeptide, and other 7 antibiotics (carbadox, furazolidone, monensin,
306 nalidixic acid, novobiocin, rifampicin, and salinomycin). Information of the antibiotics
307 is summarized in [Table S1](#).

308 The annual emission of Σ antibiotics (the sum of the 80 antibiotics) used in livestock
309 and aquaculture under the BG scenario ranged from 23,110 to 40,850 tonnes during
310 2010–2020, peaking in 2015 ([Figure 1a](#)). Annual emissions ranged from 13,870 to
311 27,560 tonnes in the LE scenario, whereas it reached 63,620 to 93,780 tonnes in the HE
312 scenario. According to the HE scenario, there would be a rebound in emissions after
313 2019, if there was not any restriction on use, mainly due to a 30% increase in the
314 quantity of livestock. The LE and HE scenarios only provide insight to potential
315 emission ranges, considering uncertainties in parameters. The following results and
316 discussion are based on the BG scenario. The emission increased by 10% from 2010 to

317 2015 and decreased by 43% from 2015 to 2020. The total emission was around 23,110
318 tonnes in 2020. The increase in the numbers of swine and poultry resulted in the
319 increasing emissions between 2010 and 2015. The decrease in emissions after 2015 was
320 mainly driven by fluctuations in total livestock numbers, freshwater aquaculture area
321 and proportions of livestock farming types. The populations of swine, cattle and poultry
322 declined by 24%, 21% and 5% from 2015 to 2020, while the freshwater aquaculture
323 area reduced by 96%. The reduction in farming volumes contributed 84% of the
324 emission decline after 2015. Meanwhile, a 12% increase in the proportion of intensive
325 farms versus a corresponding decrease in the proportion of family farms contributed
326 12% of the emission decline after 2015. The more dramatic decline after 2018 was
327 probably attributed to (1) the outbreak of African swine fever (ASF) in China with a
328 mortality rate of nearly 100%, (2) the prohibition of veterinary use of chloramphenicol,
329 carbadox, and vancomycin in 2020, leading to 14% emission reduction compared to
330 2019, and (3) a 74% reduction in pen culture area from 2017 to 2018.^{45,56}

331 On average 47% of the excreted antibiotics by livestock were removed in manure
332 treatment processes in the BG scenario every year. Due to the temporal variation in
333 proportions of intensive farms and family farms, the percentage varied from 36 to 50%
334 during 2010–2020. Therefore, manure treatment has been effective at removing
335 antibiotics before being discharged to environments. Livestock farming accounted for
336 the majority of veterinary antibiotic emissions (80–88%), while aquaculture only
337 accounted for 12–20% of total emissions. Excretion from swine accounted for >60% of
338 the total livestock emission in the 11 years. This is because China is one of the world's
339 largest producers and consumers of pork, and farms more swine than cattle and
340 poultry.⁵⁶ Cattle and poultry contributed, on average, 22% and 14% to total livestock
341 emissions, respectively. Freshwater and marine aquaculture contributed 44% and 56%
342 of the total aquaculture emission, respectively.

343 Emissions projected under the BG scenario in this study are considered reliable, after
344 being compared with estimates from other studies. Our estimated livestock emission is
345 higher than the antibiotic consumption by livestock in China estimated by Van Boeckel
346 et al. for 2010 (29,720 versus 14,525 tonnes), but is generally comparable with the

347 emission derived from their newly estimated consumption for corresponding years
348 using the updated data, if excretion rates at 30–90% by livestock are applied.^{3,23,24}
349 Furthermore, our estimated emission (39,900 tonnes for the 80 antibiotics) is lower than
350 that estimated from food animals by Zhang et al., which was 45,360 tonnes from 36
351 antibiotics in 2013.¹⁷ This is because removal of antibiotics during manure treatment
352 processes before release to the environment was considered in this study, whereas
353 Zhang et al.'s study assumed that all antibiotics excreted by livestock were directly
354 released into the environment. The estimated excretion of 80 antibiotics in 2013 reached
355 92,720 tonnes in this study, which would be directly discharged if no removal before
356 release was assumed. The aquaculture emission estimated by our study is also
357 comparable with the antibiotic consumption in aquaculture in China estimated by Schar
358 et al. for 2017 (5,143 versus 5,940 tonnes) and Yang et al. for 2014 (3,226 versus 2052
359 tonnes).^{16,57}

360 The average annual emissions during 2010–2020 varied from 0.04 to 7,048
361 tonnes/year for individual antibiotics (Table S8). There were 31 veterinary antibiotics
362 with an annual average emission >100 tonnes/year. Tetracyclines, sulfonamides,
363 chloramphenicols, and quinolones were four predominant antibiotic categories
364 throughout the 11 years (except 2020 due to the ban of chloramphenicol), accounting
365 for 94% on average of the total emissions (Figure 1b). The proportion of individual
366 antibiotics in emissions remained generally steady over the 11 years, without
367 considering the prohibition of use. Although only six substances were in the
368 tetracyclines category, they were the predominant component constituting 58% (13,290
369 tonnes) of the total emission in 2020. Tetracyclines accounted for 66%, 50%, 67%, and
370 18% of the emissions from swine, cattle, poultry, and aquaculture, respectively, in 2020.
371 Oxytetracycline, tetracycline, and chlortetracycline were the top three tetracyclines,
372 with an emission of 5,539, 4,242 and 2,237 tonnes, respectively, in 2020. Tetracyclines
373 are extensively used as feed additives to promote animal growth, primarily due to their
374 low cost, wide availability, and broad-spectrum antibacterial activity.⁵⁸ The average
375 residual concentration of individual tetracyclines in swine faeces reached 9,890 µg/kg,
376 much higher than that for other categories (126–1,230 µg/kg) (SI Table S2). Before

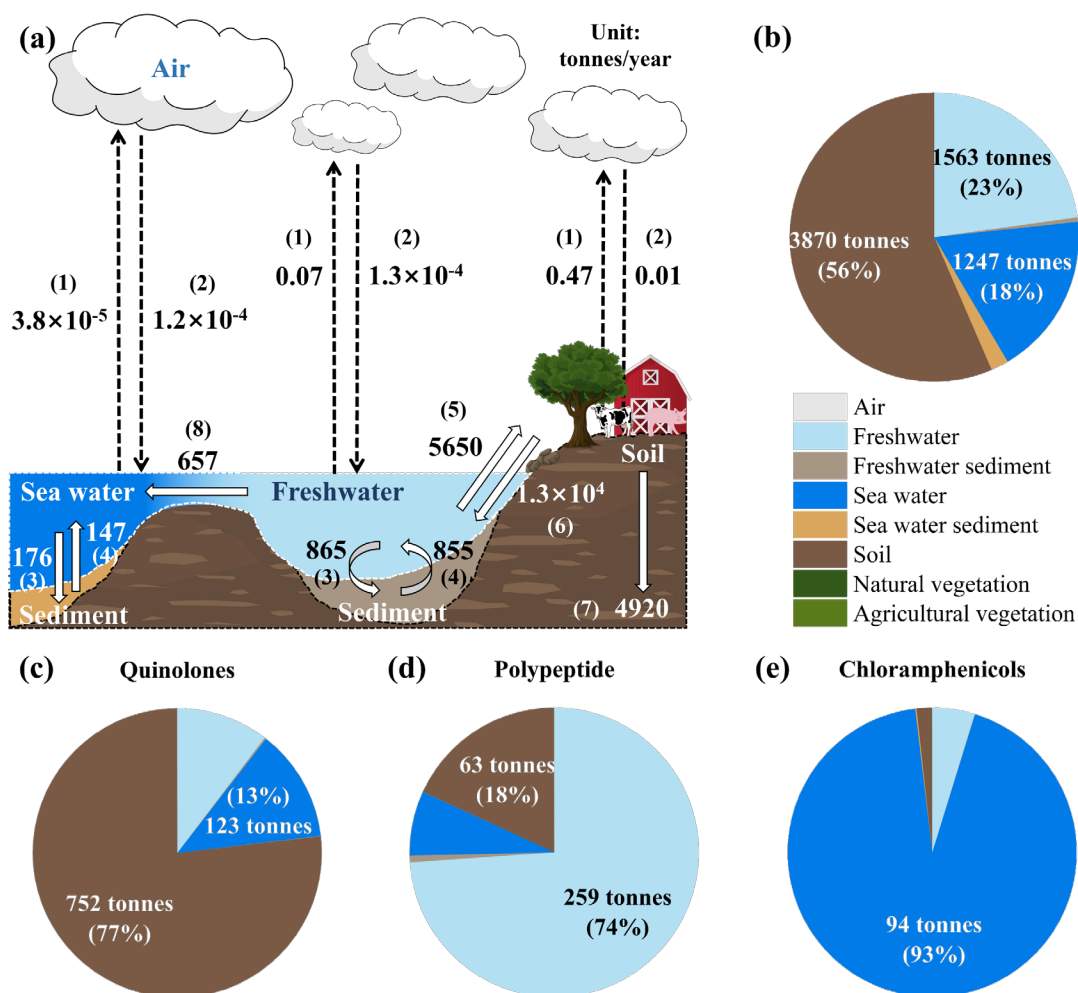
377 chloramphenicol was banned, chloramphenicols comprised the primary component
378 (53%) of emissions from cattle, although only constituting 14% of the total emission in
379 2019. Chloramphenicol exhibited the highest emission within this category at 3,762
380 tonnes in 2019. In contrast to livestock, sulfonamides were the main antibiotic
381 categories, which accounted for 63% and 29% of the emissions from freshwater and
382 marine aquaculture, respectively, in 2020. The AUO antibiotics contributed only 8.0%
383 of the total emission in 2020 (SI Table S8). Enrofloxacin is the single most emitted
384 AUO antibiotic, with an emission of 1,135 tonnes in 2020, 1.6 times higher than the
385 summed emission of all other AUO antibiotics.

386 The spatial pattern of emissions was mainly ascribed to distribution of livestock
387 farming volumes and aquaculture areas, while geographically varied residues in faeces
388 and aquaculture water (reflecting use variations), as well as proportions of two major
389 livestock farm types with distinct manure treatment rates are also influential factors.
390 Seven provinces, namely Hebei, Henan, Shandong, Guangxi, Guangdong, Yunnan, and
391 Shaanxi, exhibited elevated emissions compared to the other regions, in all contributing
392 49–61% of the total emissions during 2010–2020 (Figure 1d). In the livestock sector,
393 Henan province generally exhibited the highest antibiotic emissions at 2,410–3,490
394 tonnes/year due to the consistently highest swine and poultry farming volumes
395 throughout 2010–2020, contributing to 10–12% of the emission from livestock farming.
396 Sichuan was also among the provinces with a high swine farming volume, almost
397 comparable with Henan. However, it had substantially lower emissions (490–870
398 tonnes/year), probably due to the lower antibiotic use (lower residual concentrations in
399 faeces). In contrast to livestock farming, Anhui and Shandong showed the highest
400 antibiotic emissions at 48–2,180 tonnes/year and 954–1,282 tonnes/year, respectively,
401 from aquaculture during 2010–2020, which was driven by the high aquaculture area in
402 the two provinces.

403 As the majority of counties experienced extremely limited temporal changes (<10
404 tonnes) in emissions, with ca. 95% of counties from 2010 to 2015 and ca. 80% of
405 counties from 2015 to 2020, a relatively constant spatial pattern was found for
406 emissions during 2010–2020 (Figure S2). However, exceptionally large variations

407 occurred in certain regions (Figure 1e-f). For instance, in contrast to most regions,
 408 emissions in Gongzhuling city (Jilin province) dramatically increased by ca. 67 times
 409 (120 tonnes) from 2010 to 2015, meanwhile Tibet and Jilin provinces showed a
 410 significant decrease in emissions by 50% (Figure 1e). From 2015 to 2020, Susong
 411 county in Anhui province exhibited a quicker decline by 96% (188 tonnes), while
 412 counties in Hebei and Shaanxi provinces showed the largest increase at 30% in
 413 emissions across China (Figure 1f). In comparison to 2010, emissions of veterinary
 414 antibiotics were reduced in 82% of China's counties, but primarily increased in Shanxi,
 415 Gansu and Fujian provinces in 2020, with a summed increase of 143 tonnes.

416 **Antibiotic budget**



417
 418 **Figure 2.** (a) Antibiotic budget in the multimedia environment at the steady state in
 419 2020 (Unit, tonnes/year), and (b-e) multi-media distribution of all antibiotics,
 420 quinolones, polypeptide and chloramphenicols at the steady state in 2020. (The inter-

421 media processes in [Figure 2a](#) are explained as follows: (1) volatilization; (2) gas
422 absorption and deposition (dry deposition and wet deposition); (3) absorption and
423 sedimentation; (4) desorption and resuspension; (5) irrigation; (6) runoff and erosion;
424 (7) leaching; (8) discharges from inland freshwater to the ocean)

425 In 2020, we estimated that around 85% (19,650 tonnes) of the total emissions in
426 China were released to soils, with the remainder discharged into freshwaters and
427 seawaters. When reaching the steady state (i.e. when all mass transfer and degradation
428 fluxes became constant), there were only 56% (3,870 tonnes) of residual antibiotics in
429 the environment distributed in soils, and another 23% (1,563 tonnes) and 18% (1,247
430 tonnes) were in freshwaters and seawaters, respectively ([Figure 2b](#)). Therefore, on
431 average, soil was generally the sink of veterinary antibiotics at the national level in
432 China. Surface runoff and soil water erosion were the major processes transporting
433 1.3×10^4 tonnes veterinary antibiotics from soils to freshwaters per annum ([Figure 2a](#)).
434 This far surpassed the direct source emission to freshwaters, and became the major
435 route of antibiotics into freshwater systems. Irrigation transported 5,650 tonnes/year
436 veterinary antibiotics back to soils from freshwaters. Meanwhile, soil leaching
437 transported 4,920 tonnes/year of antibiotics into deeper soil layers, posing a potential
438 risk of groundwater contamination. Approximately 1.1×10^4 tonnes/year of veterinary
439 antibiotics were degraded, with freshwater and soil experiencing the most significant
440 degradation at 5,290 and 3,850 tonnes/year, respectively. The exchange between
441 freshwaters and sediments was both around 850 tonnes/year, with slightly more
442 transport from freshwaters to sediments. The antibiotic exchange between air and land
443 (soils and freshwaters) was minimal compared to the fluxes at the interface between the
444 other compartments, ranging 10^{-5} – 10^{-1} tonnes/year. Antibiotics are hardly volatilized to
445 air. Furazolidone and flumequine were the most volatile antibiotics, contributing almost
446 the 79% of the annual flux from land to air.

447 However, different antibiotics demonstrated different multimedia distribution
448 patterns, due to their distinct physico-chemical properties ([Figure 2](#) and [Table S9](#)). For
449 instance, quinolones and lincomycins were mainly distributed in soils (77–92%), while
450 polypeptide (74%) was primarily in freshwaters, aminoglycosides, chloramphenicols,

451 nalidixic acid, novobiocin, and monensin were mainly distributed in seawaters (93–
452 99%) (Figure 2c-e and S3). Macrolides have a higher percentage in freshwater
453 sediments (0–13%) than the other categories (0–1.0%). Macrolides have lower water
454 solubility (7.8×10^{-3} – 107 mg/L) and higher $\log K_{ow}$ (0.6–4.75) than other antibiotic
455 categories, so are more likely to remain in soils after being released. In contrast,
456 aminoglycosides and chloramphenicols have relatively high water solubility (2.5×10^3 –
457 4.1×10^5 mg/L) and lower $\log K_{ow}$ (-1.48–1.14), while the $\log K_{ow}$ of polypeptide are
458 extremely low at only -3.10 to -6.83 (Table S1). Some quinolones are different to above
459 antibiotics. For example, ciprofloxacin, difloxacin, enrofloxacin, ofloxacin and
460 norfloxacin also have high water solubility (850 – 1.8×10^5 mg/L) and low $\log K_{ow}$ (-
461 1.03–2.32), but have a low proportion in freshwaters, ranging from 0.37–13%. This is
462 because their half-lives in freshwaters are much shorter than those in soils. Many
463 antibiotics in other categories have comparable proportions in both freshwaters and
464 soils, with slightly higher percentages typically found in soil (Figure S3).

465 **Predicted environmental concentrations**

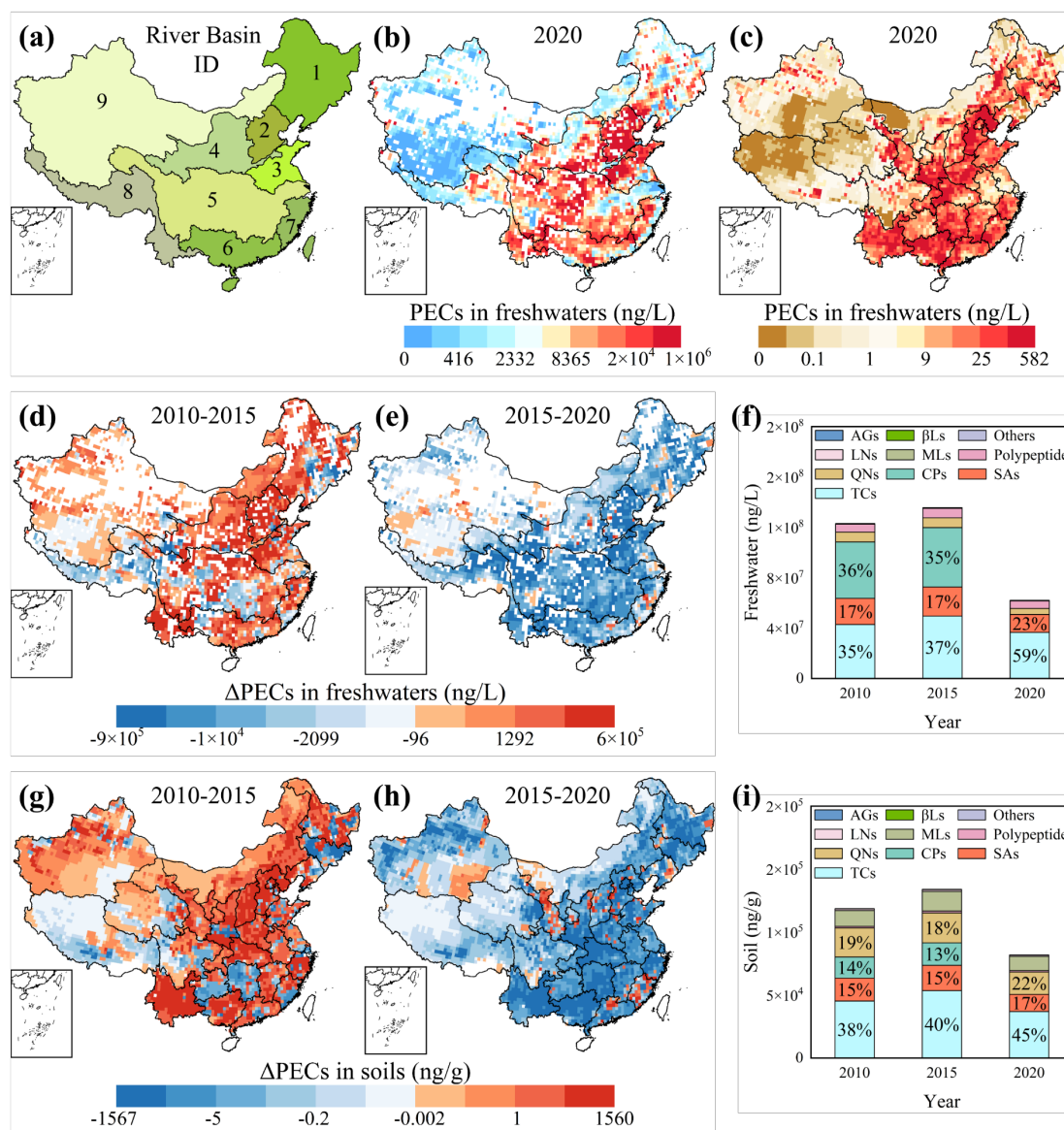
466 Although only ca. 15% of veterinary antibiotics was released to freshwaters, the
467 concentration therein was not low. The median PECs (5th–95th percentile range in
468 brackets) of Σ antibiotics in freshwaters were 1.2×10^4 (137 – 1.2×10^5), 1.3×10^4 (126 –
469 1.4×10^5) and 4.7×10^3 (48 – 6.6×10^4) ng/L in China in 2010, 2015 and 2020, respectively.
470 The median PECs of Σ antibiotics in soils were 5.3 (0.03–91), 6.0 (0.02–100) and 2.9
471 (0.02–61) ng/g in China in 2010, 2015 and 2020, respectively. Statistical characteristics
472 of PECs for individual antibiotics are shown in Table S10–S11. Elevated PECs of
473 Σ antibiotics in freshwaters in 2020 were found in the Haihe River Basin (Hebei
474 province), Huaihe River Basin (Henan province) and the section of Yangtze River Basin
475 in Henan province ($2,044$ – 1.3×10^6 , 600 – 3.0×10^5 and 1.8×10^4 – 2.6×10^5 ng/L) (Figure
476 3a-b). Meanwhile, Northwest Inland River Basin, Songhua and Liaohe River Basin,
477 and Southwest River Basin demonstrated lower PECs (0.30 – 8.4×10^5 , 0.01 – 2.4×10^5 ,
478 and 0 – 1.0×10^5 ng/L) compared to other basins. The highest PECs in freshwaters were
479 found in the Haihe River Basin in Hebei province, while the lowest PECs were
480 predicted in Southern Tibet Rivers (Southwest River Basin) in Tibet. This is consistent

481 with the findings of Zhang et al.¹⁷ The geographical pattern is also generally consistent
482 with that of emissions, but with regional contrast driven by environmental parameters,
483 such as dilution of freshwater discharge flows (SI Figure S2). In addition, Hebei,
484 Guangxi, and Henan provinces exhibited higher PECs in agricultural soils (11–582,
485 0.74–360, and 4.5–194 ng/g) than other areas. The highest PECs of antibiotics in soils
486 were found in Hebei province and the lowest PECs were found in Liaoning province
487 (Figure 3c). Resembling emissions, the provinces with elevated soil concentrations
488 were typically the seven provinces with the higher emission levels than other regions.
489 In line with the temporal pattern of emissions, most regions showed an increasing PECs
490 of Σ antibiotics from 2010 to 2015 followed by a decline after 2015. The PECs increased
491 0.02–8,150% in 46% of freshwater area and 0.001–4,130% in 75% of soil area (Figure
492 3d and 3g). However, variations of PECs in around 70% of regions was actually <30%.
493 Contrarily, a high percentage of regions (84% for freshwaters and 96% for soils)
494 exhibited a decline PECs of Σ antibiotics over 80% from 2015 to 2020 (Figure 3e and
495 3h).

496 Generally consistent with emissions, tetracyclines, chloramphenicols (except for
497 2020) and sulfonamides were major components in both freshwaters and soils, with
498 tetracyclines contributing the most (Figure 3f and 3i). Tetracyclines are a group of more
499 stable antibiotics in environments among the three highly emitted antibiotic categories
500 identified in this study.⁵⁹ The average half-life of tetracyclines is 1.2-6.0 times that of
501 chloramphenicols and sulfonamides in freshwaters and sediments, while it is
502 comparable with or 1.2 times that of chloramphenicols and sulfonamides in soils (Table
503 S1). Contrary to emissions, the proportion of chloramphenicols in freshwaters reached
504 35–36% (2015 and 2010), compared to around 14% in emissions, resulting from their
505 higher water solubility and lower log K_{ow} than the other categories - as indicated above
506 (Table S1). While chloramphenicol was banned for veterinary use in 2020, tetracyclines
507 became the major components, contributing 59% in freshwaters. Meanwhile in soils,
508 besides the above three major components, macrolides and quinolones were extra
509 essential components with proportions ranging from 11–14% and from 18–22%,
510 respectively. This is much higher compared to <1% and around 2% of macrolides, as

511 well as 6–8% and around 11% of quinolones in freshwaters and emissions, respectively.
512 The higher proportion of macrolides in soils than in freshwaters was attributed to its
513 hydrophobicity as stated above. The rising proportion of quinolones in soils was mainly
514 driven by the high soil PECs of enrofloxacin, due to its relatively higher $\log K_{ow}$ (2.32)
515 compared to other quinolones.⁶⁰

516 Before the ban, chloramphenicol (median, 2,959 ng/L, and the 5th–95th percentile
517 range, 34– 3.6×10^4 ng/L), should be the antibiotic with the highest PECs in freshwaters
518 in 2019, which is consistent with the findings of Zhang et al. (2015). In contrast to
519 Zhang et al.'s study, certain tetracyclines, specifically chlortetracycline (577 ng/L, 9.0–
520 1.4×10^4 ng/L) and oxytetracycline (310 ng/L, 3.1– 1.4×10^4 ng/L) have the second
521 highest PECs in freshwaters.¹⁷ Chloramphenicol (0.60 ng/g, 3.8×10^{-3} –10 ng/g) also
522 showed the highest concentration in soils in 2019. However, after it was banned,
523 oxytetracycline (0.23 ng/g, 2.4×10^{-3} –14 ng/g) and enrofloxacin (0.23 ng/g, 2.2×10^{-3} –
524 12 ng/g) became the antibiotics with the highest concentrations in soils in 2020 ([Table](#)
525 [S11](#)).



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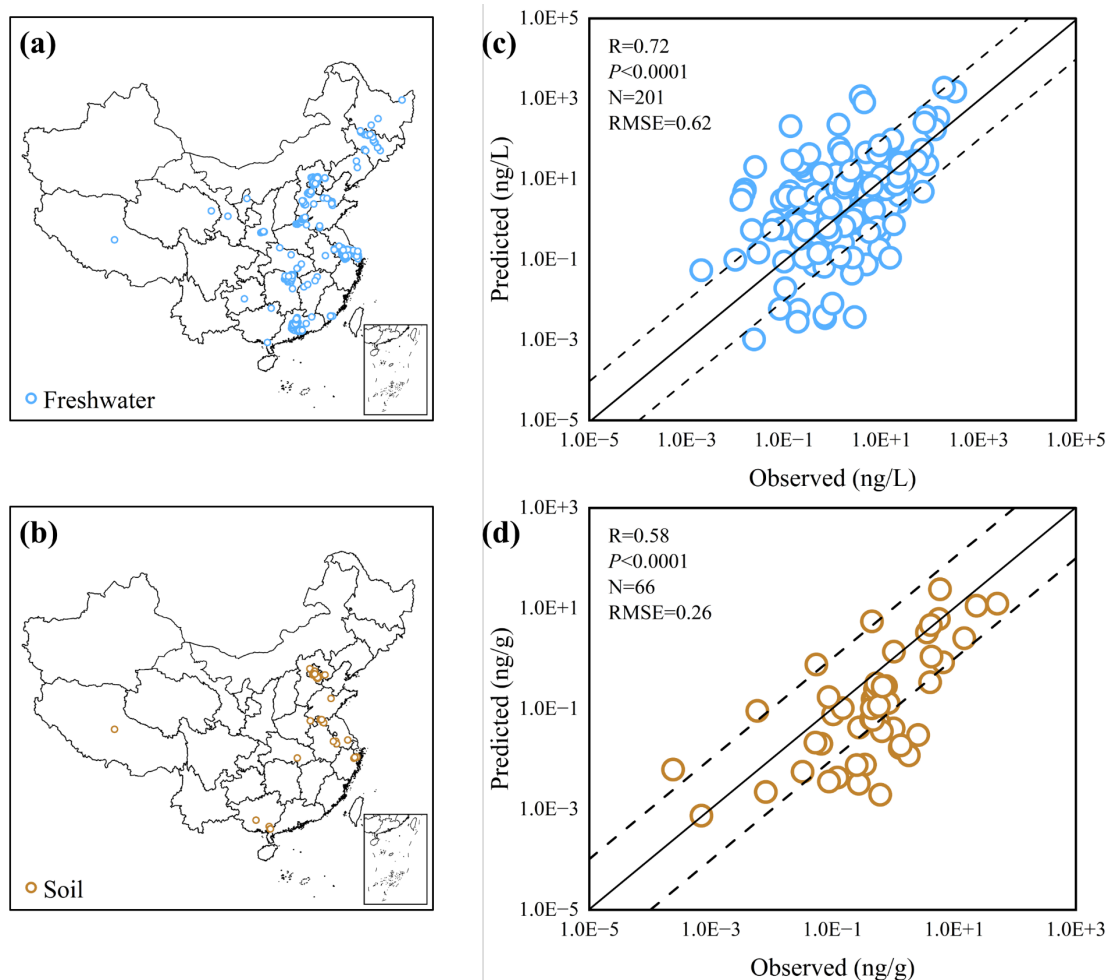
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Figure 3. (a) River Basin IDs: 1. Songhua and Liaohe River Basin; 2. Haihe River Basin; 3. Huaihe River Basin; 4. Yellow River Basin; 5. Yangtze River Basin; 6. Pearl River Basin; 7. Southeast River Basin; 8. Southwest River Basin; 9. Northwest Inland River Basin. Predicted environmental concentrations (PECs) of antibiotics in freshwaters (b) and soils (c) across China in 2020; variations in PECs of antibiotics in freshwaters (d-e) and soils (g-h) from 2010 to 2015 and 2015 to 2020; composition of antibiotics in freshwaters (f) and soils (i) in 2010, 2015 and 2020.



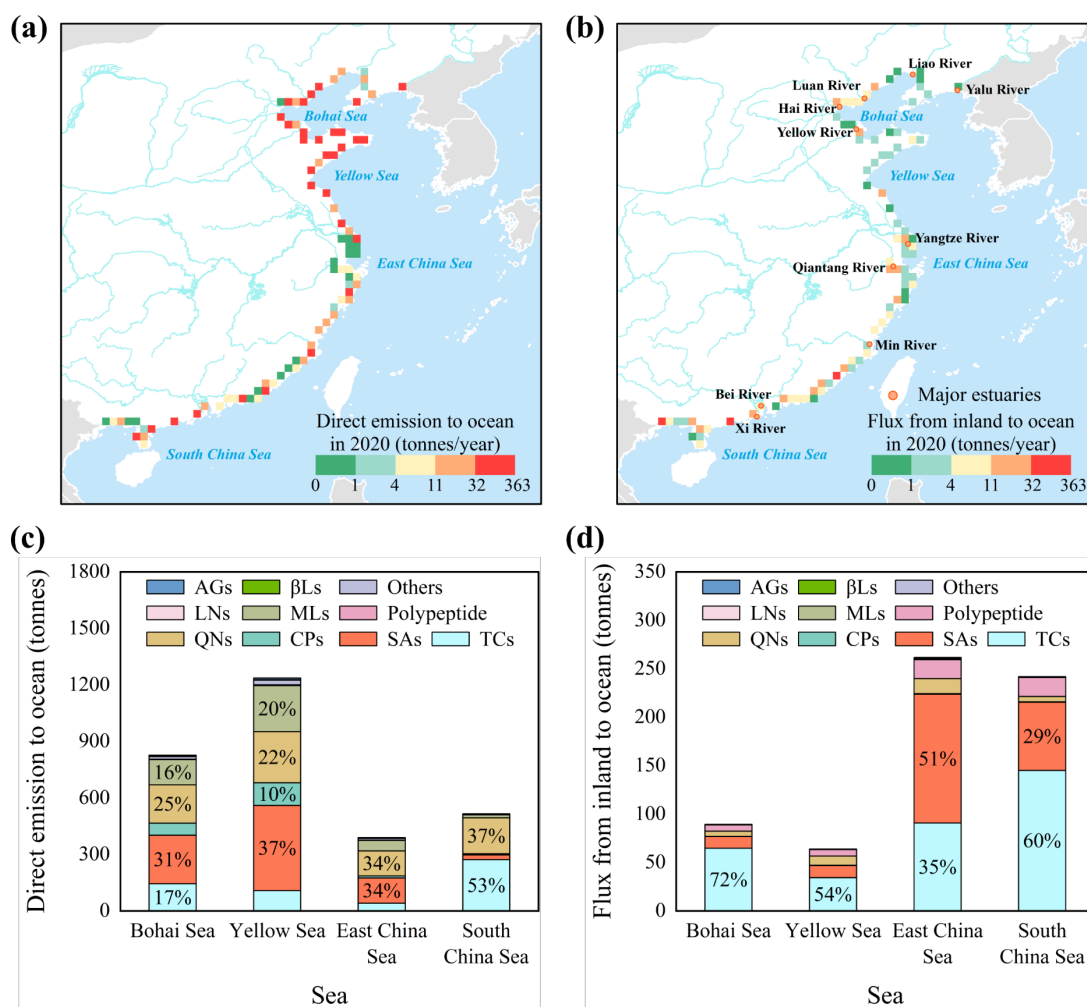
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535 **Figure 4.** (a-b) Sampling sites of observed data in freshwaters and soils; (c-d) point-to-
 536 point comparison of predictions and observations in freshwaters and soils for the nine
 537 animal-use-only (AUO) antibiotics. The solid line is 1:1 line. The dashed lines are 10:1
 538 and 1:10 lines. The root mean square error (RMSE) is logarithmic scaled.

539 Model performance was evaluated by comparing predictions with measurements of
 540 the nine AUO antibiotics with relatively sufficient monitoring data in freshwaters and
 541 soils. As our predictions are merely derived from veterinary sources, it is not reasonable
 542 to validate the model using field measurements of the veterinary antibiotics having
 543 human excretion as additional emission sources, even though some of them have
 544 abundant measured data. Sampling sites for all validated antibiotics in each
 545 compartment are shown in [Figure 4a](#) and [4b](#). Most major catchments in China are
 546 covered, whilst soil samples are mainly located in Beijing and Jiangsu province. A
 547 generally good agreement was found between predictions and measurements for both
 548 freshwaters and soils with a root mean square error (RMSE) at 0.62 and 0.26 log units,

549 respectively (Figure 4c-d). Significant and strong correlation (Pearson correlation
550 coefficient (R) > 0.5 , $P < 0.0001$) was found between predictions and measurements.
551 The model exhibits better performance on soils than freshwaters, demonstrated by the
552 lower RMSE for soils. Because soil is less mobile and exhibits more stable
553 characteristics compared to freshwaters, and the residue in soils can better reflect a
554 long-term situation. This better aligns with the nature of modelling results, which are
555 annual average concentrations. The underestimation in soils is mostly for the regions
556 with sampling sites close to livestock farms but having limited spatial coverage within
557 the grid cell to reflect an average concentration. Freshwater samples reflect momentary
558 concentrations, which can be easily affected by various factors, such as river discharge
559 flow rates and instantaneous release of pollutants. They usually cannot reflect a long-
560 term average contamination level. Therefore, although measurements may exhibit
561 generally consistent spatial patterns and contamination levels with predictions in
562 freshwaters, large local uncertainties may exist. This caused an under- or over-
563 estimation over one order of magnitude at several sites, with the corresponding points
564 falling out of the 1:10 lines. Therefore, the uncertainty falls within the acceptable range.
565 Considering the previous external validation of the model on other substances as
566 indicated in the method section, it is feasible and reliable to extrapolate the model on
567 other antibiotics having human sources or without sufficient measurements for model
568 validation. This is a common practice in modelling approaches especially for large scale.
569 The result of the uncertainty analysis conducted by Monte Carlo simulation is shown
570 in Figure S4–S5.

571 **Discharges to the ocean**



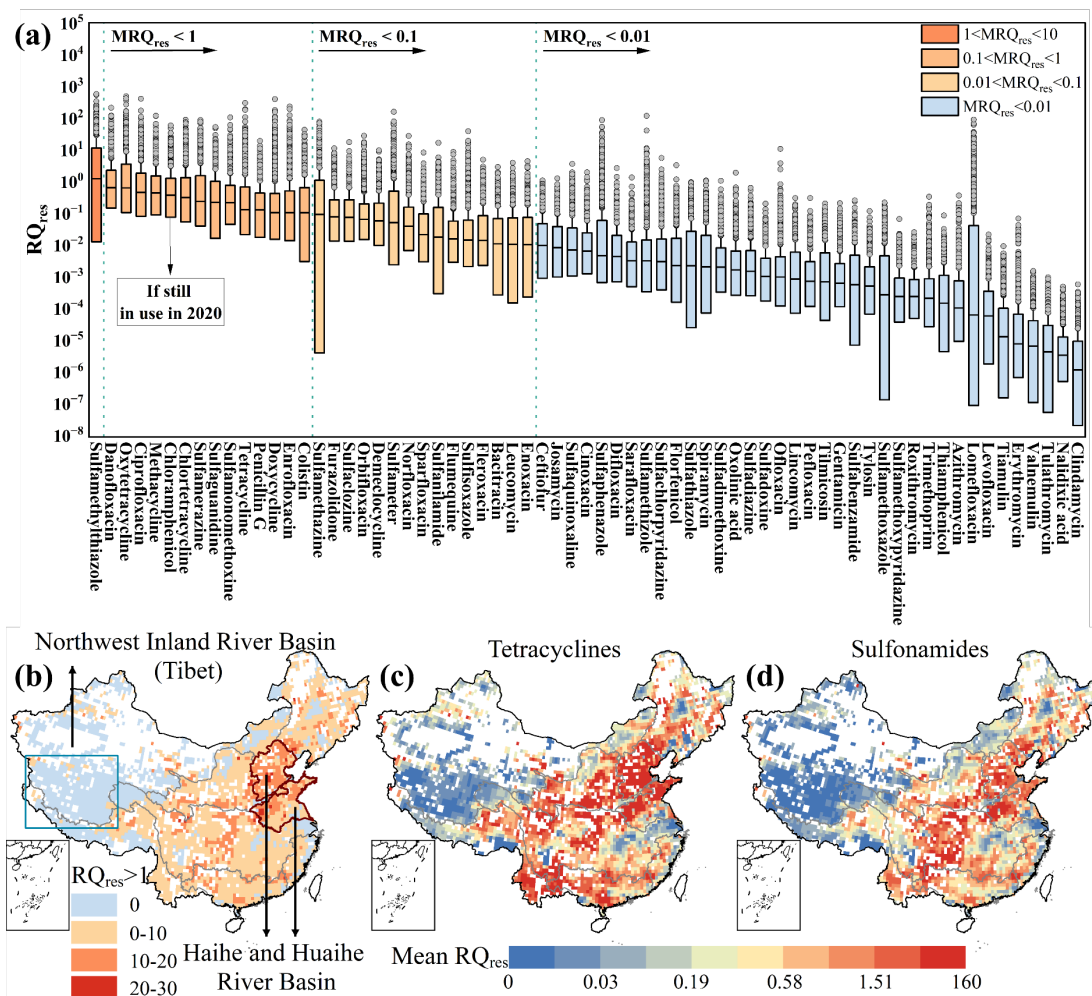
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573 **Figure 5.** Antibiotic direct emission to ocean from marine aquaculture (a) and fluxes
 574 from inland freshwaters to ocean (b) in 2020; composition of antibiotics in (c) marine
 575 aquaculture and (d) fluxes from inland freshwaters entering the Bohai Sea, Yellow Sea,
 576 East China Sea, and South China Sea, respectively.

577 The total discharges to the seawater along coastal regions of the Chinese Mainland
 578 were estimated, which comprised of source emissions from marine aquaculture and
 579 discharges from ten major inland rivers. In 2020, the total discharge to the ocean was
 580 3,626 (2,457–4,961) tonnes, resulting in a median PEC of Σ antibiotics reaching 1,094
 581 ng/L in seawaters. Direct emissions from marine aquaculture accounted for 82% of total
 582 discharges to the ocean. Around 42% and 28% of marine aquaculture emissions entered
 583 the Yellow Sea (1,237 tonnes) and Bohai Sea (827 tonnes), respectively (Figure 5a),
 584 due to the high marine aquaculture area in Shandong and Liaoning provinces. This
 585 resulted in a higher median PECs of Σ antibiotics in Yellow Sea (8,310 ng/L) and Bohai

586 Sea (4,119 ng/L) than in East China Sea (581 ng/L) and South China Sea (653 ng/L).
587 In contrast, the 657 (319–1470) tonnes of veterinary antibiotics delivered from inland
588 rivers primarily entered the East China Sea (262 tonnes) and South China Sea (242
589 tonnes), due to high discharge flow rates at the river estuaries here (Figure 5b). For
590 example, the Yangtze River, with an annual freshwater discharge flow of 8.5×10^{11}
591 m^3/year , delivered at least 22 tonnes of veterinary antibiotics to the East China Sea in
592 2020. Sulfonamides, tetracyclines, quinolones, and macrolides were the primary
593 components in the source emissions from marine aquaculture into all the four seas
594 (Figure 5c), which accounted for 87–98% of the source emissions to individual seas in
595 2020. However, tetracyclines and sulfonamides were the primary components in
596 discharges from inland freshwaters into all the four seas (Figure 5d), which accounted
597 for 35–72% and 14–51% of discharges to individual seas in 2020. Sulfonamides were
598 the predominant components in the freshwater discharge to the East China Sea, while
599 tetracyclines were the predominant component for the other three seas. The distinct use
600 pattern of veterinary antibiotics in varying regions may influence the composition in
601 the discharge to different seas.

602 **Risks**



603

604 **Figure 6.** (a) Boxplot of predicted RQ_{res} in freshwater for 67 antibiotics in China; the
 605 horizontal solid line in the box is the median RQ_{res} (MRQ_{res} as shown, the MRQ_{res} of
 606 all antibiotics are greater than 0); the top and bottom of the box are the 75th (Q3) and
 607 25th (Q1) percentiles, respectively; the top and bottom of the whisker are the highest
 608 and lowest values within 1.5 times of the interquartile range (IQ, i.e., [$Q1-1.5IQ$, $Q3+$
 609 $1.5IQ$]). The circles are outliers with RQ_{res} out of the range of the whiskers. (b) Spatial
 610 distribution of the number of antibiotics in freshwaters at high risk ($RQ_{res} > 1$) of
 611 antibiotic resistance development; and spatial distribution of mean RQ_{res} for
 612 tetracyclines (c) and sulfonamides (d). The other 13 antibiotics only occur in a few cells
 613 due to the only application in aquaculture, so the MRQ_{res} across China is zero.

614 The PECs in freshwaters in 2020 under the BG scenario were applied to be compared
 615 with $PNEC_{res}$, to assess the risk of antibiotic resistance development (SI Table S7). A
 616 total of 44 antibiotics sourced from veterinary use had a high risk of antibiotic resistance

617 development ($RQ_{res} > 1$) in freshwaters in at least one grid cell (Figure 6a). There were
618 seven of them exhibiting the high risk in >10% of the total freshwater area (ranging
619 from 1.6×10^4 – 5.0×10^4 km²), namely sulfamethylthiazole, danofloxacin,
620 oxytetracycline, chlortetracycline, methacycline, sulfamerazine, ciprofloxacin, and
621 sulfamethazine. Sulfamethylthiazole was the antibiotic demonstrating the highest risk
622 with a median RQ_{res} (MRQ_{res}) > 1. Over 20 antibiotics simultaneously exhibited a high
623 risk in ca. 610 km² freshwater areas across China, with most areas distributed in the
624 Haihe River Basin (130 km²) as well as the Huaihe River Basin (310 km²), accounting
625 for 4.2% and 3.0% of the freshwater area in the respective river basins. The highest risk
626 appeared in the Haihe River Basin in the Beijing-Tianjin-Hebei region, with 30
627 antibiotics exhibiting a high risk ($RQ_{res} > 1$) in at least one grid cell within the basin
628 (Figure 6b). Notably, the prediction was an average level within a 0.5° grid cell. An
629 identified high risk implied the existence of near sources or local areas having hotspots
630 of potentially extremely high risk. A total of 13 antibiotics demonstrated $0.1 < MRQ_{res}$
631 < 1 , while 63 antibiotics had a $MRQ_{res} < 0.1$. There were 38 out of the 63 antibiotics
632 possessing a low risk ($RQ_{res} < 0.1$) in more than 99% of the total freshwater area in
633 China. The risk associated with these antibiotics was deemed negligible. A generally
634 lower risk was observed in the Northwest Inland River Basin within the Tibet (Figure
635 6b).

636 Tetracyclines had a generally higher risk, with 5 out of 6 antibiotics having a $MRQ_{res} >$
637 0.1, followed by sulfonamides (4 antibiotics having a $MRQ_{res} > 0.1$) (Figure 6c-d). All
638 other categories of antibiotics, apart from danofloxacin, ciprofloxacin, penicillin G,
639 enrofloxacin, and colistin, had a $MRQ_{res} < 0.1$. This was consistent with the wide
640 distribution of sulfonamide (antibiotic resistance genes (ARGs) and tetracycline ARGs
641 in the aquatic environment.^{61,62} Concentrations of tetracyclines and sulfonamides
642 exhibited concordance with levels of the *tet* and *sul* genes.⁶³ Therefore, high
643 concentrations of tetracyclines and sulfonamides emitted from veterinary sources may
644 allow high levels of relevant ARGs to be transmitted in the environment, posing a risk
645 to human health. A risk screening based on aquatic toxicity is presented in Figure S6,
646 illustrating that 26 antibiotics may pose potential high risks to freshwater ecosystems.

647 **Uncertainties and limitations**

648 Limitations exist due to the lack of data, which may introduce uncertainties in the
649 estimation and assessment. (1) Livestock farming volumes were lacking in a few
650 regions for certain years, which were estimated by temporal extrapolation based on data
651 present for other years. This may introduce uncertainties to emission estimation.
652 However, the uncertainty should be extremely limited, as the data were collected on a
653 county-level basis, which has ensured the highest accuracy. The farming yield should
654 not change dramatically in a short time. (2) Spatially varied measurements of antibiotics
655 in faeces and aquaculture water were applied regionally to reflect the potential regional
656 difference in use. However, for regions without sampling sites, a national average data
657 was applied, which would possibly result in uncertainties regionally. (3) Manure
658 treatment might not be fully implemented in reality and follow the treatment rate
659 recorded in the literature. The application of literature data may lead to slight
660 underestimation. (4) The input of physico-chemical parameters largely relied on
661 predictions, which may have uncertainties especially for half-lives. More
662 measurements of these parameters are required to refine the prediction of environmental
663 concentrations. (5) $PNEC_{res}$ for more antibiotics are needed for a better assessment of
664 risks for resistance development in environments. Future improvement can be achieved
665 with the acquisition of more precise data, particularly antibiotic use data accounting for
666 spatial variations, removal rates in different manure treatment processes and $PNEC_{res}$.
667 The acquisition of such data is pivotal in enhancing the accuracy and robustness of the
668 research. Despite all the above limitations, the method and relevant data we adopted are
669 mature and the optimal choice based on current available data, so far as we know.

670 **Implications**

671 This is the first study to conduct a comprehensive assessment on emissions and
672 environmental fate for a relatively complete range of veterinary antibiotics with a high
673 spatial resolution over the past 11 years in China. The result indicates that multiple
674 measures should be taken to further control antibiotic emissions and reduce potential
675 risks of resistance development in environments, sourced from animal use. Firstly, the
676 prohibition of veterinary use of certain antibiotics, such as pefloxacin, ofloxacin and

677 norfloxacin, as well as combination therapy should be better supervised in China by
678 relevant governmental agencies.⁶⁴ Because (a) pefloxacin, ofloxacin and norfloxacin
679 can still be detected in fresh faeces samples taken even after being banned in 2016, and
680 (b) multiple antibiotics are often detected in faeces of certain livestock within one farm
681 during a single monitoring campaign.^{65,66} Techniques to efficiently remove antibiotics
682 from manure should be investigated and adopted in manure treatment processes.
683 Meanwhile, it is easier to improve and implement advanced manure treatments at
684 intensive livestock farms which can improve the removal efficiency of antibiotics
685 before release. Supervision of manure treatment could be more easily implemented in
686 intensive farms. On the other hand, considering that the risk of resistance development
687 in freshwaters is mainly caused by the transport of antibiotics from soils rather than the
688 direct source emission to freshwaters, the current manure discharge measure may not
689 be appropriate. Efforts should be made to explore optimized discharge measures aimed
690 at retaining more antibiotics in the ground, rather than allowing them to be easily
691 transported to freshwater, which is a mobile compartment more capable of spreading
692 the contamination. Finally, colistin, as the last-resort antibiotic for multidrug resistant
693 gram-negative infections in humans was detected in livestock faeces. There have also
694 been reports indicating its extensive use in livestock in China and Vietnam.⁶⁷ Veterinary
695 use of such high-end antibiotics should be heavily controlled under strict supervision
696 by the government. This is a concern because once resistance to them appears, it will
697 spread rapidly among different types of bacteria, leaving the human population at risk
698 of being exposed to a range of untreatable infections. This comprehensive assessment
699 is therefore an important reference source for policy makers.

700 **Supporting Information**

701 The Supporting Information is available at <https://pubs.acs.org/>

702 Supplemental methods, tables, and figures (PDF)

703 The raw data of livestock numbers and aquaculture areas; proportion of treatment
704 process in livestock farms; concentration of antibiotic residues in aquaculture and
705 faeces; proportion of intensive farms (Data S1-S17) ([Appendix A](#))

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