1 Supplementary File

2	Including bioconcentration of pesticide metabolites in plant uptake modeling
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23 S1. Analytical solutions of the general model

- 24 The analytical solutions of the general model are derived in this section.
- 25 In the soil compartment:

Parent compound:
$$C_{P,Soil}(t) = C_{P,Soil}(0) \exp(-k_{P,Soil}^{Diss}t)$$
 (s1)

$$Metabolite \ i: C_{M,i,Soil}(t) = \frac{k_{M,i,Soil}^{Trans}}{k_{M,i,Soil}^{Diss}} C_{P,Soil}(0) \left[\exp\left(-k_{P,Soil}^{Diss}t\right) - \exp\left(-k_{M,i,Soil}^{Diss}t\right) \right]$$
(s2)

26 In the tuber compartment:

Parent compound:
$$C_{P,Tuber}(t)$$
 (s3)

$$= \frac{k_{P,Tuber}^{S \to T}}{k_{P,Tuber}^{T \to S} + k_{P,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss}} C_{P,Soil}(0) \{ \exp(-k_{P,Soil}^{Diss}t) - \exp[-(k_{P,Tuber}^{T \to S} + k_{P,Tuber}^{Met} + k_{Tuber}^{Grow})t] \}$$

(s4a)

(s4b)

Metabolite i:
$$C_{M,i,Tuber}(t)$$

$$= \frac{k_{M,i,Soil}^{Trans}k_{M,i,Tuber}^{S \to T}}{(k_{M,i,Soil}^{Diss} - k_{P,Soil}^{Diss})} + \frac{k_{M,i,Tuber}^{Trans}k_{P,Tuber}^{S \to T}}{(k_{P,Tuber}^{T \to S} + k_{P,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss})} C_{P,Soil}(0) \exp(-k_{P,Soil}^{Diss}t)$$

$$- \frac{k_{M,i,Tuber}^{S \to T} + k_{M,i,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss}}{(k_{M,i,Soil}^{T \to S} - k_{P,Soil}^{Diss})(k_{M,i,Tuber}^{T \to S} + k_{M,i,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss})} C_{P,Soil}(0) \exp(-k_{M,i,Soil}^{Diss}t)$$

$$- \frac{k_{M,i,Tuber}^{S \to T} + k_{M,i,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss}}{(k_{M,i,Soil}^{T \to S} - k_{P,Soil}^{Diss})(k_{M,i,Tuber}^{T \to S} + k_{M,i,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{M,i,Soil}^{Diss})} C_{P,Soil}(0) \exp(-k_{M,i,Soil}^{Diss}t)$$

$$- \frac{k_{M,i,Tuber}^{T ans} + k_{P,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss}}{(k_{P,Tuber}^{T \to S} + k_{P,Tuber}^{Met} - k_{P,Soil}^{P,Soil})(k_{M,i,Tuber}^{T \to S} + k_{M,i,Tuber}^{Met} - k_{P,Tuber}^{T \to S} - k_{P,Tuber}^{Met})} C_{P,Soil}(0) \exp[-(k_{P,Tuber}^{T \to S} + k_{P,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss})(k_{M,i,Tuber}^{T \to S} + k_{M,i,Tuber}^{Met} - k_{P,Tuber}^{T \to S} - k_{P,Tuber}^{Met})t] + A \exp[-(k_{M,i,Tuber}^{T \to S} + k_{M,i,Tuber}^{Met} + k_{Tuber}^{Grow})t]$$

Integration constant: A

$$= -\frac{\frac{k_{M,i,Soil}^{Trans}k_{M,i,Tuber}^{S \to T}}{(k_{M,i,Soil}^{Diss} - k_{P,Soil}^{Diss})} + \frac{k_{M,i,Tuber}^{Trans}k_{P,Tuber}^{S \to T}}{(k_{P,Tuber}^{T \to S} + k_{P,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss})}C_{P,Soil}(0)$$

$$+ \frac{k_{M,i,Tuber}^{S \to T} + k_{M,i,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss}}{(k_{M,i,Soil}^{Diss} - k_{P,Soil}^{Diss})(k_{M,i,Tuber}^{T \to S} + k_{M,i,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{P,Soil}^{Diss})}C_{P,Soil}(0)$$

$$+ \frac{k_{M,i,Soil}^{T \to S} - k_{P,Soil}^{Diss}}{(k_{M,i,Soil}^{T \to S} - k_{P,Soil}^{Diss})(k_{M,i,Tuber}^{T \to S} + k_{M,i,Tuber}^{Met} + k_{Tuber}^{Grow} - k_{D,Soil}^{Diss})}C_{P,Soil}(0)$$

27 where A is the integration constant.

28 S2. Model application for glyphosate

In this section, we present the calculation process of the rate constants of glyphosate and its
primary toxic metabolite (i.e., AMPA).

31 In the soil compartment:

For the parent compound (glyphosate), the overall dissipation process comprises biodegradation (microorganism), volatilization, water-induced elimination (e.g., surface runoff, and vertical filtration), the uptake by plants (e.g., grasses, weeds), and photolysis (Li, 2021a; Li and Niu, 2021). For the simulation purpose, we only considered the biodegradation process, and other major elimination processes were included in the variability analysis. Thus, the $k_{P,Soil}^{Diss}$ value of glyphosate was approximated as 0.058 d⁻¹ (Fantke et al., 2015; Lewis et al., 2016; Li, 2021a), which was based on the biodegradation process in the soil.

For the primary toxic metabolite of glyphosate (AMPA), the overall dissipation process in the soil comprises processes that are similar to those of glyphosate, which is also discussed in the variability analysis. Thus, the $k_{M,i,Soil}^{Diss}$ value of AMPA was approximated as 0.0057 d⁻¹ based on the biodegradation process (European Commission, 2020). The $k_{M,i,Soil}^{Trans}$ value of AMPA was estimated as 0.38 d⁻¹, which is the product of the biodegradation rate constant (i.e., 0.058 d⁻¹) and the molecular weight ratio of AMPA to glyphosate (i.e., 0.66) (Li, 2021a).

45 In the tuber compartment:

46 The default value of 0.139 d⁻¹ was set to k_{Tuber}^{Grow} , which was based on the logistic growth dynamics

47 of the potato (Fantke et al., 2011a; Trapp et al., 2007). Due to little information on the metabolic

48 rate constants of pesticides in plant tissues (Jacobsen et al., 2015; Xiao et al., 2021a, 2021b), we

used half of the dissipation rate constant tabulated by the USEtox database (Fantke et al., 2017), 49 which was estimated as 0.15 d⁻¹ for $k_{P,Tuber}^{Met}$. To evaluate the impact of the metabolic kinetics of the 50 compound on simulation results, we conducted the variability analysis in Section S3. In addition, 51 for the same reason, the metabolic rate constant of AMPA in potatoes is unknown; thus, for the 52 risk assessment (Juraske et al., 2011; Trapp et al., 2007), the k_{M,i,Tuber} value of AMPA was set as 53 zero. The value $k_{M,i,Tuber}^{Trans}$ of AMPA can be calculated based on the same method for that in the soil, 54 which is estimated as 0.1 d⁻¹, namely the product of the metabolic rate constant of glyphosate in 55 the potato (i.e., $k_{P,Tuber}^{Met} = 0.15 \text{ d}^{-1}$) and the molecular weight ratio of AMPA to glyphosate (i.e., 56 0.66). 57

58 For the uptake and elimination routes of the compound in potatoes via diffusion, the spherical-59 shape-based point-estimate approach (Trapp et al., 2007) was adopted to simulate the uptake and 60 elimination rate constants as follows:

$$k_{Tuber}^{S \to T} = \frac{23\theta_{W,T}T_{Tuber}D_W}{r_S^2 \rho_{Tuber}K_{SW}}$$
(s5)

$$k_{Tuber}^{T \to S} = \frac{23\theta_{W,T}T_{Tuber}D_W}{r_S^2\rho_{Tuber}K_{TW}}$$
(s6)

where D_W (m² d⁻¹) is the diffusivity of the chemical in water, and the values for glyphosate and AMPA were calculated as 7.5×10^{-5} m² d⁻¹ and 9.3×10^{-5} m² d⁻¹, respectively, based on the molecular weight approach (Trapp et al., 2003). $\theta_{W,T}$ (dimensionless) is the water content of the tuber tissue, which was estimated as 0.778 (Caetano et al., 2018). T_{Tuber} (dimensionless) is the tortuosity of the tuber tissue, which was estimated as 0.72 (Li, 2021b) based on the water content and air pores of the tuber tissue (Trapp, 2007; Trapp et al., 2007). r_s (m) is the radius of the tuber, which was estimated as 0.04 m based on the medium-sized potato (Trapp et al., 2007). ρ_{Tuber} (kg 68 L^{-1}) is the density of the tuber (medulla), which was estimated as 1.1 (Li, 2020). κ_{sw} (L kg⁻¹) is 69 the soil-water partition coefficient of the compound, which can be estimated according to the major 70 (air, water, and organic matter) components of the soil (Paraíba and Kataguiri, 2008; Trapp et al., 71 2007) as follows:

$$K_{SW} = \frac{\rho_{S,Dry} f_{OC} K_{OC} + f_W + K_{AW} f_A}{\rho_{S,Wet}} \approx \frac{\rho_{S,Dry} f_{OC} K_{OC} + f_W}{\rho_{S,Wet}}$$
(s7)

where $\rho_{s,Dry}$ (kg L⁻¹) and $\rho_{s,Wet}$ (kg L⁻¹) are the densities of dry and wet soil, respectively, which 72 were estimated as 1.6 kg L⁻¹ and 1.95 kg L⁻¹, respectively (Trapp et al., 2007). f_{oc} (dimensionless) 73 is the fraction of the organic matter of the soil, which was estimated as 0.018 (Trapp et al., 2007). 74 f_W (dimensionless) is the water fraction of the soil, which was estimated as 0.28 (Trapp et al., 75 2007). f_A (dimensionless) is the air fraction of the soil, which was estimated as 0.12 (Trapp et al., 76 2007). κ_{oc} (L kg⁻¹) is the soil sorption partition coefficient of the compound, and the logarithm 77 values (log Koc) of glyphosate and AMPA were taken as 4.34 and 3.7, respectively (Daouk, 2013). 78 K_{AW} (dimensionless) is the air-water partition coefficient (or the dimensionless Henry's law 79 constant) of the compound. We note that the K_{AW} values of glyphosate and AMPA are so low (i.e., 80 non-volatile compounds) that the term ' $K_{AW}f_A$ ' in Eq. (s7) can be negligible (Bento, 2018). Then, 81 the K_{SW} values of glyphosate and AMPA were calculated based on the inputs above as 323 L kg⁻¹ 82 and 74.2 L kg⁻¹, respectively. K_{TW} (L kg⁻¹) is the tuber tissue-water partition coefficient of the 83 84 compound, which can be estimated using the major nutritional components of the tuber (Chiou et al., 2001; Trapp et al., 2007) as follows 85

$$K_{TW} = 1.22 f_{Lip,T} K_{OW}^{0.77} + f_{Carb,T} K_{Ch} + \theta_{W,T}$$
(s8)

86 where K_{ow} (dimensionless) is the octanol-water partition coefficient of the compound, and the

logarithm values of glyphosate and AMPA were estimated as -3.4 and -1.4, respectively, (National 87 Library of Medicine, 2021; Royal Society of Chemistry, 2022). $f_{Lip,T}$ (dimensionless) and $f_{Carb,T}$ 88 (dimensionless), are the massive contents of lipid and non-lipid carbohydrate, respectively, which 89 were estimated as 0.001 and 0.154, respectively (Trapp et al., 2007). K_{ch} (L kg⁻¹) is the 90 carbohydrate-water partition coefficient of the compound, and the values of glyphosate and AMPA 91 were estimated (via lipophilicity) as 0.1 (Chiou et al., 2001; Trapp et al., 2007). $\theta_{W,T}$ (L kg⁻¹) is 92 the volumetric content of water in the tuber, which was estimated as 0.778 (Trapp et al., 2007). 93 Then, the K_{TW} values of glyphosate and AMPA were calculated as 0.79. 94

95 S3. Variability analysis

96 The variability analysis was conducted by varying selected rate constants or essential 97 physicochemical properties of glyphosate and AMPA, which generated the variability intervals of 98 the simulation results (e.g., concentrations of the glyphosate and AMPA in the potato). The 99 variability analysis can help users conduct the regional-specific assessment of bioconcentration of 100 pesticides and their toxic metabolites by focusing on site-specific model inputs.

101 **S3.1 Dissipation rate constants of compounds in the soil**

In the previous section (i.e., simulating concentrations of glyphosate and AMPA), we only considered the degradation kinetics of glyphosate and AMPA in the soil to estimate the degradation rate constants (i.e., $k_{P,Soil}^{Diss}$ and $k_{M,i,Soil}^{Diss}$) for the conservative consideration (i.e., for the human health risk assessment). However, the overall dissipation of glyphosate and AMPA involves multiple processes. Thus, we varied $k_{P,Soil}^{Diss}$ and $k_{M,i,Soil}^{Diss}$ values of glyphosate and AMPA to generate the variability intervals of the simulated concentrations in the potato. The ranges of $k_{P,Soil}^{Diss}$ and $k_{M,i,Soil}^{Diss}$ values of glyphosate and AMPA were taken as $[3.5 \times 10^{-3}, 0.7] d^{-1}$ and $[7.2 \times 10^{-4}, 3.0 \times 10^{-2}] d^{-1}$, respectively (Bento, 2018). Accordingly, the $k_{M,i,Soil}^{Trans}$ value of AMPA was estimated by multiplying the half of $k_{P,Soil}^{Diss}$ and 0.66 (i.e., the molecular weight ratio between AMPA and glyphosate). The variability intervals of the simulation results were generated by using the

112 **S3.2** Koc of compounds in the soil

We also varied the K_{OC} values of glyphosate and AMPA to generate the variability intervals of the simulated concentrations of glyphosate and AMPA in the potato because the K_{OC} values can be substantially affected by soil properties (e.g., pH and charges). The ranges of K_{OC} values of glyphosate and AMPA were taken as [884, 60000] L kg⁻¹ and [1160, 24800] L kg⁻¹, respectively (Bento, 2018).

118 **S3.3 Metabolic rate constants of compounds in the potato**

In the previous section, we set the metabolic rate constant of AMPA as zero for the conservative consideration (i.e., the purpose of the human health risk assessment). However, AMPA can be further biotransformed by plant enzymes (la Cecilia et al., 2018). Thus, we varied the $k_{M,i,Tuber}^{Met}$ value of AMPA in the potato to generate the variability interval of the simulation results. Due to data limitations, we set $k_{M,i,Tuber}^{Met}$ values as 0 d⁻¹, 0.1 d⁻¹, 0.25 d⁻¹, and 0.5 d⁻¹.

124 S4. Case study

125 Maggi et al. (2020) evaluated the concentrations of glyphosate and AMPA in global surface soil.

126 We utilized the data of Maggi et al. (2020) (soil residue concentrations) to model the toxic pressure

- 127 of glyphosate and AMAP in potatoes. The results of the simulation are presented in Table S1.
- 128 Importantly, certain observed AMPA concentrations in the soil are notably higher than those of

glyphosate, while in other cases, this difference is not evident. This phenomenon arises from the fact that pesticides are usually applied in pulses, and if measurements occur shortly after pesticide application, the parent compounds might exhibit considerably greater soil concentrations than their corresponding metabolites. Additionally, various dynamic environmental factors, including precipitation, wind, sunlight, and humidity, can exert influence on the soil concentrations of chemical contaminants.

Table S1. Summary of glyphosate and AMPA concentration simulation results in potatoes based on their concentrations in global surface soil. The soil concentrations were obtained from Maggi et al. (2020), which were compiled from the scientific literature. Metabolite and non-metabolite models were utilized to calculate glyphosate and AMPA concentrations in potatoes using simulated bioconcentration factors with a 30-day time-to-harvest interval.

	Observed (means)		Metabolite model		Non-metabolite model	
	Glyphosate	AMPA	Glyphosate	AMPA	Glyphosate	AMPA
Location	concentration	concentration	concentration	concentration	concentration	concentration
	in soil	in soil	in potato	in potato	in potato	in potato
	(mg/kg)	(mg/kg)	(mg/kg)	(mg/kg)	(mg/kg)	(mg/kg)
Province of Buenos Aires,						
Argentina	0.0347	0.2993	2.31E-06	40.99	2.76E-06	34.31
Province of Buenos Aires,						
Argentina	0.53	0	3.53E-05	0	4.22E-05	0
Toholamni, Western Finland	0.1476	0.1284	9.84E-06	17.58	1.17E-05	14.72
Nntuna, Sweden	0.00965	0.0535	6.43E-07	7.33	7.68E-07	6.13
Lanna, Sweden	0.02147	0.1177	1.43E-06	16.12	1.71E-06	13.49
Sandved, Denmark	0.00081	0.01046	5.40E-08	1.43	6.44E-08	1.20
Montepaldi-San Csciano, Italy	0	0.0609	0	8.34	0	6.98
United Kindom	0.15	0.15	1.00E-05	20.54	1.19E-05	17.20
Denmark	0.11	0.14	7.33E-06	19.17	8.75E-06	16.05
Portugal	1.14	0.73	7.60E-05	99.97	9.07E-05	83.69
Italy	0.13	0.1	8.66E-06	13.70	1.03E-05	11.46
Greece	0.54	0.21	3.60E-05	28.76	4.30E-05	24.08

Spain	0.22	0.09	1.47E-05	12.33	1.75E-05	10.32
Hungary	0.1	0.23	6.67E-06	31.50	7.96E-06	26.37
Poland	0.16	0.14	1.07E-05	19.17	1.27E-05	16.05
Netherland	0.13	0.13	8.66E-06	17.80	1.03E-05	14.90
France	0.08	0.13	5.33E-06	17.80	6.36E-06	14.90
Germany	0.13	0.15	8.66E-06	20.54	1.03E-05	17.20

140

141 **S5. Supplementary figures**



142 The supplementary figures of the main text are provided in this section.

143

Figure S1. Simulated concentrations of glyphosate and aminomethylphosphonic acid (AMPA) in
the soil based on the initial concentration of glyphosate of 1 mg kg⁻¹ in the soil plotted against time





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Figure S2. The ratio of the simulated AMPA concentrations in the potato between the metabolite
and non-metabolite models plotted against time (t, d) after glyphosate application.

150 S6. Model extension

In this study, we present a modeling approach that considers the bioconcentration of pesticide 151 152 metabolites in plants. We demonstrate the effectiveness of this approach using potatoes as an example crop. While there is currently a shortage of experimental studies to evaluate the 153 simulations of pesticides including their metabolites, the plant uptake models, which underlie our 154 proposed approach, have been robustly supported by field observations (Pang et al., 2020, 2016). 155 This robustness lends credibility to our approach for predicting the uptake of organic compounds, 156 encompassing both parent compounds and their metabolites. The modeling experiments utilizing 157 158 glyphosate as an example serve as a reminder to risk and impact assessors to prioritize the consideration of pesticides' toxic metabolites in addition to the parent compounds. This will 159 contribute to a more comprehensive assessment framework. We note that different plant species 160 have varying pesticide uptake routes that depend on both plant physiology and pesticide 161

application patterns. However, our proposed modeling approach can be adapted to other crops and 162 their respective pesticide uptake routes. Table S2 provides a summary of the extension of our 163 modeling approach to other crops. Above-ground crops have more pathways for pesticide uptake 164 compared to tuber crops, including surface penetration (such as fruit, leaves, and stems) and 165 transpiration. This is because pesticide residues can be distributed in soil, air, and plant surfaces 166 167 after their application, particularly through aerial spraying. As a result, pesticide metabolites can be generated in various compartments, which increases their uptake routes by above-ground crops. 168 To simulate metabolite concentrations in above-ground crops, additional equations for metabolite 169 170 generation compartments and pesticide and metabolite uptake routes are necessary. These equations can be generated and solved using a first-order kinetics-based matrix approach (Fantke 171 et al., 2011b, 2011a). This matrix-based approach offers the capability to provide a comprehensive 172 pesticide bioconcentration or bioaccumulation modeling framework for various plant 173 compartments (including stem, leaf, fruit, and root) across diverse plant species. This allows users 174 to tailor their simulations to edible portions of the plant. For instance, risk assessors may be 175 176 concerned with pesticide bioconcentration modeling in the leaves of lettuces, while the focus might shift to the fruit of apple trees. With the integration of metabolite simulations into this matrix 177 approach, the added toxicity of metabolites to their parent compounds across all modeling 178 179 compartments-leaf, fruit, stem, and root-can be generated, which equips users to choose simulation outcomes tailored to any consumable plant parts. 180

181 Notably, the PHI for glyphosate application in crops can vary widely, such as ranging from 7 to 28
182 d (EFSA, 2013). While our simulation results have revealed that the concentration of glyphosate

in potatoes starts to decline shortly after pesticide application (around 2-3 d) and continues to 183 decrease during the potato's growth period, its main metabolite, AMPA, displays a consistent 184 increase in concentration within the potato over a longer period. Therefore, even though certain 185 suggested PHIs for glyphosate might result in reduced pesticide concentrations (e.g., glyphosate 186 itself) at harvest time, their metabolites are likely to accumulate in the edible parts of crops, 187 188 introducing potentially significant health risks to consumers. Consequently, extending the PHI can indeed help diminish pesticide residue levels in crops. However, this approach warrants careful 189 consideration, particularly for toxic compounds and their metabolites, which could pose substantial 190 health concerns. Hence, it is advisable that the proposed PHI of pesticides, in conjunction with 191 their efficacy in pest management, takes into account the potential adverse effects posed by their 192 metabolites on human and ecological health. 193

Plant growth dynamics can exert a significant influence on the bioconcentration of pesticides in plant tissues. In this study, we employed a simple yet effective strategy, utilizing the growth rate constant to capture the dilution effect on chemical fate and transport within plants. This approach, widely adopted in modeling and risk assessment endeavors, allows for a straightforward depiction of pesticide behavior (Fantke et al., 2011a; Juraske et al., 2011; Trapp et al., 2007).

The growth and cultivation of crops invariably display region-specific patterns, contributing to the variability of pesticide bioconcentration in plant tissues. As evidenced by our variability analysis, various environmental factors, including soil properties and weather conditions, yield a significant influence on pesticide and metabolite bioconcentration potentials within plants. This is largely attributed to the temperature's substantial impact on the biotransformation rate of the parent

compound. Consequently, an increase in surrounding temperature can lead to a heightened 204 production rate of metabolites. Moreover, given that soil serves as a primary source of pesticide 205 bioconcentration in plants (Fantke et al., 2011a), its composition can significantly differ across 206 regions. For instance, varying soil types possess distinct organic matter contents that directly 207 influence pesticide absorption in the soil. Soils rich in organic matter yield elevated K_{OC} values 208 209 for lipophilic pesticides, hindering the diffusion kinetics of these compounds from the soil into the tuber. Conversely, diverse pH values in the soil can impact the dissociation process of certain 210 pesticides, particularly those that are ionic or polar in nature (Trapp, 2004). Thus, the uptake of 211 212 such compounds by plants from the soil will manifest region-specific patterns. To conduct a regional assessment of pesticide and metabolite bioconcentration in plants, we recommend that 213 users modify region-specific model input variables, such as K_{OC}, to generate simulation outcomes 214 215 that accurately reflect dynamic environmental conditions across regions.

The comparison between metabolite and non-metabolite models reveals that, under specific 216 217 conditions, such as when plant tissues contain low concentrations of the parent compound or when the biotransformation rate of the parent compound within the plant tissue is notably small, the non-218 219 metabolite model can serve as a viable proxy approach. This enables the simulation of the total toxicity equivalent of both the parent compound and its metabolites, especially when the uptake 220 221 of metabolites from environmental compartments (like soil, air, and plant surfaces) is considered. Given its simplicity and ease of operation, the non-metabolite model is recommended if the 222 aforementioned conditions are met. To ensure this, we suggest performing a preliminary 223 assessment of the degradation kinetics of the parent compound within the plant tissue (Li and 224

Fantke, 2023). Should the degradation rate constant prove to be relatively low when compared to other uptake or elimination rate constants, opting for the non-metabolite model is advisable, which can facilitate uncomplicated modeling experiments.

228 Table S2. Summary of notable plant uptake models with their respective metabolite-based

229 modeling strategies.

Crop examples	Plant uptake models	Potential sources of	Potential sources of	References
		pesticides	pesticide metabolites	
Lettuce	Grass uptake model	Leaf surface; soil	Leaf surface; soil;	(Itoiz et al., 2012;
			plant tissues	Trapp and Matthies, 1995)
Tomato	Herbaceous uptake	Leaf surface; soil;	Leaf surface; soil;	(Juraske et al., 2012,
	model	fruit surface	fruit surface; plant	2007)
			tissues	
Apple	Fruit tree uptake	Leaf surface; soil;	Leaf surface; soil;	(Fantke et al., 2013;
	model	fruit surface; stem	fruit surface; stem	Mendez et al., 2018;
		surface	surface; plant tissues	Trapp, 2007; Trapp et
				al., 2003)
Carrot	Root crop uptake	Leaf surface; soil	Leaf surface; soil;	(Band and King,
	model		plant tissues	2012; Li, 2022;
				Trapp, 2002, 2000)

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