

# Does sustainable agriculture promote biodiversity and yield? A second-order meta-analysis.

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## 33 Abstract

34 Among the biggest challenges of modern society are biodiversity conservation and food  
35 security. Food security requires the increase of agricultural yields, though land use  
36 intensification is one of the main drivers of biodiversity loss. Environmentally friendly farming  
37 practices, such as organic farming, have positive effects on biodiversity, but are accompanied  
38 by yield losses. Other practices, such as diversification, result in a simultaneous increase of  
39 biodiversity and yield. In this study, we quantitatively synthesize the results of multiple meta-  
40 analyses, to identify the impact of sustainable farming practices on the biodiversity-food nexus.  
41 Our results show that sustainable farming practices have a positive effect on biodiversity  
42 without compromising productivity. Notably, when we pooled all meta-analytic means,  
43 biodiversity and yield gains were significantly correlated. In conclusion, sustainable farming  
44 practices have a positive effect on both biodiversity without significant yield losses.

45

46

## 47 Introduction

48 Land use intensification is considered one of the main drivers of biodiversity loss<sup>1-3</sup>, as it  
49 causes habitat loss, grassland degradation, reduction of landscape heterogeneity and  
50 environmental pollution. Fertilizer applications can alter soil composition and nutrient cycling,  
51 pesticides have detrimental impacts on non-target species, often disrupting food webs and  
52 lead to reduction of plant diversity, while intensive tillage disturbs the soil environment. A global  
53 meta-analysis on the effects of intensification on biodiversity and yield estimated that, on  
54 average, conventional intensification leads to a 20% increase in agricultural or silvicultural  
55 yields, but a 9% decrease in species richness<sup>4</sup>. In addition, high management intensity is  
56 associated with lower plant diversity<sup>5</sup> and reduced ecosystem functions<sup>6</sup>. Multiple studies have  
57 highlighted the need for a more sustainable agriculture that ensures food security and hinders  
58 biodiversity loss<sup>3,7-10</sup>.

59 The Target 2 of the Sustainable Development Goals (SDG) is to ensure food security, support  
60 smaller farms and achieve sustainable agriculture, while maintaining biodiversity by 2030  
61 (Goal 2: Zero Hunger. *United Nations*, 2015). However, the adoption of sustainable practices  
62 is impeded by the agro-chemical industry lobby, armed conflicts, climate change, populism  
63 and other socio-political and economic factors<sup>11,12</sup>. One of the main challenges for the adoption  
64 of sustainable farming practices is ensuring high productivity<sup>13</sup>, since sustainable practices are  
65 commonly associated with lower yields. To tackle this issue, ecological intensification has been  
66 proposed as the pathway to sustainable food security<sup>14</sup>. Ecological intensification is the  
67 process of optimizing ecosystem services to either supplement or replacing human-made  
68 inputs (e.g. pesticides, fertilizers), with the goal of sustaining or boosting agricultural  
69 productivity<sup>14,15</sup>.

70 The adoption of sustainable farming practices can lead to three possible cases for biodiversity  
71 and yield outcomes: win-win (biodiversity and yield increase), trade-offs (one increases and  
72 the other one decreases) or lose-lose (biodiversity and yield decrease). At a global scale,  
73 agricultural diversification, for example, is benefitting biodiversity, without compromising

74 yield<sup>16</sup>, but in tropical and sub-tropical regions, it is more likely to result in trade-offs<sup>17</sup>. Agri-  
75 environment schemes (AES) lead, in general, to an increase of insect diversity but a reduction  
76 of yield, whereas the presence of flower strips in orchards has been shown to favour both  
77 insect diversity and yield<sup>18</sup>. Lose-lose outcomes (also known as intensification traps) are  
78 observed mostly in conventional intensification scenarios, when they are implemented in  
79 natural communities<sup>4,19</sup>. In grasslands, spatially optimized management can lead to win-win  
80 outcomes for insect diversity and biomass<sup>20</sup>, emphasizing the high potential of grasslands in  
81 multifunctional landscapes<sup>21</sup>. Grasslands have high economic value<sup>22</sup>, as they contribute to  
82 global productivity (i.e., feed), and they are fundamental for biodiversity conservation<sup>23</sup>. The  
83 outcome, however, is not uniform across systems, biogeographic regions, crop types and  
84 taxonomic groups.

85 In order to reach generalizable conclusions, by surpassing the high heterogeneity between  
86 studies, multiple meta-analyses have synthesized the results of observational and  
87 experimental field studies on the impact of sustainable farming practices on both crop  
88 production and biodiversity<sup>4,17,24</sup>. However, these first order meta-analyses have examined the  
89 impact of either single management practices<sup>25,26</sup> or focus on specific crop systems, such as  
90 vineyards<sup>27,28</sup>.

91 The goal of this study is to systematically analyse the impact of sustainable management  
92 practices on the biodiversity-food nexus and summarize the relevant meta-analytic evidence.  
93 The synthesis of meta-analyses is important to inform policy because it integrates a larger  
94 body of evidence<sup>29</sup>. Second-order meta-analysis is a method that combines the outcomes of  
95 multiple first order meta-analyses using statistical models, while accounting for variation  
96 among them<sup>30</sup>. This method has been adopted only recently in the field of Ecology, as a result  
97 of the increasing number of first-order meta-analyses published in the last decade and the  
98 need for higher level of evidence synthesis. In the present study, second-order meta-analyses  
99 allowed for a nuanced understanding of how different management practices influence both

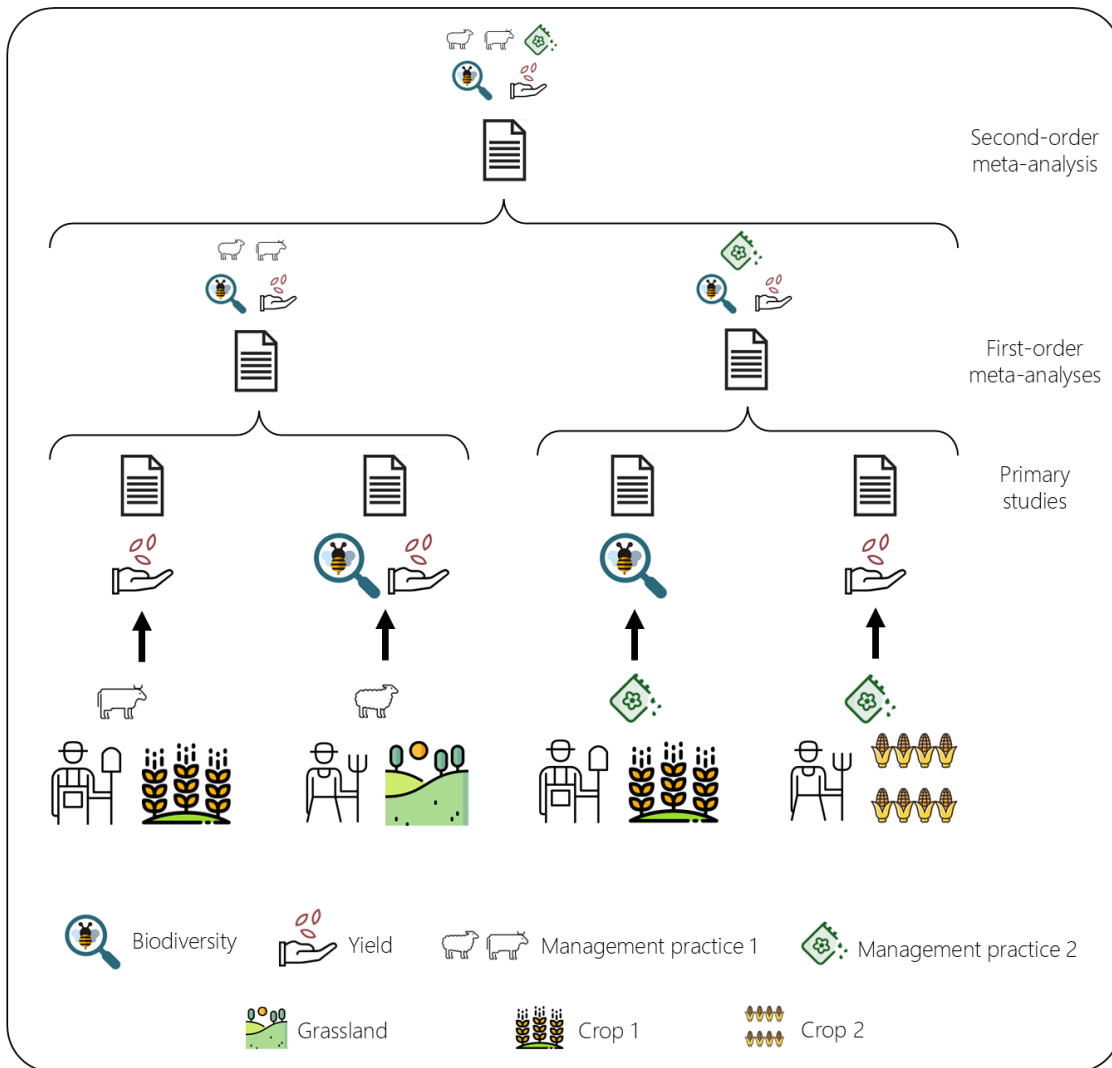
100 biodiversity and productivity across various ecosystems. In this paper, we examine the  
101 following research questions:

- 102 1) What is the impact of sustainable farming practices on both biodiversity and yield?
- 103 2) Which management practices lead to win-win scenarios for biodiversity and yield?
- 104 3) Which taxa and crop systems are benefited the most by specific agricultural  
105 management practices?

106 We conducted a systematic review to identify meta-analyses focusing on both biodiversity and  
107 agricultural production as response variables, following specific inclusion criteria outlined in  
108 [Takola et al. \(2023\)](#)<sup>31</sup>. The study material consists of studies which examined management  
109 effects on croplands, grasslands, and agroforestry systems. We conducted a second-order  
110 meta-analysis, combining multiple first-order meta-analyses to quantify the effects of various  
111 agricultural practices on biodiversity and yield (Fig. 1). We extracted overall meta-analytic  
112 means and subgroup estimates, enabling us to perform meta-regressions with moderators  
113 and analyze subgroups. We provide a comprehensive overview of the body of literature that  
114 examines the interplay of biodiversity and yield in productive ecosystems and quantify the  
115 impact of sustainable agricultural practices.

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118

119 **Figure 1.** The study framework of the present paper; from farm to our second-order meta-analysis.

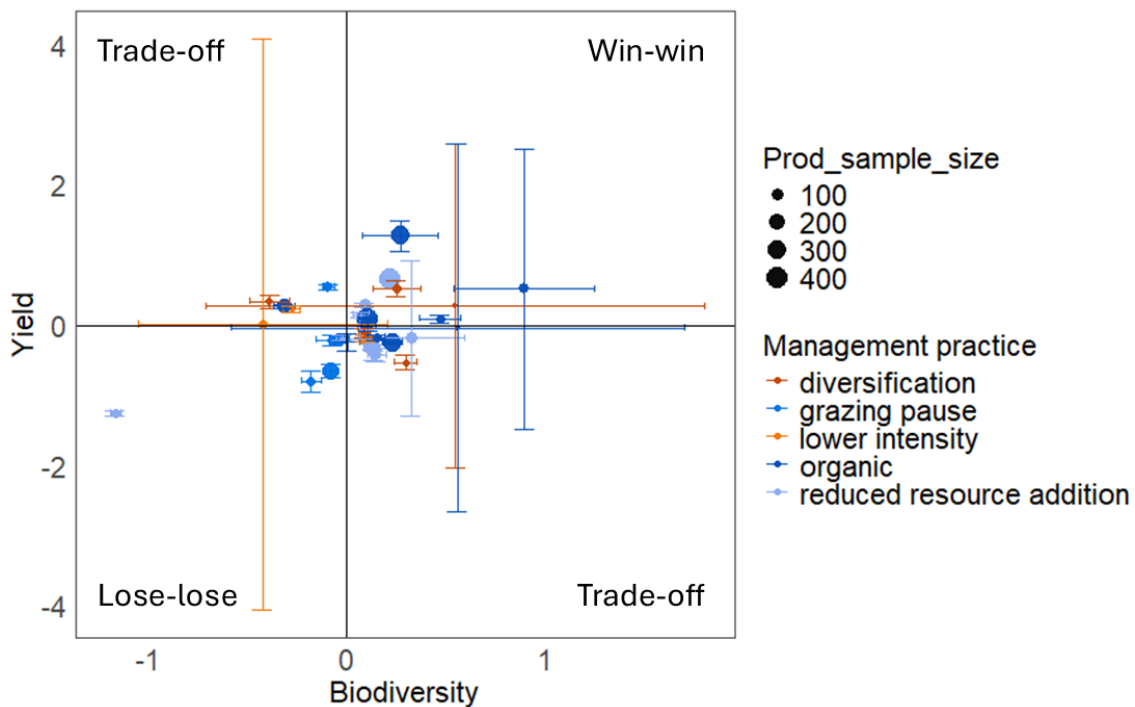
120

121 **Results**

122 Our systematic review yielded 27 meta-analyses. From these, we used 22 for the second-  
123 order meta-analysis (ca. 2,000 primary studies), which provided 41 pairs of overall means for  
124 biodiversity and yield. We also extracted subgroup estimates for taxonomic groups and crop  
125 systems, from 21 meta-analyses. The subgroup datasets contained 298 estimates for  
126 biodiversity and 51 for yield. All data are available online ([here](#)).

127 **Trade-off analysis**

128 In order to examine the relationship of biodiversity and yield within the context of sustainable  
129 farming, we plotted all the meta-analytic means in a scenario space (Fig. 2).



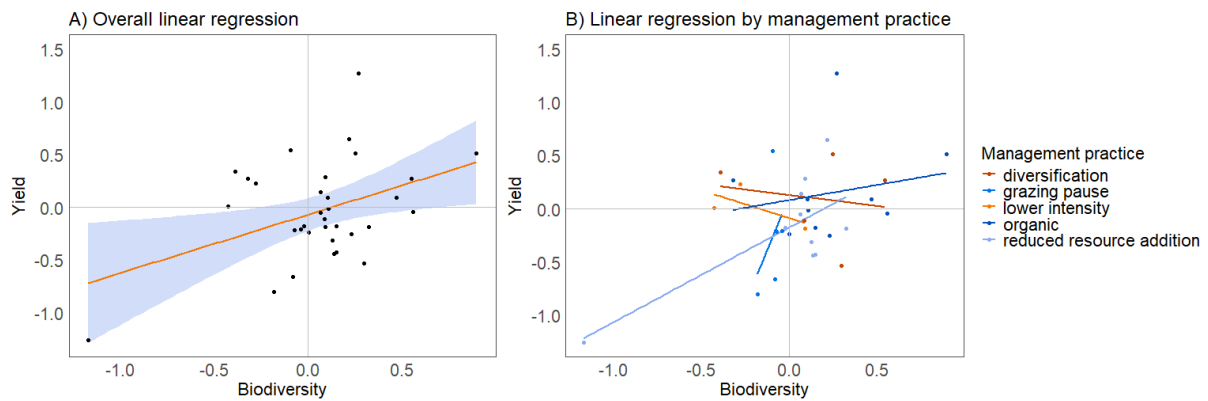
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131 **Figure 2.** Cartesian plane showing a scenario space. The top-right quadrant includes the win-win cases,  
132 bottom-left quadrant includes the lose-lose cases and the remaining two are showing trade-offs. Points  
133 represent the overall meta-analytic means for biodiversity and yields, as reported in each meta-analysis  
134 ( $n = 22$ ). Horizontal and vertical error bars represent the 95% C.I. for biodiversity and yield respectively.  
135 Colours represent management practices. Points are scaled based on the number of primary studies  
136 that was used for each meta-analytic mean.

137

138 We found that, based on 22 meta-analyses, the effect of sustainable farming on biodiversity  
139 is positively correlated to the effect of sustainable farming on yield (slope = 0.55,  $SE_{\text{slope}} =$   
140 0.216,  $p = 0.015$ ). Though, when fitting a linear regression for each management practice  
141 separately, some relationships were negative (Fig. 3).

142



143

144 **Figure 3.** A) Overall linear regression and B) Linear regression by management practice of the log  
145 response ratios (LRR) for yield and biodiversity, as reported in 22 meta-analyses included in our study.

146

### 147 Win-win scenarios for biodiversity and yield

148 We examined separately the meta-analyses that reported positive values for both biodiversity  
149 and yield (Table 1). In total, these were 11 meta-analyses which contributed 14 effect sizes to  
150 our dataset. From these, 13 effect sizes were used in the second-order meta-analysis,  
151 because they were reported as log-response ratio (LRR) or some other metric that we could  
152 convert to log-response ratio (Table S1). Two meta-analyses were focusing on invertebrates,  
153 one was focusing on plants and one was focusing on animals. Ten meta-analyses were  
154 focusing on various species groups. Regarding the metrics of biodiversity, one meta-analysis  
155 was reporting abundance, one was focusing on biocontrol species, one was not reporting the  
156 metric that was used and eight meta-analyses were reporting species richness. Regarding the  
157 metrics of productivity, five meta-analyses were reporting crop yields, another five meta-  
158 analyses were reporting biomass, and one meta-analysis was reporting productivity. Six meta-



159 analyses were using primary studies with paired control and treatment measurements of  
 160 biodiversity or yield and five meta-analyses were using unpaired measurements. Five meta-  
 161 analyses were including non-paired measurements of biodiversity and yield (i.e. measured in  
 162 the same location), while six meta-analyses were included paired measurements of  
 163 biodiversity and yield or a mix of paired and non-paired. Two meta-analyses were examining  
 164 below-ground biodiversity, eight meta-analyses were examining above-ground biodiversity  
 165 and one meta-analysis both. One meta-analysis was focusing on grapes, one on cocoa  
 166 plantations, one on herbs, three on various crops and four were referring to grasslands.

167

Crop type	Reference	Win-win scenario	Sample size
Meadow, forest, grassland	Ma <i>et al.</i> (2020)	Reduced resource addition (nitrogen, phosphorus and both) leads to an increase of Shannon's Index of arbuscular mycorrhizal fungi (AMF) and the biomass of above-ground plants in alpine meadows, forests and grasslands.	Biodiversity: 127 Yield: 100
Agroforestry	Neither <i>et al.</i> (2020)	When compared to cocoa monoculture, cocoa agroforestry schemes have more animal species richness and higher yields.	Biodiversity: 5 Yield: 8
Agroforestry	Winter <i>et al.</i> (2018)	Organic farming increases all species richness (plants, insects and birds) and yields in vineyards.	Biodiversity: 24 Yield: 45
Cropland	Bai <i>et al.</i> (2018)	When compared to monocultures, crop rotations result to higher abundance of earthworms and higher yields.	Biodiversity: 2 Yield: 14
Cropland	Iverson <i>et al.</i> (2014)	Biocontrol invertebrate species and crop yields are higher in crop rotation schemes, than in monocultures.	Biodiversity: 54 Yield: 39
Cropland	Bai <i>et al.</i> (2018)	When adding organic matter to crops, the abundance of earthworms and the crop's yield are higher.	Biodiversity: 6 Yield: 54
Cropland	Shu <i>et al.</i> (2022)	Microbial species richness and crop yield are increasing when using organic fertilizer instead of mineral fertilizer in agricultural crops.	Biodiversity: 484 Yield: 379
Grassland	Guldmond <i>et al.</i> (2017)	When elephants are removed (through fences) from grasslands, there is an increase of plant and animal species richness diversity and an increase of herb and tree abundance.	Biodiversity: 18 Yield: 132
Grassland	Su <i>et al.</i> (2022)	Reduced resource addition increases species richness and biomass of plants in grasslands.	Biodiversity: 12 Yield: 35
Grassland	Ploughe <i>et al.</i> (2020)	The addition of biosolids increases plant species richness and productivity in grasslands.	Biodiversity: 159 Yield: 269
Grassland	Wang <i>et al.</i> (2020)	Reduced resource addition (nitrogen, phosphorus and both) leads to an increase of plant species richness and the above-ground biomass in grasslands.	Biodiversity: 133 Yield: 412

168 **Table 1.** Win-win scenarios in the database identified in the current study.

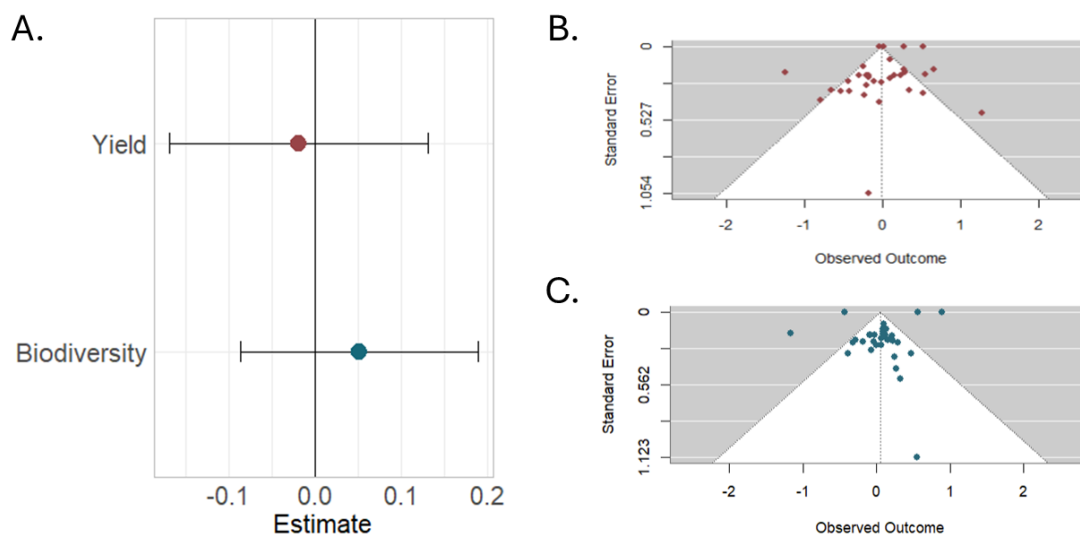
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170 **Second-order meta-analysis**

171 *Overall estimates on the impact of sustainable management on biodiversity and yield*

172 The random effects model of the second order meta-analysis showed that sustainable farming  
173 has an overall positive effect on biodiversity but neutral on yield (Table S2, Fig. 4). Meta-  
174 analyses examining reduced resource addition, organic farming and grazing pause showed  
175 positive relationships between biodiversity and yield, while lower land use intensity and  
176 diversification showed a negative relationship (Figure 3).

177



178

179 **Figure 4.** (A) Overall means of random effects models for biodiversity and yield, (B) Funnel plot for  
180 biodiversity means of first-order meta-analyses, (C) Funnel plot for yield means of first-order meta-  
181 analyses.

182

183 *Meta-regressions to examine the impact of different management practices*

184 To examine the effect of sustainable management on biodiversity and yield, we performed  
185 meta-regressions using management practices as a moderator. Overall, diversification and  
186 organic farming lead to win-win outcomes for biodiversity and yield, while grazing pause, lower  
187 intensity and reduced resource addition have moderately negative effects on the two response  
188 variables (Table 2). However, none of the aforementioned estimates was significant.

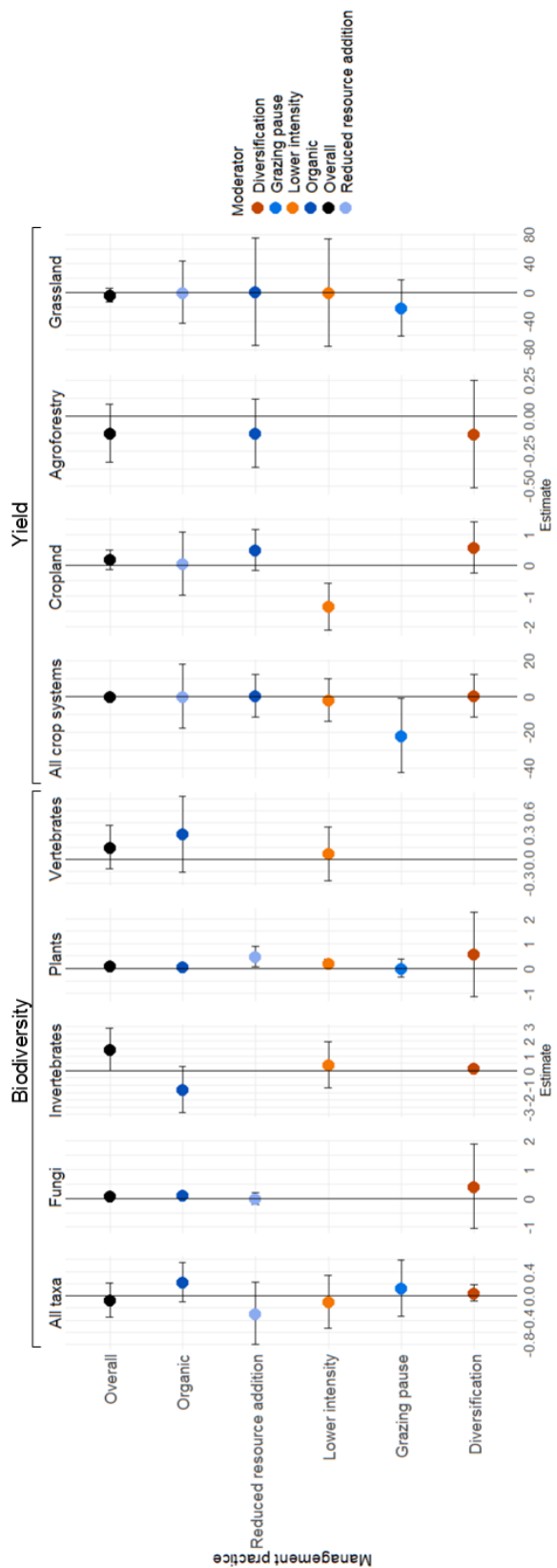
Response variable	Moderator	Estimate	CI low	CI up
Biodiversity	Diversification	0.13	-0.33	0.59
	Grazing pause	-0.09	-0.50	0.34
	Lower intensity	-0.29	-0.80	-0.22
	Organic	0.23	-0.09	0.55
	Reduced resource addition	-0.06	-0.35	0.22
Yield	Diversification	0.56	-11.23	12.42
	Grazing pause	-21.61	-42.1	-1.12
	Lower intensity	-1.72	-13.6	10.15
	Organic	0.56	-11.23	12.35
	Reduced resource addition	0.23	-17.66	18.13

190 **Table 2.** Moderator estimates of each management practice on the overall meta-analytic estimates  
 191 biodiversity and yield.

192

193 We fitted a meta-regression with yield as a response variable and biodiversity as a moderator.  
 194 The moderator estimate was positive and statistically significant (estimate = 0.42, 95% C.I.:  
 195 [0.11, 0.72], SE = 0.16, z-value = 2.66, p-value = 0.008).

196 Overall, diversified farming had positive effects on the diversity of all taxa but this was not the  
 197 case for crop systems (Table S3, S4). However, the meta-analytic models that we fitted for  
 198 each taxonomic group showed positive meta-analytic means for all taxa (Table 2). Regarding  
 199 crop types, the meta-analytic means were positive for productivity of agroforestry schemes,  
 200 but negative for croplands and grasslands. Reduced resource addition had either a positive  
 201 or neutral effect on all taxa and crop systems, while lower management intensity, similarly to  
 202 grazing pause, had a negative or neutral effect on biodiversity and yield (Fig. 5).



203

204 **Figure 5.** Moderator estimates of different management practices for meta-regression models per taxon  
 205 and crop system subgroup. The category Overall refers to the estimates of the random-only second  
 206 order meta-analytic models

207

208 *Subgroup analysis; taxa and crop systems*

209 The overall meta-analytic means for each subgroup of taxonomic groups and crop systems  
210 showed that overall, sustainable farming practices are associated with increased biodiversity  
211 and decreased yields in croplands and grasslands, but increased yield in agroforestry  
212 schemes (Table 3).

213

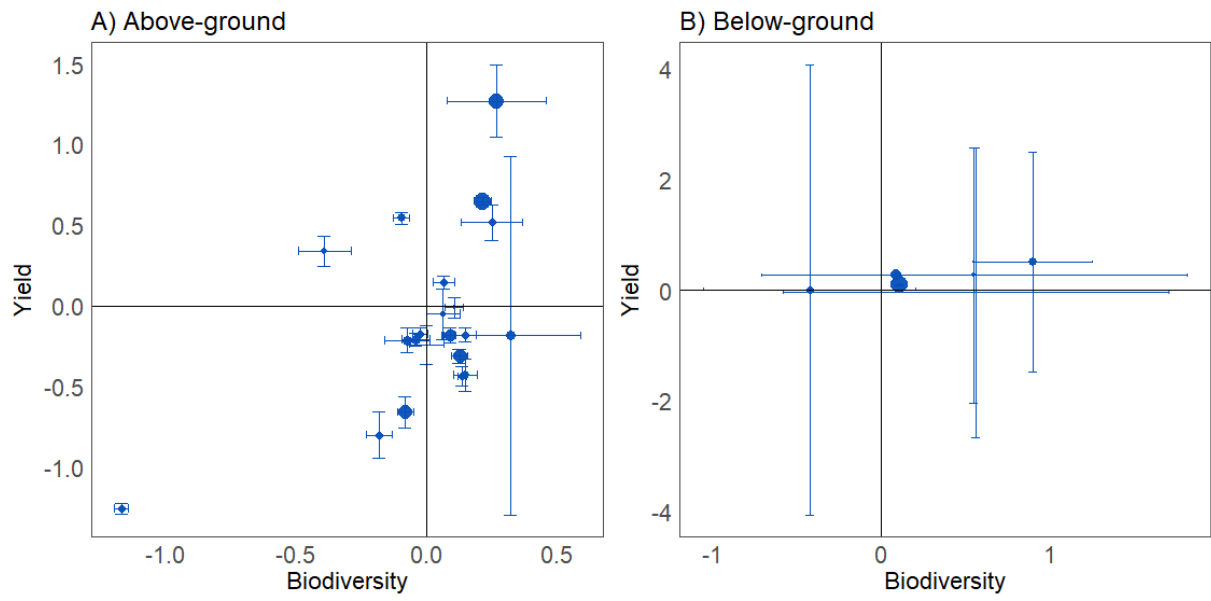
Response variable	Subgroup	Estimate	CI low	CI up	Prediction Interval low	Prediction Interval up
Biodiversity	Invertebrates	0.16	-0.005	0.33	-0.005	0.33
	Plants	0.11	0.02	0.19	0.02	0.19
	Fungi	0.09	0.01	0.16	0.01	0.16
	Vertebrates	0.15	-0.12	0.42	-0.12	0.42
Yield	Cropland	0.19	-0.13	0.51	-1.03	1.41
	Agroforestry	-0.12	-0.33	0.08	-0.33	0.08
	Grassland	-3.87	-13.10	5.36	-46.18	38.44

214 **Table 3.** *Meta-analytic means of random-only effect models for all subgroups of taxa and crop systems.*

215

216 Interestingly, we observed different patterns when separating the data into above- and below-ground  
217 biodiversity and productivity. Some studies did not distinguish between the two groups, so they were  
218 excluded. Above-ground effect sizes show significantly higher heterogeneity (based on the Q-test),  
219 while below-ground effect sizes have wide confidence intervals (Fig. 6).

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**Figure 6.** Visualization of effect sizes referring to (A) Above-ground and (B) Below-ground biodiversity and productivity.

## 224 Discussion

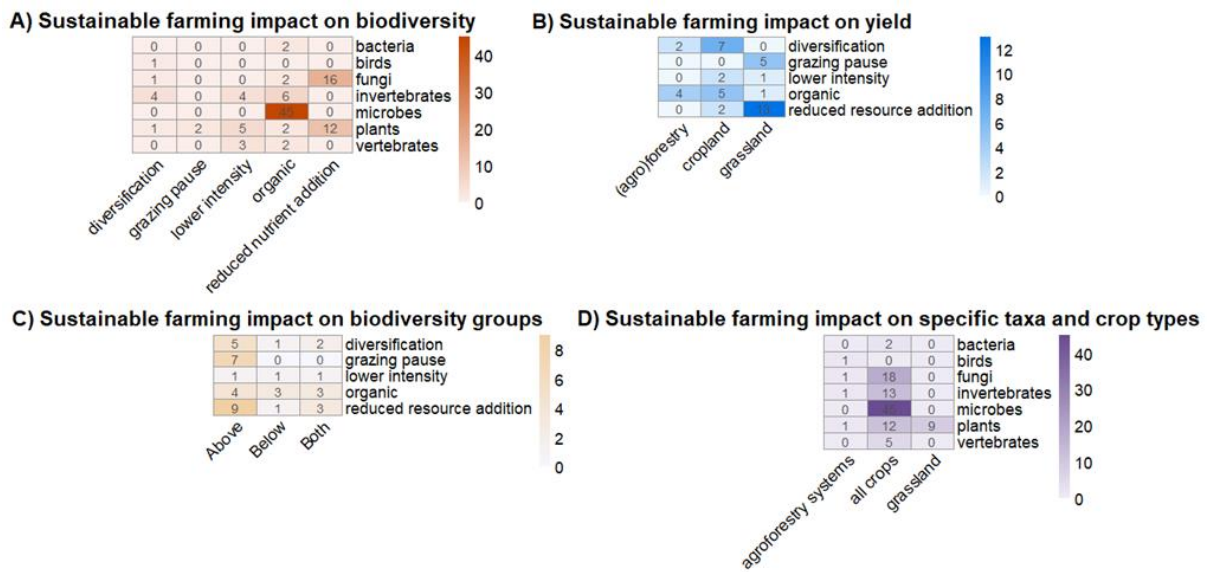
225 We conducted a systematic review and a second-order meta-analysis of ca. 2,000 primary  
226 studies (22 meta-analyses), to summarize the effect of sustainable farming on both  
227 biodiversity and yield. Birds, other vertebrates and trees are the most underrepresented  
228 taxonomic groups in our meta-analyses database (Fig. 7). This does not mean that there are  
229 no relevant studies, but it means that these taxa are rarely studied jointly with yield outcomes.

230

### 231 Biodiversity is mediating yield gains in sustainable farming

232 Our second-order meta-analysis on sustainable farming practices showed that there was a  
233 positive relationship between the change in biodiversity and yield caused by different  
234 sustainable farming practices. This relationship has been described in literature<sup>32</sup>, but it is  
235 highly context-dependent. Indeed, our analysis showed that this effect is not uniform across  
236 taxa<sup>33</sup> and in our dataset we even discerned above- and below-ground differences. Organic  
237 farming practices, such as the addition of biosolids, organic matter and the replacement of  
238 mineral fertilizers with organic fertilizers, as well as diversified farming, were associated with  
239 an increase in biodiversity and yield. For example, nutrient enrichment increases invertebrate  
240 herbivory and pathogen damage in grasslands<sup>34</sup> and it can lead to eutrophication. Reduced  
241 resource addition and lower intensity do not lead to win-win outcomes, possibly due to slower  
242 processes of ecosystem restoration after the removal of the disturbances.

243



244

245 **Figure 7.** Heatmaps of frequencies representing (A) taxonomic groups per management practice, (B)  
 246 crop types per crop systems, (C) biodiversity groups per management practice, (D) crop types per  
 247 taxonomic groups, based on the dataset of the present study.

248

249 The relationship between biodiversity and yield within the context of sustainable farming is  
 250 complex and multi-factorial<sup>35,36</sup>, thus we cannot assume causality in every aspect. There are  
 251 different underlying mechanisms that can be best studied through well-designed long-term  
 252 experiments rather than through second-order meta-analyses. Although multiple meta-  
 253 analyses study sustainable farming practices, there is a lack of evidence i) on the effectiveness  
 254 of the existing yield enhancement measures in relationship to biodiversity and ii) on the  
 255 specific ways this relationship is mediated by the structure of the surrounding landscape.

256 The analysis of taxonomic and crop system subgroups resulted in both positive and negative  
 257 effects. For instance, organic farming having either positive or negative results for different  
 258 subgroups and a negative effect on productivity for all crop systems, except for agroforestry  
 259 schemes. Although the test for heterogeneity (Q) was not significant for most subgroups, we  
 260 performed meta-regressions aiming at examining the effect of management practices. We  
 261 found that diversification positively affects biodiversity while having a neutral effect on yield.  
 262 Additionally, within the context of diversification, there is a negative relationship between yield



263 and biodiversity. These findings are in accordance with findings from previous meta-  
264 analyses<sup>16,17</sup>, that focused on the impact of diversification.

265 Translating international commitments (i.e. SDGs and COP21 INDCs) into actionable plans  
266 requires a focus on environmental contexts due to the highly context-dependent nature of  
267 sustainable farming success. Significant knowledge gaps remain regarding farm size and  
268 landscape heterogeneity, both crucial for assessing spillover effects. Smaller farms tend to  
269 have higher yields and species richness<sup>37</sup>, while landscape heterogeneity is known to enhance  
270 biodiversity<sup>38</sup>. However, the influence of surrounding landscapes on yields is complex and  
271 context-dependent. Additionally, these factors contribute to the adoption of agri-environmental  
272 schemes<sup>39</sup> (AES). Developing policies that address contextual challenges and are tailored to  
273 fit national, regional or even local scales could be supported by spatially-explicit typologies  
274 that capture archetypal patterns of agri-environmental systems<sup>40</sup>.

275 Based on our dataset, win-win scenarios are more likely to be achieved through the following  
276 actions: i) reduced addition of nutrients in grasslands is benefitting fungi and plant diversity  
277 and biomass production ii) organic farming (such as addition of organic matter, addition of  
278 biosolids and substituting mineral with organic fertilizers) is promoting species richness of  
279 plants, invertebrates, vertebrates and microbes as well as productivity in vineyards, crops and  
280 grasslands iii) crop rotation is increasing the richness of invertebrates (biocontrol species and  
281 earthworms) and yields iv) cocoa agroforestry schemes are hosting more animals and higher  
282 yields than cocoa monocultures v) removal of large herbivores (i.e. elephants) from grasslands  
283 is improving plant and animal richness while increasing herb and tree abundance. More  
284 detailed and targeted studies across different crop systems will help to shed light on the  
285 synergies between biodiversity and productivity. We don't just "need more data", though. What  
286 we need is coordinated global efforts of carefully designed long-term experiments, which can  
287 then be used in evidence syntheses and causal inference<sup>41</sup>, with the aim to understand  
288 processes, instead of having to conform to "snapshots"<sup>42-44</sup>. When coupled with simulations  
289 and models, carefully designed experiments can inform us very well<sup>45</sup>. There has been a long

290 history of efforts to reverse biodiversity loss<sup>46</sup>. Conclusions from the Biodiversity Strategy for  
291 2030 and COP16 re-iterated the need for more political will and legally binding legislation as  
292 well as adequate funding<sup>47</sup>, because it is linked to motivations for farmers<sup>48</sup>.

293

## 294 Methods

### 295 Literature search and systematic review

296 We systematically screened literature (Fig. S1) in order to identify meta-analyses that examine  
297 the effect of an agricultural management practice on biodiversity and yield. We leveraged  
298 references from three published studies<sup>16,49,50</sup> and Web of Science.

299 The query we used to search Web of Science was:

300 *(diversity OR species richness OR biodiversity OR (taxonomic AND richness) OR (abundance\* AND species) OR even\*ess OR*  
301 *shannon OR simpson) AND (provisioning OR producti\* OR food OR fodder OR feed OR fibre OR logg\* OR fuel OR commodit\**  
302 *OR harvest\* OR wood OR timber OR coffee OR cacao OR crop\* OR yield\* OR oil OR abundanc\* OR biomass\*) AND (diversifi\**  
303 *OR intensif\* OR fertili\* OR nutrient\* OR organic OR manag\* OR pest\* OR insectic\* OR graz\*) AND ("meta analys\*" OR "meta-*  
304 *analys\*" OR "metaanalys\*")*

305 The inclusion criteria were: i) the study had to be a first-order meta-analysis ii) the meta-  
306 analysis had to use biodiversity and yield as response variables iii) the meta-analysis had to  
307 examine the effect of some management practice on the aforementioned response variables.  
308 We did not restrict the studies regarding language, time period or geographical area. We  
309 identified 27 meta-analyses and extracted all reported results (overall estimates and subgroup  
310 estimates). We used *metaDigitise*<sup>51</sup> to extract data from figures, whenever necessary. All  
311 analyses were conducted in R.

### 312 Data grouping and pre-processing

313 We grouped management practices in five categories (Table 4). All categories have been  
314 (re)phrased in order to ensure that the sustainable practice (or lower management intensity)  
315 is the treatment, while the intensive management is the control (see below for statistical  
316 explanation).

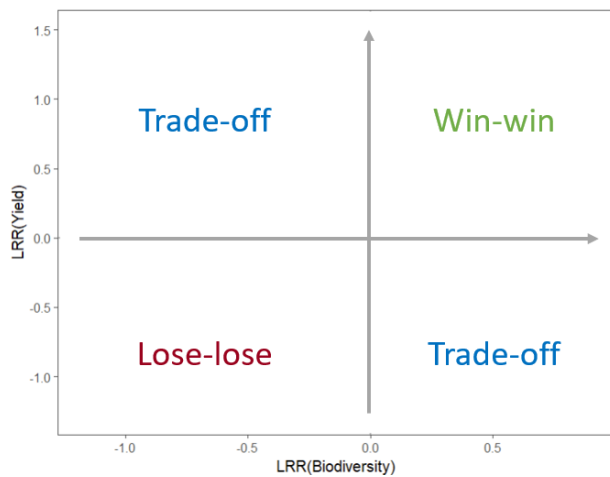
Sustainable Management Group (second-order analysis)	First-order groups of practices (first-order meta-analysis)	Crop system
Diversification	Monocultures vs. Polycultures Monoculture vs. Agroforestry Bare Soil vs. Cover Crops Agriculture vs. Agroforestry Monoculture vs. Crop rotation Monoculture vs. Polyculture Bare/Unmanaged Edges vs. Edge Plantings	Cropland, agroforestry
Grazing management	Grazing vs. No grazing Other grazing regimes vs. Communal grazing Continuous grazing vs. Strategic grazing Strategic grazing vs. No grazing	Grassland
Lower intensity	Mowed field vs. Abandoned field Intensification vs. No intensification Tillage vs. No tillage	Cropland, grassland
Organic	Conventional vs. Organic agriculture No Biosolids vs. Biosolids No organic matter vs. Organic matter Conventional vs. Mixed organic Mixed organic vs. Pure organic Mineral-only fertilizer vs. Organic fertilizer Intensive management vs. Organic agriculture	Cropland, grassland, agroforestry
Reduced addition	Nitrogen vs. No nitrogen Phosphorus vs. No phosphorus Nitrogen and Phosphorus vs. No Nitrogen and Phosphorus Water addition vs. No water addition	Cropland, grassland

318 **Table 4.** Grouping of management practices.

319

## 320 Trade-off analysis

321 We plotted the reported meta-analytic results in a scenario space, which is depicted as a  
 322 Cartesian plane with biodiversity in the x axis and yield in the y axis. This scenario space (Fig.  
 323 8) allows further quantitative evidence synthesis and the exploration of general patterns,  
 324 potentially identifying synergies and trade-offs.



325

326 **Figure 8.** Cartesian plot showing the scenario space for the present study.

327

328 We visualized the results of meta-analyses in a Cartesian plot representing a scenario space.  
 329 Every pair of yield and biodiversity overall meta-analytic means (reported as log-response  
 330 ratio or some other convertible metric) and confidence intervals was placed in this scenario  
 331 space. We then fitted a regression line through the points to examine the overall relationship  
 332 between yield and biodiversity.

333 We also created a “win-win table”, which is a tabular representation of the effect sizes that  
 334 were located in the top-right panel of the scenario space. We then fitted a regression line on  
 335 these values, to assess the effect of biodiversity on yield change in the context of sustainable  
 336 farming.

337

### 338 Effect size calculations

339 It is common practice for meta-analyses to convert log response ratios (LRR) to percentages  
 340 of change, to facilitate interpretation. We converted percentages and response ratios to log  
 341 response ratios (LRRs).

342 We standardized all effect sizes based on the reference scenario of each meta-analysis. Some  
 343 meta-analyses used the intensified management as an intervention, while other meta-

344 analyses used the sustainable management as the intervention. For studies that reported the  
345 intensified management as an intervention, we used the inverse of the log response ratio (by  
346 inverting the control and treatment in the LRR equation). We did the same for confidence  
347 intervals but not with standard errors and standard deviations.

348

## 349 **Second-order meta-analysis**

350 We conducted the second-order meta-analysis using first-order meta-analyses that used log  
351 response ratio (LRR) as an effect size metric. We excluded studies reporting Hedges' g,  
352 Hedges' d, or Cohen's D, because they cannot be converted in LRR. We analyzed log-  
353 response ratio data using a multi-level meta-analytic model and meta-regressions without an  
354 intercept (with the `rma.mv()` and `rma()` functions from the R package `metafor`) with random  
355 and fixed effects. As random effects we used the effect size and study ID. As fixed effects we  
356 used management practice (grouping is explained in Table), taxonomic group, crop type and  
357 geographic region. We also fitted a meta-regression with biodiversity as a predictor and yield  
358 as a response variable.

359 We analysed separately the taxonomic groups that had enough effect sizes (fungi, insects,  
360 vertebrates and plants), as well as the three crop systems (cropland, grassland, agroforestry).  
361 In our study, the taxonomic group of plants is not referring to the plants of the crop, but wild  
362 plant communities that are related to the crop (e.g. the understory of an agroforestry scheme).  
363 An overview of the models used is provided in Tab S5.

364

## 365 **Statistical independence of meta-analyses**

366 First and second order meta-analyses assume independence among studies. Non-  
367 independence can be addressed by fitting a correlation matrix (R) in the meta-analytic model.  
368 This practice is common in phylogenetic meta-analyses<sup>52</sup>. In our dataset, some meta-analyses  
369 used the same primary studies in their sample size, therefore the assumption of independent

370 samples in meta-analytic models was violated. In order to account for non-independence  
371 among individual meta-analyses, we fitted a correlation matrix with the percentage of overlap  
372 among primary studies from each pair of meta-analyses.

373 We calculated the percentage of overlap for each pair of meta-analyses using the formula:

374

$$375 \quad \text{Percentage overlap} = \frac{C}{N_i + N_j - C} * 100$$

376 where

377  $N_i, N_j$ : are the total number of primary studies included in meta-analysis i and j respectively

378  $C$ : is the number of primary studies that  $N_i$  and  $N_j$  have in common

379

### 380 *Heterogeneity analysis*

381 Due to the inherent differences between studies in the ecological literature, true effect sizes  
382 were presumed to differ between studies.

383 In order to test for heterogeneity between studies, the weighted sum of squares,  $Q$ , was  
384 calculated as:

$$385 \quad Q = \sum_{f=1}^k W_f g^2 - \frac{\left( \sum_{f=1}^k W_f g \right)^2}{\sum_{f=1}^k W_f}$$

386 If the  $Q$  statistic is significant, then the meta-analyst should proceed with meta-regressions, to  
387 explore the sources of heterogeneity.

388 We calculated the amount of heterogeneity for each dataset and subgroup using the I-squared  
389 ( $I^2$ ) statistic:

$$390 \quad I^2 = 100\% * \frac{Q - df}{Q}$$

391  $I^2$  is a metric of variability between studies and it was calculated as:

$$392 \quad I^2 = \frac{Q - df}{C}$$

393 where

394  $df = \text{number of independent datasets} - 1$

$$395 \quad C = \sum W_f - \frac{W_f^2}{W_f}$$

395

396

397

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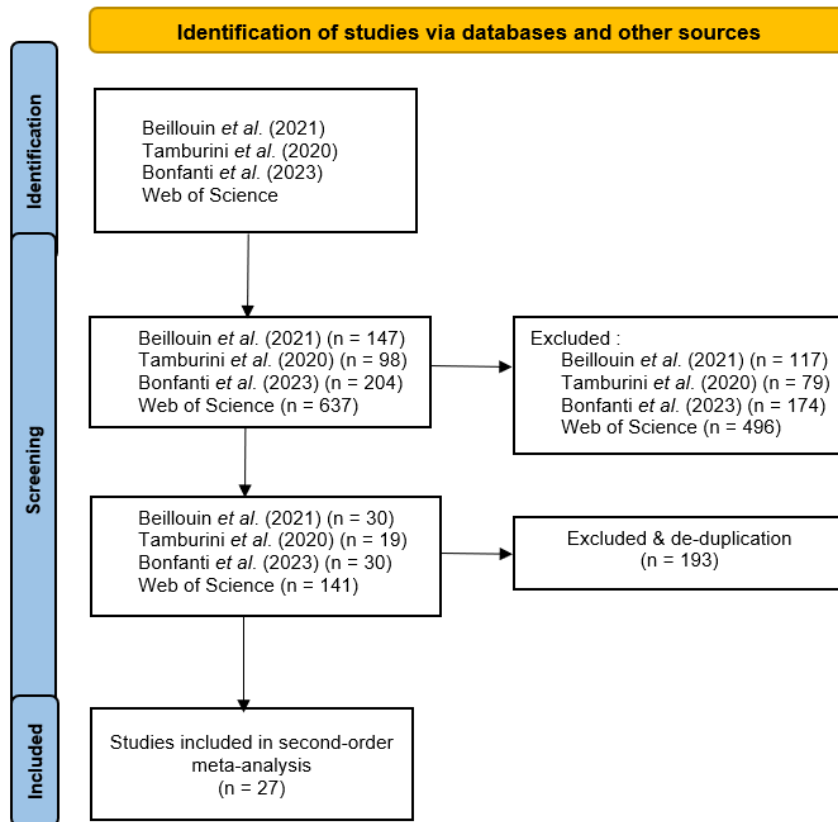


Table S1. Data for the studies that reported win-win outcomes for biodiversity and yield.

No.	Reference	Title	Year	DOI	StudyID	ES_ID	Taxonomic_group	Scenario_raw	Biodiv_ES	Biodiv_ES_Metric	Biodiv_messure	Biodiv_size	Management_group	Prod_ES	Prod_ES_Shem	ControlT	LandSH	BiodivYieldPair	CropType	Sustain	AboveB	Biodiv_C	Biodiv_CProd_C	Prod_CI					
1	Ma et al. (2020)	Global negative effects of nutrient enrichment on arbuscular mycorrhizal fungi, plant diversity and ecosystem multifunctionality	2020	<a href="https://doi.org/10.1111/nph.17077">https://doi.org/10.1111/nph.17077</a>	FEB23_13_2	E5005	fungi	No nutrient enrichment	-0.09	LRR	Sh	127	reduced resource addition	-0.29	LRR	biomass	100	Y	Sh	Y	all crops	N	Below	0.09	0.06	0.12	0.29	0.22	0.35
2	Niether et al. (2020)	Cocoa agroforestry systems versus monocultures: a multi-dimensional meta-analysis	2020	<a href="https://doi.org/10.1088/1748-1748/39326/abb005">https://doi.org/10.1088/1748-1748/39326/abb005</a>	FEB23_22_4	E5011	animals	Monocult ure vs Agroforestry	4.67	HedgesG	SR	5	diversification	9.06	HedgesG	yield	8	Y	Sh	N	cocoa	Y	Above	4.67	8.13	1.22	9.06	30.49	-12.37
3	Guidemond et al. (2017)	A systematic review of elephant impact across Africa	2017	<a href="https://doi.org/10.1371/journal.pone.0177893">https://doi.org/10.1371/journal.pone.0177893</a>	FEB23_30_3	E5020	all	Presence vs absence of elephants from plots	-0.18	CohensD	Diversity	18	grazing pause	-0.48	Cohen abund	ance	132	Y	Sp	Yeome	herbs/trees	N	Above	0.18	-2.22	2.58	0.48	0.28	0.68
4	Su et al. (2022)	Multiple global changes drive grassland productivity and stability: A meta-analysis	2022	<a href="https://doi.org/10.1111/1365-2745.13983">https://doi.org/10.1111/1365-2745.13983</a>	FEB23_50_2	E5030	plants	fertilized vs non-fertilized	-6.57	Percentage change	SR	12	reduced resource addition	-13.48	Percentage change	biomass	35	Y	Sh	Yeome	grassland	N	Above	0.07	0.00	0.14	0.14	0.21	21.35
5	Wang et al. (2020)	Effects of nutrient addition on degraded alpine grasslands of the Qinghai-Tibetan Plateau: A meta-analysis	2020	<a href="https://doi.org/10.1016/j.agee.2020.106870">https://doi.org/10.1016/j.agee.2020.106870</a>	FEB23_59_7	E5036	all	no nutrient addition	-0.22	LRR	SR	133	reduced resource addition	-0.65	RR++	biomass	412	Y	Sh	N	all crops	N	Above	0.22	0.15	0.28	0.65	0.60	0.70
6	Ploughe et al. (2020)	Revegetation of degraded ecosystems into grasslands using bioislands as an organic amendment: A meta-analysis	2020	<a href="https://doi.org/10.1111/1365-2745.12558">https://doi.org/10.1111/1365-2745.12558</a>	FEB23_95_2	E5040	all	Bioislands vs no bioislands	0.27	LRR	SR	159	organic	1.27	LRR	productivity	269	N	Sh	Yeome	grassland	Y	Above	0.27	0.65	-0.10	1.27	1.71	0.83
7	Bai et al. (2018)	Effects of agricultural management practices on soil quality: A review of long-term experiments for Europe and China	2018	<a href="https://doi.org/10.1016/j.agee.2018.05.028">https://doi.org/10.1016/j.agee.2018.05.028</a>	JB005	E5043	invertebrates	crop rotation vs no crop rotation	1.73	RR	Ab	2	diversification	1.31	RR	yield	14	Y	Sh	N	all crops	Y	Below	0.55	8.65	-5.19	0.27	1.50	1.12
8	Bai et al. (2018)	Effects of agricultural management practices on soil quality: A review of long-term experiments for Europe and China	2018	<a href="https://doi.org/10.1016/j.agee.2018.05.028">https://doi.org/10.1016/j.agee.2018.05.028</a>	JB005	E5041	invertebrates	organic matter input vs no input	2.45	RR	Ab	6	organic	1.67	RR	yield	54	Y	Sh	N	all crops	Y	Below	0.90	3.82	1.08	0.51	1.94	1.40
9	Iverson et al. (2014)	Do polycultures promote win-wins or trade-offs in agricultural ecosystem services? A meta-analysis	2014	<a href="https://doi.org/10.1111/1365-2684.12334">https://doi.org/10.1111/1365-2684.12334</a>	JB073	E5045	invertebrates	polyculture vs monoculture	0.25	LRR	Biocontrol	54	diversification	0.52	LRR	yield	39	Y	Sh	Y	all crops	Y	Above	0.25	0.49	0.02	0.52	0.74	0.30
10	Shu et al. (2022)	Organic amendments enhance soil microbial diversity, microbial functionality and crop yields: A meta-analysis	2022	<a href="https://doi.org/10.1016/j.scirev.2022.154627">https://doi.org/10.1016/j.scirev.2022.154627</a>	JB158	E5051	all	no treatment vs organic amendment	0.11	LRR	SR	484	organic	0.09	LRR	yield	379	N	Sh	Yeome	all crops	Y	Below	0.11	0.12	0.09	0.09	0.11	0.08
11	Winer et al. (2015)	Effects of vegetation management intensity on biodiversity and ecosystem services in vineyards: A meta-analysis	2015	<a href="https://doi.org/10.1111/1365-2684.13124">https://doi.org/10.1111/1365-2684.13124</a>	JB188	E5053	all	conventional vs organic	60.11	Percentage change	Biodiversity	24	organic	9.46	Percentage change	yield	45	N	Sh	N	grapes	Y	Both	0.47	0.65	0.29	0.09	0.00	-1.04

530 **Methods**

531 We converted the percentage change back into log response ratios using the formula:

532 
$$LRR = \ln\left(\frac{\text{Percentage change}}{100} + 1\right)$$

533

534 *Reference scenarios*

535 In our second-order meta-analysis we used only results that were reported as log response  
536 ratios (LRR). The formula for the log response ratio is

537 
$$LRR = \ln\left(\frac{\bar{X}_T}{\bar{X}_C}\right)$$

538 Where:

539  $\ln$  is the natural logarithm

540  $\bar{X}_T$  is the mean of the treatment group

541  $\bar{X}_C$  is the mean of the control group

542 The assignment of control and treatment variables depends on the research question. For  
543 example, if a meta-analysis is examining the effect of a sustainable agricultural practice, such  
544 as replacement of mineral fertilizers with organic amendments, then the control would be the  
545 plots with mineral fertilizers, while the treatment would be the organic amendments. However,  
546 if a meta-analysis is examining the effect of intensification practices, such as tillage, on a  
547 response variable, then the control would be the no-tilled plots and the treatment would be the  
548 tilled plots. As expected, the reference scenarios were not homogeneous across our sample.

549 We standardized reference scenarios by examining closely the phrasing of the research  
550 questions of each meta-analysis and keeping a record of the study design. For our analysis,  
551 we used all sustainable management practices as treatments. In essence, the control should  
552 be an intensified and non-diversified system and the treatment should be a less intensive,

553 diversified system. Since not all meta-analyses were abiding to this design, we inverted some  
554 of the effect sizes by multiplying the reported log response ratio by -1, because:

555 
$$LRR_{reversed} = \frac{1}{LRR} = \ln \frac{1}{\frac{\bar{X}_T}{\bar{X}_C}} = -\ln \frac{\bar{X}_T}{\bar{X}_C} = -LRR$$

556

557 Table S2. The overall means of the random effects second-order meta-analysis based on first-  
558 order meta-analytic means.

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<b>Response variable</b>	<b>Estimate</b>	<b>CI low</b>	<b>CI up</b>	<b>Prediction Interval low</b>	<b>Prediction Interval up</b>	<b>Sample size (studies)</b>
Biodiversity	0.05	-0.08	0.19	-0.63	0.73	22
Yield	-0.019	-0.17	0.13	-0.76	0.72	22

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570 Table S3. Moderator estimates per taxonomic group.

Moderator	Estimate	CI.LB	CI.UB	Taxon
Diversification	0.05	-0.09	0.19	All taxa
Grazing pause	0.13	-0.33	0.59	All taxa
Lower intensity	-0.10	-0.53	0.34	All taxa
Reduced resource addition	-0.29	-0.80	0.22	All taxa
Organic	-0.07	-0.35	0.22	All taxa
Overall	0.23	-0.09	0.55	All taxa
Diversification	0.16	0.00	0.33	Invertebrates
Lower intensity	0.41	-1.16	1.99	Invertebrates
Organic	-1.30	-2.85	0.26	Invertebrates
Overall	1.46	0.02	2.90	Invertebrates
Diversification	0.58	-1.13	2.28	Plants
Grazing pause	0.01	-0.35	0.38	Plants
Lower intensity	0.19	0.02	0.37	Plants
Reduced resource addition	0.05	-0.05	0.16	Plants
Organic	0.48	0.08	0.89	Plants
Overall	0.12	0.01	0.23	Plants
Diversification	0.42	-1.05	1.89	Fungi
Reduced resource addition	0.10	0.00	0.20	Fungi
Organic	-0.01	-0.22	0.20	Fungi
Overall	0.09	0.01	0.16	Fungi
Lower intensity	0.07	-0.26	0.40	Vertebrates
Organic	0.31	-0.17	0.79	Vertebrates
Overall	0.15	-0.12	0.42	Vertebrates

571

572

573 Table S4. Moderator estimates per crop type.

Moderator	Estimate	CI.LB	CI.UB	Crop_system
Diversification	-0.44	-15.75	14.87	All crop systems
Grazing pause	-11.92	-38.47	14.62	All crop systems
Lower intensity	-2.76	-18.08	12.56	All crop systems
Organic	-0.44	-15.70	14.82	All crop systems
Reduced resource addition	0.25	-22.93	23.44	All crop systems
Overall	-0.02	-0.17	0.13	All crop systems
Diversification	-2.93	-8.63	2.76	Cropland
Lower intensity	-5.27	-10.95	0.41	Cropland
Organic	-2.99	-8.66	2.67	Cropland
Reduced resource addition	0.02	-9.33	9.37	Cropland
Overall	-2.56	-7.69	2.58	Cropland
Diversification	-0.33	-7.08	6.43	Agroforestry
Organic	4.66	-2.18	11.49	Agroforestry
Overall	2.13	-2.79	7.06	Agroforestry
Grazing pause	-11.94	-61.44	37.57	Grassland
Lower intensity	-0.26	-95.34	94.82	Grassland
Organic	1.27	-93.82	96.36	Grassland
Reduced resource addition	0.35	-54.76	55.47	Grassland
Overall	-4.86	-30.81	21.09	Grassland

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576 Table S5. Structure of models fitted in the study

Model type	Model name	Structure
<b>Linear regression</b>		
	reg	Yield ~ Biodiversity
<b>Second-order meta-analysis</b>		
<b>Random effects models</b>		
	meta_yield	rma(yi = Prod_ES_homog, vi = Prod_SE3, method = "REML", data = df_Irr)
	meta_bio	rma(yi = Biodiv_ES_homog, vi = Biodiv_SE3, method = "REML", data = biodiv)
	meta_invertebrates	rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_invertebrates), data = invertebrates)
	meta_plants	rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_plants), data = plants)
	meta_fungi	rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_fungi), data = fungi)
	meta Vertebrates	rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1 Vertebrates), data = vertebrates)

meta_crop	rma.mv(yi = Prod_ES_homog, V = Prod_SE3, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_crop), data = crop)
meta_agrof	rma.mv(yi = Prod_ES_homog, V = Prod_SE3, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_agrof), data = agrof)
meta_grass	rma.mv(yi = Prod_ES_homog, V = Prod_SE3, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_grass), data = grass)

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### Meta-regressions

metareg_bioyield	rma.mv(yi = Prod_ES_homog, V = Prod_SE3, mods = ~ 0 + Biodiv_ES_homog, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_df_lrr), data = df_lrr)
metareg_bio	rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_biodiv_lrr), data = biodiv)
metareg_invertebrates	rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML",

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	<pre>R = list(StudyID = A1_invertebrates), data = invertebrates)</pre>
metareg_plants	<pre>rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_plants), data = plants)</pre>
metareg_fungi	<pre>rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_fungi), data = fungi)</pre>
metareg_vertebrates	<pre>rma.mv(yi = Biodiv_ES_homog, V = Biodiv_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_vertebrates), data = vertebrates)</pre>
metareg_yield	<pre>rma.mv(yi = Prod_ES_homog, V = Prod_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_yield_Irr), data = yield)</pre>
metareg_crop	<pre>rma.mv(yi = Prod_ES_homog, V = Prod_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_crop), data = crop)</pre>

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metareg_agrof	rma.mv(yi = Prod_ES_homog, V = Prod_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_agrof), data = agrrof)
metareg_grass	rma.mv(yi = Prod_ES_homog, V = Prod_SE3, mods = ~ 0 + Management_grouped, random = list(~1 StudyID, ~1 ES_ID), method = "REML", R = list(StudyID = A1_grass), data = grass)

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