

## Supplementary Information

### **Biodiversity stabilizes plant communities through statistical-averaging effects rather than compensatory dynamics**

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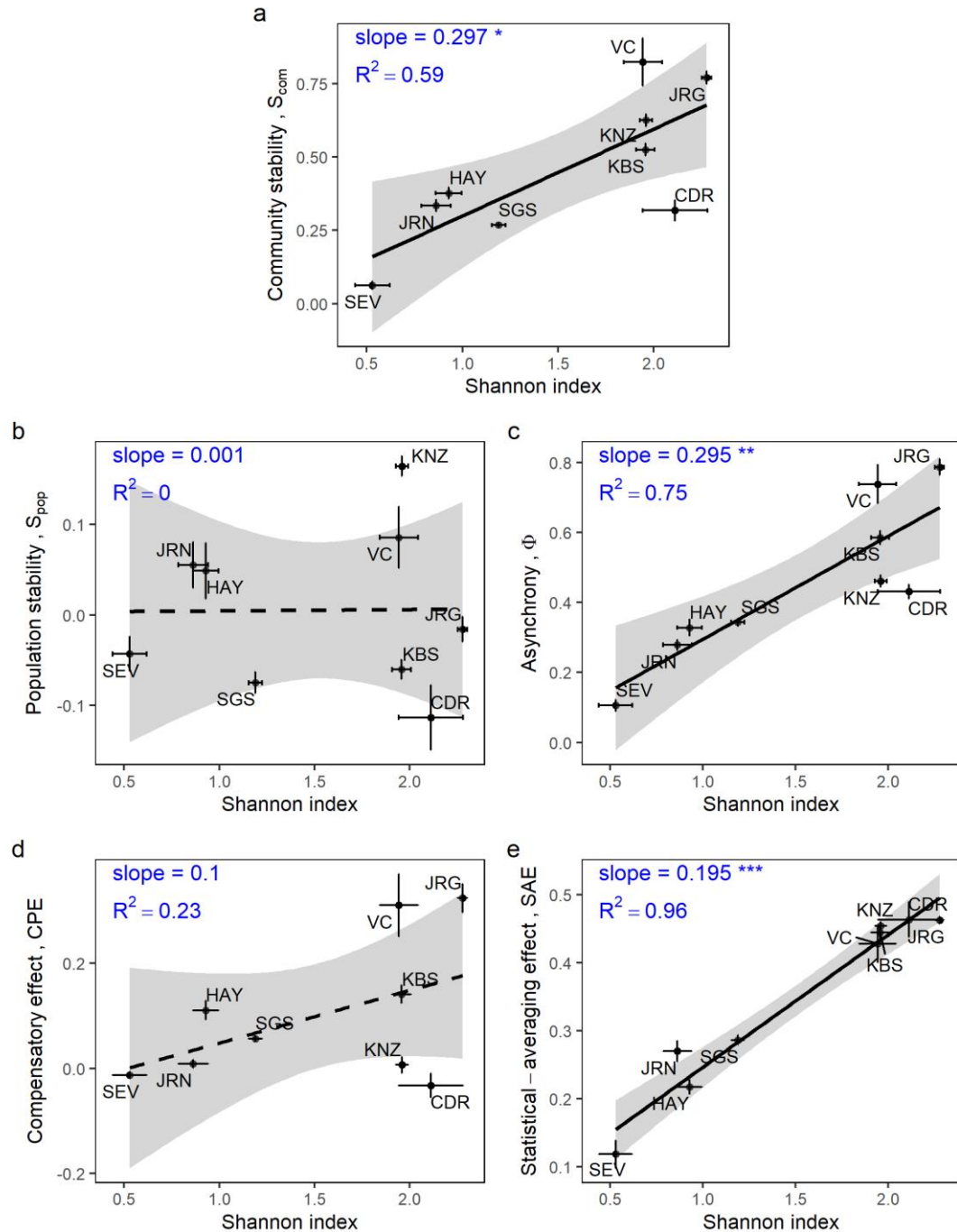
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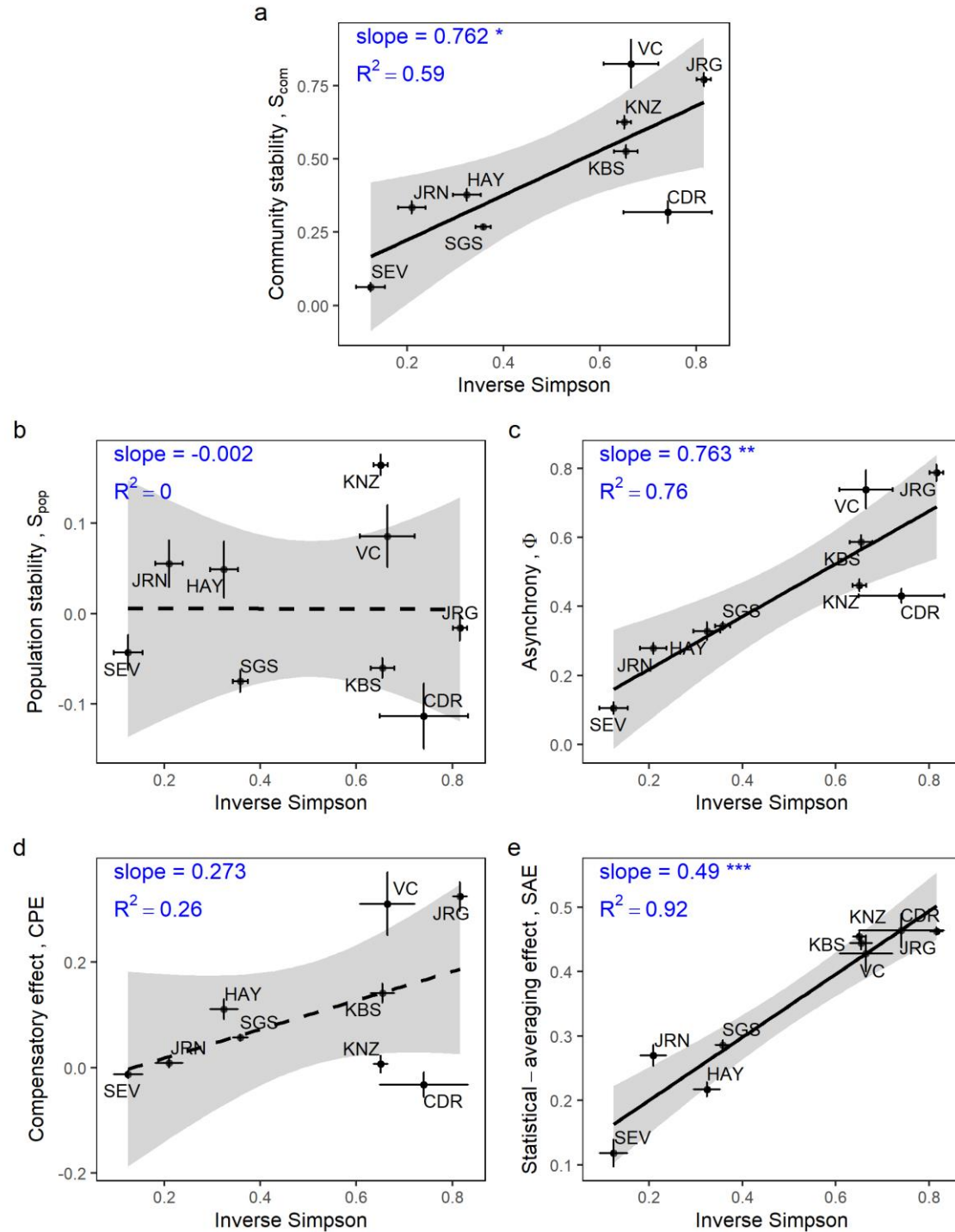
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**Supplementary Table 1: Significance of compensatory effects (*CPE*).** Within each plot, the significance of *CPE* was assessed as compared to asynchronous surrogates (Methods). Each plot was deemed to have significantly compensatory dynamics (*CPE* significantly greater than 1), or significantly synchronous dynamics (*CPE* significantly less than 1), or neither.

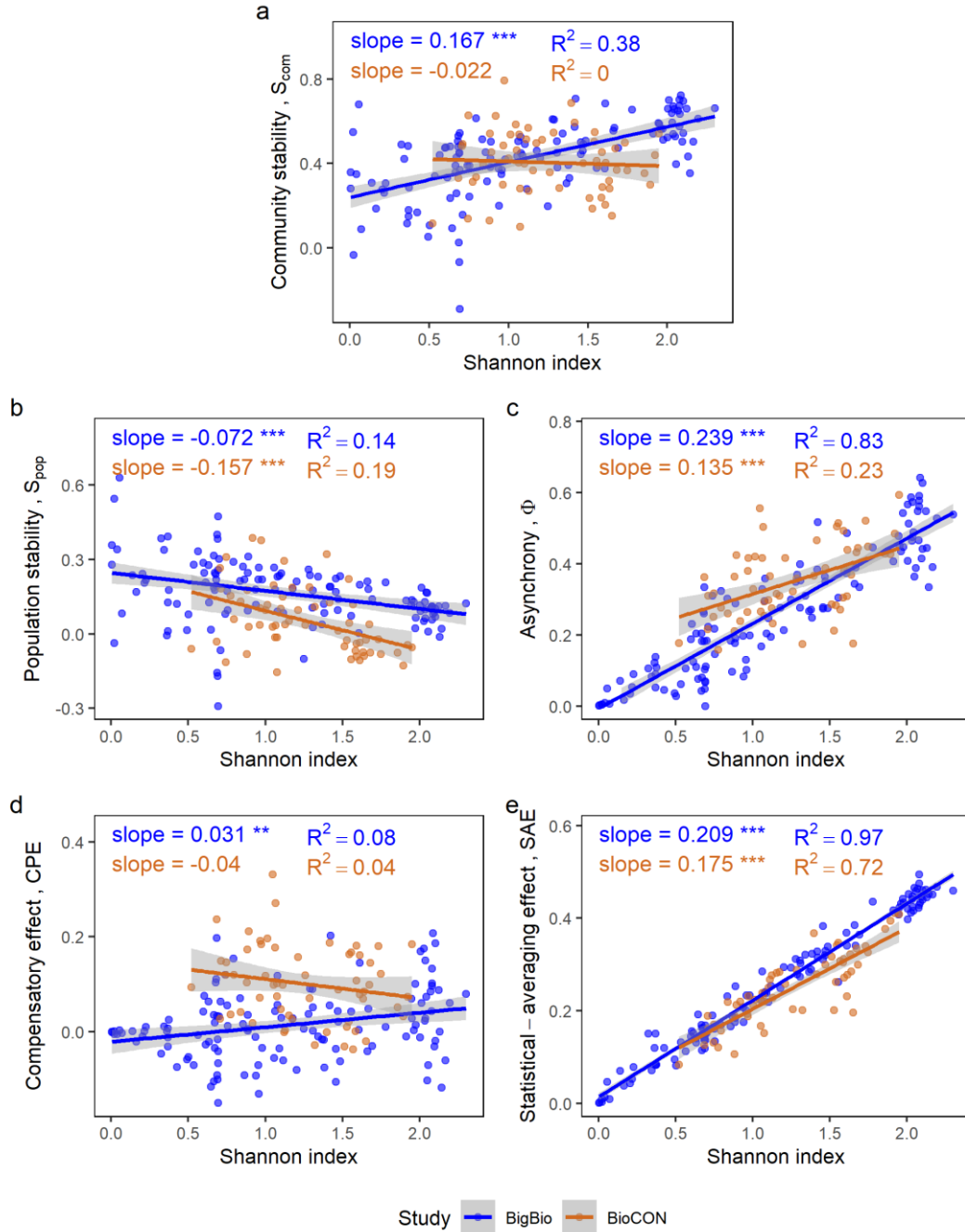
Dataset	Site	Plots	Proportion of plots showing significantly compensatory dynamics	Nonsignificance	Proportion of plots showing significantly synchronous dynamics
Plant Survey Dataset	JRG	18	100%	0%	0%
	KBS	30	30%	70%	0%
	HAY	13	46.2%	53.8%	0%
	JRN	47	12.5%	83.3%	4.2%
	KNZ	20	5%	90%	5%
	SEV	22	0%	90.9%	9.1%
	CDR	5	0%	100%	0%
	SGS	100	14%	86%	0%
	VC	6	66.7%	33.3%	0%
Biodiversity-manipulated Experiment Dataset	BigBio	122	4.8%	86.4%	8.8%
	BioCON	42	30.5%	69.5%	0%



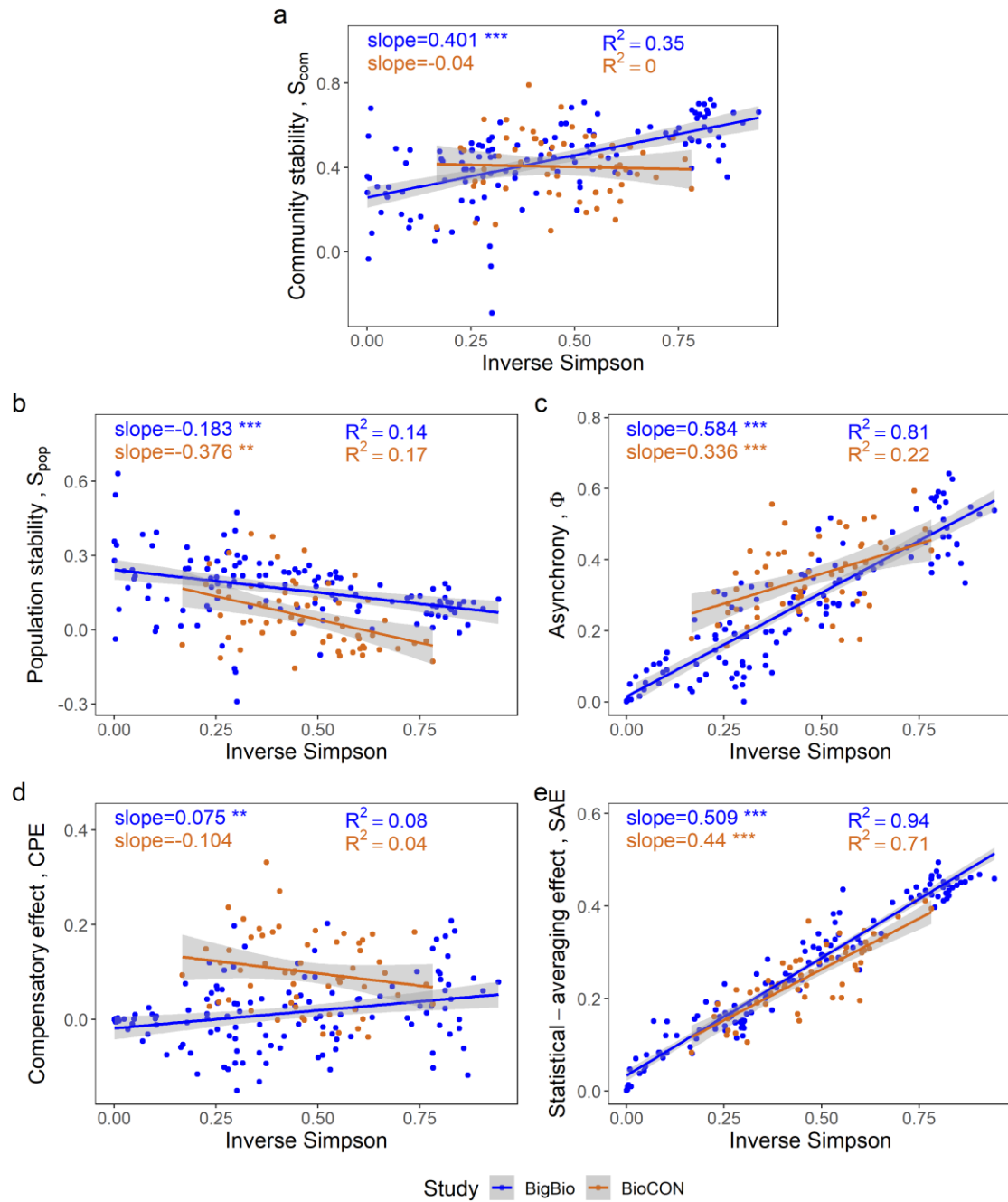
**Supplementary Fig. 1: Effects of the Shannon index on community stability and its components across the nine sites in the plant survey dataset.** Relationships are shown between the Shannon index and (a) community stability ( $p = 0.015$ ), (b) population stability ( $p = 0.979$ ), (c) asynchrony ( $p = 0.003$ ), (d) the compensatory effect ( $p = 0.189$ ), and (e) the statistical-averaging effect ( $p < 0.001$ ). All values (including standard error) are  $\log_{10}$  transformed except for the Shannon index. Regression and significance results formatted as in Fig. 2.



**Supplementary Fig. 2: Effects of the inverse of Simpson's index on community stability and its components across the nine sites in the plant survey dataset.** Relationships are shown between the inverse of Simpson's index and (a) community stability ( $p = 0.016$ ), (b) population stability ( $p = 0.991$ ), (c) asynchrony ( $p = 0.002$ ), (d) the compensatory effect ( $p = 0.160$ ), and (e) the statistical-averaging effect ( $p < 0.0001$ ). All values (including standard error) are  $\log_{10}$  transformed. Regression and significance results formatted as in Fig. 2.



**Supplementary Fig. 3: Effects of the Shannon index on community stability and its components in the biodiversity-manipulated experimental dataset.** Relationships are shown between the Shannon index and (a) community stability ( $p < 0.001$  for BigBio and  $p = 0.675$  for BioCON), (b) population stability ( $p < 0.001$  for both studies), (c) asynchrony ( $p < 0.001$  for both studies), (d) the compensatory effect ( $p = 0.002$  for BigBio and  $p = 0.147$  for BioCON), and (e) the statistical-averaging effect ( $p < 0.001$  for both studies) within two studies (indicated by different colours). All values are  $\log_{10}$  transformed except for the Shannon index. Regression and significance results formatted as in Fig. 2.



**Supplementary Fig. 4: Effects of inverse Simpson's index on community stability and its components in the biodiversity-manipulated experimental dataset.** Relationships are shown between the inverse Simpson's index and (a) community stability ( $p < 0.001$  for BigBio and  $p = 0.769$  for BioCON), (b) population stability ( $p < 0.001$  for BigBio and  $p = 0.001$  for BioCON), (c) asynchrony ( $p < 0.001$  for both studies), (d) the compensatory effect ( $p = 0.002$  for BigBio and  $p = 0.147$  for BioCON), and (e) the statistical-averaging effect ( $p < 0.001$  for both studies) within two studies (indicated by different colours). All values are log10 transformed. Regression and significance results formatted as in Fig. 2.

Asynchrony

Compensatory effect

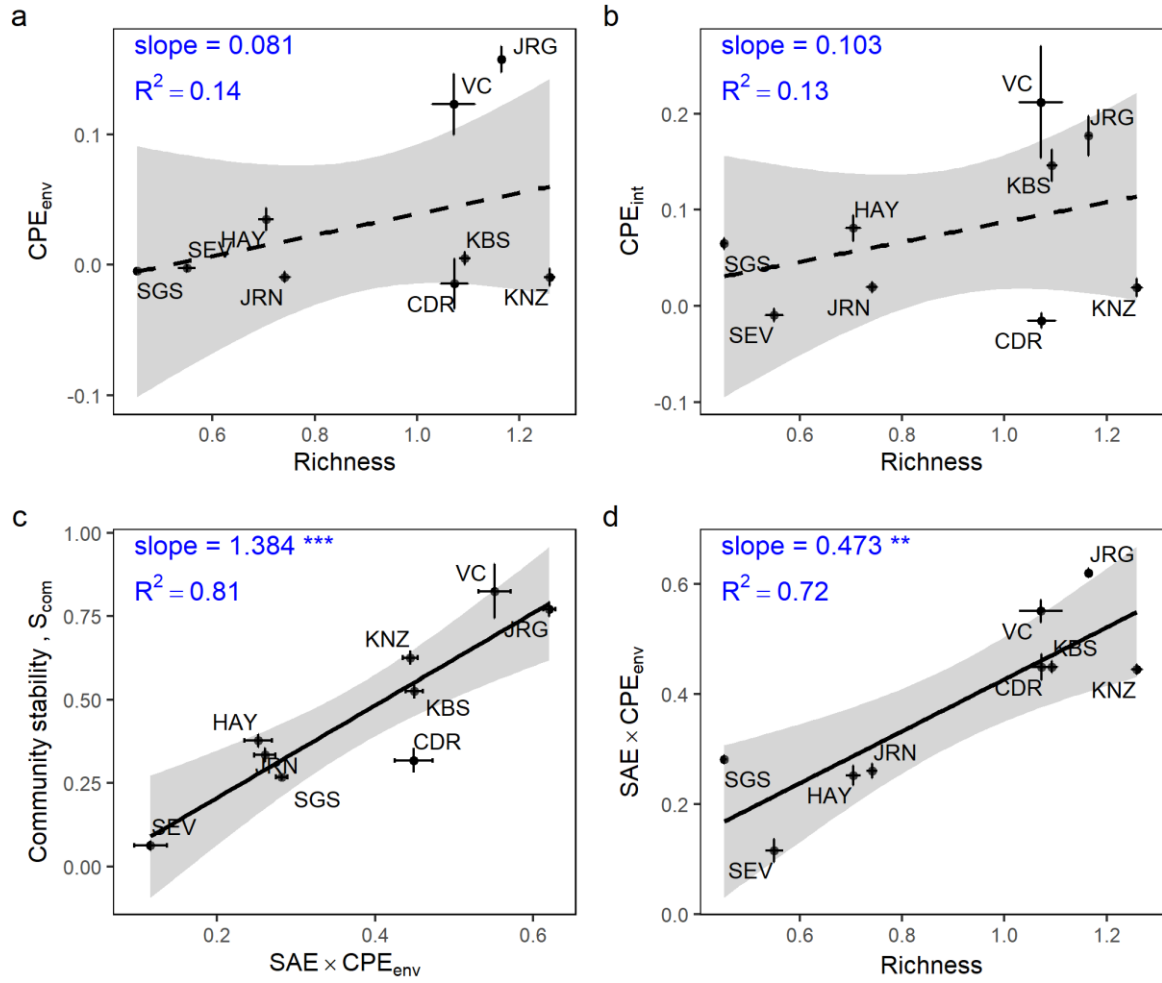
$$S_{\text{com}} = CPE_{\text{int}} \times CPE_{\text{env}} \times SAE \times S_{\text{pop}}$$

Portfolio effect as defined by Doak *et al.*<sup>19</sup>

Portfolio effect as defined by Tilman<sup>20</sup>

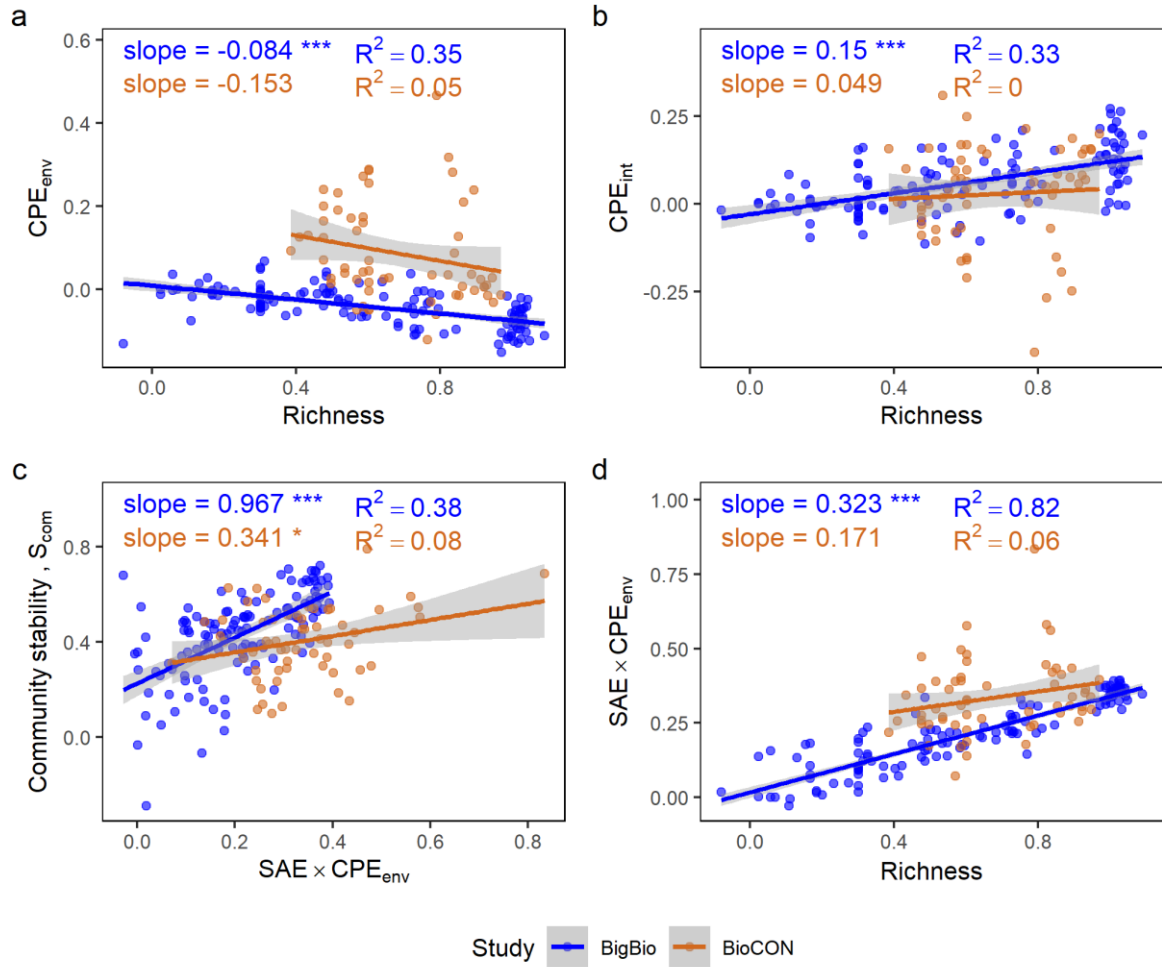
The diagram illustrates the decomposition of community stability ( $S_{\text{com}}$ ) into four components. A green bracket above the equation groups  $CPE_{\text{int}}$  and  $CPE_{\text{env}}$  under the label 'Asynchrony'. A blue bracket below the equation groups  $CPE_{\text{int}}$  and  $CPE_{\text{env}}$  under the label 'Compensatory effect'. A red bracket below the equation groups  $CPE_{\text{env}}$  and  $SAE$  under the label 'Portfolio effect as defined by Doak *et al.*<sup>19</sup>'. A purple line points from  $SAE$  to the label 'Portfolio effect as defined by Tilman<sup>20</sup>'.

**Supplementary Fig. 5: Summary equation of our full decomposition of community stability, and ecological meanings of components.**



**Supplementary Fig. 6: Relationships involving compensatory effects due to joint responses to environment fluctuations ( $CPE_{env}$ ) and due to species interactions ( $CPE_{int}$ ) across the nine sites in the plant survey dataset.** Relationships are shown between (a)  $CPE_{env}$  and richness ( $p = 0.329$ ), (b)  $CPE_{int}$  and richness ( $p = 0.342$ ), (c) community stability and the product of statistical averaging effect and  $CPE_{env}$  ( $p < 0.001$ ), and (d) richness and the product of statistical averaging effect and  $CPE_{env}$  ( $p = 0.004$ ). All values (including standard errors) are  $\log_{10}$  transformed. Regression and significance results formatted as in Fig. 2.





**Supplementary Fig. 7: Relationships involving compensatory effects due to joint responses to environment fluctuations ( $CPE_{env}$ ) and due to species interactions ( $CPE_{int}$ ) within studies of the biodiversity-manipulated experimental dataset.** Relationships are shown between (a)  $CPE_{env}$  and richness ( $p < 0.001$  for BigBio and  $p = 0.094$  for BioCON), (b)  $CPE_{int}$  and richness ( $p < 0.001$  for BigBio and  $p = 0.653$  for BioCON), (c) community stability and the product of statistical averaging effect and  $CPE_{env}$  ( $p < 0.001$  for BigBio and  $p = 0.029$  for BioCON), and (d) richness and the product of statistical averaging effect and  $CPE_{env}$  ( $p < 0.001$  for BigBio and  $p = 0.072$  for BioCON). All values (including standard errors) are  $\log_{10}$  transformed. Regression and significance results formatted as in Fig. 2.