

Correction

POPULATION BIOLOGY

Correction for “Fast–slow continuum and reproductive strategies structure plant life-history variation worldwide,” by Roberto Salguero-Gómez, Owen R. Jones, Eelke Jongejans, Simon P. Blomberg, David J. Hodgson, Cyril Mbeau-Ache, Pieter A. Zuidema, Hans de Kroon, and Yvonne M. Buckley, which was first published December 22,

2015; 10.1073/pnas.1506215112 (*Proc Natl Acad Sci USA* 113: 230–235).

The authors note that the symbols “greater than” and “lower than” of the survivorship curve type (H) in Table 1 appeared incorrectly. The corrected table appears below. This error does not affect the results or conclusions of the article.

Table 1. Loadings of the life-history traits grouped by their relation to turnover, and strategies to longevity, growth, and reproduction onto the first two PCA axes

Life-history trait	Symbol	Definition	PCA 1	PCA 2
Turnover		Number of years necessary for the individuals of a population to be fully replaced by new ones	0.85	0.17
Generation time	T			
Longevity		Shape of the age-specific survivorship curve l_x as quantified by Keyfitz' entropy (H). $H < 1, = 1, > 1$ correspond to survivorship curves types I, II and III, respectively	0.55	0.23
Survivorship curve type	H			
Age at sexual maturity	L_a	Number of years that it takes an average individual in the population to become sexually reproductive	0.71	0.29
Growth		Mean probability of transitioning to a larger/more developed stage in the life cycle of the species, weighted by the stable stage distribution (SSD)	−0.73	−0.05
Progressive growth	γ			
Retrogressive growth	ρ	Mean probability of transitioning to a smaller/less developed stage in the life cycle of the species, SSD-weighted	0.07	−0.77
Reproduction		Mean per-capita number of sexual recruits across stages in the life cycle of the species, SSD-weighted	−0.83	0.30
Mean sexual reproduction	Φ			
Degree of iteroparity	S	Spread of reproduction throughout the lifespan of the individual as quantified by Demetrius' entropy (S). High/low S values correspond to iteroparous/semelparous populations	−0.23	0.51
Net reproductive rate	R_0	Mean number of recruits produced during the mean life expectancy of an individual in the population	0.04	0.75
Mature life expectancy	L_w	Number of years from the mean age at sexual maturity (L_a) until the mean life expectancy (l_e) of an individual in the population	0.15	0.27
Explained variation, %			34.06	21.23
Cumulative percentage of explained variation			34.06	55.38

Loadings in bold indicate a high contribution (greater than ± 0.50) of the life-history trait to the PCA axis.

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