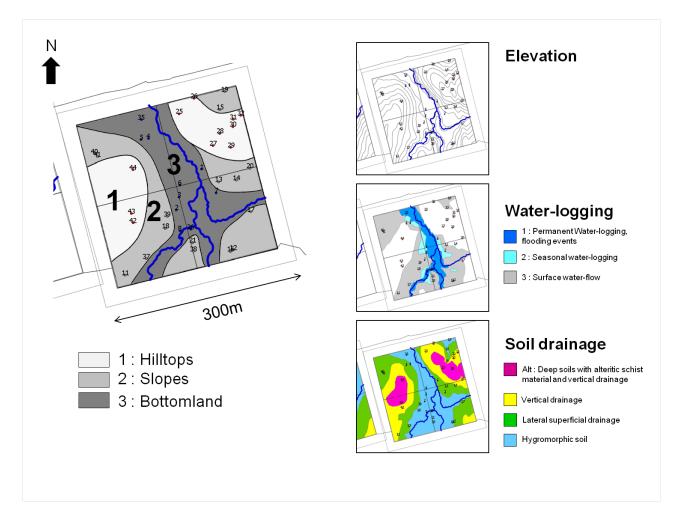
SUPPLEMENTARY DATA

FIG. S1: Three habitats were defined according to elevation, water-logging and soil drainage conditions. Sampled mother trees are indicated by red circles and numbers.



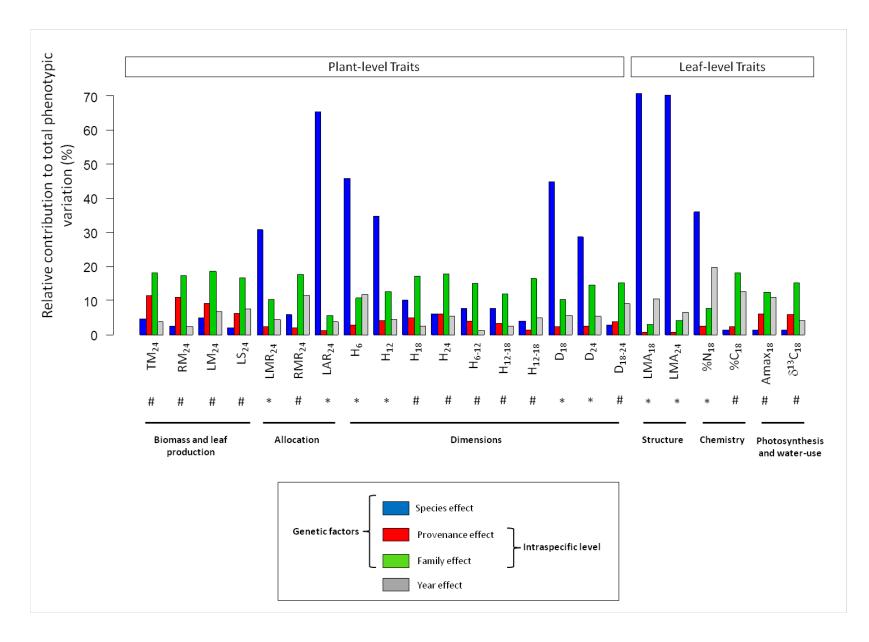


FIG. S2: relative size of different effects on trait variability (seed mass effect is not displayed here and therefore the bars do not sum to 100%).

FIG. S3: Pearson's correlation between phenotypic traits at seedling and maternal family level for *Eperua falcata*. Colours indicate the sign and strength of significant correlations (blue = negative correlation; red = positive correlation; deeper colours indicate stronger correlation). Empty cells correspond to non significant correlations ($\alpha = 5\%$). Significance levels: * = 5%; ** = 1% or less. FDR: seedling level: 0.07% ($\alpha = 5\%$) and 0.01% ($\alpha = 1\%$); maternal family level: 0.4% ($\alpha = 5\%$) and 0.06% ($\alpha = 1\%$).

	Н6	H12	H18	H24	D18	D24	H612	H1218	H1824	D1824	LMA18	LMA24	Amax18	d13C18	N18	C18	۲	LM	RM	ΜT	LAR	LMR	RMR
H6		0.91**	0.86**	0.83**	0.88**	0.77**	0.77**	0.62**	0.47**				0.55**	0.82**			0.75**	0.77**	0.71**	0.79**	-0.46**	-0.42**	-0.57**
H12	0.78**		0.97**	0.87**	0.93**	0.81**	0.97**	0.8**	0.49**				0.71**	0.9**			0.77**	0.79**	0.74**	0.82**	-0.47**	-0.41*	-0.62**
H18	0.61**	0.88**		0.88**	0.93**	0.84**	0.96**	0.9**	0.51**	0.34*			0.73**	0.9**			0.81**	0.83**	0.78**	0.85**	-0.4*	-0.34*	-0.63**
H24	0.57**	0.8**	0.92**		0.88**	0.92**	0.82**	0.74**	0.75**	0.48**			0.69**	0.78**			0.92**	0.92**	0.84**	0.92**			-0.76**
D18	0.67**	0.84**	0.81**	0.73**		0.9**		0.76**					0.76**	0.81**			0.81**	0.85**	0.85**	0.9**	-0.5**	-0.47**	-0.63**
D24		0.79**					0.78**		0.65**	0.59**			0.7**	0.73**					0.88**			-0.37*	
H612						0.71**		0.84**						0.87**			0.72**				-0.43**	-0.36*	
H1218						0.63**			0.38*	0.36*				0.81**			0.73**						-0.49**
H1824						0.39**		0.3**		0.62**			0.52**				0.71**						-0.54**
D1824			0.49**	_		0.72**	0.34**	_					0.35*				0.46**	0.38*	0.38*	0.37*			
LMA18	0.19**			-0.08**				-0.22**	-0.1**	-0.17**		0.35*									-0.48**		
LMA24		0.07**				0.15**			-0.23**		0.36**											-0.5**	
Amax18						0.52**			0.39**			-0.07*		0.58**			0.56**					-0.41*	
d13C18					_	0.54**											0.72**	0.73**	0.62**	0.71**	-0.38*		-0.64**
N18						0.26**			_		0.09**	-0.18**				0.36*							
C18						0.16**								0.11**									
LA						0.79**		0.58**	0.6**	0.56**		-0.1**	0.53**			0.14**	0.0000	0.98**		0.94**			-0.68**
LM						0.81**				0.55**		-		0.48**		0.14**			0.9**	0.97**			-0.7**
RM	0.59**					0.84**							0.51**			0.2**	0.82**	0.85**		0.97**		-0.44**	
TM		_			_	0.89**				0.55**			0.54**	_		0.18**	0.9**	0.93**	0.95**	0.0011	-0.39*	-0.33*	-0.62**
LAR						-0.34**							-0.06*							-0.26**	0.0000	0.93**	
LMR					•	-0.31**				0.5000			-0.06*	_				0.570		-0.24**		0.057	
RMR	-0.25**	-0.38**	-0.5**	-0.59**	-0.3**	-0.49**	-0.38**	-0.49**	-0.47**	-0.52**	0.2**	0.11**	-0.47**	-0.24**	-0.32**		-0.58**	-0.57**	-0.29**	-0.49**	-0.31**	-0.35**	

FIGURE S4: Pearson's correlation between phenotypic traits at seedling and maternal family level for *Eperua grandiflora*. Colours indicate the sign and strength of significant correlations (blue = negative correlation; red = positive correlation; deeper colours indicate stronger correlation). Empty cells correspond to non significant correlations ($\alpha = 5\%$). Significance levels: * = 5%; ** = 1% or less. FDR: seedling level: 0.2% ($\alpha = 5\%$) and 0.04% ($\alpha = 1\%$); maternal family level: 1.2% ($\alpha = 5\%$) and 0.3% ($\alpha = 1\%$).

