

Table S5 Female (co)variance components. Within-sex variances (diagonal) covariances (lower off diagonal) and correlations (upper off diagonal) between female life history traits ($\pm 1SE$) from a multivariate model of all female traits simultaneously. Genetic parameters do not have associated significances or standard errors as they are calculated from a fourth order FA model where significance values and errors are given on factor estimates, not on the subsequently recovered \hat{G} (see Table S8 for the non-FA estimate of G_f with associated errors on each element). Non-genetic matrices were estimated as variance-correlation matrices as these models proved more stable than unstructured variance-covariance models (Gilmour *et al.* 2009) and thus there are no standard errors on covariances. The permanent env/residual section of the table presents permanent environment (PE) variances in the upper row and residual variances in the lower row. AFR is multiplied by -1 to make any trade-offs negative in sign. Variances are presented for comparison with univariate models.

	SBA	AFR	L	ABS
Genetic				
SBA	0.165	0.220	0.147	-0.300
AFR	0.0360	0.163	-0.574	0.787
L	0.0161	-0.0624	0.0727	-0.696
ABS	-0.0257	0.0669	-0.0396	0.0444
Permanent env/ Residual				
SBA	<i>X</i> <i>0.660±0.057</i>	<i>NA</i> <i>NA</i>	<i>NA</i> <i>NA</i>	<i>NA</i> <i>NA</i>
AFR	<i>NA</i> <i>NA</i>	0.624±0.083 <i>X</i>	0.206±.0110* <i>NA</i>	0.999^{NE*} <i>NA</i>
L	<i>NA</i> <i>NA</i>	0.152 <i>NA</i>	0.867±0.121 <i>X</i>	0.504±0.172* <i>NA</i>
ABS	<i>NA</i> <i>NA</i>	0.168 <i>NA</i>	0.0997 <i>NA</i>	0.0452±0.0123 0.720±0.0171
Birth year				
SBA	0.0634±0.0228	0.381±0.249	<i>NA</i>	<i>NA</i>
AFR	0.0270	0.0792±0.0322	<i>NA</i>	<i>NA</i>
L	<i>NA</i>	<i>NA</i>	<i>X</i>	<i>NA</i>
ABS	<i>NA</i>	<i>NA</i>	<i>NA</i>	<i>X</i>
Maternal				
SBA	0.0689±0.0333	0.0145±0.360	<i>NA</i>	<i>NA</i>
AFR	0.0123	0.105±0.043	<i>NA</i>	<i>NA</i>
L	<i>NA</i>	<i>NA</i>	<i>X</i>	<i>NA</i>
ABS	<i>NA</i>	<i>NA</i>	<i>NA</i>	<i>X</i>
Year of measurement				
SBA	<i>X</i>	<i>NA</i>	<i>NA</i>	<i>NA</i>
AFR	<i>NA</i>	<i>X</i>	<i>NA</i>	<i>NA</i>
L	<i>NA</i>	<i>NA</i>	<i>X</i>	<i>NA</i>
ABS	<i>NA</i>	<i>NA</i>	<i>NA</i>	0.0293±0.0090

Bold values are significantly different from 0 ($P < 0.05$). *X* term not fit, see methods for details. *NA* covariance or correlation not applicable. *covariance is between PE for ABS and residual for other traits, estimated by forcing residual variance into permanent environment variance as detailed in the methods. ^{NE}Standard errors not estimable.