

**Table S3 Model showing variances (diagonal), covariances (below diagonal) and correlations (above diagonal) ( $\pm 1SE$ ) for the minimal models of life history traits in both sexes.** Genetic parameters do not have associated significances or standard errors as they are calculated from a fourth order FA model (see Methods). Underlined values highlight between-sex covariances and correlations. Non-genetic matrices were estimated as variance-correlation matrices and thus there are no standard errors on covariances. The permanent environment/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.

<b>Genetic</b>	Female SBA	Female AFR	Female L	Female ABS	Male SBA	Male AFR	Male L	Male ABS
Female SBA	0.187	0.294	0.00844	-0.221	<u>0.516</u>	<u>-0.138</u>	<u>-0.0639</u>	<u>0.0927</u>
Female AFR	0.0521	0.167	-0.368	0.829	<u>0.408</u>	<u>-0.0658</u>	<u>-0.350</u>	<u>-0.0668</u>
Female L	0.00108	-0.0443	0.0868	-0.510	<u>0.419</u>	<u>0.844</u>	<u>0.622</u>	<u>0.925</u>
Female ABS	-0.0206	0.0730	-0.0323	0.0464	<u>0.180</u>	<u>-0.226</u>	<u>-0.220</u>	<u>-0.21</u>
Male SBA	<u>0.0572</u>	<u>0.0426</u>	<u>0.0316</u>	<u>0.00993</u>	0.0655	0.197	0.625	0.703
Male AFR	<u>-0.0482</u>	<u>-0.0216</u>	<u>0.200</u>	<u>-0.0392</u>	0.0405	0.648	0.190	0.765
Male L	<u>-0.0118</u>	<u>-0.0608</u>	<u>0.0781</u>	<u>-0.0202</u>	0.0682	0.0650	0.181	0.722
Male ABS	<u>0.0101</u>	<u>-0.00684</u>	<u>0.0683</u>	<u>-0.0113</u>	0.0451	0.154	0.0772	0.0629

  

<b>Permanent env/ Residual</b>	Female SBA	Female AFR	Female L	Female ABS	Male SBA	Male AFR	Male L	Male ABS
Female SBA	X	NA	NA	NA	Male SBA	X	NA	NA
	<b>0.637<math>\pm</math>0.009</b>	NA	NA	NA		<b>0.707<math>\pm</math>0.047</b>	NA	NA
Female AFR	NA	<b>0.625<math>\pm</math>0.080</b>	0.188 $\pm$ 0.099*	0.999 <sup>NE*</sup>	Male AFR	NA	<b>0.456<math>\pm</math>0.208</b>	0.315 $\pm$ 0.235*
	NA	X	NA	NA		NA	X	NA
Female L	NA	0.137	<b>0.849<math>\pm</math>0.089</b>	<b>0.484<math>\pm</math>0.172*</b>	Male L	NA	0.19	<b>0.804<math>\pm</math>0.145</b>
	NA	NA	X	NA		NA	NA	X
Female ABS	NA	0.164	<b>0.0924</b>	<b>0.0429<math>\pm</math>0.0115</b>	Male ABS	NA	0.0723	0.0647
	NA	NA	NA	<b>0.721<math>\pm</math>0.017</b>		NA	NA	<b>0.648<math>\pm</math>0.023</b>

  

<b>Maternal</b>	Female SBA	Female AFR	Male SBA
Female SBA	<b>0.0740<math>\pm</math>0.0330</b>	-0.0359 $\pm$ 0.326	<b>0.835<math>\pm</math>0.362</b>
Female AFR	-0.00313	<b>0.103<math>\pm</math>0.042</b>	<u>0.285<math>\pm</math>0.328</u>
Male SBA	<b>0.0542</b>	<u>0.0218</u>	<b>0.0570<math>\pm</math>0.0282</b>

  

<b>Year of birth</b>	Female SBA	Female AFR	Male SBA
Female SBA	<b>0.0729<math>\pm</math>0.0252</b>	0.353 $\pm$ 0.250	<b>0.956<math>\pm</math>0.083</b>
Female AFR	0.0263	<b>0.0765<math>\pm</math>0.0312</b>	<u>0.386<math>\pm</math>0.235</u>
Male SBA	<b>0.0808</b>	0.0334	<b>0.0980<math>\pm</math>0.0314</b>

  

<b>Year of measurement</b>	Female SBA

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Female SBA	0.0291
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Bold values are significantly different from 0 ( $P < 0.05$ ), X term not fitted: see methods for details. *NA* covariance or correlation not applicable. <sup>NE</sup>Standard errors not estimable. \*covariance is between PE for ABS and residual for other traits (see Methods).