**S2 Table**. High posterior density (HPD) intervals for the genetic and phenotypic correlations between pairs of harvests (M1 to M6). The asterisk represents those estimates that are significant by HPD intervals.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Pair | Genetic | | Phenotypic | |
| Inferior | Superior | Inferior | Superior |
| **M1:M2** | -0.1313 | 0.9849 | -0.0517 | 0.2636 |
| **M1:M3** | -0.1262 | 0.8024 | -0.0349 | 0.2121 |
| **M1:M4** | 0.3113\* | 0.9995\* | 0.0475\* | 0.2583\* |
| **M1:M5** | 0.3401\* | 0.9821\* | 0.0153\* | 0.1493\* |
| **M1:M6** | 0.1305\* | 0.9561\* | -0.0104 | 0.0984 |
| **M2:M3** | -0.5061 | 0.7285 | -0.4366 | 0.6299 |
| **M2:M4** | -0.2176 | 0.9931 | -0.0836 | 0.3141 |
| **M2:M5** | 0.3408\* | 0.9983\* | 0.0651\* | 0.2806\* |
| **M2:M6** | 0.2762\* | 0.9985\* | 0.0124\* | 0.1720\* |
| **M3:M4** | 0.1384\* | 0.9988\* | 0.0249\* | 0.3305\* |
| **M3:M5** | 0.3185\* | 0.9217\* | 0.0803\* | 0.2870\* |
| **M3:M6** | 0.2270\* | 0.9402\* | 0.0416\* | 0.2445\* |
| **M4:M5** | 0.4933\* | 0.9997\* | 0.0708\* | 0.2505\* |
| **M4:M6** | 0.5092\* | 0.9997\* | 0.1257\* | 0.3444\* |
| **M5:M6** | 0.7612\* | 0.9999\* | 0.2619\* | 0.4395\* |