Table S5: Heritability estimates from data simulated over even and odd chromosomes of 8,000 individuals from the decode cohort.

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | s.e. |  | s.e. |  |  |
| 0.84 | 0.04 | 0.20 | 0.03 | 0.20 | 0.80 |
| 0.75 | 0.04 | 0.19 | 0.03 | 0.20 | 0.80 |
| 0.80 | 0.04 | 0.29 | 0.03 | 0.30 | 0.80 |
| 0.76 | 0.04 | 0.28 | 0.03 | 0.30 | 0.80 |
| 0.82 | 0.04 | 0.41 | 0.03 | 0.40 | 0.80 |
| 0.77 | 0.04 | 0.42 | 0.03 | 0.40 | 0.80 |
| 0.76 | 0.04 | 0.51 | 0.03 | 0.50 | 0.80 |
| 0.80 | 0.04 | 0.48 | 0.03 | 0.50 | 0.80 |
| 0.77 | 0.04 | 0.64 | 0.03 | 0.60 | 0.80 |
| 0.77 | 0.04 | 0.61 | 0.03 | 0.60 | 0.80 |