

Table A1 The means and standard errors of GEBV accuracy for ungenotyped individuals comparing different validation generations at different QTL scenarios and different single-step models (supplemental table for Figure 2).

| Models | QTL scenarios | Validation generations | | |
|-----------|---------------|------------------------|------------|------------|
| | | 5003 | 5004 | 5005 |
| SSGBLUP | 5 QTL | 0.43±0.020 | 0.31±0.027 | 0.27±0.017 |
| | 50 QTL | 0.41±0.018 | 0.34±0.015 | 0.27±0.029 |
| | 500 QTL | 0.44±0.020 | 0.34±0.018 | 0.23±0.008 |
| SS-BayesA | 5 QTL | 0.61±0.012 | 0.53±0.014 | 0.48±0.019 |
| | 50 QTL | 0.45±0.020 | 0.40±0.018 | 0.35±0.028 |
| | 500 QTL | 0.44±0.020 | 0.34±0.018 | 0.24±0.008 |
| SS-BayesB | 5 QTL | 0.63±0.012 | 0.55±0.013 | 0.47±0.039 |
| | 50 QTL | 0.51±0.018 | 0.48±0.014 | 0.40±0.031 |
| | 500 QTL | 0.40±0.033 | 0.31±0.027 | 0.22±0.018 |

Table A2 The means and standard errors of GEBV accuracy for genotyped individuals comparing different validation generations at different QTL scenarios and different single-step models (supplemental table for Figure 2).

| Models | QTL scenarios | Validation generations | | |
|-----------|---------------|------------------------|------------|------------|
| | | 5003 | 5004 | 5005 |
| SSGBLUP | 5 QTL | 0.48±0.015 | 0.41±0.020 | 0.34±0.021 |
| | 50 QTL | 0.45±0.019 | 0.39±0.008 | 0.34±0.016 |
| | 500 QTL | 0.46±0.020 | 0.36±0.017 | 0.32±0.021 |
| SS-BayesA | 5 QTL | 0.95±0.006 | 0.95±0.006 | 0.95±0.008 |
| | 50 QTL | 0.58±0.029 | 0.58±0.027 | 0.55±0.028 |
| | 500 QTL | 0.46±0.019 | 0.37±0.016 | 0.32±0.021 |
| SS-BayesB | 5 QTL | 0.98±0.002 | 0.98±0.003 | 0.93±0.050 |
| | 50 QTL | 0.74±0.019 | 0.75±0.017 | 0.66±0.065 |
| | 500 QTL | 0.40±0.025 | 0.33±0.023 | 0.28±0.016 |

Table A3 The means and standard errors of GEBV accuracy for ungenotyped individuals comparing different single-step models at different QTL scenarios and different validation generations (supplemental table for Figure 3).

| Validation generations | QTL scenarios | Models | | |
|------------------------|---------------|------------|------------|------------|
| | | SSGBLUP | SS-BayesA | SS-BayesB |
| 5003 | 5 QTL | 0.43±0.020 | 0.61±0.012 | 0.63±0.012 |
| | 50 QTL | 0.41±0.018 | 0.45±0.020 | 0.51±0.018 |
| | 500 QTL | 0.44±0.020 | 0.44±0.020 | 0.40±0.033 |
| 5004 | 5 QTL | 0.31±0.027 | 0.53±0.014 | 0.55±0.013 |
| | 50 QTL | 0.34±0.015 | 0.40±0.018 | 0.48±0.014 |
| | 500 QTL | 0.34±0.018 | 0.34±0.018 | 0.31±0.027 |
| 5005 | 5 QTL | 0.27±0.017 | 0.48±0.019 | 0.47±0.039 |
| | 50 QTL | 0.27±0.029 | 0.35±0.028 | 0.40±0.031 |
| | 500 QTL | 0.23±0.008 | 0.24±0.008 | 0.22±0.018 |

Table A4 The means and standard errors of GEBV accuracy for genotyped individuals comparing different single-step models at different QTL scenarios and different validation generations (supplemental table for Figure 3).

| Validation generations | QTL scenarios | Models | | |
|------------------------|---------------|------------|------------|------------|
| | | SSGBLUP | SS-BayesA | SS-BayesB |
| 5003 | 5 QTL | 0.48±0.015 | 0.95±0.006 | 0.98±0.002 |
| | 50 QTL | 0.45±0.019 | 0.58±0.027 | 0.74±0.019 |
| | 500 QTL | 0.46±0.020 | 0.46±0.016 | 0.40±0.025 |
| 5004 | 5 QTL | 0.41±0.020 | 0.95±0.006 | 0.98±0.003 |
| | 50 QTL | 0.39±0.008 | 0.58±0.027 | 0.75±0.017 |
| | 500 QTL | 0.36±0.017 | 0.37±0.016 | 0.33±0.023 |
| 5005 | 5 QTL | 0.34±0.021 | 0.95±0.008 | 0.93±0.050 |
| | 50 QTL | 0.34±0.016 | 0.55±0.028 | 0.66±0.065 |
| | 500 QTL | 0.32±0.021 | 0.32±0.021 | 0.28±0.016 |

Table A5 The means and standard errors of GEBV accuracy for ungenotyped individuals comparing different QTL numbers at different validation generations with different single-step models (supplemental table for Figure 4).

| Models | Validation generations | QTL scenarios | | |
|-----------|------------------------|---------------|------------|------------|
| | | 5 | 50 | 500 |
| SSGBLUP | 5003 | 0.43±0.020 | 0.31±0.027 | 0.27±0.017 |
| | 5004 | 0.41±0.018 | 0.34±0.015 | 0.27±0.029 |
| | 5005 | 0.44±0.020 | 0.34±0.018 | 0.23±0.008 |
| SS-BayesA | 5003 | 0.61±0.012 | 0.53±0.014 | 0.48±0.019 |
| | 5004 | 0.45±0.020 | 0.40±0.018 | 0.35±0.028 |
| | 5005 | 0.44±0.020 | 0.34±0.018 | 0.24±0.008 |
| SS-BayesB | 5003 | 0.63±0.006 | 0.55±0.006 | 0.47±0.008 |
| | 5004 | 0.51±0.029 | 0.48±0.027 | 0.40±0.028 |
| | 5005 | 0.40±0.019 | 0.31±0.016 | 0.22±0.021 |

Table A6 The means and standard errors of GEBV accuracy for genotyped individuals comparing different QTL numbers at different validation generations with different single-step models (supplemental table for Figure 4).

| Models | Validation generations | QTL scenarios | | |
|-----------|------------------------|---------------|------------|------------|
| | | 5 | 50 | 500 |
| SSGBLUP | 5003 | 0.48±0.015 | 0.41±0.020 | 0.34±0.021 |
| | 5004 | 0.45±0.019 | 0.39±0.008 | 0.34±0.016 |
| | 5005 | 0.46±0.020 | 0.36±0.017 | 0.32±0.021 |
| SS-BayesA | 5003 | 0.95±0.006 | 0.95±0.006 | 0.95±0.008 |
| | 5004 | 0.58±0.029 | 0.58±0.027 | 0.55±0.028 |
| | 5005 | 0.46±0.019 | 0.37±0.016 | 0.32±0.021 |
| SS-BayesB | 5003 | 0.98±0.002 | 0.98±0.003 | 0.93±0.050 |
| | 5004 | 0.74±0.019 | 0.75±0.017 | 0.66±0.065 |
| | 5005 | 0.40±0.025 | 0.33±0.023 | 0.28±0.016 |