Web table M Net reclassification improvement based on the addition of the genetic risk function to the Framingham Offspring risk score.

| Predicted | No. of individuals | | | | Reclassified | | Net |
|--------------------------------------------------------------------------------|--------------------|-------|--------|------|--------------|-----------|------------------------|
| diabetes risk | <5% | 5-10% | 10-15% | >15% | Increased | Decreased | correctly reclassified |
| Framingham | | | | | Risk | Risk | recrassified |
| Offspring | | | | | | | |
| risk score | | | | | | | |
| + genetic risk function Individuals without diabetes during follow-up (n=3324) | | | | | | | |
| 0 - <5% | 2245 | 98 | 1 | 0 | 221 | 166 | -1.7% |
| 5 - 10% | 80 | 409 | 70 | 2 | | | |
| 10 - 15% | 1 | 51 | 127 | 51 | | | |
| ≥ 15% | 0 | 2 | 32 | 156 | | | |
| + genetic risk function Individuals with diabetes during follow-up (n= 202) | | | | | | | |
| 0 - <5% | 51 | 9 | 0 | 0 | 21 | 24 | -1.5% |
| 5 - 10% | 4 | 32 | 6 | 0 | | | |
| 10 - 15% | 0 | 12 | 13 | 6 | | | |
| ≥15% | 0 | 1 | 7 | 61 | | | |
| Net reclassification improvement (95% CI) -3.2 (-9.9 to 3.5) P=0.35 | | | | | | | |