

**No signs of decline in genetic
variation in racing performance
traits in Swedish standardbred**

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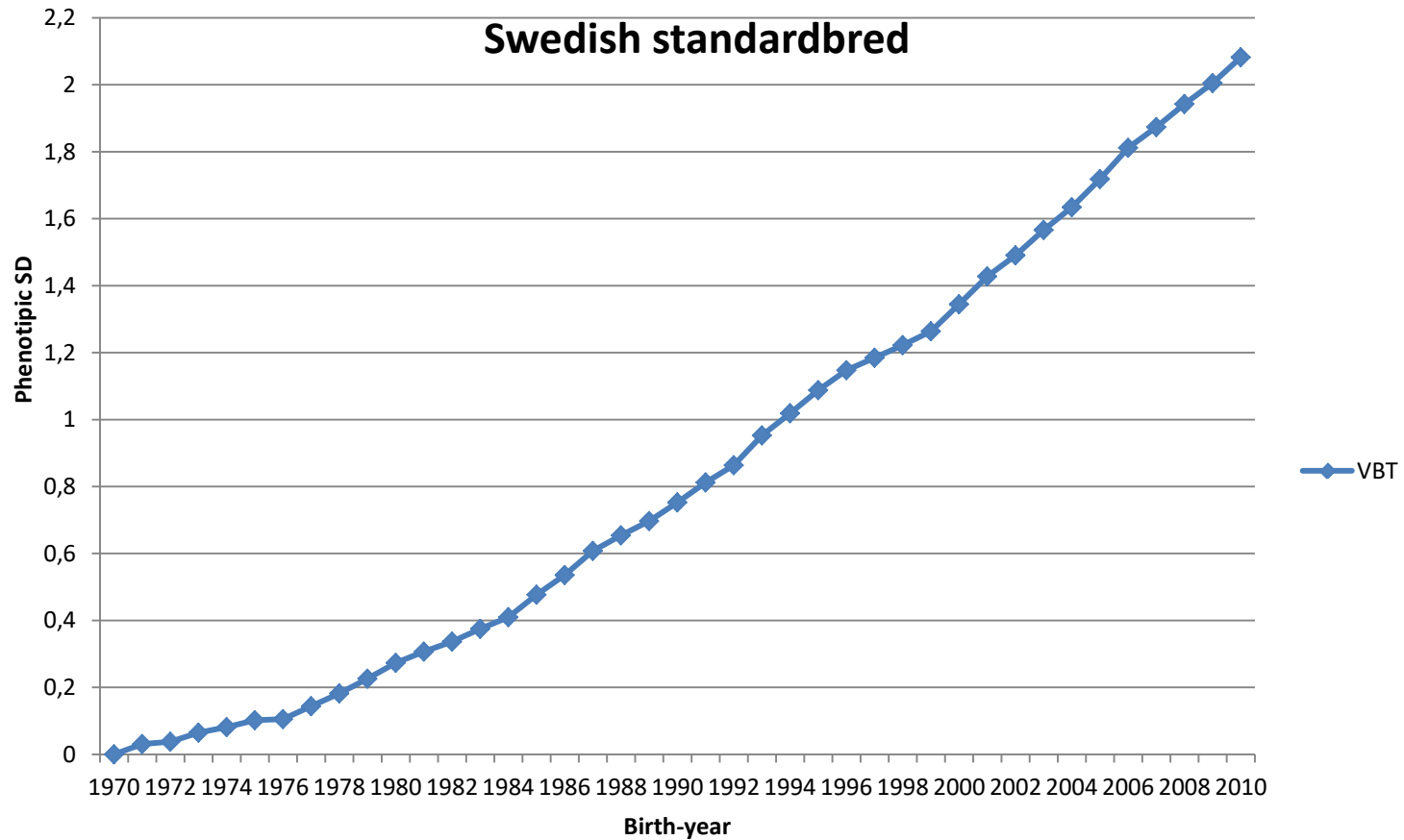
Background

- **Breeding Goal:** *fast, sound, sustainable, well tempered trotter of international standard with regular gaits and strong ability to win races*
- **Gene pool:** 94% American Standardbred, 6% French trotter
- **EBVs:** SireM-BLUP 1984-1991, AM-BLUP since 1992

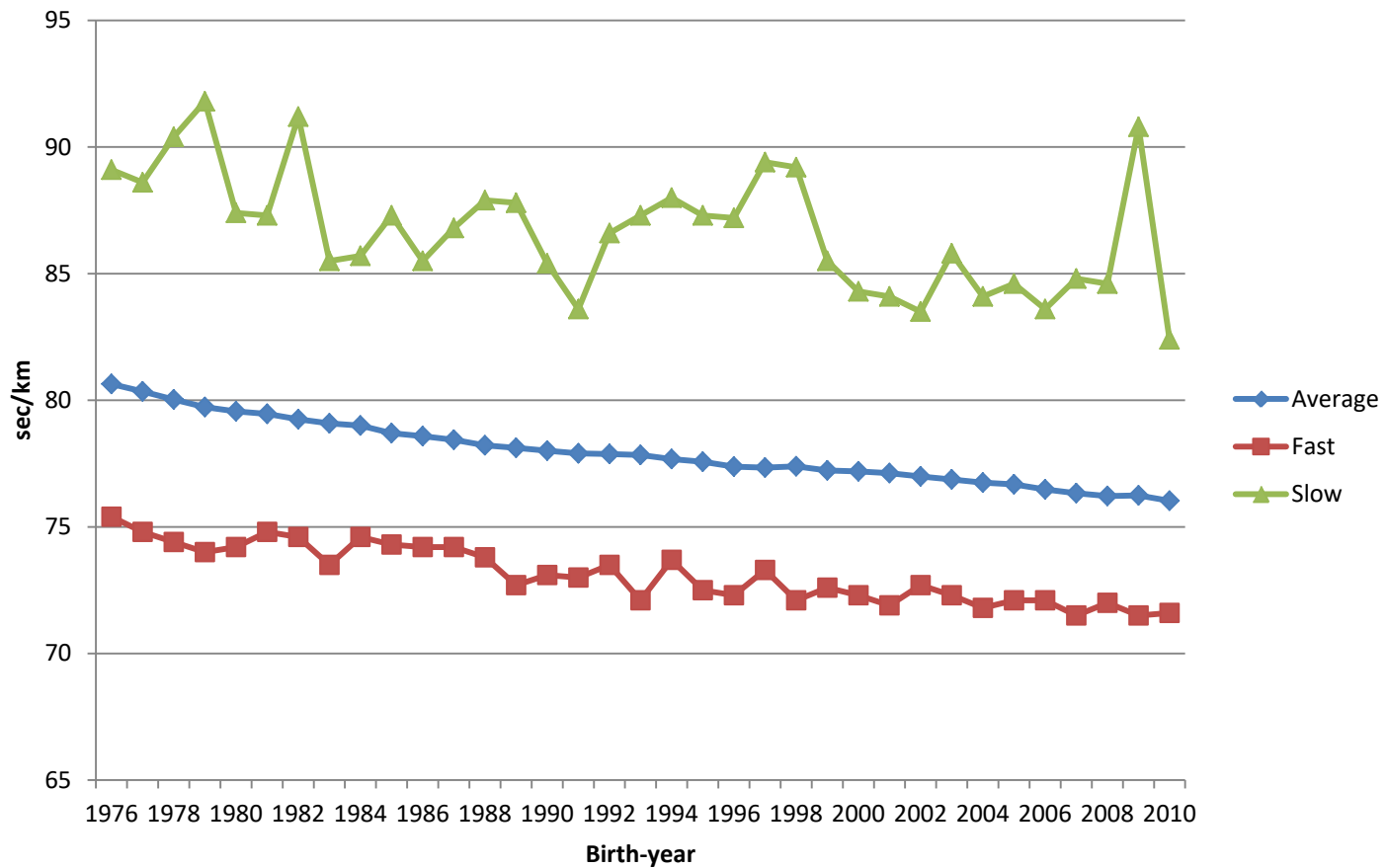
Racing performance traits

- **Accumulated 2-5 year-old-results (transformed):**
- $\text{Ln}(\text{Earnings} + 1000 \text{ SEK})$
- $\text{Ln}((\text{Earnings} + 1000 \text{ SEK})/(\text{number of races}))$
- $\text{Ln}(\text{Best racing time}(\text{sec/km}) - 68.2)$

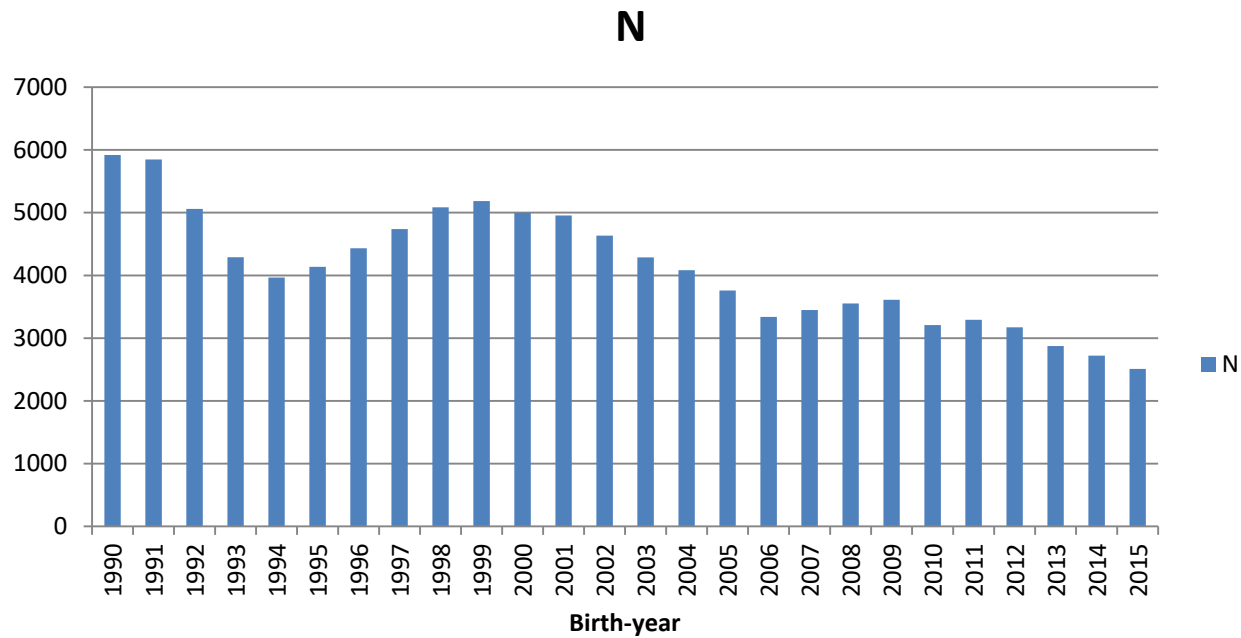
Genetic progress in racing performance



Trend in racing times (males)



Actual population size (foals registered)



Paternal and maternal lineage

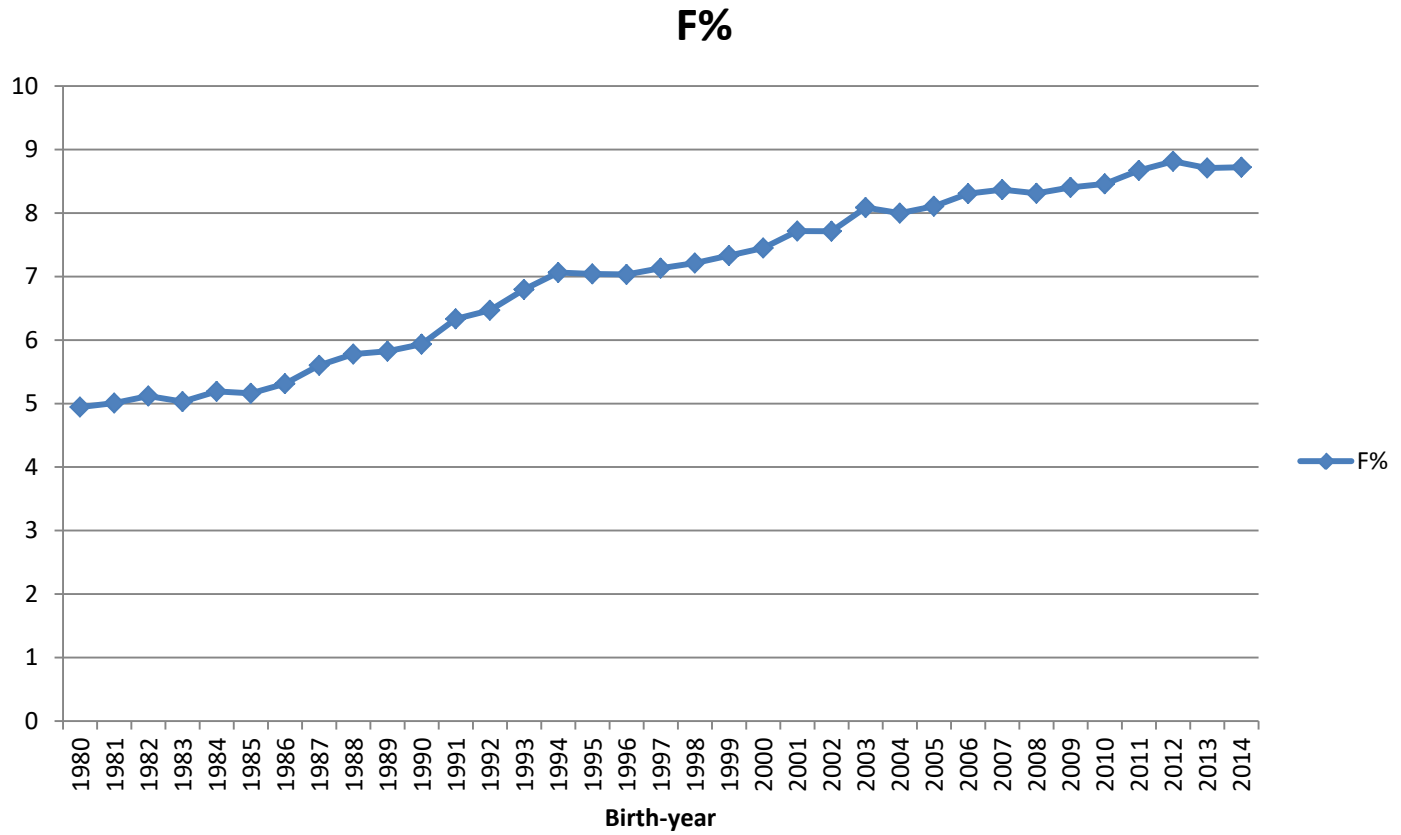
Paternal lineage founders of the current generation (2000-2009) :

| Stallion | Country | Birth-year | % |
|--------------|---------|------------|------|
| Happy Medium | USA | 1863 | 92,5 |
| Fuschia | France | 1883 | 1,8 |
| Axtell | USA | 1886 | 1,7 |
| McKinney | USA | 1887 | 1,2 |
| Beaumanour | France | 1901 | 2,8 |

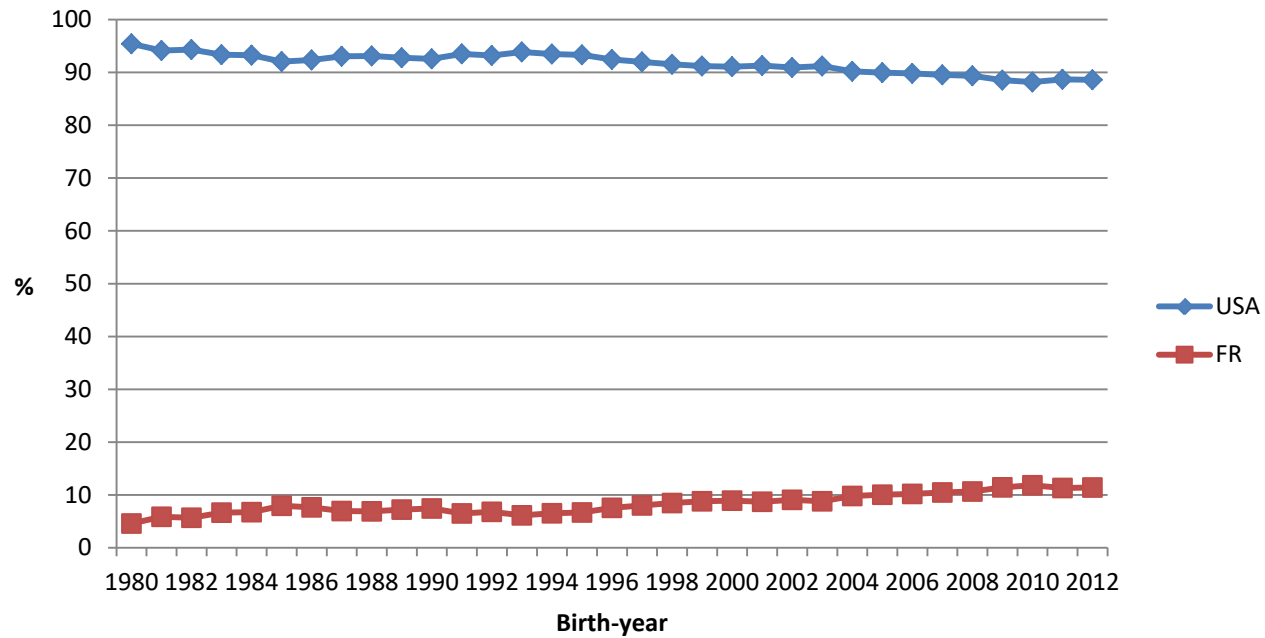
Maternal lineage founders of the current generation (2000-2009) :

781 maternal lines in total; 495 maternal lines with ≥ 5 direct decendants in the current generation; 87 maternal lines with ≥ 100 direct decendants in the current generation

Inbreeding



Influence of French genes



Effect of inbreeding and genetic progress in racing performance

- Rate of inbreeding: $\Delta F=0.0132 \approx 1,3\%$ *per* generation
- **$N_e = 35-40$ animals**
- Inbreeding depression : ca 0,5% of σ_p/yr
- Genetic progress: ca 7% of σ_p/yr
- -> **Adjusted genetic progress: 6,5% of σ_p/yr**
- **Has genetic variance reduced over time?**

Material

| Time period | No records (FR>0 excluded) | Pedigree file (5 generations) | Base average |
|--------------|----------------------------|-------------------------------|--------------|
| A. 1978-1990 | 33,099 (24,233) | 54,467 | 1932.6 |
| B. 1991-2000 | 28,753 (17,879) | 54,919 | 1942.8 |
| C. 2001-2009 | 21,994 (10,818) | 47,822 | 1953.8 |

| Time period | No records | Pedigree file (2 generations) | Base average |
|--------------|------------|-------------------------------|--------------|
| A. 1978-1990 | 33,099 | 44,348 | 1970.9 |
| B. 1991-2000 | 28,753 | 41,368 | 1982.4 |
| C. 2001-2009 | 21,994 | 33,027 | 1992.1 |

Methods

- AI-REML, DMU
- Model: $y = \text{sex/birth-year (fixed)} + \text{animal} + e$
- Single trait analysis
- Sire model tested
- Inclusion of maternal lineage (fixed or random effect) in the animal model was tested

Results (AM 5 generations traced, (FR>0 excluded))

| Trait | Period | h^2 | Var(A) | Var(P) |
|-------------------|--------|--------------------|------------------------|-----------------|
| Earnings | A | 0.34 (0.35) | 0.7888 (0.7756) | 2.2896 (2.2240) |
| | B | 0.37 (0.37) | 0.9796 (0.9402) | 2.6590 (2.5476) |
| | C | 0.37 (0.38) | 0.9782 (0.9729) | 2.6577 (2.5776) |
| Earnings per race | A | 0.46 (0.46) | 0.3737 (0.3543) | 0.8118 (0.7773) |
| | B | 0.46 (0.47) | 0.4851 (0.4703) | 1.0453 (0.9902) |
| | C | 0.48 (0.50) | 0.5123 (0.5247) | 1.0761 (1.0446) |
| Racing time | A | 0.30 (0.30) | 0.0110 (0.0110) | 0.0368 (0.0363) |
| | B | 0.34 (0.35) | 0.0129 (0.0129) | 0.0381 (0.0369) |
| | C | 0.34 (0.35) | 0.0161 (0.0163) | 0.0479 (0.0472) |

Results (AM 2 generations traced)

| Trait | Period | h^2 | Var(A) | Var(P) |
|-------------------|--------|-------------|---------------|--------|
| Earnings | A | 0.33 | 0.7411 | 2.2486 |
| | B | 0.36 | 0.9440 | 2.6177 |
| | C | 0.35 | 0.9148 | 2.5962 |
| Earnings per race | A | 0.44 | 0.3474 | 0.7905 |
| | B | 0.45 | 0.4632 | 1.0227 |
| | C | 0.45 | 0.4692 | 1.0399 |
| Racing time | A | 0.29 | 0.0101 | 0.0362 |
| | B | 0.33 | 0.0125 | 0.0375 |
| | C | 0.32 | 0.0151 | 0.0469 |

Results (Sire model)

| Trait | Period | t | Var(S) | Var(P) |
|-------------------|--------|--------------|---------------|--------|
| Earnings | A | 0.132 | 0.3115 | 2.3642 |
| | B | 0.120 | 0.3222 | 2.6740 |
| | C | 0.122 | 0.3180 | 2.6169 |
| Earnings per race | A | 0.192 | 0.1649 | 0.8580 |
| | B | 0.160 | 0.1680 | 1.0507 |
| | C | 0.163 | 0.1716 | 1.0506 |
| Racing time | A | 0.129 | 0.0050 | 0.0385 |
| | B | 0.110 | 0.0042 | 0.0382 |
| | C | 0.105 | 0.0050 | 0.0480 |

Conclusions

- No decline in h^2 or in genetic variance observed over time
- Estimated heritability of racing performance traits in Swedish standardbred trotters is high (0.3 to 0.48). I.e. 30-50% of the phenotypic variation is controlled by nuclear DNA!
- Inclusion of maternal lineage analyses indicated that variation in mitochondrial DNA contributed very little (0 to 0.5% of σ^2_p) to the variance in racing performance in Swedish standardbred

Implications and questions

- Creation of new genetic variation in the racing performance traits seems to balance loss due to inbreeding!
- Migration – DMRT3?
- Is there a critical actual population size?
- “Steady state” and Bulmer effect?

Acknowledgements

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