

Development of variation in a random bred mouse population

W. Schlote¹, A. Wolc², T.A. Schmidt³ and T. Szwaczkowski²

¹Populationsgenetik, Institut für Nutztierwissenschaften, Humboldt-Universität zu Berlin, 10099 Berlin, Germany,

²Department of Animal Genetics and Breeding, August Cieszkowski Agricultural University of Poznan, 60627 Poznan, Poland,

³T.A. Schmidt, Institute for Animal Breeding (FAL), 31535 Neustadt-Mariensee, Germany



1. Material:

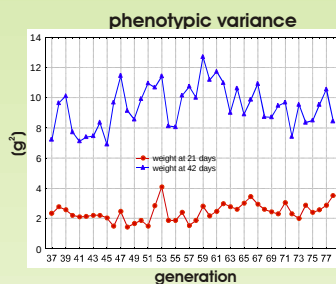
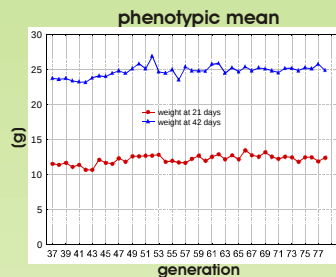
- 42 generations (1989 - 2005)
- 24 363 individual records
- varied population size
- 1- 3 halfsib families per sire



Table 1: Overall heritability

	W21(g)	W42(g)	G42(g)
s^2_A	1.006	2.761	.00329
s^2_P	1.793	4.676	.00664
h^2 (sh^2)	.563 (0.12)	.591 (.014)	.496 (.015)

2. Trend of phenotypic mean and variance ?



4. Trend of h^2 ?

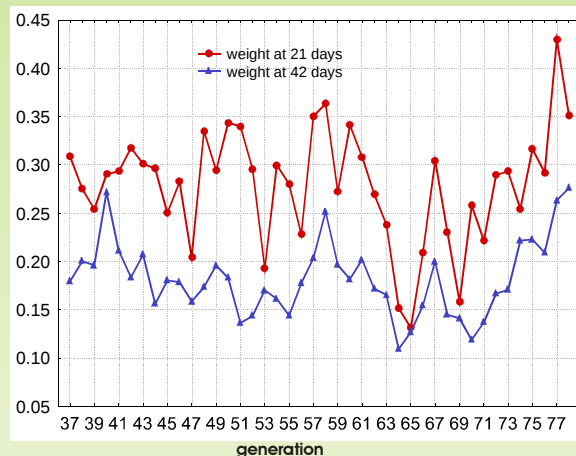
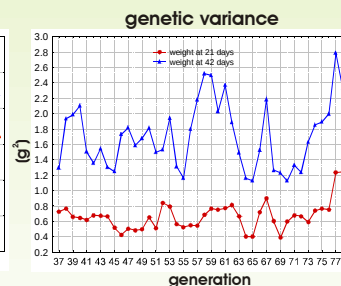
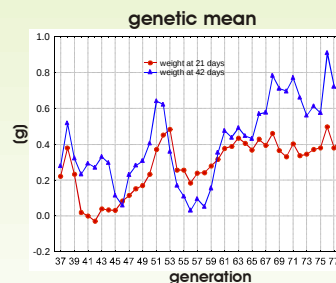


Table 2: Overall correlation (r_A/r_P)

	W21	W42	G42
W21	1	.654	.062
W42	.518	1	.782
G42	-.022	.894	1

3. Trend of genetic mean and variance ?



5. Conclusions

- stable phenotypic mean and variance
- increased genetic mean and variance
- high variation and increase in generation estimate for heritability
- marked difference between generation and overall estimates

W21 = weight at 21 days W42 = weight at 42 days G42 = growth 21- 42 days