

PREDICTION OF BREEDING VALUE AND ESTIMATION OF VARIANCE
COMPONENTS IN SPAIN BY MIXED MODELS.

1983. Sire Model (unpublished results).
1986. Animal Model (unpublished results).
1987. REML - EM (Sire Model).
1988. Animal Model (Official catalogue).
1992. Changes in the model:
- New definition of Contemporary Groups.
 - Days-open effect.
 - Heterogeneity of variance correction.
 - P.E.V. (modified Meyer, 1989).
 - Genetic grouping (origin by sex and age).

CONTEMPORARY GROUP (CG). TREATMENT AND DEFINITION.

RESULTS FROM SIMULATION (UGARTE et al., 1992).

C.G. RANDOM IMPLIES SOME GOOD PROPERTIES:

- ↓ PREDICTION ERROR VARIANCE.
- ↓ MSE WHEN G.C. > 4.
- ↑ GENETIC CORRELATION BETWEEN
PREDICTED AND TRUE BREEDING VALUE.

RESULTS FROM REAL DATA. (BEN GARA AND ALENDA, 1992).

G.C. RANDOM IMPLIES:

* GOOD PROPERTIES:

- 1 CORRELATION PEDIGREE INDEX - YD
- 1 EFFECTIVE NUMBER (N_e)/REAL NUMBER (N_r)
- 1 N_e/σ^2 .

* BAD PROPERTIES:

- 1 CORRELATION PREDICTED B.V. - C.G. EFFECT.
- B.V. OF BULLS USED IN ^{good environment} POSITIVE C.G. EFFECTS ARE OVERESTIMATED.

ERROR VARIANCE ($\sigma^2_{\bullet}$), COEFFICIENT OF DETERMINATION (CD) AND RATIO OF EFFECTIVE SIZE ($N_{\bullet}$) AND ERROR VARIANCE FOR THE MODELS THAT CONSIDERED HERD-YEAR AND HERD-YEAR-MOBILE SEASON AS FIXED OR RANDOM.

MODEL		$\sigma^2_{\bullet}$	CD	$N_{\bullet}/\sigma^2_{\bullet}$
HERD-YEAR	FIXED	1,465,300	.42	.044
	RANDOM	1,310,610	.48	.050
HERD-YEAR-SEASON (MOBILE)	FIXED	1,026,250	.62	.057
	RANDOM	1,026,500	.62	.061

WITHIN HERD CORRELATION BETWEEN PEDIGREE INDEX ($PI = 1/2 \text{ B.V. SIRE} + 1/2 \text{ B.V. DAM}$) AND YIELD
 DEVIATION (YD) FROM HERD CONTEMPORARY.

MODEL		CORRELATION PI - YD
HERD-YEAR	FIXED	.28
	RANDOM	.38
HERD-YEAR-SEASON (MOBILE)	FIXED	.27
	RANDOM	.40

EXPECTED WITHIN HERD CORRELATION PI - YD = .42

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CORRELATION BETWEEN SIRE PREDICTED BREEDING VALUE (BV) AND
CONTEMPORARY GROUP (CG) ESTIMATED EFFECTS.

MODEL		CORRELATION BV - CG
HERD-YEAR	FIXED	.07
	RANDOM	.34
HERD-YEAR-SEASON (MOBILE)	FIXED	.14
	RANDOM	.36

CONCLUSIONS:

1. USE OF C.G. AS RANDOM EFFECTS IN H-FRIESIAN
EVALUATION IN SPAIN IS NOT RECOMMENDED
 $(H_f + YS(H), ?)$.

2. MOBILE HYS DEPENDING ON HERD SIZE IMPLEMENTED
FOR NEXT EVALUATION.

HETEROGENEITY OF VARIANCE CORRECTION

- MANAGEMENT GROUPS (MG) WITH AT LEAST 30 OBSERVATIONS WERE CREATED FROM SIMILAR CONTEMPORARY GROUPS (C.G.) WITH A MAXIMUM OF 3 C.G.

- STANDARD DEVIATION FOR MG (σ_{MG}):

$$\sigma_{MG_i} = \left[\frac{N_i V_i + 30 PV_i + .5 N_{i-1} V_{i-1} + .5 N_{i+1} V_{i+1}}{N_i + 30 + .5 N_{i-1} + .5 N_{i+1}} \right]^{1/2}$$

N_i = NUMBER OF OBSERVATIONS IN MG_i .

V_i = VARIANCE OF MG_i .

PV_i = PREDICTED VARIANCE OF MG_i .

$$PV_i = (CV_{pop} * \mu_i)^2$$

CV = COEFFICIENT OF VARIATION (0.19 IN SPANISH POPULATION)

μ_i = PHENOTYPIC MEAN OF MG_i

(BAGNATO AND JENSEN PERSONAL COMMUNICATION)

IMPROVEMENT OF COMPUTING ALGORITHMS

- G.S. (100 ITERATIONES) CURRENTLY USED TO SOLVE BLUP-ANIMAL MODEL MME.
 - G.S. SLOW TO CONVERGE
 - UNCERTAINTY ABOUT STOPPING POINT
- SIMULATION APPLIED TO:
 - COMPARE NUMERICAL ACCURACY OF 6 ITERATIVE PROCEDURES:
 - GAUSS-SEIDEL (GS)
 - JACOBI-CONJUGATE GRADIENT (JCG)
 - SUCCESSIVE OBERRELAXATION (SOR)
 - SYMMETRIC SOR - CONJUGATE GRADIENT (SSORCG)
 - BLOCK (200 ECUATIONS) (BLO)
 - BLOCK (FIXED EFFECTS + 200 EQ) (BLOF).
 - STUDY "ACCURACY OF PREDICTIONS" DURING ITERATION UNDER THE 6 PROCEDURES
 - DETERMINE STOPPING CRITERIA BY COMPARING PROGRESS OF ACCURACY OF PREDICTIONS AND ESTIMATED GENETIC TREND WITH NUMERICAL CRITERIA USED TO MONITOR CONVERGENCE.

- DATA SETS:

- 2 SIZES (5000, 20000)

- SELECTED vs UNSELECTED

- NUMERICAL CRITERIA: $Cd = \frac{\|b^{n-1} - b^n\|}{\|b^n\|}$ (Cb = y)

$$Md = \max |b^{n-1} - b^n|$$

$$Cp = \frac{\|b^n - b\|}{\|b\|}$$

- ACCURACY OF EVALUATION: - ρ_{rr}

$$- MSD = \frac{\sum (\hat{u}_i - u_i)^2}{n}$$

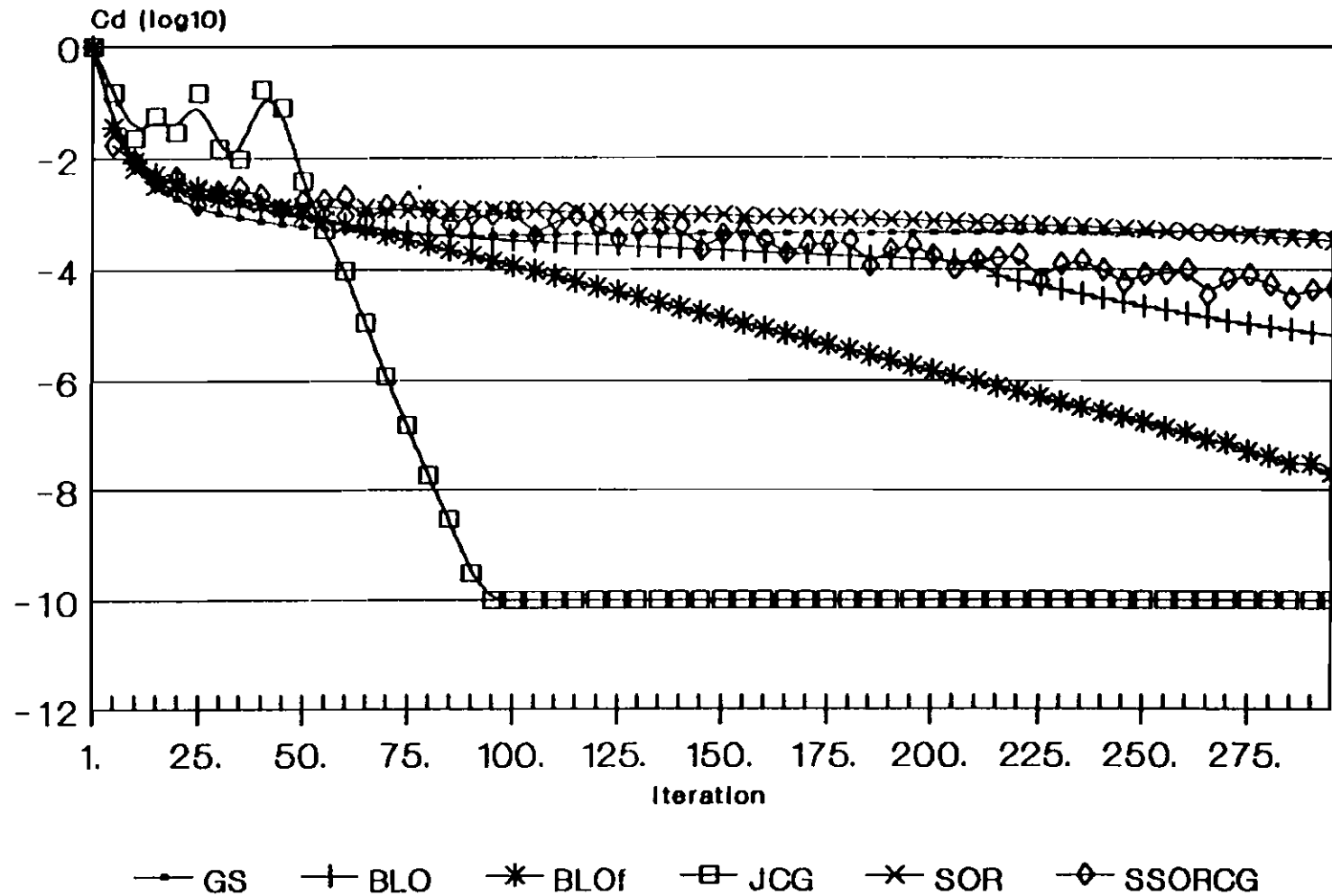
(u_i = additive genetic value)
(n = number of animals evaluated)

- ESTIMATED GENETIC TREND (b)

CONCLUSIONS:

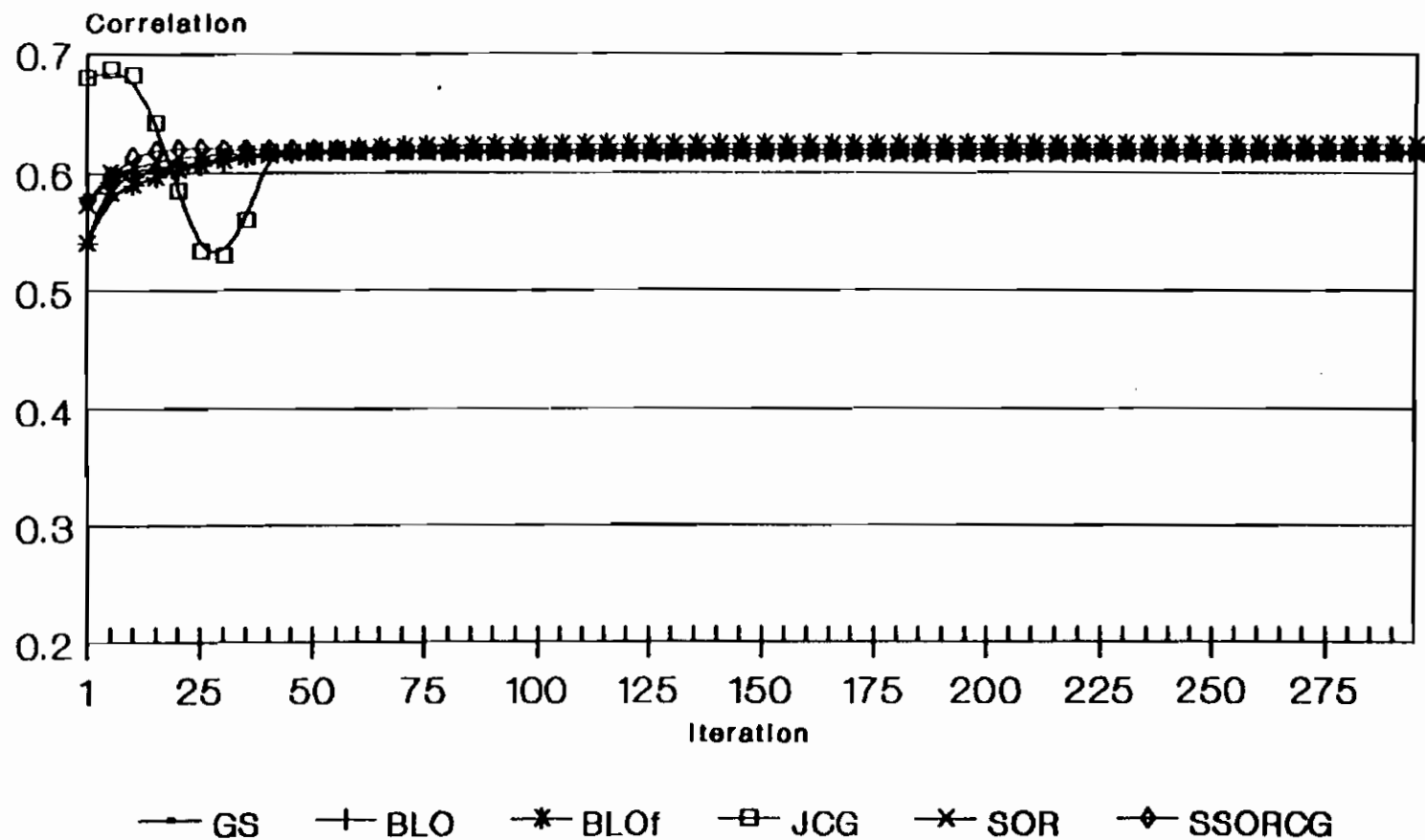
- JCG AND BLOCKING ALLOW MORE NUMERICAL ACCURACY BUT LARGER MEMORY (4x) AND CPU TIME (4-5x) REQUIRED.
- FIXING RANKING DOES NOT REQUIRE HIGH NUMERICAL ACCURACY ($C_d \geq 10^{-3}$).
- OBTAINING STABLE MSD AND b REQUIRED MORE NUMERICAL PRECISION ($C_d \geq 10^{-4}$).
- SLIGHTLY SLOWER CONVERGENCE FOR SMALLER AND SELECTED DATA SETS.
- POSSIBLE SHIFT TO JCG FOR FUTURE EVALUATIONS.

Changes in Cd during iteration



(size: 5000; selected pop'n)

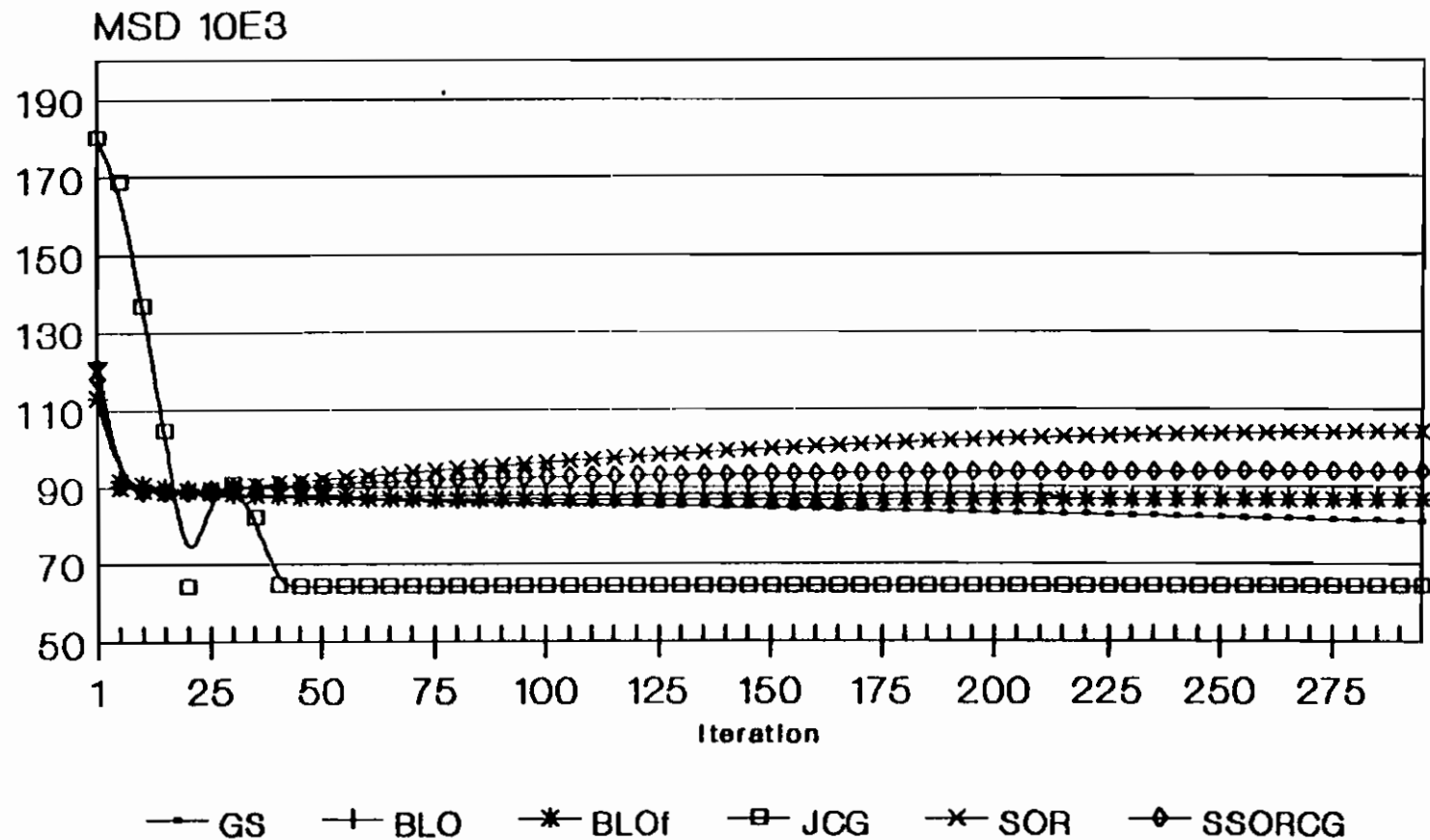
Correlation between real and predicted breeding value for bulls during iteration



(size: 5000; selected pop'n)

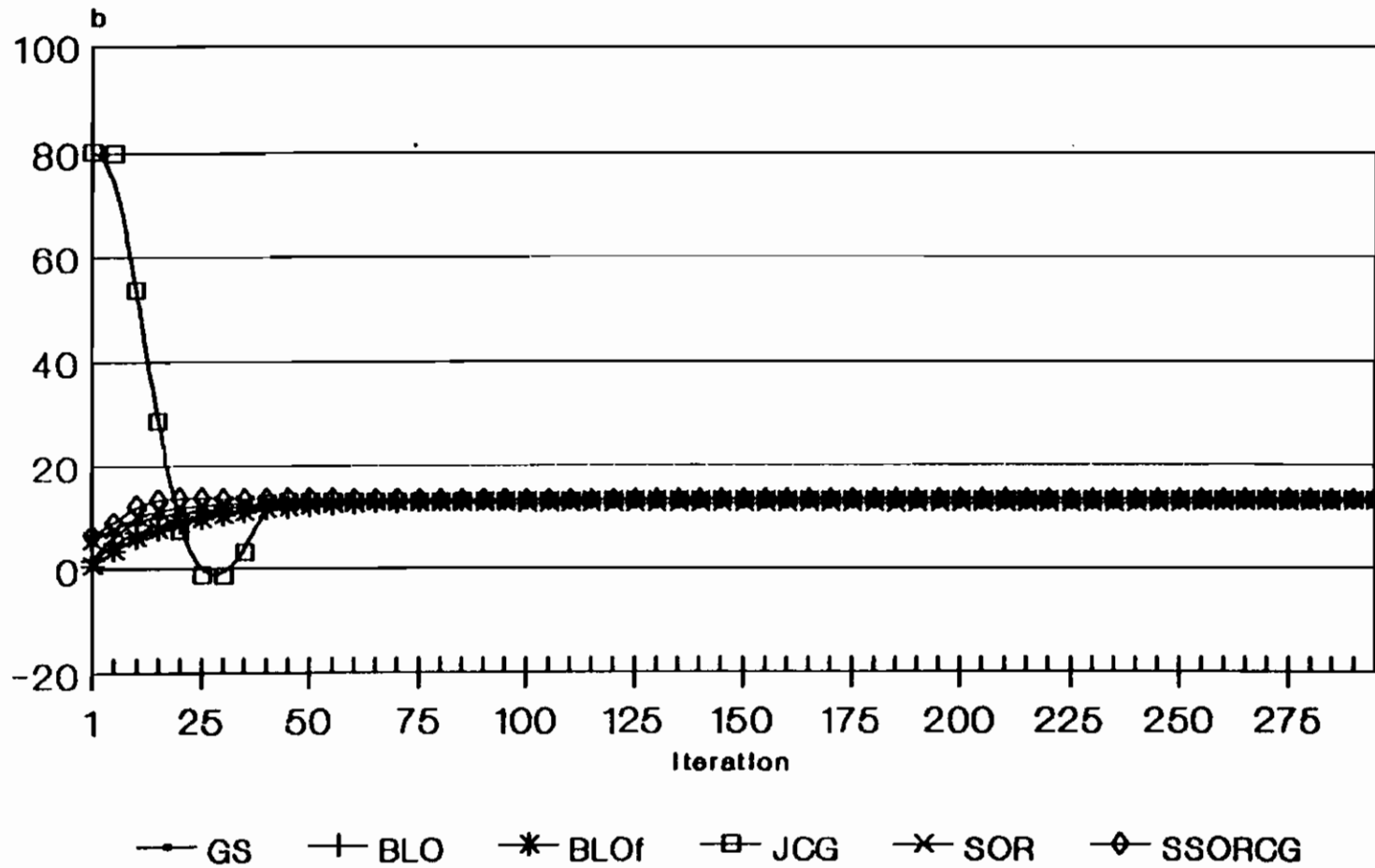
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Mean Squared Difference between real and predicted breeding value (MSD) for bulls during iteration



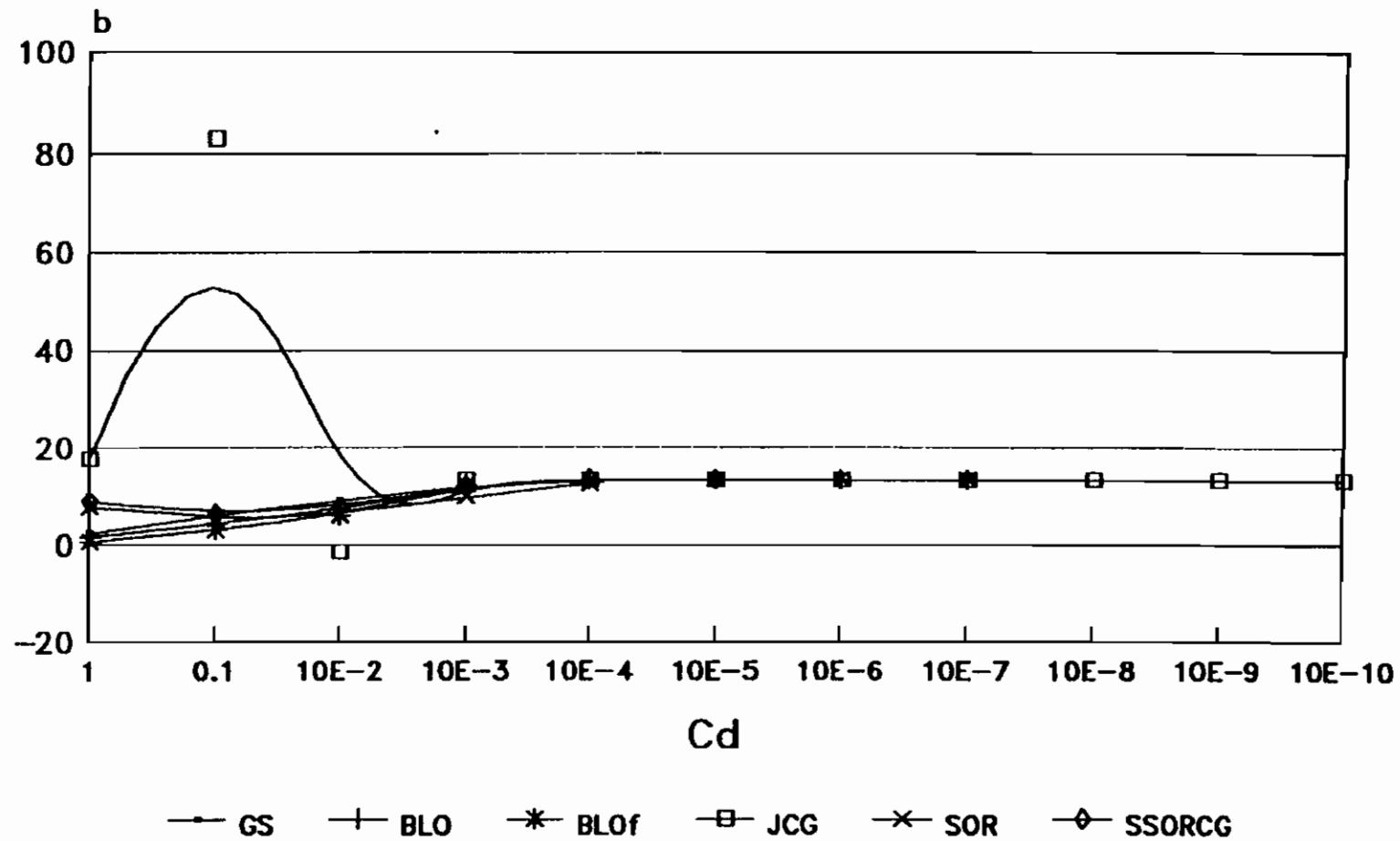
(size:5000; selected pop'n)

Estimated genetic trend (b) during iteration



(size: 5000; selected pop'n)

Estimated genetic trend (b) when Cd changed one digit of accuracy



(size: 5000; selected pop'n)