

Muller, H.J. Insertion of Foreign Chromosome into Homozygous Host Stock

(Stocks f and h need not contain any visible mutants)

- (1) For substituting 1^f for 1^h in $1^h; 2^h; 3^h; 4^h$ stock

P_1 $f \sigma \sigma \times ClB/+; T(2-3)S \text{ Cy}/+; ey^D/+ \text{ } \sigma \sigma$
 F_1 $1^f/ClB; 2^f; 3^f/T(2-3)S \text{ Cy}; 4^f/ey^D \text{ } \sigma \sigma$
 F_2 one σ $1^f; 2^h; 3^h/T(2-3)S \text{ Cy}; 4^h/ey^D$ $\times ClB/+; Cy/+; CC$
 (save for remating!) $D/+; ciD/+ \text{ } \sigma \sigma$
 (Different markers; Inversions)

F_3 Father (remated) $\times 1^f/ClB; 2^h/Cy; 3^h/CC, D; 4^h/ciD \text{ } \sigma$

F_4 $1^f; 2^h; 3^h; 4^h$ completely homozygous $\sigma \sigma$ $\times \sigma \sigma$

- (2) For substituting 2^f for 2^h in $1^h; 2^h; 3^h; 4^h$ stock

P_1 $f \sigma \sigma \times Pm/+; H/+; ey^D/+ \text{ } \sigma \sigma$ AND $h \sigma \sigma \times ClB/+; Cy/+; CC, D/+; ciD/+ \text{ } \sigma \sigma$
 F_1 $1^f; 2^f/Pm; 3^f/H; 4^f/ey^D \text{ } \sigma \sigma$ $\times 1^h/ClB; 2^h/Cy; 3^h/CC, D; 4^h/ciD \text{ } \sigma \sigma$
 F_2 one σ $1^h; 2^f/Cy; 3^h/H; 4^h/ey^D$ $\times 1^f/ClB; 2^f/Cy; 3^f/CC, D; 4^f/ciD \text{ } \sigma \sigma$
 (Recombination; save!) $ciD \text{ } \sigma \sigma$

F_3 Father $\times 1^h/ClB; 2^f/Cy; 3^h/CC, D; 4^h/ciD \text{ } \sigma \sigma$

F_4 $1^h; 2^f; 3^h; 4^h$ completely homozygous $\sigma \sigma$ and $\sigma \sigma$

- (3,4) Analogous schemes apply for 3 and 4 (selecting appropriate $F_2 \sigma$)

If desired, foreign autosome may first be made like that in homozygous stock except in regard to particular region. Method for this is shown below in scheme for combining invisible genes in the same chromosome.

Muller, H. J. Insertion of Desired Genes into Attached X's

P_1 $y \sigma \sigma \times T(1-4)B^S \sigma$ (Austin T; left part X attached to 4; rt end X carrying "Bar-super")

F_1 $y/XR, B^S \sigma \sigma \times m \text{ } g^2 \text{ } sd \text{ } f \text{ } \sigma \sigma$ (Let m, g^2 , sd, f be the genes to be inserted)

F_2 (Crossing-over) $y \cdot B^S/m \text{ } g^2 \text{ } sd \text{ } f \text{ } \sigma \sigma \times m \text{ } g^2 \text{ } sd \text{ } f \text{ } \sigma \sigma$

F_3 (Crossing-over) $m \text{ } g^2 \text{ } sd \text{ } f \cdot B^S/m \text{ } g^2 \text{ } sd \text{ } f \text{ } \sigma \sigma \times \sigma \sigma$

F_4 (Crossing-over) $m \text{ } g^2 \text{ } sd \text{ } f \text{ } Y \sigma$

In earlier method, used by L.V. Morgan et. al, genes were inserted by crossing-over, from separate X-chromosome containing them, into attached X's, in triploids.

Muller, H. J. Combination of Invisible Genes

Nearly same methods apply for setting desired chromosome

combinations of recessives that always must be worked with in the heterozygous condition.

1. Two invisibles in different autosomes (a in 2; b in 3)

P_1 $a/2 \sigma \times S/+; CC, D/+ \text{ } \sigma$ AND $b/3 \sigma \times Cy/+; CC, Dfd/+ \text{ } \sigma$

F_1 $a/S; CC, D/+ \text{ } \sigma$ $\times Cy/+; b/CC, Dfd \text{ } \sigma$

F_2 $a/Cy; b/CC, D \text{ } \sigma$ $\times a/Cy; b/CC, D \text{ } \sigma$

(balanced stock, if recessives have low viability or low fertility)

F_3 $a/a; b/b \text{ } \sigma \sigma$ and $\sigma \sigma$ (if homozygotes are viable and fertile)

F_4 (for chromosome 4, markers ciD and ey^D may be used in same way.)

2. Getting combination of invisibles in 1 and 2 (a in 1; b in 2)

P₁ a/1 ♂ X Cy/+ ♂ AND b/2 ♂ X ClB/+;Pm/+ ♀.
(double mated) y;Pm/+ ♀

F₁ a;Cy/+ ♂ (double-mated) X ClB/+;b/Pm ♀ and Xy;b/Pm ♀

F₂ a/ClB;b/Cy ♀ X a;b/Cy ♂ (balanced ♂)

F₃ a/a;b/b ♀ and a;b/b ♂ (if homo. viab. and fert.)

Analogous scheme using CC;D and CC,Dfd or Mc-2 for 3 applies for chromosomes 1 and 3.

3. Getting combination of two invisible genes (a and b) in same chromosome (in known positions). Let map order be a, v₁, v₂, b where v₁ and v₂ are visible genes between a and b.

P₁ a/+ X v₁v₂ AND b/+ X v₁v₂

F₁ a/+v₁v₂ ♀ X v₁v₂ ♂ AND b/+v₁v₂ ♀ X v₁v₂ ♂

F₂ (by crossing-over) a/+v₂/v₁v₂ X v₁/v₁v₂ b/+v₁v₂ ♀
(chance of losing a or b from ♀ obviated by interpolating cross of ♂ by individual containing inversions)

F₃ a/+v₂/v₁v₂ ♀ X v₁v₂ ♂

F₄ (by crossing-over) a/+b/v₁v₂ X D₁v ♀
(where D₁ is dominant linked and where v is either v₁ or v₂)

F₅ a b/D₁v ♂ X C,D₂ ♀ (where D₂ is a different linked dominant)

(lethal when homozygous, and C is an inversion preventing crossing-over).

F₆ a b/C,D₂ ♂ and ♀ (balanced ♂)

F₇ a b/a b ♂ and ♀ (homozygous ♂)

Combination of one invisible with one visible is correspondingly simpler.

Muller, H. J. and C. B. Bridges
Balanced Stocks

Given "1", a gene (or combination of linked genes) with low or no

viability or fertility when homozygous in ♂ or ♀ (or both, and "C" a genetic complex that is non-perpetuating or lethal when homozygous, that contains gene rearrangements which prevent crossing-over between "1" and "C" and that preferably contains a dominant marker, then the most usual balanced stock for maintenance of "1" without selection is:

1/C ♂ X 1/C ♀
1/1 (fails to perpetuate); C/C (dies); 1/C and C/1 (continue as before).

To construct simple balanced autosomal stocks (1/C)
Given 1 a/+ (or 1/a) ♂, where a = recessive marker and M = a dominant linked marker subject to crossing-over. Then:

P₁ M/+ X a/a

F₁ M/a ♀ X a/a ♂

F₂ M a/a ♀ X 1 a/+ ♂ (or 1/a ♂)

F₃ M a/1 a ♂ X C/+ ♀ (or M a/1 X C/+ ♀)

F₄ 1 a/C ♂ and ♀ (or 1/C ♂ and ♀)

The more used complexes represented above by "C" are as follows: