

Catalogue of Stocks

Chromosome 1

sr₁ z_b₄ P^{WW}
 sr₁ P^{WR}
 sr₁ P^{WR} an₁ gs₁ bm₂
 sr₁ P^{WR} an₁ bm₂
 sr₁ P^{RR} gs₁ bm₂
 sr₁ P^{WR} bm₂
 vp₅
 z_b₄ ms₁₇ P^{WW}
 z_b₄ ts₂ P^{WW} br₁ f₁ bm₂
 z_b₄ ts₂ P^{WW} bm₂
 z_b₄ P^{WW}
 z_b₄ P^{WW} br₁
 z_b₄ P^{WW} br₁ f₁ bm₂
 z_b₄ P^{WW} bm₂
 ts₂ P^{RR}
 ts₂ P^{WW} br₁ bm₂
 ts₂ P^{WW} bm₂
 P^{CR}
 P^{RR}
 P^{RW}
 P^{CW}
 P^{MO}
 P^{VV}
 P^{RR} as br₁ f₁ an₁ gs₁ bm₂

Chromosome 1 (Continued)

P^{RR} br₁ f₁ an₁ gs₁ bm₂
 P^{RR} an₁ ad₁ bm₂
 P^{RR} an₁ gs₁ bm₂
 P^{RR} ad₁ bm₂
 P^{WR} an₁ Kn bm₂
 P^{WR} an₁ ad₁ bm₂
 P^{WR} an₁ bm₂
 P^{WR} ad₁ bm₂
 P^{WR} br₁ vg
 P^{WR} br₁ f₁ gs₁ bm₂
 P^{WW} rs₂
 P^{WW} rs₂ br₁ f₁
 P^{WW} as br₁ f₁ bm₂
 P^{WW} hm₁ br₁ f₁
 P^{WW} br₁ f₁ ad₁ bm₂
 P^{WW} br₁ f₁ bm₂
 P^{WW} br₁ f₁ an₁ gs₁ bm₂
 as
 as rs₂
 rd-Hy
 br₁ f₁
 br₁ f₁ Kn
 br₁ f₁ Kn Ts₆
 br₁ f₁ Kn bm₂

Chromosome 1 (Continued)br₁ bm₂

Vg

Vg an₁ bm₂Vg br₂ bm₂bz₂^m m ; A₁ A₂ C₁ R Prbz₂^m M ; A₁ A₂ C₁ R Pran₁ bm₂an₁ bz₂ 6923 (apparent deficiency
including an₁ and bz₂)br₂br₂ bm₂tb₈₉₆₃

Kn

Kn Ts₆lw₁vp₈gs₁ bm₂Ts₆bm₂

id

nec₈₁₄₇ms₉ms₁₂ms₁₄mi₈₀₄₃ = mi₁D₈Chromosome 1 (Continued)

TB-1a (1L.20)

TB-1b (1S.05)

Chromosome 2ws₃ lg₁ gl₂ Bws₃ lg₁ gl₂ B skws₃ lg₁ gl₂ B sk fl₁ v₄ws₃ lg₁ gl₂ B ts₁ws₃ lg₁ gl₂ bws₃ lg₁ gl₂ b sk fl₁ v₄ws₃ lg₁ gl₂ b fl₁ v₄ws₃ lg₁ gl₂ b ts₁ws₃ lg₁ gl₂ b v₄

al

al lg₁al lg₁ gl₂ B sk v₄al lg₁ gl₂ b sk v₄lg₁lg₁ gl₂ Blg₁ gl₂ B gl₁₁lg₁ gl₂ B gs₂lg₁ gl₂ B gs₂ v₄lg₁ gl₂ B gs₂ Chlg₁ gl₂ B sk v₄lg₁ gl₂ B v₄lg₁ gl₂ blg₁ gl₂ b gs₂

Chromosome 2 (Continued)lg₁ gl₂ b gs₂ sk Chlg₁ gl₂ b gs₂ v₄lg₁ gl₂ b gs₂ v₄ Chlg₁ gl₂ b sklg₁ gl₂ b sk fl₁ v₄lg₁ gl₂ b sk v₄lg₁ gl₂ b wt₁ v₄lg₁ gl₂ b fl₁ v₄lg₁ gl₂ b fl₁ v₄ Chlg₁ gl₂ b v₄lg₁ gl₂ b v₄ Chlg₁ gl₂ wt₁lg₁ gl₂ w₃lg₁ gl₂ w₃ Chlg₁ gl₂ Chlg₁ b gs₂ v₄lg₁ Chd₅ = d₀₃₇₋₉B gl₁₁B ts₁gl₁₁ = gl₈₇₁₂wt₁mn₁fl₁ts₁v₄Chromosome 2 (Continued)w₃w₃ Ht₁w₃ ChHt₁ A sourceHt₁ B sourceba₂R₂; r₁ A₁ A₂ C₁

Ch

TB-2₆₂₇₀ (2S)TB-2₄₄₆₃ (2L)

Primary Trisomic 2

Chromosome 3cr₁cr₁ d₁cr₁ d₁ Lg₃cr₁ ts₄ na₁d₁ Tall = d₆₀₁₆ = tnd₁ rt₁ Lg₃d₁ Rf₁ lg₂d₁ ys₃d₁ ys₃ Rgd₁ Lg₃d₁ Rg ts₄ lg₂d₁ pmd₁ ts₄ lg₂d₁ ts₄ lg₂ a₁^m; A₂ C₁ R Dt₁

Chromosome 3 (Continued)ra₂ra₂ ys₃ Lg₃ Rgra₂ ys₃ Rgra₂ Rg lg₂ra₂ pm lg₂ra₂ lg₂

Cg

cl₁cl₁ Cl₂cl₁ Cl₃clp Cl₄rt₁ys₃ys₃ Lg₃ys₃ gl₆ lg₂ a₁^m et; A₂ C₁ R Dt₁ys₃ ts₄Lg₃Lg₃ Rggl₆ lg₂ A₁; A₂ C₁ Rgl₆ lg₂ A^b et; A₂ C₁ R Dt₁gl₆ lg₂ a₁^m et; A₂ C₁ R dt₁gl₆ lg₂ a₁^m et; A₂ C₁ R Dt₁ts₄ts₄ ba₁ na₁ts₄ lg₂ a₁^m; A₂ C₁ R Dt₁ts₄ lg₂ gl₇Chromosome 3 (Continued)ts₄ na₁ a₁^m et; A₂ C₁ R Dt₁ts₄ a₁^m; A₂ C₁ R Dt₁ba₁lg₂ A^b et; A₂ C₁ R Dt₁lg₂ a₁^m sh₂ et; A₂ C₁ R Dt₁lg₂ a₁^m et; A₂ C₁ R dt₁lg₂ a₁^m et; A₂ C₁ R Dt₁lg₂ a₁st sh₂ et; A₂ C₁ R Dt₁lg₂ a₁st et; A₂ C₁ R Dt₁na₁A₁ sh₂; A₂ C₁ R B Pl dt₁A₁^d-31; A₂ C₁ RA₁^d-31; A₂ C₁ R pr dt₁A₁^d-31; A₂ C₁ R B Pl dt₁A₁^d-31; A₂ C₁ R Dt₁A₁^d-31; A₂ C₁ R pr Dt₁A₁^d-31 sh₂; A₂ C₁ R B Pl dt₁A₁^d-31 sh₂; A₂ C₁ R Dt₁A₁^d-31 sh₂; A₂ C₁ R B Pl Dt₁A₁^d-31 et; A₂ C₁ R Dt₁a₁^m; A₂ C₁ R dt₁a₁^m; A₂ C₁ R B Pl dt₁a₁^m; A₂ C₁ R Dt₁a₁^m; A₂ C₁ R B Pl Dt₁a₁^m sh₂; A₂ C₁ R B Pl dt₁a₁^m sh₂; A₂ C₁ R B Pl Dt₁

Chromosome 3 (Continued)

a_1^m et; $A_2 C_1 R D t_1$
 a_1^{st} ; $A_2 C_1 R D t_1$
 $a_1^{st} sh_2$; $A_2 C_1 R D t_1$
 $a_1^{st} sh_2$ et; $A_2 C_1 R D t_1$
 a_1^{st} et; $A_2 C_1 R D t_1$
 a_1^p et; $A_2 C_1 R dt_1$
 a_1^p et; $A_2 C_1 R B P l D t_1$
 $a_1 - xl$
 $a_1 Ga_7$; $A_2 C_1 R$
 $sh_2 = bt_{60-156} = sh_{Garwood}$

 $v p_1$ $R p_3$ $g l_{12}$

TB-3a (3L.10)

TB-3b (3S.50)

Primary Trisomic 3

Chromosome 4 $R p_4$ $G a_1$ $G a_1 su_1$ $G a_1^s$ $G a_1^s bt_2$

st

st $T s_5$ st $f l_2$ st $T s_5 su_1$ Chromosome 4 (Continued)

$T s_5$
 $T s_5 f l_2$
 $T s_5 su_1$
 $T s_5 su_1 z b_6$
 $T s_5 su_1 z b_6 o_1$
 $T s_5 Tu$
 $l a su_1 Tu g l_3$
 $l a su_1 g l_3$
 $l a su_1 g l_3 c_2$; $A_1 A_2 C_1 R$
 $l a su_1 g l_3 o_1$

 $f l_2$ $f l_2 su_1$ su_1 su_1^{am} $su_1 b m_3$ $su_1 z b_6$ $su_1 z b_6 Tu$ $su_1 z b_6 C_2^{Idf (Active-1)}$; $A_1 A_2 C_1 R$ $su_1 g l_4$ $su_1 g l_4 Tu$ $su_1 g l_4 j_2$ $su_1 g l_4 o_1$ $su_1 j_2$ $su_1 g l_3$ $su_1 g l_3 o_1$ $su_1 o_1$

Chromosome 4 (Continued)

$bt_2 = bt_4 = bt_{60-158} = bt_{Williams}$

$bt_2\ gl_4$

$bt_2\ gl_4\ j_2$

$gl_4 = gl_{16} = gl_{Stadler}$

Tu

Tu¹ 1st

Tu¹ 2nd

Tu^d

Tu^{md}

Tu gl₃

j₂

j₂ c₂; A₁ A₂ C₁ R

j₂ c₂; A₁ A₂ C₁ R

v₈

gl₃

gl₃ dp

c₂; A₁ A₂ C₁ R

c₂; A₁ A₂ C₁ R

c₂ Idf (Active-1); A₁ A₂ C₁ R

v₁₇

gl₇

o₁

ra₃

TB-4a (4S.20)

TB-4₄₆₉₂ (4L)

Primary Trisomic 4

Chromosome 5

lu₁

lu₁ sh₄

ms₁₃

gl₁₇

gl₁₇ A₂ pr; A₁ C₁ R

gl₁₇ a₂; A₁ C₁ R

A₂ vp₇ pr; A₁ C₁ R

A₂ bm₁ pr; A₁ C₁ R

A₂ bm₁ pr ys₁; A₁ C₁ R

A₂ bm₁ pr ys₁ eg; A₁ C₁ R

A₂ bt₁ pr; A₁ C₁ R

A₂ sh₃ pr ys₁; in A₁ C₁ R

A₂ v₃ pr; A₁ C₁ R

A₂ pr na₂; A₁ C₁ R

A₂ pr ys₁; A₁ C₁ R

a₂; A₁ C₁ R

a₂; A₁ C₁ R B Pl

a₂ bm₁ bt₁ bv₁ pr; A₁ C₁ R

a₂ bm₁ bt₁ pr; A₁ C₁ R

a₂ bm₁ bt₁ pr ys₁; A₁ C₁ R

a₂ bm₁ pr ys₁; A₁ C₁ R

a₂ bm₁ pr v₂; A₁ C₁ R

a₂ bt₁ v₃ pr; A₁ C₁ R

a₂ bt₁ pr; A₁ C₁ R

a₂ bt₁ v₂; A₁ C₁ R

a₂ v₃ pr; A₁ C₁ R

Chromosome 5 (Continued)a₂ pr; A₁ C₁ Rvp₂vp₂ gl₈vp₇bm₁ yg₁bt₁ = bt_{Alex-Krug} = bt_{Krug6-1303-2}= bt_{Vineyard} = bt₆₋₇₈₃₋₇ =sh_{Eldridge} = bt_{C103} = sh₃ = sh₅ms₅v₃ = v₈₉₈₃

td ae

ae

sh₄gl₈ = gl₁₀na₂lw₂ys₁

eg

v₂yg₁ms₁₃v₁₂lw₃ lw₄

Primary Trisomic 5

Chromosome 6rgd po y₁rgd Y₁po = ms₆po y₁ plpo Y₁ ply₁ = pb₁ = w^my₁ l₁₀y₁ l₄₉₂₀y₁ w₈₈₉₆y₁ pb₄y₁ pb₄ ply₁ pb₄ Ply₁ ms-siy₁ at-si = ms-siy₁ wi Ply₁ pg₁₁; wx pg₁₂y₁ pg₁₁; wx pg₁₂Y₁ pg₁₁; wx pg₁₂Y₁ pg₁₁; wx pg₁₂y₁ ply₁ Ply₁ Pl Bh; c₁ sh₁ wx A₁ A₂ Ry₁ su₂y₁ l₄₁₂₀Y₁ l₁₀Y₁ pb₄

Chromosome 6 (Continued)

$\text{Y}_1 \text{ wi pl}$
 $\text{Y}_1 \text{ wi Pl}$
 $\text{Y}_1 \text{ su}_2$
 wi
 $\text{pg}_{48-040-8} = \text{pg}_{11} \text{ pg}_{12}$
 $\text{pg}_{6656} = \text{pg}_{11} \text{ pg}_{12}$
 $\text{yg}_{6853} = \text{pg}_{11} \text{ pg}_{12}$
 $\text{Pl Dt}_2; \text{a}_1 \text{ A}_2 \text{ C R}$
 $\text{pl sm; P}^{\text{RR}}$
 $\text{Pl sm; P}^{\text{RR}}$
 $\text{Pl sm py; P}^{\text{RR}}$
 Pt
 w_1
 $\text{w}_{8657} = \text{w}_{025-12} = \text{w}_{035-2} =$
 $\text{w}_{5946} = \text{w}_{8050} = \text{w}_{6853} =$
 $\text{w}_{1-74302}$

Primary Trisomic 6

Chromosome 7

$\text{Hs o}_2 \text{ v}_5 \text{ ra}_1 \text{ gl}_1$
 In^D
 $\text{In}^D \text{ o}_2 \text{ v}_5 \text{ ra}_1 \text{ gl}_1$
 $\text{In}^D \text{ gl}_1$
 o_2
 $\text{o}_2 \text{ v}_5$
 $\text{o}_2 \text{ v}_5 \text{ ra}_1 \text{ gl}_1$

Chromosome 7 (Continued)

$\text{o}_2 \text{ v}_5 \text{ ra}_1 \text{ gl}_1 \text{ Tp}_1$
 $\text{o}_2 \text{ v}_5 \text{ ra}_1 \text{ gl}_1 \text{ ij}$
 $\text{o}_2 \text{ v}_5 \text{ gl}_1$
 $\text{o}_2 \text{ ra}_1 \text{ gl}_1 \text{ ij}$
 $\text{o}_2 \text{ gl}_1$
 $\text{o}_2 \text{ gl}_1 \text{ sl}_1$
 $\text{o}_2 \text{ bd}$
 in
 in gl_1
 v_5
 vp_9
 $\text{vp}_9 \text{ gl}_1$
 $\text{ra}_1 \text{ gl}_1 \text{ ij bd}$
 $\text{gl}_1 = \text{gl}_9$
 gl_1^m
 $\text{gl}_1 \text{ Tp}_1$
 $\text{gl}_1 \text{ o}_5$
 $\text{gl}_1 \text{ g}_2$
 Tp_1
 ij
 Bn
 bd
 Pn
 o_5
 g_2
 va_1

Chromosome 7 (Continued)Dt₃; a₁ A₂ C₁ Rv₈₆₄₇yel₇₇₄₈

TB-7b (7L.30)

Primary Trisomic 7

Chromosome 8gl_gv₁₆ = v₈₆₆₁v₁₆ j₁v₁₆ ms₈ j₁nec₆₆₉₇ = sie₇₇₄₈ = nec₀₂₅₋₄v₁₆ ms₈ j₁ gl_g

TB-8a (8L.70)

Primary Trisomic 8

Chromosome 9yg₂ C₁ sh₁ bz₁; A₁ A₂ Ryg₂ C₁ sh₁ bz₁ wx; A₁ A₂ Ryg₂ C₁^I sh₁ bz₁ wx; A₁ A₂ Ryg₂ C₁ sh₁ bz₁ wx K^L9; A₁ A₂ Ryg₂ C₁ bz₁ wx; A₁ A₂ Ryg₂ c₁ sh₁ bz₁ wx; A₁ A₂ Ryg₂ c₁ sh₁ wx; A₁ A₂ Ryg₂ c₁ sh₁ wx gl₁₅; A₁ A₂ Ryg₂ c₁ sh₁ wx gl₁₅ K^L9; A₁ A₂ R^Gyg₂ c₁ bz₁ wx; A₁ A₂ Rwd-Ring C₁^I; A₁ A₂ RChromosome 9 (Continued)C₁ sh₁ bz₁; A₁ A₂ RC₁ sh₁ bz₁ wx; A₁ A₂ RC₁ sh₁ bz₁ wx gl₁₅ bm₄; A₁ A₂ RC₁ sh₁; A₁ A₂ RC₁ sh₁ wx; A₁ A₂ RC₁ wx ar; A₁ A₂ RC₁^I sh₁ wx v₁; A₁ A₂ RC₁ sh₁ wx K^L9; A₁ A₂ RC₁ sh₁ ms₂; A₁ A₂ RC₁ bz₁ wx; A₁ A₂ RC₁ Ds wx; A₁ A₂ R y₁C₁ Ds wx; A₁ A₂ R prC₁^I Ds wx; A₁ A₂ RC₁^I; A₁ A₂ RC₁; A₁ A₂ RC₁; A₁ A₂ R B PlC₁ wx; A₁ A₂ RC₁ wx; A₁ A₂ R B PlC₁ wx; A₁ A₂ R b PlC₁ wx; A₁ A₂ R B plC₁^I wx; A₁ A₂ R y₁C₁^I wx; A₁ A₂ R y₁ B plC₁ wx ar da; A₁ A₂ RC₁ wx v₁; A₁ A₂ RC₁ wx v₁; A₁ A₂ R PlC₁ wx gl₁₅; A₁ A₂ R

Chromosome 9 (Continued)

c₁ wx gl₁₅; A₁ A₂ R pr
 c₁ wx Bf₁; A₁ A₂ R
 c₁ sh₁ bz₁ wx; A₁ A₂ R y₁
 c₁ sh₁ wx; A₁ A₂ R
 c₁ sh₁ wx v₁; A₁ A₂ R
 c₁ sh₁ wx gl₁₅; A₁ A₂ R
 c₁ sh₁ wx gl₁₅ bk₂; A₁ A₂ R
 c₁ sh₁ wx gl₁₅ Bf₁; A₁ A₂ R
 c₁ sh₁ wx bk₂; A₁ A₂ R
 c₁; A₁ A₂ R
 c₁ wx; A₁ A₂ R y₁
 c₁ wx v₁; A₁ A₂ R
 c₁ wx gl₁₅; A₁ A₂ R
 c₁ wx Bf₁; A₁ A₂ R
 c₁ wx bk₂; A₁ A₂ R
 sh₁ = sh₆₃₄₉ = sh₆₀₋₁₅₅ = sh_{67-Vineyard}
 sh₁ bp₁ wx; P^{RR}
 sh₁ bp₁ wx; P^{RW}
 sh₁ wx v₁
 bp wx; P^{RR}
 bp wx; P^{RW}
 bp wx; P^{WW}
 lo₂
 wx = wx^a
 w₁₁
 wx d₃

Chromosome 9 (Continued)

wx pg₁₂; y₁ pg₁₁
 wx pg₁₂; y₁ pg₁₁
 wx pg₁₂; y₁ pg₁₁
 wx pg₁₂; y₁ pg₁₁
 wx v₁
 wx bk₂
 wx bk₂ bm₄
 wx Bf₁
 wx Bf₁ bm₄
 d₃ = d₀₁₅₋₁₂ = d₀₇₂₋₇ = d_{fg} =
 d₈₀₅₄ = d_{x-ray}
 v₁ = v₈₅₈₇
 gl₁₅
 gl₁₅ bm₄
 bk₂ wc
 wc
 bm₄
 l₆
 l₆; l₁
 l₇
 l₇; l₁
 yel₀₃₄₋₁₆
 yg z_b₅₅₈₈
 w₄₈₈₉
 w₈₈₈₉
 w₈₉₅₁

Chromosome 9 (Continued)w₈₉₅₀w nl₀₃₄₋₅w₉₀₀₀

TB-9a (9L.40)

TB-9b (9S.40)

Primary Trisomic 9

Chromosome 10

oy

oy bf₂oy bf₂ R; A₁ A₂ C₁oy bf₂ ms₁₀oy du R; A₁ A₂ C₁oy du r; A₁ A₂ C₁oy zn₁

Og

Og du R; A₁ A₂ C₁bf₂bf₂ li g₁ r; A₁ A₂ C₁bf₂ g₁ R sr₂; A₁ A₂ C₁bf₂ g₁ r sr₂; A₁ A₂ C₁nl g₁ R; A₁ A₂ C₁y₉li zn₁ g₁ r; A₁ A₂ C₁li g₁ R; A₁ A₂ C₁li g₁ r; A₁ A₂ C₁li g₁ r v₁₈; A₁ A₂ C₁Chromosome 10 (Continued)

du

du g₁ r; A₁ A₂ C₁zn₁zn₁ g₁ r; A₁ A₂ C₁Tp₂ g₁ r; A₁ A₂ C₁g₁ R sr₂; A₁ A₂ C₁g₁ r; A₁ A₂ C₁g₁ r sr₂; A₁ A₂ C₁g₁ R^E sr₂; A₁ A₂ C₁g₁ R^E sr₂ v₁₈; A₁ A₂ C₁g₁ R^E KLO; A₁ A₂ C₁g₁ R^r sr₂; A₁ A₂ C₁g₁ R^r KLO; A₁ A₂ C₁g₁ r^r sr₂; A₁ A₂ C₁E^j r^r; A₁ A₂ C₁E^j r^r sr₂; A₁ A₂ C₁r sr₂ l₁; A₁ A₂ C₁R^E; A₁ A₂ C₁r^E sr₂; A₁ A₂ C₁r KLO; A₁ A₂ C₁r^E; A₁ A₂ C₁r^r; A₁ A₂ C₁R^{mb}; A₁ A₂ C₁R^{nj}; A₁ A₂ C₁R^r; A₁ A₂ C₁

Chromosome 10 (Continued)R^r
Boone; A₁ A₂ C₁R^{lsk}; A₁ A₂ C₁R^{sk mc.2}; A₁ A₂ C₁R^{sk}; A₁ A₂ C₁Rst; A₁ A₂ C₁

Lc

w₂w₂ l₁l₁v₁₈

Mt

yel₈₉₆₂l₁ yel₅₃₄₄yel₈₇₂₁yel₈₄₅₄yel₈₇₉₃w₇₇₄₈ = w₈₉₀₅

TB-10a (1OL.35)

Primary Trisomic 10

Unplaced Genes

dv

dy

el

g^l₁₄

h

l₃Unplaced Genes (Continued)l₄Rs₁v₁₃ws₁ ws₂

ub

zb₁zb₂zb₃zn₂l₄₉₂₃

"necrotic 8376" (seedling)

Multiple Gene StocksA₁ A₂ C₁ R^g Pr B PlA₁ A₂ C₁ R^r Pr B PlA₁ A₂ C₁ R PrA₁ A₂ C₁ R Pr wxA₁ A₂ C₁ R Pr wx g^l₁

Multiple Gene Stocks (Continued)

$A_1 A_2 C_1 R Pr wx y_1$
 $A_1 A_2 C_1 R pr$
 $A_1 A_2 C_1 R pr y_1 gl_1$
 $A_1 A_2 C_1 R pr y_1 wx$
 $A_1 A_2 C_1 R pr y_1 wx gl_1$
 $A_1 A_2 c_1 R Pr y_1 wx$
 $A_1 A_2 C_1 r Pr y_1 wx$
 $a_1 su_1 A_2 C_1 R$
 $bm_2 lg_1 a_1 su_1 pr y_1 gl_1 j_1 wx g_1$
 colored scutellum
 $lg_1 gl_2 wt_1 ; a_1 Dt_1 A_2 C R$
 $lg_1 su_1 bm_2 y_1 gl_1 j_1$
 $su_1 y_1 wx a_1 A_2 C_1 R^E pr$
 $y_1 wx gl_1$
 $hm_1 hm_2$
 $ts_2; sk$
Popcorns
 Amber Pearl
 Argentine
 Black Beauty
 Hulless
 Ladyfinger
 Ohio Yellow
 Red
 South American
 Strawberry

Popcorns (Continued)

Supergold
 Tom Thumb
 White Rice
Exotics and Varieties
 Black Mexican Sweet Corn
 (with B-chromosomes)
 Black Mexican Sweet Corn
 (without B-chromosomes)
 Knobless Tama Flint
 Knobless Wilbur's Flint
 Gaspe Flint
 Gourdseed
 Maiz chapolote
 Papago Flour Corn
 Parker's Flint
 Tama Flint
 Zapaluta chica
Tetraploid Stocks
 P^{RR}
 P^{VV}
 Ch
 B Pl
 $a_1 A_2 C_1 R Dt_1$
 su_1
 $pr; A_1 A_2 C_1 R$
 y_1
 gl_1

Tetraploid Stocks (Continued)

ij

 $y_1\ sh_1\ wx$ $sh_1\ bz_1\ wx$

wx

 g_1 $A_1\ A_2\ C_1\ R$ $A_1\ A_2\ C_1\ R\ B\ Pl$ Cytoplasmic Steriles and RestorersWF9 - (T) $rf_1\ rf_2$

N6 (S)

WF 9 $rf_1\ rf_2$ N6 $rf_1\ Rf_2$ R213 $Rf_1\ rf_2$ Ky21 $Rf_1\ Rf_2$

These combinations are also available
in other inbred backgrounds.

Chromosome rearrangements

The following rearrangements are being maintained primarily for use in determining the chromosome locations of new traits. All are marked with closely-linked endosperm or seedling traits.

The cytological positions of Inv 2a were determined by Dr. Morgan; those of Inv 9a were determined by Dr. Li. The indicated interchange points of the reciprocal translocations are taken from published work of Dr. Longley.

Inversions

*gl ₂	Inv 2a (also available with Ch)	2S.7; 2L.8
*wx ²	Inv 9a	9S.7; 9L.9

Reciprocal translocations

*wx 1-9c	1S.48; 9L.22
*wx 1-9 4995	1L.19; 9S.20
*wx 1-9 8389	1L.74; 9L.13
*wx 2-9b	2S.18; 9L.22
*wx 3-9c	3L.09; 9L.12
wx 3-9 5775	3L.09; 9S.24
*wx 4-9b	4L.90; 9L.29
*wx 4-9 5657	4L.33; 9S.25
*wx 4-9g	4S.27; 9L.27
*wx 5-9a	5L.69; 9S.17
*wx 5-9c	5S.07; 9L.10
*wx 5-9d	5L.14; 9L.10
wx 5-9 4817	5L.06; 9S.07
*wx 6-9a	6S.79; 9L.40
*wx, y 6-9b	6L.10; 9S.37
wx 6-9 4505	6L.13; 9 cent
wx 6-9 4778	6S.80; 9L.30
*wx 7-9a	7L.63; 9S.07
*wx or gl ₁ 7-9 4363	7 cent; 9 cent
*wx 8-9d	8L.09; 9S.16
*wx 8-9 6673	8L.35; 9S.31
*wx 9-10b	9S.13; 10S.40

*These constitute a basic series of twenty rearrangements for use in locating unplaced genes.