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## SUMMARY:

TITLE: A Plan for breeding *Pinus radiata* in Western Australia.

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The long term nature of forest breeding implies a commitment by the organisation for its financing and staffing priorities and pressures for research will always lead to speculation about the need to continue and to the actual economic value of the research. Published results show increases in yield and improvements in quality for our *Pinus* *pinaster* and increased disease resistance of *P. radiata* but although these results are acknowledged, they go unnoticed because they are not quantified in terms of money.

The management model for *P. pinaster*, embracing the concepts of intensive management and applied genetics for more production and greater efficiency shows that tree improvement is a paying proposition. And there is an extra payoff in the Sunklands where increased disease resistance of *P. radiata* to Phytophthora *cinnamomi* is a uniquely genetic effect that could not be achieved by any other means. International investment, "perhaps even the best of all possible forestry investment", "tree improvement as a good economic future support to the best of all possible forestry investments" (Zobel, 1978).

Most pine plantations in Australia are currently grown from orchard seed and the main task is to now improve the genetic quality of this seed. Most of the future improvement in the genetic quality of Australia's orchard seed will come from a new generation of orchards begun in the early 1980's. These orchards will contain mostly 1.5 or 2nd generation clones.

The solution of many of the problems facing forestry, particularly the conflicts between alternative uses of the same forest lies in intensive management, encompassing genetic improvement, of part of the forest area for wood production. There is an opportunity to concentrate wood production on a smaller land base permitting the larger remaining areas to be managed more flexibly toward other objectives.

## INTRODUCTION

At this time when our organisation is undergoing major restructuring and Land management functions are being discussed, it is opportune to review the Department's pine tree breeding programmes. A plan for the breeding of radiata pine is presented for discussion and endorsement.

Nearly all of the Australian programmes breeding *Pinus radiata* have reached the end of the forest generation. The lack of an overall plan and particularly the uncertainty of the breeding methods for future generations, has resulted in a 20 year generation interval for this first generation. A promised mass production of seed to establish plantations of fast-growing, fine-branched, narrow-crowned trees seems unlikely until at least the third generation of seed orchards (Eldridge, 1983). To maximize commercial gains, structured breeding plans incorporating reduced generation intervals are required.

Early results from first generation seed orchard progeny are clearly superior to that of non-orchard sources. Pedreick and Eldridge (1983) reported that initial orchards have yielded plants that grow at least 10 per cent faster in volume than unimproved plants and the gains many even increase with age. Trees of orchard origin were also straighter and had thinner and more frequent branches than the controls.

As soon as progeny information is available about the clones used in the first generation orchard, the genetic quality of the orchard can be improved by removing (culling) the poor performance clones. For example, Pederick and Eldridge (1983) calculated an increased average gain in volume of young trees derived from the rogued (culled) orchard from 7.4% to 10.2% by removing 12 poor performance clones from a 50 clone seed orchard in Victoria. They were also able to increase this gain to about 16% (i.e. double) by the establishment of a "1.5 generation" orchard (knowledge of breeding values of clones).

Data from Western Australian progeny tests and yield trials are in accord with reported gains from Australian tests. This is discussed in the section on Advances in the first phase of the W.A. breeding plan. Some data is shown in Appendix 1.

The first generation of breeding radiata pine in Australia has made a distinct improvement in growth rate and tree form. In a recent economic analysis of the gains achieved from a radiata pine orchard, Wright and Eldridge (1983) calculated that, by spending \$10 to \$20 per hectare on improving seed, net present values of plantations would be several hundred dollars higher and internal rates of return would exceed 20%. Continued investment in breeding seems certain to result in further genetic improvement in second, third and subsequent generations.

A plan for the breeding of radiata pine in Western Australia is presented. It is logically divided into five segments that detail the historical time frame, major objectives and progress of the plan. The plan has, and will continue to rely upon considerable co-operation and co-ordination with other Australian states breeding radiata pine to achieve maximum increasing gains and to provide a broad genetic base to ensure these advances in perpetuity.

The major segment of the plan is phase 5 (research and development). Objectives are to continue the cumulative development). Objectives are to maintain genetic diversity in established plantations, and to maintain genetic diversity in the ability to adapt to changing environmental and economic circumstances. Shortened generation intervals of about 14 years are proposed.

111 southern Australian states began tree breeding programmes for *Pinus radiata* during this period. Programme objectives were for increased yields of high quality wood through multiple-trait selection. The foundation of the programme was large areas were searched for very few plus trees on the plus trees. From this limited area, 20 plus trees were selected. Section material was collected for grafting and establishment at the Mundaring clone bank. Grafts were planted at the Mundaring clone bank and Victoria (13), South Australia (11) and New Zealand (3). As sections from Canberra (10), New South Wales (9), 1962-64; W.A. Plantations at Mundaring, Margaret River and Grimwade over 20 years of age were searched for and additional plus trees were selected. Section material was collected for grafting and establishment at the Mundaring clone bank. Grafts were planted at the Mundaring clone bank and Victoria (13), South Australia (11) and New Zealand (3). 1964-65; additional plus tree selections were imported as sections from Canberra (10), New South Wales (9), 1969-74; progeny tests were mainly based on open-pollinated seed in seed orchards. \* The first tests, planted in 1969 (32 families - Collie) and 1971 (60 families - Collie and Blackwood Valley) used polycross seed of Canberra, South Australian and Victorian origin. Test objectives were to rank parents for breeding values, demonstrate genetic gains and examine genotype x environment interactions.

#### Breeding population:

1964-65; additional plus tree selections were imported as sections from Canberra (10), New South Wales (9), 1969-74; progeny tests were mainly based on open-pollinated seed in seed orchards. \* The first tests, planted in 1969 (32 families - Collie) and 1971 (60 families - Collie and Blackwood Valley) used polycross seed of Canberra, South Australian and Victorian origin. Test objectives were to rank parents for breeding values, demonstrate genetic gains and examine genotype x environment interactions.

#### Gene pool:

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Yield trials were planted in 1973 at Grimwade, Blackwood Valley and Yanchep. Test objective was to evaluate the potential of improved seed on the establishment and management of plantations. Treatments included orchard, seed production stand, crop tree and routine seed planted on large plots at either 1000 or 2000 spha.

1967-71; seed production stands were developed at Grimmelwade A section. Together with third thinnings come collection, they were to provide an interim source of improved seed prior to the availability of orchard seed.

1965-68; a 4.5 ha seed orchard using 40 clones from the gene bank was established at Glenelgote. This was later abandoned, in 1976, because of very low cone yields.

### Production population:

Local selections were included in comprehensive progeny tests for the first time in 1974. The eleven W.A. clones used in the Glenelgale orchard were collected as polycross seed and trials were planted on major plantation site types at Donnybrook sunkenlands, Blackwood Valley (later destroyed by rabbits) and yellow sands at Yanchep. Principal objective was the ranking of parents for breeding value, as well as demonstration of gains and evaluation of genotype x environment interactions.

1972 was devoted to the planting of the International gene pool at Grimwade and Kirup. Seed of 319 seed lots, mainly wind-pollinated progenies collected in seed orchards, was provided from 13 different breeding programmes. Test objectives were the establishment of a combined genetic variability pool, a source for second generation selections, evaluation of the genotypic environment interaction and development of seedling seed orchards.

A third objective was the evaluation of the importance of genotype x environment interactions. Generally family performance has been similar whether grown on former pasture or bushland sites and also on the sunkenland site, although rankings may be changed on the latter site due to variable susceptibility to Phytophthora cinnamomi. Matheson and Raymond (1984), reported a loss of potential gain using a single

Another major objective was the ranking of breeding values of clones in the Australian population for W.A. afforestation. This has been achieved (Appendix 2). Information has been used to upgrade the genetic quality of the Manjimup orchard by the removal of the poorer performing clones. The information is also being used in the creation of the new "1.5" and 2 generation seed orchards proposed in phase 4 of this breeding plan (Appendix 2a).

a) Increased volume production (diameter and height) of the order of 10%; will lead to a shortening of the rotation and earlier returns from intermediate yields.

b) Reduction of planting and thinning costs due to the better quality and more uniform plants i.e. orchard seedlings will yield a greater number of straight, smaller limbed trees thus requiring the planting of fewer seedlings for the selection of the growing crop of sawlog trees. Cheaper systematic thinning techniques can be applied in the more homogeneous stands established at a wider spacing.

c) An increase in the percentage recovery and in the grade of the processed product due to improvement in stem form and branching properties.

\* One of the major objectives of the early progeny tests was to show benefits from tree breeding. Data from

### Advances:

Australian populations; regionalization of breeding programmes was suggested as one solution but the better solution would be to omit interactive families which seem to be particularly susceptible to environmental variation.

\* Poor quality areas at Grimwade (32 ha) were searched in 1971, yielding 14 plus trees. These new selections were planted in the seed orchard and clone bank.

\* The Pimelia plantation (125 ha) was intensively searched for plus trees in 1970. The search revealed 74 good trees, 21 of which were considered candidate plus trees. Because of the high standards imposed, only 5 were classified as plus trees and added to the clone bank and orchard.

Western Australia's major radiata pine seed orchard was planted at Manjimup on a site that should have greater flowering potential than the poor Glenelg site.

Local plantations at Pemberton (Pimelia), Grimwade and Harvey (coast) were intensively searched for plus trees; 27 plus trees were selected and added to the breeding population.

The provenance collections made by Eldridge (1978) were planted at the Sunklands in 1979 and will provide a useful resource of plants for breeding in the future.

During this second decade the Donnybrook Sunklands was proposed for a major P. radiata afforestation programme. This project envisaged the conversion of some 60,000 ha of degraded dieback-infested native forest to pine planting over a period of 30 years. A limitation to the seed source adapted to this hostile environment. Progeny trials of the Australian breeding population were established in the Sunklands to identify suitable clones for orchard development.

\* Harvey coastal plantations (SN 1899) were searched in 1974; this yielded 7 plus trees from 31 ha searched.

\* Progeny tests in 1976 used half-sib and full-sib seed supplied from the NSW programme. Objectives were to rank the breeding value of these clones and to provide a pedigree source for second generation selections.

\* Full pedigree seed provided by CSIRO was planted out in progeny tests in 1977 and 1978. Principal objective was to provide a source for second generation selections.

\* The major planting in 1979 was the natural populations of *P. radiata*. Objectives were to provide a long-term and variable stock of genetic variation, a new source for selection of plus trees and the identification of environments. A progeny test to evaluate early WA.

\* In 1980, the Grimwade marginal site selections were planted out in a large progeny trial at Baudin. Test objective is to rank the breeding value of these parent trees. Again the objective was to rank parental breeding populations from the A.C.T. and Tasmania open pollinated seedlots from the A.C.T. and Tasmania parent trees.

\* Parents from the Victorian and NSW breeding populations were planted in 1983, in a trial in the Vasse plantation. Objectives are to rank parents on their breeding value.

#### Breeding population:

of between 1700 - 2200 ha per year. In the field, the orchard will only cover the planting nursery and a planting density of 1 000 per hectare 11 000 plants raised per kilogram of seed sown in the i.e. orchard yield of 160-200 kg. Based on average to reduce the average orchard yield to about 12-15 kg/ha, pollarding to facilitate cone collection is expected however culturing to enhance the genetic quality and orchards in Australia.

hectare and makes it one of the higher yielding radiata to an orchard production of 25 kg seed per orchard (SN 8301) was achieved in 1983; this was equivalent seed (SN 5099). A maximum harvest of 327 kg seed orchard was in full production in 1978 yielding 195 kg was in 1976 when 64 kg seed (SN 5087) was collected. \* First cone harvest from the West Manjimup seed orchard was in 1976 when 64 kg seed (SN 5087) was collected.

#### Advances:

The remaining 6 ha (P71-72) was culled in 1982 of Phytophthora cinnamomi susceptible clones to develop orchard, stocking is 130 s.p.ha. form, vigour and P.C. tolerance now constitute the a P.C. tolerant seed source. 40 clones with good Western Australia and other Australian tests and on ramet shape in the orchard. Additional culling was based on progeny performance in was carried out in 1983 reducing the stocking to 100 s.p.ha.

7 ha of the P69-70 area was culled to retain 36 clones at a stocking of 155 s.p.ha in 1979; this culturing planted at 370 s.p.ha. The orchard was established using grafts of 92 clones 1969-72 West Manjimup seed orchard (13 ha). \*

#### Production population:

If the radiata programme is to exceed 2 000 ha per year an additional planting source is immediately required. Cone collection from unimproved forests would be a backward step and is not advocated. Instead the use of cuttings of selected *radiata* pine is recommended. Proposals are discussed in Phase 4 of the breeding plan for the immediate production of 30 000 woody cuttings (60 ha) leading to future production of 100 000 woody and fascicle cuttings (200 ha).

Scheduling of seed orchard populations is documented in Appendix 3. As there most likely will be a shortage of improved seed every endeavour should be made to purchase orchard seed over this period from elsewhere in Australia e.g. Saxtons seed orchard.

extensive planting of *Pinus radiata* in the Sunbelt and commerce in 1974. Seeding and young tree deaths were found to be consistently associated with root infection by *Phytophthora* spp. These deaths were scattered and could be due to the distribution of the pathogen in the surface soil or there could be genetic variation in resistance. A pot trial was initiated in 1978 to see if there was variation and if it was heritable. This initial work, reported in Butcher et al. (1984), established that resistance in *Pinus radiata* to *Phytophthora cinnamomi* was under strong genetic control and that the genetic control of resistance also acted in the field.

objectives of this phase were to identify genotypes that were tolerant to P.C. for the future development of a tolerant seed source, and to show that genetic tolerance is effective against a range of P.C. isolates and stable over the growing rotation of the tree.

\* Series I - 1979/80; 18 parents from West Manjimup seed orchard. Test objective was to determine genetic variation and heritability as well as a ranking of parents on disease resistance. Split-plot design with inoculation (1 isolate) treatments applied to single family pots.

\* Series II - 1980/81; 31 parents from West Manjimup seed orchard. Test objective was to determine genetic variation and heritability as well as a ranking of parents on disease resistance. Split-plot design with inoculation (1 isolate) treatments applied to single family pots.

Breeding populations: glasshouse screening tests.

parents on disease resistance. Split-plot design with inoculation (1 isolate) treatments applied to single family pots.

\* Series II - 1980/81; 31 parents from West Manjimup seed orchard. Test objective was to determine genetic variation and heritability as well as a ranking of parents on disease resistance. Split-plot design with inoculation (1 isolate) treatments applied to single family pots.

RADIATA PHASE 3: 1979-85 (*Phytophthora cinnamomi* disease resistance).

\* Series III - 1981/82; 35 parents from West Manjimup seed orchard. Test objective was to rank parents on disease resistance and to look at host x pathogen interaction. Treatments applied to single family pots.

\* Series IVA - 1982/83; Wanneroo - parents from W.A. (13), A.C.T. (17), Tasmania (40) and Victoria A.P.M. (55); multi-family pots and 2 P.C. isolate treatments, 5x5x5 cubic lattice design.

\* Series IVB - 1982/83; Como - parents from W.A. (12), A.C.T. (17), Tasmania (36) and Victoria A.P.M. (35); single-family pots, 2 P.C. isolate, 10x10 triple lattice design.

\* Series IVC - 1982/83; Como - host x pathogen interaction; the Australian breeding population on disease resistance. Series IV A and B objectives were to rank parents from

\* Series VA - 1983/84; Como - parents from W.A. (10), Victoria (19) and N.S.W. (20); multi-family pots, 8 P.C. isolates to each pot; 7 x 7 balanced lattice.

\* Series VB - 1983/84; Como - as for VA except that single family pots were used; 7 x 7 triple lattice.

\* Series VA and VB objectives were to rank parents from the Victorian and N.S.W. breeding population on disease resistance.

\* Series VC - 1983/84; Como - host x pathogen interaction; 8 parents (4 tolerant, 4 susceptible) and 10 P.C. isolate treatments including the Al mating type and control, multi-family pots with a split-plot design.

Series VIA - 1983/84; Como - natural populations; Ano Nuevo (10), Monterrey coast (10), Monterrey inland (9), Cambria (10) and Cedros Island (10); multi-family pots, 8 different P.C. isolates per pot  $7 \times 7$  balanced lattice design.

Series VIB - 1983/84; Como - natural populations; Ano Nuevo (12), Monterey coast (12), Monterey inland (12), Cambria (12), Guadalupe Island (12) and Cedros (12); multi-family pots, 8 different P.C.

\* Series VIC - 1983/84; Wanneroо - domesticated populations; W.A. (21), S.A. (21), New Zealand (21) and South Africa (21); multi-family pots, split-plot design of control and inoculated (8 P.C. isolates/pot) treatments.

Series VI objectives were to evaluate variances within and between populations, heritabilities, made of gene action.

Series VII - 1984/85; Como - parents from W.A. (4), S.A. (18 + 5), South Africa (22 + 4) and New Zealand (5 + 1). Series VIIA multi-family pots with 4 P.C. isolates, 7 x 7 balanced lattice design. Series VIIB single-family pots with 4 P.C. isolates, 7 x 7 triple lattice. Objectives are to evaluate some of the parents from the South Australian, New Zealand and South African breeding population for disease resistance.

Objectives are to evaluate some of the parents from the South Australian, New Zealand and South African breeding population for field screening tests.

Breeding population: field screening tests.

RS 17(1980) - Baudin II, *Pinus radiata* families (27), orchard sources (3), *P. pinaster* and *P. taeda* planted as single tree plots on a severely infested dieback site; split-plot treatments of natural and natural plus applied inoculum. Objective was to correlate the field and glasshouse test results on genetic variability.

- \* RS. 18 (1981) - Baudin 12, parents from the West Manjimup seed orchard (35) and the hybrid *P. attenuata* dieback site; split-plot treatment of natural and natural plus applied inoculum. An additional, control series was planted on a healthy banksia site high in the landscape. Plus objectives were the same as the first field test.
- \* There is genetic variation on *P. radiata* to *P.C.* and this is strongly heritable.
- \* 239 plus tree selections from the Australasian breeding population have been screened for resistance to *P.C.* of these 61 were classified as tolerant, 75 as moderately tolerant, 57 as moderately-susceptible and 46 as high or low. This includes an isolate of type A1.
- \* Tolerance is stable against all *P.C.* isolates tested so far, irrespective of whether their virulence levels are high or low. Results of the glasshouse screening trials have been reproduced on field trials in the Sunkland.
- \* Infection by *P.C.* leads to depressed height growth in the survivors but it has little effect on the growth of some of the tolerant families.
- \* Sub-lethal infections can have a major effect on the productive potential of a *P.C.* infested site. For example, four year-old trees of susceptible genotype 60017 were 20% taller than tolerant genotype 80007 on a dieback free Blackwood valley site, but they were 20% smaller on a Sunkland site.

#### Advances:

to study disease expression with time and to evaluate sub-lethal infections on growth rates.

Cedros Island, Ano Nuevo and Guadalupe Island populations were generally highly susceptible to P.C. while the Cambria population was tolerant. The Monterey population, on which the Australian breeding population is based, was moderately-tolerant. Usable variation exists in each population.

Susceptible genotypes were culled from half the West Manjimup seed orchard in 1982 to develop a P.C. tolerant seed source.

A "specialty" (P.C. tolerant) seed orchard is to be established over the 1985-1990 period, using Phytophthora cinnamomi tolerant clones.

This phase is the culmination of the early breeding programme. Breeding values of the plus tree selections and the imported Australian parents have been ascertained for the W.A. environment. The interaction of the genotype with different environments has not been strong, generally the top performers have been universally good and the converse also applies. It was noted that sub-lethal infections in the Sunland tests can have a significant effect on the genotype x environment interaction.

While considerable gains in growth rates, stem quality and branch quality have been reported earlier (Appendix 1) this has resulted from the simple mass selection. Family selection, based on our test results will give further significant benefits but large gains are still achievable from selection of the best individuals within the good families and using these as breeding trees. However to do this, complete pedigree records are required to prevent any future seed orchards. The cone parent is known but the pollin based on family seed lots that were open-pollinated in seed orchards. The cone parent is known but the pollin parent is not and will be drawn from only a very limited gene pool (n = 50). Progeny trials have shown that parents from different families may be related through a common best tree is an open-pollinated (orchard) family (1 in 50 or 100), there is an increasing likelihood that selections breeding population but not from the production population, provided that the possibility of relatedness is recognized. It can be accepted by restricting the number of reselections by the positioning of grafts when planting the orchard.

Most of the Western Australian tests of radiata pine are based on family seed lots that were open-pollinated in seed orchards. The cone parent is known but the pollin parent is not and will be drawn from only a very limited gene pool (n = 50). Progeny trials have shown that parents with high general combining ability (gca) provide a larger number of outstanding offspring. Then, if selecting the best tree is an open-pollinated (orchard) family (1 in 50 or 100), there is an increasing likelihood that selections from different families may be related through a common best tree of outstanding offspring. Then, if selecting the best tree is an open-pollinated (orchard) family (1 in 50 or 100), there is an increasing likelihood that selections breeding population but not from the production population, provided that the possibility of relatedness is recognized. It can be accepted by restricting the number of reselections by the positioning of grafts when planting the orchard.

character need not be included in the genetic makeup. Other for planting on disease safe sites where the tolerance for planting on all potential P.C. infection sites and the developed, one to include the character of P.C. tolerance for this reason that two P. radiata seed sources are being coastal plain where dieback does not impose a threat. It is of the river valleys and deep yellow sands on the southern however, there are plantation sites such as the fertile loams occurrence in all intensively managed afforestation areas. In Western Australia, Phytophthora cinnamomi has a potential

and Western Australia is proposed. However, direct transfer of material between South Australia quarantine period to guard against Ditrichostroma introduction. can be made available, there would then need to be a programmes. This is still being explored. If the material within specified genotypes growing in other Australian another immediate option for orchard use is to make selections and Western Australia is proposed.

new orchards.

otherwise elite trees (certified plus trees) will be used in selections in the open-pollinated families have been made. tests have this. Rather than forgo the added gain, some in the International gene pool and the 1976, 1977 and 1978 objective of within-family selections; only some families family selection but they were not designed to meet the selection. Our trials have provided the information for genetic gain through family selection and within-family

using seed orchards for seed propagation and tree improvement, there is a minimum lead time of some eight to ten years between tree selection and large scale seed production. With vegetative propagation selected trees can be established in the field in a much shorter time. Advantages of clonal propagation of improved material are a greater genetic gain, more rapid realization of this gain and the development of more uniform forests than is possible with seedlings.

For example, cuttings from more mature radiata pines typically produce trees with less volume but they develop better stem form, have fewer and smaller knots and take less time to prune compared to juvenile cuttings or seedlings. If objective is high quality timber production, then management may be willing to sacrifice some per-tree volume growth for the better role characteristics. This would not be critical on W.A. sites where agroforestry management is practised. The optimum maturation state will be a balance between the easy rooting and rapid growth rates of the juvenile age, to the declining root regeneration, improved branching habit, reduced taper and more vertical growth of the aging state.

Production population: seed orchards

straight stems, fine branching and vigour characters.

Genotypes with good expression of P.C. tolerance,

\* "Specialty" (P.C. tolerance) orchard, 8.5 ha, P85-90.

Clones will be represented by grafts and cuttings. A reversed - interlocking - duplicate orchard design will be used with ramets spaced at 8 x 4 m. Ramet crowns will be manipulated to facilitate efficient cone collection.

Intention is to plant the orchard over a 5 year period, establishing approximately 600 ramets each year. The genotype mix need not be the same each year; the best available genetic material at the time will be used.

Stock is available for the planting of 1 ha in 1985, using 40 clones selected from W.A. progeny trials.

Genetic content is made up of: 23 second generation selections (genetic gain component of phenotypic + between-family + within-family selection); 6 have full pedigree, 2 were open-pollinated selection; 6 have full pedigree, 2 were open-pollinated on ortet and 15 were open-pollinated in seed orchards (5 ex Taltaganda, 3 ex NSW and Victoria, 2 ex WA and 1 ex SA and NZ).

: 19 first generation selections (phenotypic + between-family selection). 19 first generation selections (phenotypic + between-family selection).

"General" orchard, 10 ha, p 85-90. Genotypes with good expression of straight stems, fine branching and vigour characters. The orchard established plant is the same as the "specialty" orchard.

In 1985, using 50 clones selected from W.A. progeny trials. Sufficient stock is available for the planting of 1 ha

will be imported through South Australia.

CSIRO and top selections from the New Zealand programme from the Australian population will be available from genotypes will be required in 1985. Pedigree seed available as seed but extra controlled crossings of nursery facilities. Some created pedigree trees are more initial development of donor cutting hedges and fascicle shoot cuttings - favoured method but requires stock available for p 86.

\* Annual programme of 30 000 cuttings is proposed, first

orchard stock plantings in the Blackwood valley.

2 - 5 year old families in progeny trials and from stem cuttings - branches from selected trees in

\* production population: clonal propagation.

between-family selection).

: 14 first generation selections (phenotypic + ex WA, SA, Victoria, Queensland and South Africa).

24 were open-pollinated in seed orchards and clone banks (6 ex Tallaganda, 5 ex NSW, 3 ex NZ, and 2

fully pedigree, 3 were open-pollinated on root and between-family + within-family selection); 9 have

: 35 second generation selections (phenotypic + genetic content is made up of:

Techniques for the rooting of fascicle shoots and needles are being refined at Wanneroo. Hedgerows will be developed in 1986, leading to operational propagation of 50 000 cuttings in 1987/88.

\* first clonal afforestation in 1986 of 60ha leading to 200 ha in 1988.

\* first seed collection from the 1.5 to 2 generation seed orchards in 1993.

#### Advances:

Techniques for the rooting of fascicle shoots and needles are being refined at Wanneroo. Hedgerows will be developed in 1986, leading to operational propagation of 50 000 cuttings in 1987/88.

It is proposed to conduct an intensive plus tree search in 1985 (Appendix 7). Priority areas for the search have been chosen for maximum expression of crown character. The search will be concentrated in plantations growing on high fertility sites on trees that are 25 years of age or older.

#### Gene pool:

The Western Australian proposal is to adopt the basic principles of the South Australian Plan (Cottrell, 1984) with some added simplifications to reduce generation intervals, to reduce the work load in progeny testing and have a more effective selection within families. The description of the proposed WA Plan is summarised in the hierarchy of populations.

The main conflict in tree breeding is between achieving rapid gains in the genetic quality of seed orchards, while at the same time attempting to maintain genetic diversity. The South Australian solution to this is to gradually improve one large breeding population over repeated generations while concurrently developing from this population a more highly selected seed orchard population. This strategy creates a hierarchy of populations with tree levels: (1) gene pool, (2) breeding population and (3) seed orchard population considered "best" by industry at that time. This individual selection is to maintain genetic diversity. The Western Australian proposal is to adopt the basic principles of the South Australian Plan (Cottrell, 1984) with some added simplifications to reduce the work load in progeny testing and have a more effective selection within families. The description of the proposed WA Plan is summarised in the hierarchy of populations.

The earlier phases of this tree breeding plan are the short-term operational or utilisation phases. They consist of mass-producing seed or vegetative propagules for operational planting. The result of selection is to reduce genetic variability. This is needed if gain is to be achieved. However, if selection is intensive on an initial base, genetic variability can be eroded to an extent that it will jeopardise the long-term improvement in future generations. Care must be taken to keep a broad and flexible base in the breeding programme from which outstanding trees can be developed in the future for use in operational planting programmes.

The intention is to initially improve the breeding population as a single large population by selecting for traits which are most likely to remain economically important in the long-term, such as rapid growth, straight stems and fine branching. Wood density is an important trait but only needs to be maintained at its present level in the breeding population.

The intention is to initially improve the breeding population and so on.

new and improved breeding population for the next generation, and crossed in a particular mating pattern to regenerate a superior individual are selected from the breeding population and crossed in a particular mating pattern to regenerate a cumulative improvement of this population. Most tree breeding programmes rely on recurrent selection where breeding individuals are selected from the breeding population where and so on.

#### Breeding population:

The WA radiata gene pool will be created by mixing an equal number of seed from all selections. This will then be raised, planted at normal spacing on a 20 ha typical afforestation site, and managed by operations. The site need not be isolated as fire protection has priority. The gene pool area will be routinely thinned leaving about 250 stems per hectare. Cones will be collected from a large number of trees (stratified random sample) at an approximate age of 25 years to perpetuate the gene pool.

minimum of 8 cones will be collected from each selection. Selection intensity of 1 in 1500 will be used (Burdon and Shetbouren, 1971); 600 - 800 plus trees will be selected from this search. Wood quality characters such as wood density and spiral grain will be later assessed. A number of seed from all selections. This will then be raised, planted at normal spacing on a 20 ha typical afforestation site, and managed by operations. The site need not be isolated as fire protection has priority. The gene pool area will be routinely thinned leaving about 250 stems per hectare. Cones will be collected from a large number of trees (stratified random sample) at an approximate age of 25 years to perpetuate the gene pool.

The main search will be at Nannup (600 ha) and Collie (100 ha) covering the P 56 to P 59 plantations. These plantings include six separate serial lots, four from South Australia and two from New Zealand. Complementary searches will be carried out at Grindmade (23 ha), Margaret River (47 ha) and Harvey Weir (105 ha), on relatively small areas of a greater diversity of seed origins.

Selection intensity of 1 in 1500 will be used (Burdon and Shetbouren, 1971); 600 - 800 plus trees will be selected from this search. Wood quality characters such as wood density and spiral grain will be later assessed. A minimum of 8 cones will be collected from each selection.

The essence of the breeding plan is a low intensity selection among families in the first generation, followed by selection within families in successive generations of the plan. The initial family selection is intended mainly as a preliminary screening to rid the breeding population, as far as possible, of families that have an unacceptable level of malformations. The intensity of family selection (retaining 400 parents) should be sufficient to reduce long-term gains. One phenotypic open-pollinated (ortet) families retained and these second generation selections will be mated in single-pairs to generate a second generation breeding population of 200 unrelated full-sib families. Selection in the second and subsequent breeding populations will be entirely within families and based on individual performance. Successive breeding population sizes of 200 full-sib families will be maintained by selecting the best two trees in each family for the next single pair mating and perpetuation of the plan.

The first generation breeding population will comprise about 100-200 plus trees selected in earlier phases of the programme and an additional 600-700 plus trees selected from unimproved plantations in Western Australia in 1985/86.

RECAUSE of the ateback disease, a separate breeding population may be developed when the second generation is constituted.

The benefit of the plan relates on reduced generation time intervals. This will be maximised by immediately using the nine year-old selections in the family blocks for single-pair mating rather than establishing clone banks for the mating and two-stage selection proposed in the South Australian plan. Selections in the family blocks will be pollarded, thinned to a radius of 12m and heavily fertilized to stimulate flower production.

A net-merit index or complementary mating system can be used in the deliberate culling of individuals for the next generation. Highly selected individuals can be used in the deliberate culling of individuals for the next generation. Highly selected individuals can be created for certain traits, for example, the perfect agroforestry tree with desirable wood properties. These super trees can be immediately commercially propagated using fascicle cuttings.

The intention is to develop a separate population having tolerance to Phytophthora cinnamomi as a primary character. Each of the 400 selections from the first breeding population will be screened for tolerance to P.C.

A timetable of operations for the development of the breeding population is given in Appendix 5.

Major differences to the South Australian plan can be summarised as:

\* definition of objectives; intensive progeny tests to define parent breeding values and large family blocks selected and maintained by operations, for efficient planting and selection of individuals for the next generation of the plan. In the South Australian plan, all of this is done in intensive progeny tests and it is questionable if the next generation selections are really the best.

The upper level of the hierarchy is the highly selected production population, which is a subset of the breeding population. Superior families in the breeding population can be cloned for use in seed orchards or for cutting propagation and thereby achieve substantial short-term gains in the plantation whilst genetic diversity is maintained over the long-term in a cumulative improving breeding population and a large gene pool.

#### Production population:

generation time interval will be further lengthened. Selection bias in the first selection, or the cones at the age of 9 years and this could give a non-experience indicates that not all trees would have ripe selections with an arguable base of selection intensity. Selections are done at a very early age, on trees originating from a very scarce pollen cloud and on reselection is done at a very early age, on trees the two-stage approach but this is questionable. The (1983) expects an increase of 40% in genetic gain using is to use the normal single stage selection. Cotterill each set of four parents tested per family. The WA plan year 4 are used to choose the best two parents from the results of the open-pollinated progeny tests at second stage of selection. At stage 2 of selection, clone banks and open pollinated cones collected for the progeny tests. Selection material is collected to establish no two-stage selection in the WA plan. In the South Australian plan, four individuals having superior phenotypes are chosen in year 9 from each of the 200 full-sib families growing in single tree plot clone banks and open pollinated cones collected for the progeny tests. Selection material is collected to establish no two-stage selection in the WA plan. In the South Australian plan is 14 years compared with a 18 year interval for the South Australian plan. This is largely due to the two-stage selection proposed in the South Australian plan and controlled pollinations being made in clone banks. The WA proposal is to shorten the generation interval by manipulating the selections in the family block and performing the pollinations on these trees.

\* generation interval of the WA plan is 14 years compared with a 18 year interval for the South Australian plan. This is largely due to the two-stage selection proposed in the South Australian plan and controlled pollinations being made in clone banks. The WA proposal is to shorten the generation interval by manipulating the selections in the family block and performing the pollinations on these trees.

The production population will be supplemented by some of the outstanding families from other Australian breeding populations. This can also include material from specialized populations, for example clones with high wood density. Genotype x environment interaction for growth has been found not to be strong and inclusion of different superior genotypes should add to the potential gains. However this will be limited to 25% of the orchard content.

What size should the orchard population be? With a few highly selected clones the potential productivity of plantations will be increased but a narrow genetic base would increase the risk of plantation failure due to extremes of environment (drought) or invasions by pests or diseases. The South Australian plan will have at least 20 clones in the orchard population in each generation. We will be aiming for a minimum of 40 clones until there is information on the flowering habits of individual clones.

For clonal plantations, Libby (1980) has calculated that a safe minimum number of clones is 7 to 25 but larger numbers should be preferred. For example, in Norway spruce programme in Germany 50 to 100 clones are being planted in a random mixture (Kleinischmidt and Schmidt, 1977).

If the timetable for the breeding plan is followed it will take 10 years until the progeny tests are assessed to evaluate breeding values. Superior selections from the family blocks could be used as cuttings to establish a new series of orchards twelve years from now, in 1997. Single-pair matings could be used for clonal propagation with first cuttings could be used for second generation with first population of superior selections for the second generation population available from 2001.

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Progeny trial	routine	pedigree	difference	top quartile*
RS.1(1969) age 8	12.8	13.2	+3%	14.0
RS.2(1971) age 7	17.7	18.6	+5%	19.6
RS.3(1971)	16.7	17.9	+7%	19.0
RS.4(1972)	15.4	15.8	+3%	16.7
RS.5(1972)	13.8	13.6	-1%	14.8
RS.8(1974) age 10	12.3	13.8	+12%	14.5
RX.1(1973) age 6	16.0	16.6	+4%	17.2
RX.2(1973) age 7	11.6	12.0	+3%	12.6
r/o (1974) age 10	12.3	13.2	+7%	14.2
* average for the top ranking 25% of families in the trial.				

1. Character: Diameter (bhd) in cm).

Appendix 1: Results from W.A. *Pinus radiata* progeny and yield trials.

2. Character: Height (m).

Progeny trial	routine	pedigree	difference	top quartile
RS.1(1969)	2.2	2.1	-5%	2.1
RS.2(1971)	7.5	7.5	=	8.1
RS.3(1971)	3.1	3.5	+11%	3.8
RS.4(1972)	5.9	5.9	=	6.3
RS.3(1974)	3.8	4.2	+11%	4.5
Yield trial				
RX.3(1973) age 4	4.2	4.4	+5%	4.4
r/o (1974) age 4	3.8	3.8	+1%	3.8
Yield trial				

3. Character: Stem straightness (% of trees in trial that are straight).

Progeny trial	routine	pedigree	difference	top quartile
RS.1(1969)	2.39*	2.59	+8%	2.79
RS.2(1971)	52	58	+12%	70
RS.3(1971)	76(12**)	83(22)	+10% (+83%)	91(34)
RS.4(1972)	33	41	+24%	64
RS.8(1974)	32	63	+97%	72
Yield trial				

\* mean point score

Progeny trial	routine	pedigree	difference	top quartile
RS.1(1969)	2.82*	2.83	+1%	3.1
RS.2(1971)	50	56	+12%	69
RS.3(1971)	85(15**)	89(27)	+4%	97(41)
RS.4(1972)	56	80	+43%	94
RS.8(1974)	52	66	+27%	75
Yield trial				
RX.1(1973)	26	29	+12%	
r/o (1974)	52	58	+12%	

\*\* % of trees in trial with right angle branching.

5. Character: branch angle (% of trees in trial with flat  
angle branches).

\*\* % of trees in trial with very small branches.

\* mean point score

Progeny trial	routine	pedigree	difference	top quartile
RS.1(1969)	2.57*	2.67	+4%	2.92
RS.2(1971)	44	59	+34%	70
RS.3(1971)	71(8**)	85(23)	+20%(+200%)	95(38)
RS.4(1972)	36	69	+92%	88
RS.8(1974)	30	49	+63%	63
Yield trial				
RX.1(1973)	87	88	=	
r/o (1974)	30	42	+40%	

4. Character: branch thickness (% of trees in trial with  
thin branches).

\*\* % of trees in trial that are very straight

\* mean point score

Yield trial	routine	pedigree	difference	top quartile
RX.1(1973)	49	52	+6%	
r/o (1974)	32	64	+100%	

Appendix 1 (continued).

P. radiata	Height	Diameter	Stem	Striagneness	Branch	Angle	Total	Wood	Density
12038	**	**	***	**	**	*	**	**	80055
12349	**	**	**	**	***	*	**	*	80007
12374	*	**	**	**	**	*	**	*	70053
10957	**	**	***	*	**	-	**	*	70052
10956	*	**	*	*	**	*	*	*	60002
12112	*	**	*	*	**	*	**	*	60015
12130	*	**	*	*	**	*	**	*	60004
30016	*	**	**	*	*	*	**	*	50269
30026	*	**	*	*	**	*	**	*	50266
30001	*	**	**	*	**	*	**	*	50015
30012	*	**	**	*	**	*	**	*	50048
30012	*	**	**	*	**	*	**	*	50012
50006	*	**	**	*	**	*	**	*	50006
50001	*	**	**	*	**	*	**	*	50016
30026	*	**	**	*	**	*	**	*	30016
30012	*	**	**	*	**	*	**	*	30012
30011	*	**	**	*	**	*	**	*	30007
30002	*	**	*	*	**	*	**	*	12130
12112	*	**	*	*	**	*	**	*	12112
10956	*	**	*	*	**	*	**	*	10957
10957	*	**	***	*	**	-	**	*	12374
30007	-	-	*	*	**	*	**	*	12349
30002	*	*	*	*	**	*	**	*	10956
30011	*	**	**	*	**	*	**	*	10957
30012	*	**	**	*	**	*	**	*	30011
30016	*	**	**	*	**	*	**	*	30012
50001	*	**	**	*	**	*	**	*	50001
50006	*	**	**	*	**	*	**	*	50012
50015	*	**	**	*	**	*	**	*	50048
50016	*	**	**	*	**	*	**	*	50015
50266	*	*	*	*	**	*	**	*	50269
50269	-	-	*	*	**	*	**	*	50004
60004	*	-	*	*	*	*	**	*	60015
60002	*	-	*	*	*	*	**	*	70052
70052	-	-	*	*	***	*	**	*	70053
70053	*	**	**	*	*	*	**	*	80007
80007	***	***	**	**	**	*	*	*	80055
80055	--	--	**	*	*	*	**	*	--

(a) some Clones with high gca.

Appendix 2: Breeding values of clones in W.A. tests.

(b) some Clones with poor gca.

Appendix 2 (continued).

P. radiata	Clone	Height	Diameter	Stem	Branch	Branchiness	Thickness	Angle	Tolerance	Density
12236	60017	---	---	---	---	---	---	*	***	***
	60003	*	---	---	---	---	---	-	-	-
	60001	---	---	---	---	---	---	-	*	*
	50024	**	*	---	---	---	---	-	*	*
	30048	-	-	---	---	---	---	-	-	-
	30043	*	**	---	---	---	---	-	-	*
	30040	**	**	---	---	---	*	---	---	**
	30036	**	**	---	---	---	*	---	---	**
	30017	**	*	---	---	---	-	---	---	**
	30004	*	---	---	---	---	*	*	*	*
	12408	-	*	---	---	---	*	-	-	-
	12378	-	-	---	---	---	*	**	---	-
	12247	-	-	---	---	---	*	*	-	*
	12197	-	-	---	---	---	*	-	-	*
	12001	-	-	---	---	---	-	-	-	-

Year	Phase 2 W.M.S.O.	Age yield	Phase 4 "General"	Age yield	Phase 5 gen.	Gen. Spec.	Gen. Spec.	Gen. Spec.	Gen. Spec.
1982	11 205kg								
1983	12 327								
1984	13 310								
1985	14 200	Plant	Plant	Plant	Plant	Plant	Plant	Plant	Plant
1986	15 160	Plant	Plant	Plant	Plant	Plant	Plant	Plant	Plant
1987	16 200	Plant	Plant	Plant	Plant	Plant	Plant	Plant	Plant
1988	17 200	Plant	Plant	Plant	Plant	Plant	Plant	Plant	Plant
1989	18 250	Plant	Plant	Plant	Plant	Plant	Plant	Plant	Plant
1990	19 250								
1991	20 250								
1992	21 250								
1993	22 250	6 60kg	6 50kg	6 50kg	6 50kg	6 50kg	6 50kg	6 50kg	6 50kg
1994	23 250	7 100	7 80	8 150	8 130	8 130	8 130	8 130	8 130
1995	24 250	7 100	7 80	8 150	8 130	8 130	8 130	8 130	8 130
1996	1996	9 150	9 130	9 130	9 130	9 130	9 130	9 130	9 130
1997	1997	10 180	10 150	10 150	10 150	10 150	10 150	10 150	10 150
1998	1998	11 180	11 150	11 150	11 150	11 150	11 150	11 150	11 150
1999	1999	12 180	12 150	12 150	12 150	12 150	12 150	12 150	12 150
2000	2000	13 180	13 150	13 150	13 150	13 150	13 150	13 150	13 150
2001	2001	14 180	14 150	14 150	14 150	14 150	14 150	14 150	14 150
2002	2002	15 180	15 150	15 150	15 150	15 150	15 150	15 150	15 150
2003	2003	16 180	16 150	16 150	16 150	16 150	16 150	16 150	16 150
2004	2004	17 180	17 150	17 150	17 150	17 150	17 150	17 150	17 150
2005	2005	18 180	18 150	18 150	18 150	18 150	18 150	18 150	18 150
2006	2006	19 180	19 150	19 150	19 150	19 150	19 150	19 150	19 150
2007	2007	20 180	20 150	20 150	20 150	20 150	20 150	20 150	20 150
2008	2008	21 180	21 150	21 150	21 150	21 150	21 150	21 150	21 150
2009	2009	22 180	22 150	22 150	22 150	22 150	22 150	22 150	22 150
2010	2010	200	150	200	150	200	150	200	150
2011	2011	200	150	200	150	200	150	200	150
2012	2012	200	150	200	150	200	150	200	150
2013	2013	200	150	200	150	200	150	200	150
2014	2014	200	150	200	150	200	150	200	150
2015	2015	200	150	200	150	200	150	200	150

Appendix 3: Schedule of seed orchard populations.

## PROGENY TEST

- \* objective - to define breeding values of parents.
- \* information - (a) screen initial selections for parents to go into the breeding population. (b) provide a ranking of parents for use in design - incomplete block, 30-50 individuals in each family planted in single tree plots. standard management prescriptions. repeat on principal planting sites. measure - height at 4 years, diameter, height, tree form and wood quality at 9 years. conversion to seedling seed orchard after 9 years. isolation not necessary, maximum fire protection essential.

## FAMILY BLOCK

- \* objective - selection of phenotypically superior individuals within a family. design - single family planting. standard management prescriptions. repeat on principal planting sites. within a family planting. of 100-200 seedlings to facilitate selection of the best individuals in family groups.

- \* release selected trees by removing adjacent trees to a distance of 12 m, Pollard selected tree and apply fertilizer and hormone treatments to stimulate flower for single-pair mating.

- \* collect scions for establishment in clone bank and pollen be selected at age 9 years. Evaluate wood properties, phenotypically superior individuals within a family will no measurements will be carried out.

- \* standard management prescriptions.

repeat on principal planting sites.

the best individuals in family groups.

of 100-200 seedlings to facilitate selection of

- \* design - single families will be planted in compact blocks no measurements will be carried out.

phenotypically superior individuals

- \* repeat on principal planting sites.

production.

fertiliser and hormone treatments to stimulate flower

distance of 12 m, Pollard selected tree and apply

release selected trees by removing adjacent trees to a

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distance of 12 m, Pollard selected tree and apply

Appendix 4: Progeny test and family block features of the breeding plan.

Appendix 4 (continued).

- \* single-pair mating of selected trees will be performed in the family block to decrease the time interval between generations.

Appendix 5: Timetable of operations in Phase 5.

Year 0 (1985): SEARCH 1985, search 700 ha of plantation type 1 for 700 plus trees.

Year 1 (1986): Sow family seed lots in nursery.  
Mix some seed from each family to create the gene pool and sow in nursery.

Year 2 (1987): Plant the 1st generation breeding population in progeny tests and family blocks at two or three major plantation sites.

Year 6 (1991): Progeny test - measure heights.  
Develop progeny test as a seedling seed orchard.  
best 400 parents to use in the breeding population.  
all parents on these characters and select the best 400 families. Collect scions for cutting the phenotypically superior individual for each of the 400 families. Collect sections for cutting propagation and establishment of clone banks and collect pollen. Release selected trees and treat to stimulate flowering. Use cuttings of the best 40 clones to plant in seed orchards.

Year 11 (1996): Progeny test - measure heights, diameter and wood quality, assess tree and branching form. Rank best 400 parents to use in the breeding population.  
all parents on these characters and select the best 400 families. Collect scions for cutting propagation and establishment of clone banks and collect pollen. Release selected trees and treat to stimulate flowering. Use cuttings of the best 40 clones to plant in seed orchards.  
the phenotypically superior individual for each of the 400 families. Collect sections for cutting propagation and establishment of clone banks and collect pollen. Release selected trees and treat to stimulate flowering. Use cuttings of the best 40 clones to plant in seed orchards.  
Family block - in the chosen 400 families, select 40 clones to plant in seed orchards.  
the phenotypically superior individual for each of the 400 families. Collect sections for cutting propagation and establishment of clone banks and collect pollen. Release selected trees and treat to stimulate flowering. Use cuttings of the best 40 clones to plant in seed orchards.  
Clone bank - collect open-pollinated seed of the 400 chosen parents, raise seedlings and carry out glasshouse screening tests for Phytophthora cinnamonomi tolerance.

: Establish a yield trial.

Year 12 (1997): General seed orchard (11 ha) - plant cuttings of the best 50 second generation selections, 3 ha a year for 4 years. This orchard will be developed at the West Mangimpup Forestry Centre.

Year 14 (1999): Family block - collect control pollinated seed. chosen clones to create 200 unrelated families.

Year 15 (2000): Sow family seed lots in the nursery.

Year 16 (2001): Plant the 2nd generation breeding population in progeny tests and family blocks at two or three major plantation sites.

Year 17 (2002): Specialty seed orchard (8.5 ha) - plant cuttings of P.C. tolerant seedlings over 3 years. This orchard will be developed on part of the original West Mangimpup seed orchard site (cleared in 1998).

Year 18 (2003): Plant the 2nd generation breeding population in progeny tests and family blocks at two or three major plantation sites.

Year 19 (2004): *Phytophthora cinnamomi* tolerant population - complete the ranking of the chosen families for P.C. tolerance and recommend a strategy for the development of this population.

Year 20 (2005): Progeny test - measure heights.

Year 25 (2010): Progeny test - measure heights, diameter and wood quality, assess tree and crown form. Rank breeding values of parents for use in seed orchards and clonal propagation. Develop a seedling seed orchard.

: Family block - select two phenotypically superior individuals in each family ( $200 \times 2 = 400$ ). Collect scions for cutting propagation and establishment of clone banks, and collect pollen. Release selected trees and treat to stimulate flowering. Use cuttings from the best 40 clones to plant in seed orchards.

Year 26 (2011): General seed orchard (10 ha) - plant cuttings of the best 50 third generation selections, 3 ha a year for 4 years. This orchard will be developed on the Manjimup general orchard site planted 1985-90 (clearfelled 2009).

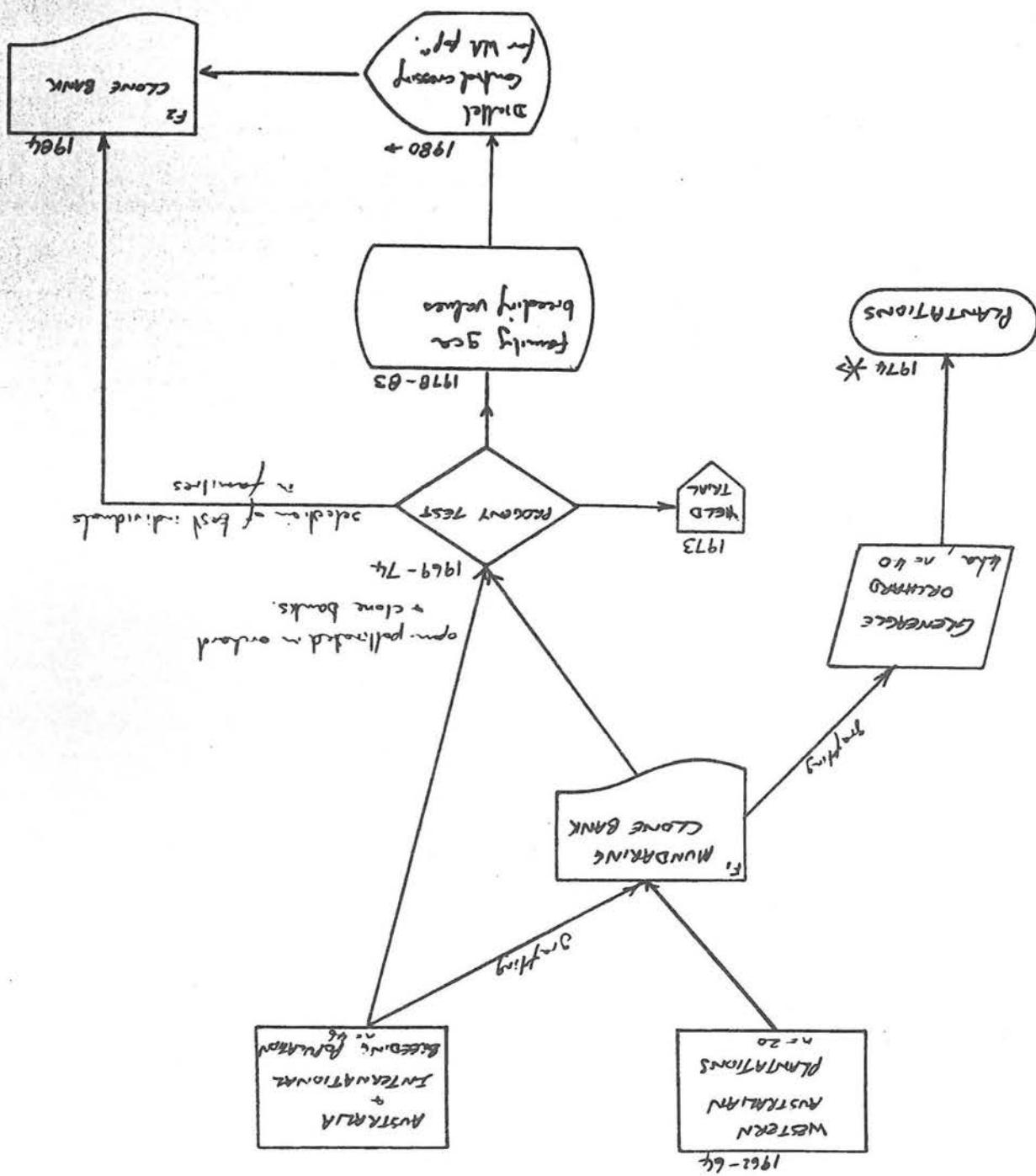
: Family block - single-pair mating of the 400 selections to create 200 new families.

Year 28 (2013): Family block - collect cones at random from a large gene pool - plant the gene pool. Gene pool - collect cones at random from a large number of trees (planted in 1987), mix seed to regenerate and form the new gene pool.

: Speciality seed orchard (8.5 ha) - plant cuttings of the best third generation selections, 3 ha a year for 3 years. This orchard will be developed on the speciality orchard site planted 1985-89 (clearfelled 2011). Speciality seed orchard (8.5 ha) - plant cuttings of the best third generation selections, 3 ha a year for 3 years. This orchard will be developed on the speciality orchard site planted 1985-89 (clearfelled 2011).

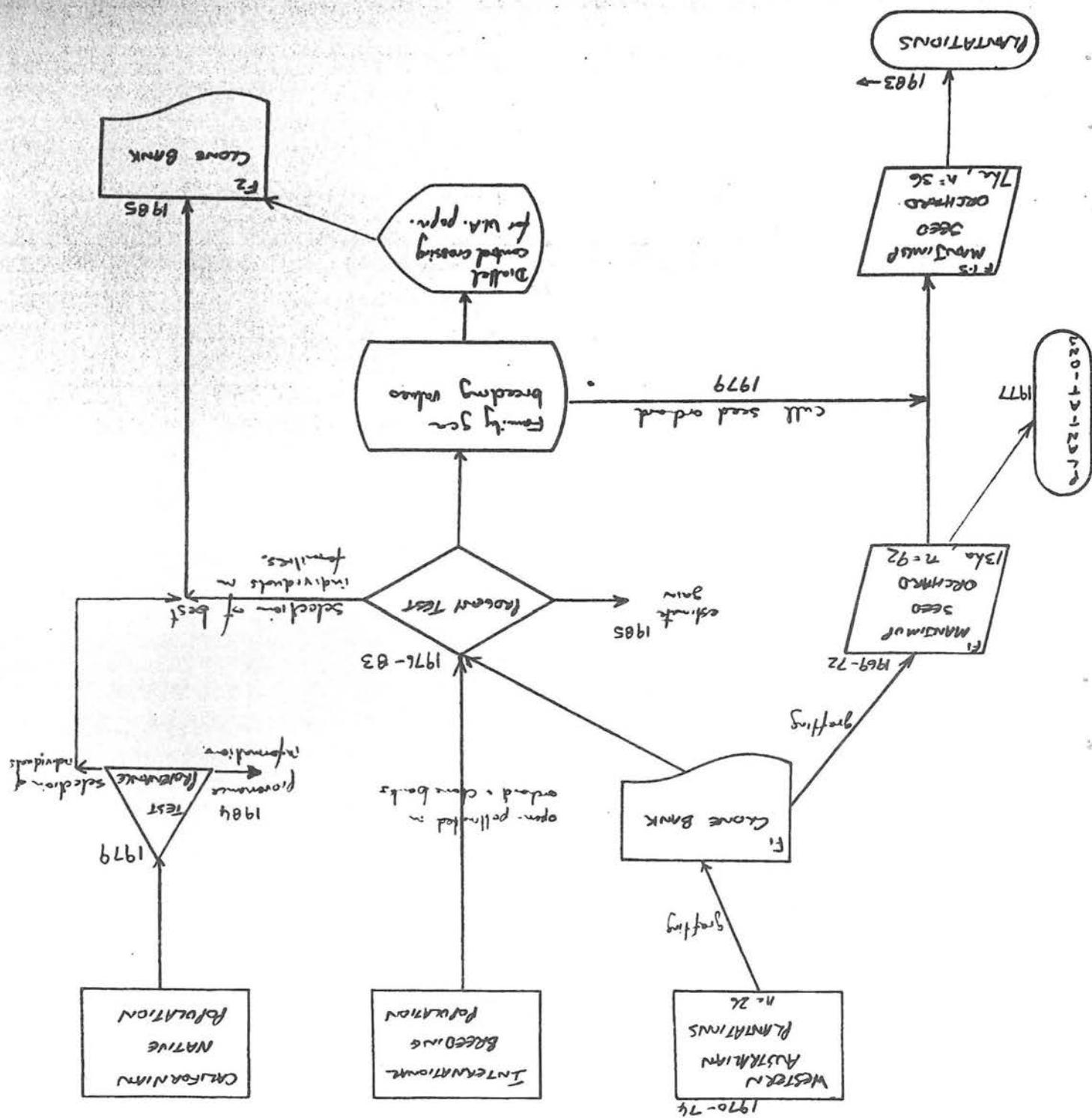
Year 29 (2014): Sow family seed lots in the nursery. General seed orchard, final planting of 3 ha.

Year 30 (2015): Plant the 3rd generation breeding population in year 30 (2015): Plant the 3rd generation breeding population in this population for clonal afforestation. Family shoot propagation of outstanding families major plantation sites.

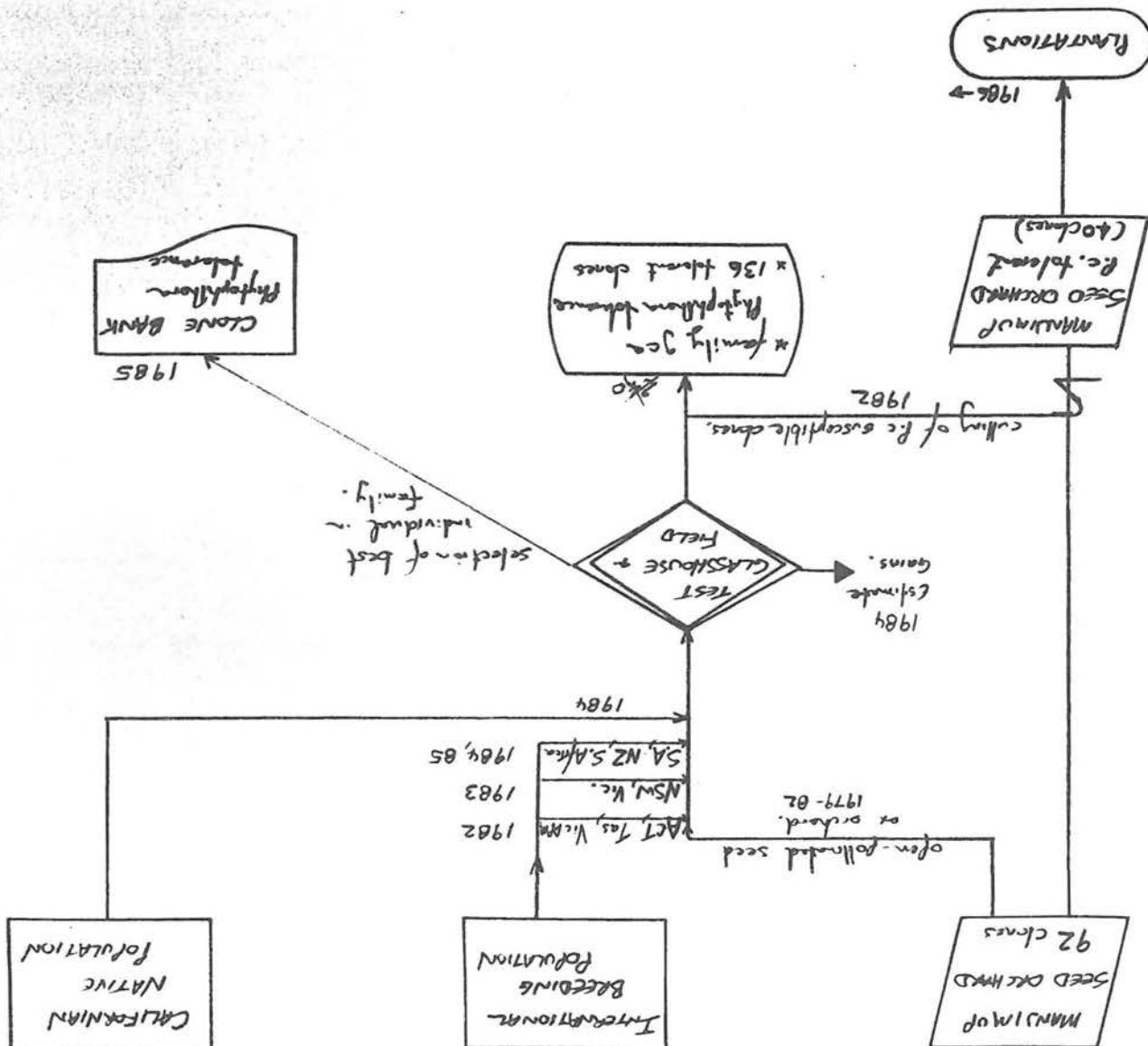


KADIRIA Phase 1: 1960-1970 (commencement)

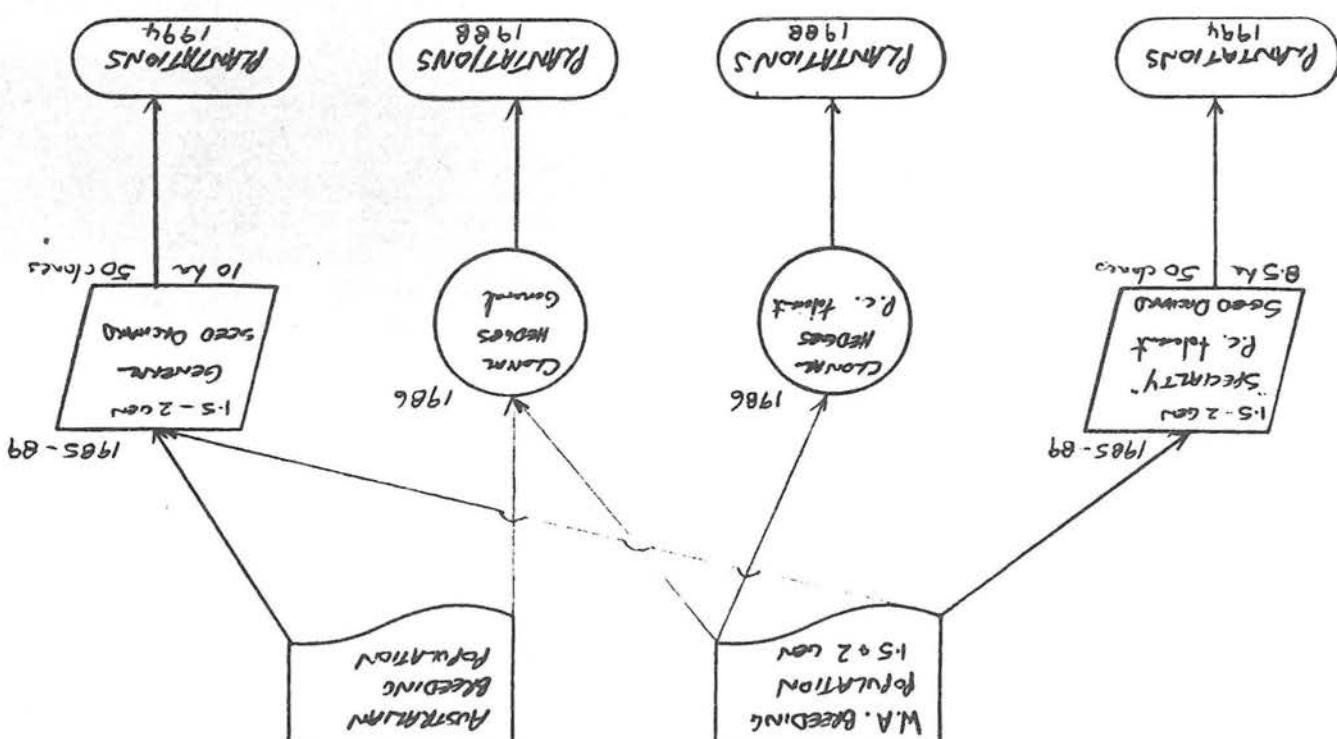
Appendix 6.



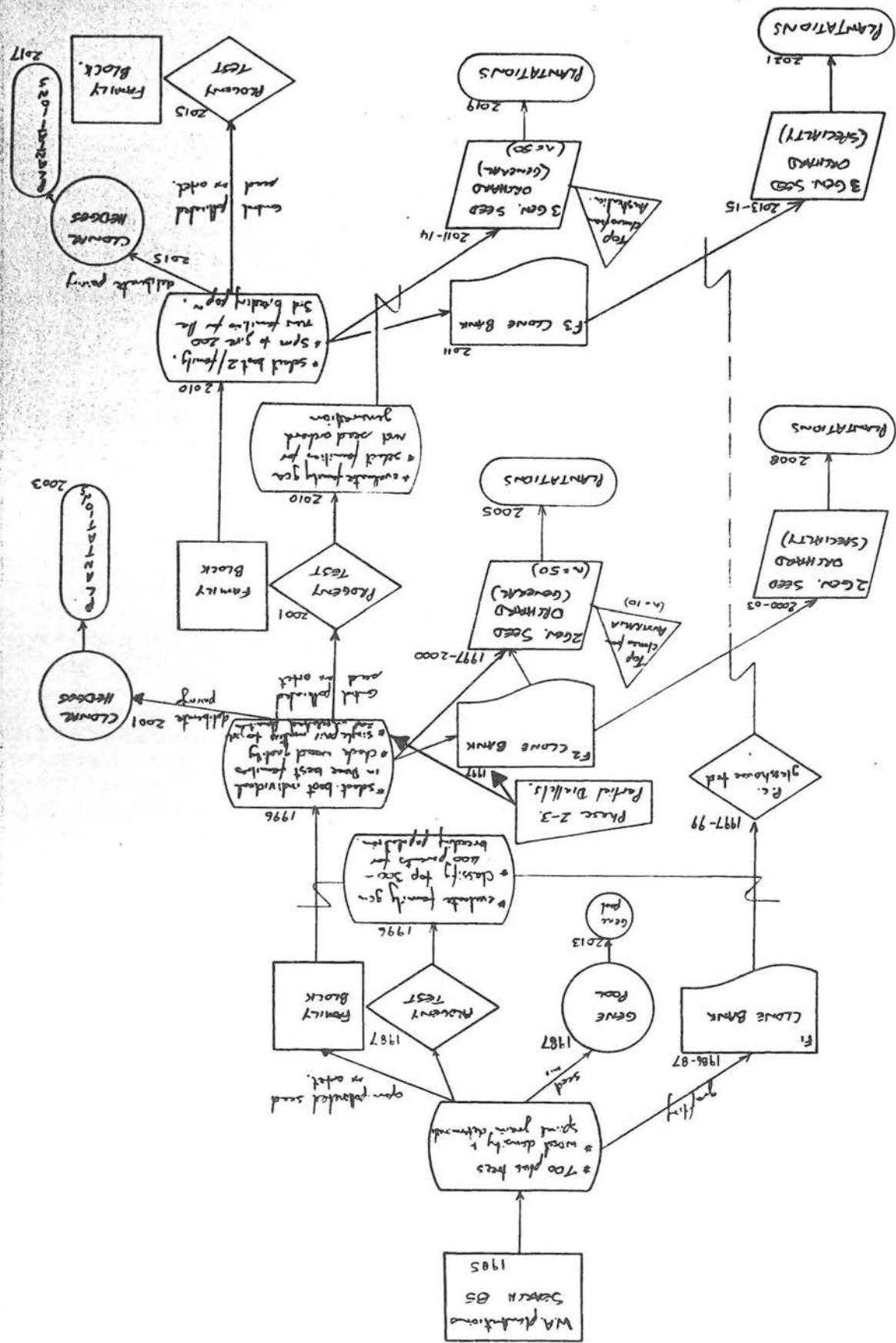
RADIATA. PHASE 2 = 1970-1980 (enrichment)



RADIATA Phase 3 : 1980-1985 (Hybridization, inbreeding, disease resistance)



DATA FILE 4 : 1985-1990 (Commercial Operations)



DATAITA Phase 5 : 1985 → (Development)

Attach.

3 December 1984

Como Research

TBB:KG

T.B. BUTCHER

S.D.F.O.

Attention: Mr. Have1

Mr. Underrwood  
Mr. Van Noort

Loc. 6/3  
H.O. 400/65

A/CONSERVATOR OF FORESTS  
STATE HEADQUARTERS

## PINUS RADIATA PLUS TREE SELECTION

Plus tree selection in Pinus radiata plantations in Western Australia is expanding rapidly. It is essential that the base breeding population of Pinus radiata in Western Australia is expanded immediately. It is proposed to conduct an intensive plus tree search in 1985. The priority is for "agroforestry" plus trees that are straight with small limbs, deep crowns. The 1985 search will be concentrated in plantations growing on high fertility sites on trees that are 25 years of age or older. Selection intensities of 1 plus tree hectare (SI, 1:1000-2000 trees) will be used. The proposal is to search 600 ha in Folly/Milwards plantation at Nanup and 100 ha in the Mungalup plantation, 600 plus trees will be selected from this search. The project is beyond the capacity of technical staff. As this project is of vital concern to the Central Region, the region could provide 4 recent graduates for three weeks based at Nanup to assist in the plus tree selection. Failing Central Region, the region could provide 4 recent graduates for three weeks based at Nanup to assist in the plus tree selection. A 2 year period. This is to be carried out over a 2 year period.

1. W.A. Plantation site types for selection.
2. Intensity of selection.
3. Pinus radiata plus tree selection traits, notes for searchers.
4. Plus tree classification.
5. Method of plus tree selection.
6. P.O.C.S sheets for the 1985 search.

Appendices to this proposal include:

3 December 1984  
Como Research  
Mr Underrwood  
Mr. Have1  
Attention: T.B. BUTCHER  
S.D.F.O.

Plantation	Group	P Year	Area	Type
Nanup	Folly	1956-1963	734 ha	1a
	Lewana	1961-1965	387 ha	1a
	Milward	1957-1965	498 ha	1a
Kirup	Grimwade	1933-1960	689 ha	1b
	Ferndale	1961-1965	549 ha	1b
	Kelly	1965	50 ha	1b
Collie	Mungalup	1957-1961	260 ha	1a & 1b
	Bussells	1957-1965	428 ha	1b
	Wellington	1960-1962	118 ha	1b
Harvey	Harvey Wet	1939-1963	186 ha	1b
	Brunswick	1959-1964	149 ha	1b
	Myalup	1930-1965	100 ha	2
	McLarty	1953-1965	38 ha	2
	Tallanalla	1955-1965	205 ha	2
Busselton	Margaret	1933-1957	64 ha	1b
	Ludlow	1910-1965	79 ha	2
Mundaring	Greystones	1922-1960	28 ha	1a
	Heleena	1923-1965	30 ha	1a
	Gortie	1959-1963	117 ha	1a
	Berakine	1931-1965	210 ha	1a
	Wellbuckett	1953-1965	68 ha	24 ha
	Ciffords	1957-1965		16 ha
	Carinyaah	1925-1965		

Twenty year and older plantation areas are tabulated:

1. River valley - high fertility sites
2. Coastal plain sands - low fertility sites.
3. Donnybrook sunkland - infertile, poor drainage, shallow soils and dieback infested.

(a) former pasture

(b) former bushland

*Pinus radiata* plantations in Western Australia can be divided into:

#### 1. PLANTATION SITE TYPES FOR SELECTION

These are the base populations; they can be searched to locate the best adapted trees for the particular site type. "Plus" trees are selected for their distinctive superiority to the average of neigbhouring trees; they are phenotypes (interactions of environment with genotype). Heritability of the character will determine how closely the phenotype resembles the genotype.

heritability	character	high	intermediate	low
	branching type	Phytophthora cinnamomi tolerance	stem straightness	forking and ramification branches
	"	"	branch quality	branch diameter
	"	"	height	wood density
	"	"	height	future.

The 1985 search will concentrate on the plantations:

Nannup	Folly	P56	116 ha	Folly	P57	92 ha	Milward	P58	166 ha	Milward	P59	247 ha	Mungalup	P57	75 ha	Mungalup	P58	86 ha
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Selection standards for plus trees were set very high for all breeding programmes when initiated in the late 1950s; large areas were searched for very few outstationing plus trees on the assumption that heritability and additive genetic variance would be fairly high for all characters. Now that some preliminary estimates of heritability are available the tree breeders relied on intensive phenotypic selection has decreased. The Pimeliá plantation plus tree selection carried out by Butcher, Fremlin and Kruger in 1970 can be used as an example. The 125 ha of 36-42 year old P. radiflata were crissed on a 100 m grid, resulting in:

initial search	= 74 trees; SI, 1:3800.
candidate plus trees	= 21 ; SI, 1:13000.
plus trees	= 5 ; SI, 1:56000.

of the 5 plus trees selected for the breeding population, only 1 appears to be good in all characters viz. stem straightness, branching quality, vigour and Phytosphaera tolerance. It is obvious now that all of the 21 candidates plus trees (SI, 1:13000) if not the 74 trees from the initial cruise should have been included in the initial breeding population and progress testing then would have revealed more trees suitable for inclusion in the breeding population. The breeding population requirements for very large gains in productivity.

Selection intensity of 1 plus tree per hectare or 1:1000 to 2000 trees will be used in the 1985 selection series.

3. *PINNUS RADIATA* PLUS TREE SELECTION TRAITS

1. STEM FORM:
  - \* The tree must be straight. The stem should have no bends or lean and it should be circular in cross-section. Slight bends can be only accepted if the tree is better than its neighbors.
2. BRANCH QUALITY:
  - \* Multi-nodal trees with regular spaced, small diameter, flat angled branches. There should be 2-3 branch whorls per year, preferably at an even distance apart. Avoid trees with long branches; in wide spaced plantations, a tree with short branches is particularly desirable. Branches coming out at right angles to the stem are ideal; avoid trees with very steep angled branches.
3. TREE VIGOUR:
  - \* The tree must be at least as tall as the biggest tree in the immediate vicinity. Preferably, it should also be as big (diameter) as the biggest tree.
4. TREE HEALTH:
  - \* Crowns should be dense, healthy and vigorous; avoid thin crowns of light green needles, any signs of dead topping or rusty crowns.
  - \* Avoid any stems exuding resin or showing signs of insect attack.
5. STEM CONES:
  - \* Avoid trees with large numbers of cones on the stem.

Each candidate plus tree will be tested for wood quality, principally basic density and grain angle. The acceptance standards will be set after the population values have been determined.

Candidate plus trees should be good in all characteristics and possibly outlasting in one or two. Trees inferior in an important character should not be selected.

Following the initial search and location of candidate plus trees, quantification of the phenotypic values and classification of the trees is necessary. This is done by the comparison tree method. Nine other crop trees (rate 250 s.p.h.a) in the immediate vicinity of the candidate plus trees group of ten trees forms the micro-site population; candidate plus tree characteristics are rated against this to determine superiority.

Characteristic to be rated as:

- 1 = **superior** (better than 2 standard deviations from the mean or the best of 10 trees).
- 2 = **better than average** (between 1 and 2 Std. Dev. from the mean or the best or second best of 10 trees).
- 3 = **average** (within 1 Std. Dev. from mean, or intermediate).
- 4 = **below average** (between -1 and -2 Std. Dev. from the mean, or last or second last of 10 trees).
- 5 = **very poor** (more than -2 Std. Dev. from the mean, or poorest of 10 trees).

### External characteristics - measured

**Tree:**

- \* measure diameter over bark at height of 1.3 m.

**Height (tree):**

- \* measure height of first green crown whorl; calculate green crown length.

**Crown width:**

- \* measure crown width (diameter) along row and across row (bay); calculate crown area and the ratio of crown area/stem area.

**Tree volume:**

- \* calculate  $DBH_{UB} = 0.506 + 0.862 DBH_{OB}$  cm.
- \* calculate  $Vol_{UB} = 0.472 DBH_{UB}^2 \times Ht \text{ m}^3$ .

- 1 - trunk cones cast shortly after maturity.
- 2 - scattered persistent trunk cones.
- 3 - numerous persistent trunk cones.

stem cones:

- 1 - dark green, dense, healthy crown.
- 2 - dark green, healthy crown.
- 3 - dark green, sparse crown.
- 4 - light green, sparse crown.
- 5 - dead topping.
- 6 - rusty crown.
- 7 - resinous stem.

tree health:

- 1 - normal
- 2 - ramicorn
- 3 - fork
- 4 - multiple or repeated forking.

tree form:

- 1 - very flat, right angles to stem.
- 2 - flat angle, 70-80° to stem.
- 3 - low angle, 60-70° to stem.
- 4 - moderate angle, 50-60° to stem.
- 5 - high angle, 40-50° to stem.
- 6 - very steep angled, <40° to stem.

branch angle:

- 1 - very small, short branches.
- 2 - small, short branches.
- 3 - acceptable, average diameter and length of branches.
- 4 - moderate, slightly larger and long branches.
- 5 - large diameter, long branches.
- 6 - very large diameter, long branches.

branch thickness and length (to be related to tree vigour):

- 1 - uni-nodal.
- 2 - bi-nodal.
- 3 - multi-nodal.

branch type (3 to 10 m above ground):

- 1 - perfectly straight.
- 2 - straight, very slight deviation.
- 3 - straight, with slight deviations.
- 4 - reasonably straight, obvious deviations.
- 5 - not straight, marked kinks, sweeps or lean.
- 6 - crooked, extreme deviation.

stem straightness:

## 5. METHOD OF PLUS TREE SELECTION

1. OFFICE

- \* Collect the plantation maps.
- \* P.O.C.S. sheet for the particular age class.
- \* Aeriel photographs - Elgicht age 6-7 years to show extraction tracks, compartment crosses breaks etc. May be able to use most recent photographs as well.
- \* Aeriel photographs - Elgicht age 6-7 years to show extraction tracks, compartment crosses breaks etc.
- \* Aeriel photographs - Elgicht age 6-7 years to show extraction tracks, compartment crosses breaks etc.
- \* Photocopy aerial photographs and draw up a 200 m x 200 m grid system; index the grid with compartment number and grid unit number. These selection units will be 4-5 ha in area.

## 2. FIELD

- \* Mark out the end of the 200 m row to delineate the selection unit, using yellow plastic ribbon (need to mark the 6 trees at the end of the job).

### 3. OFFICII

Workking in teams of two assessors, traverse the selection grid as illustrated in diagram. Each assessor looks towards the other through 16 rows of trees. Either walk through or use horses; each assessor has radio communication. Search for trees that have the specified characteristics; mark any worthwhile tree in the first run, using white ribbon. At the end of each unit, several trees will have been marked; these are then closely scrutinised by both assessors and the best 4 trees selected are marked with additional white plastic ribbon; these are then located with reference to the compartment boundary for permanent marking and reforestation. Of the 4 selected trees/unit, a ranking of the best 1 and 2 trees should be given.

Cover the plantation group specified.

COLLATE all data from the initial plus tree cruise.

ALLOCATE plus tree candidate numbers to each of these trees.

Note that these are not the plus tree numbers, these will only be allocated after the candidate trees have (a) scored for external characteristics (b) screened for wood properties (c) cones collected from.

#### 4. FIELD

- Do the candidate plus tree exceed final characteristic assessment on the standard form.
- Take wood core for spiral grain and wood density determination.
- Determine wood properties of candidate. Decision then as to if to accept as a plus tree.

7. CONTINUE THE SELECTION PROCESS TO EXPAND THE BASE BREEDING POPULATION.

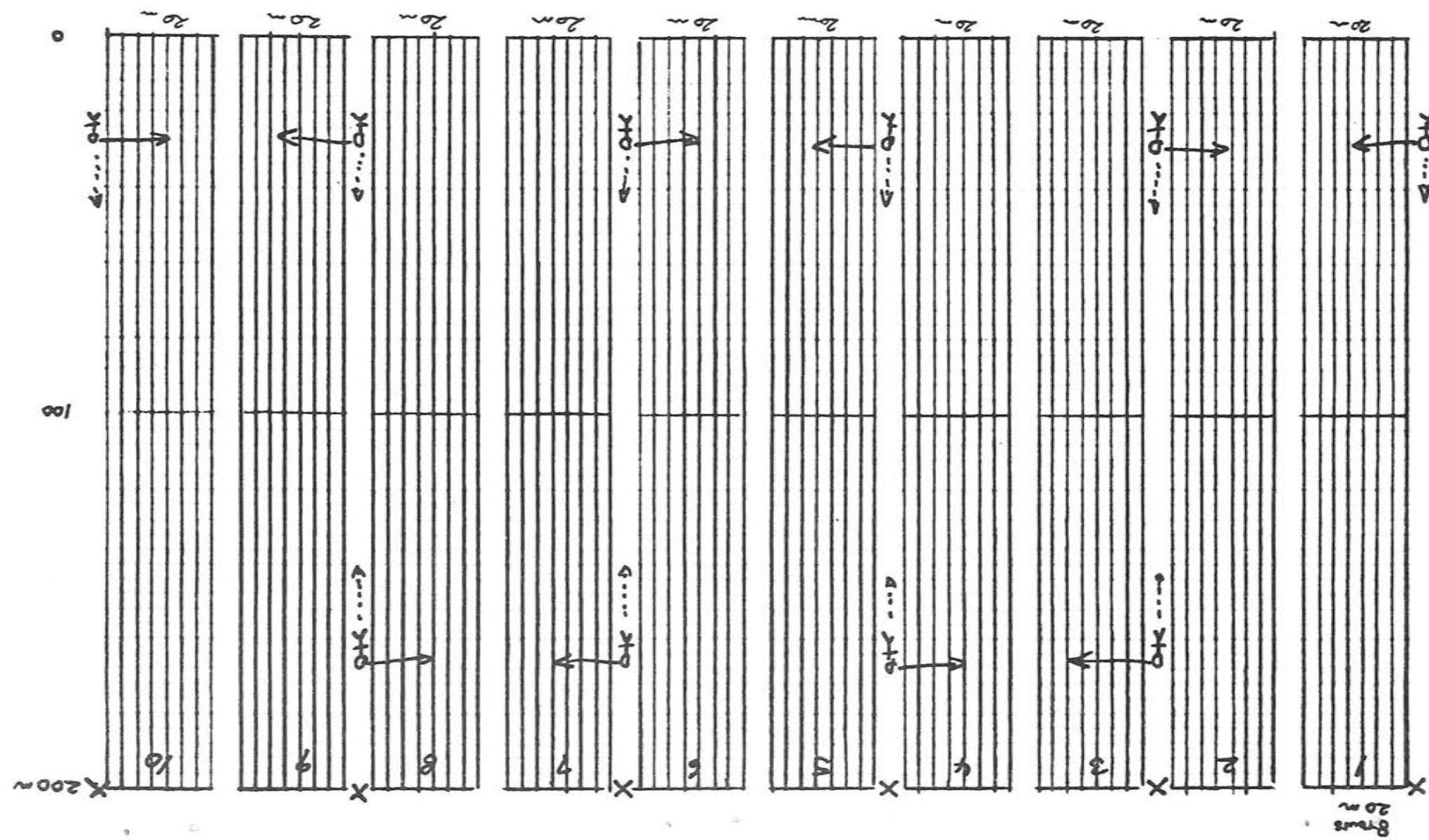
- \* Establish clone bank for all plus trees.
- \* Introduce the best of the plus trees into the general seed orchard.
- \* Raise progeny and plant the op progeny test and the family blocks on 2 representative sites.

6. FIELD

- \* Design comprehensive progeny tests and arrange for co-operative trials. Implement these screening tests as soon as possible.
- \* Register plus trees on plantation plus tree map; register plus trees on P.O.C.S.
- \* Calculate phenotypic values for each of the traits and rank plus trees for potential immediate use; top ranked trees to go into general seed orchard.

5. OFFICE

- \* Using rifle, collect minimum of 8 cones/plus tree; in each compartment where plus trees were collected randomly choose two other trees and collect 8 cones/tree to represent the bulk average population.
- \* Take photographs (3) of each plus tree to show bole, crown and total tree.
- \* Using rifle, in July/August collect scion material for grafting and establishment of clone banks.
- \* If 'plus' trees were not carrying cones, return in the following years to complete the cone collection.

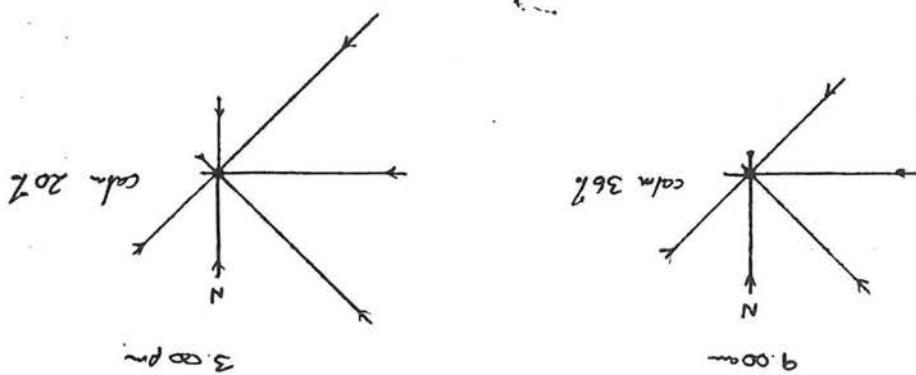




July 22, 1984 - I

FINE SEED ORCHARDS

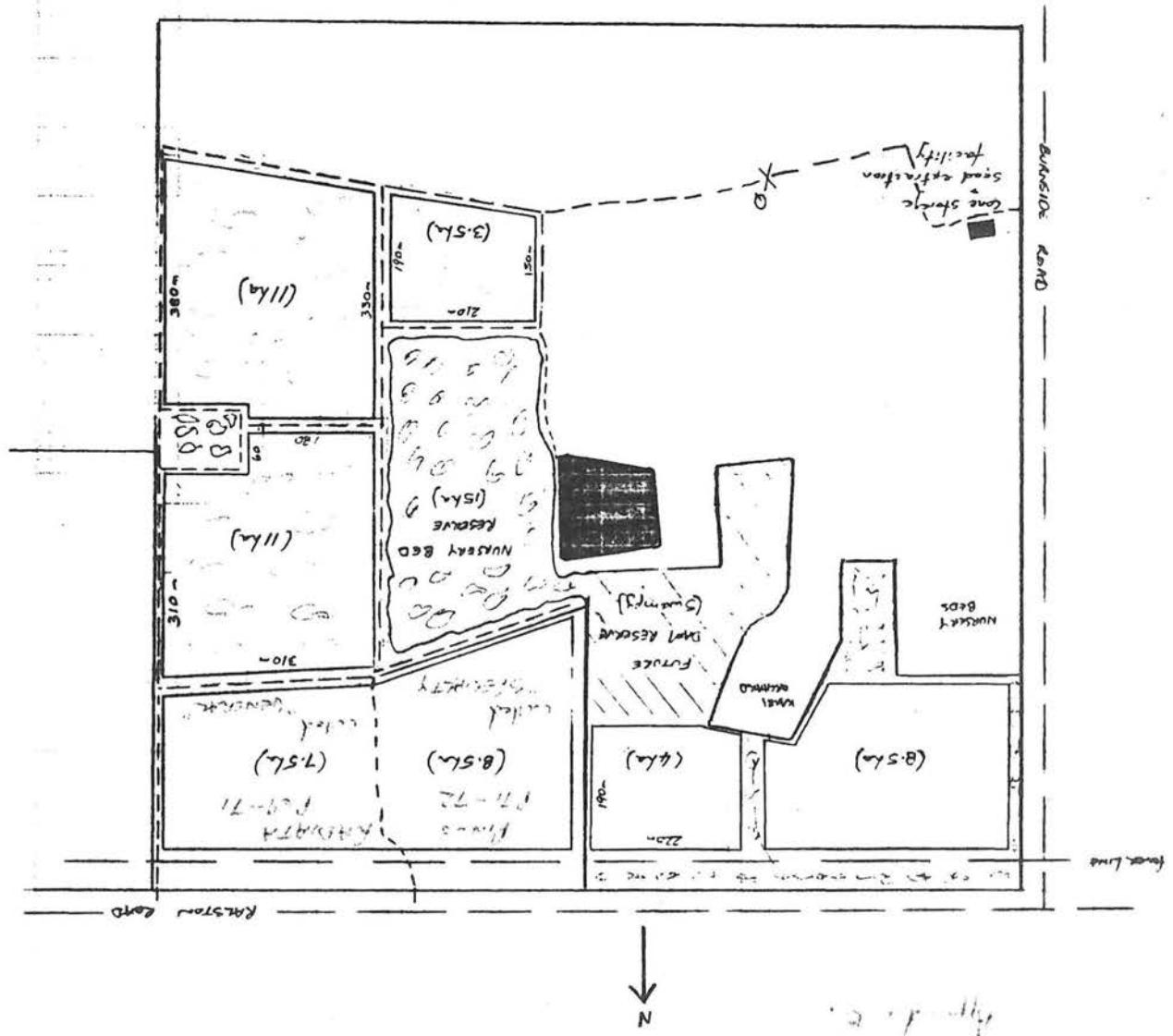
## WEST MOUNTAIN FORESTRY CENTRE



**AUGUST FREQUENCY PERCENTAGE**

## Alino J. Acetioan at Manjimup (1957-61)

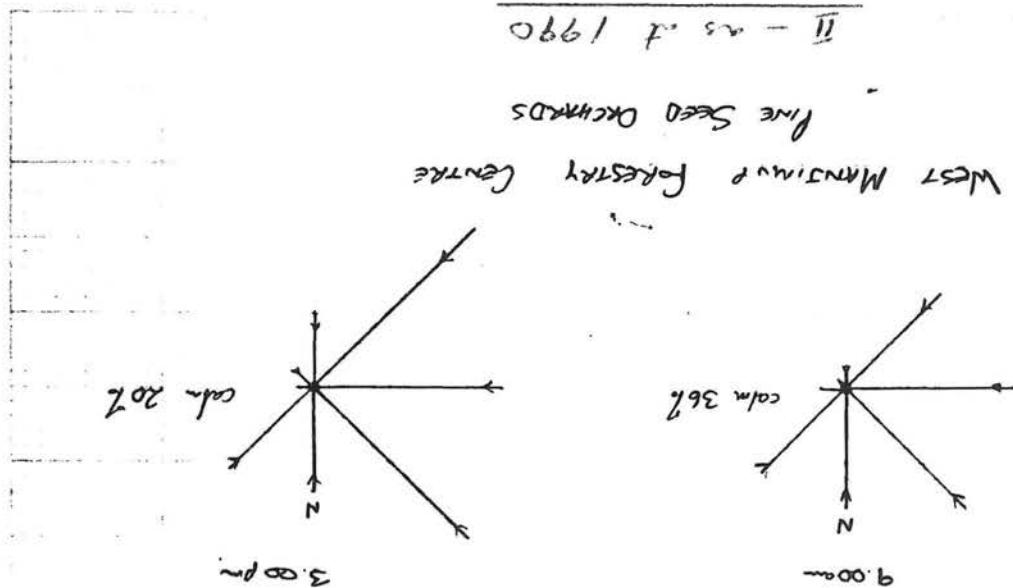
SCALE (MM) 1 cm = 84 m



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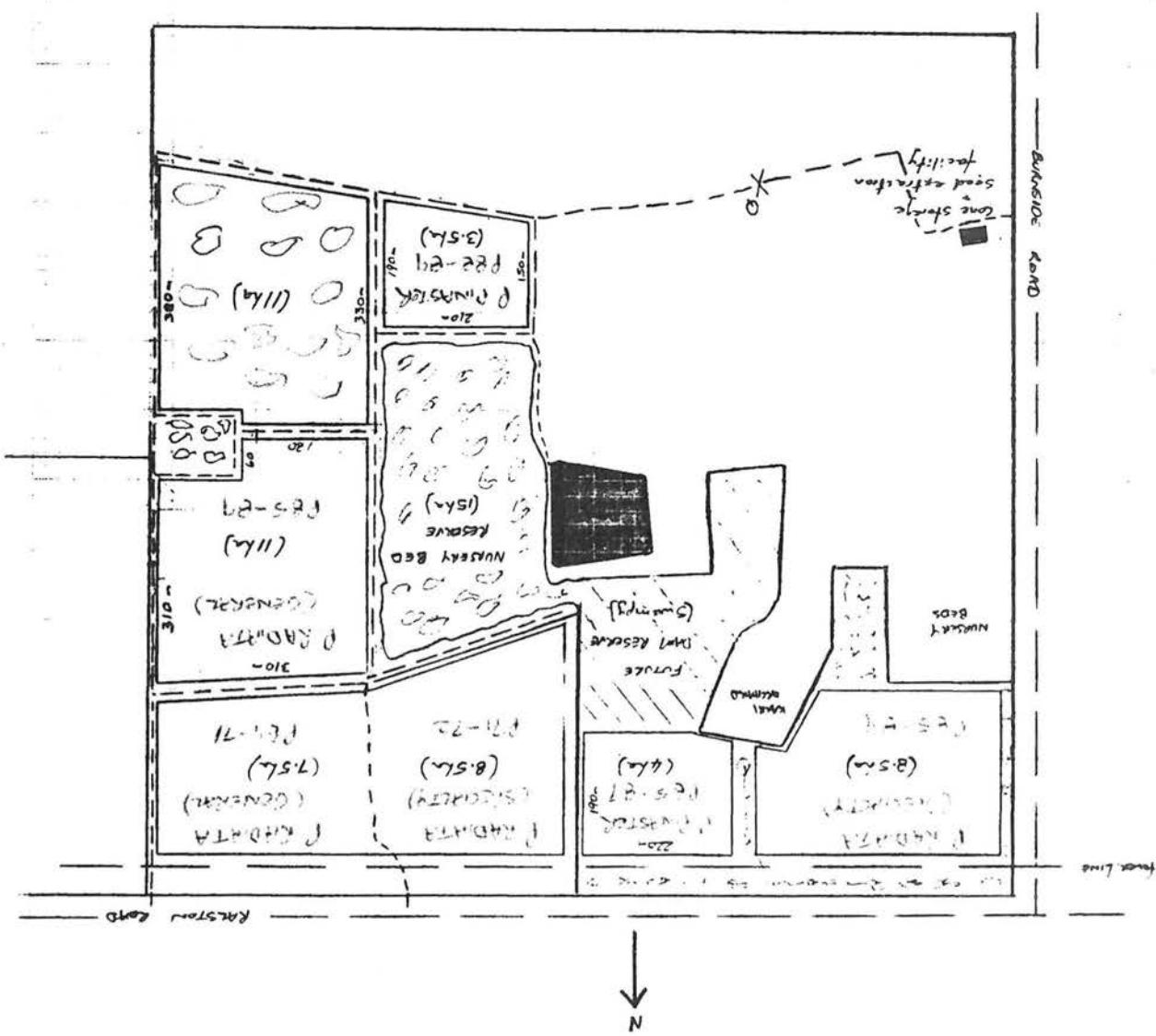
FINE SEED ORCHARDS

## WEST MOUNTAIN FOREST CENTRE



WINO D'ACCETTOU AT MANGIMUP (1957-61)  
AUGUST PRECIPITATION PRECIPITATION

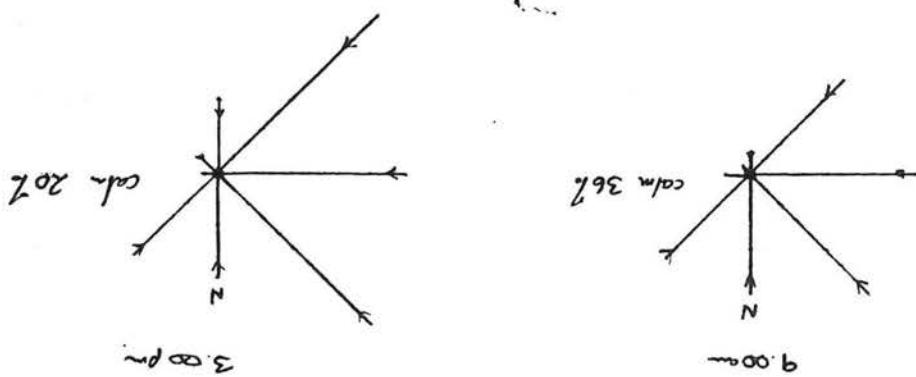
SCALE (mm) 1 cm = 84



III - 2003

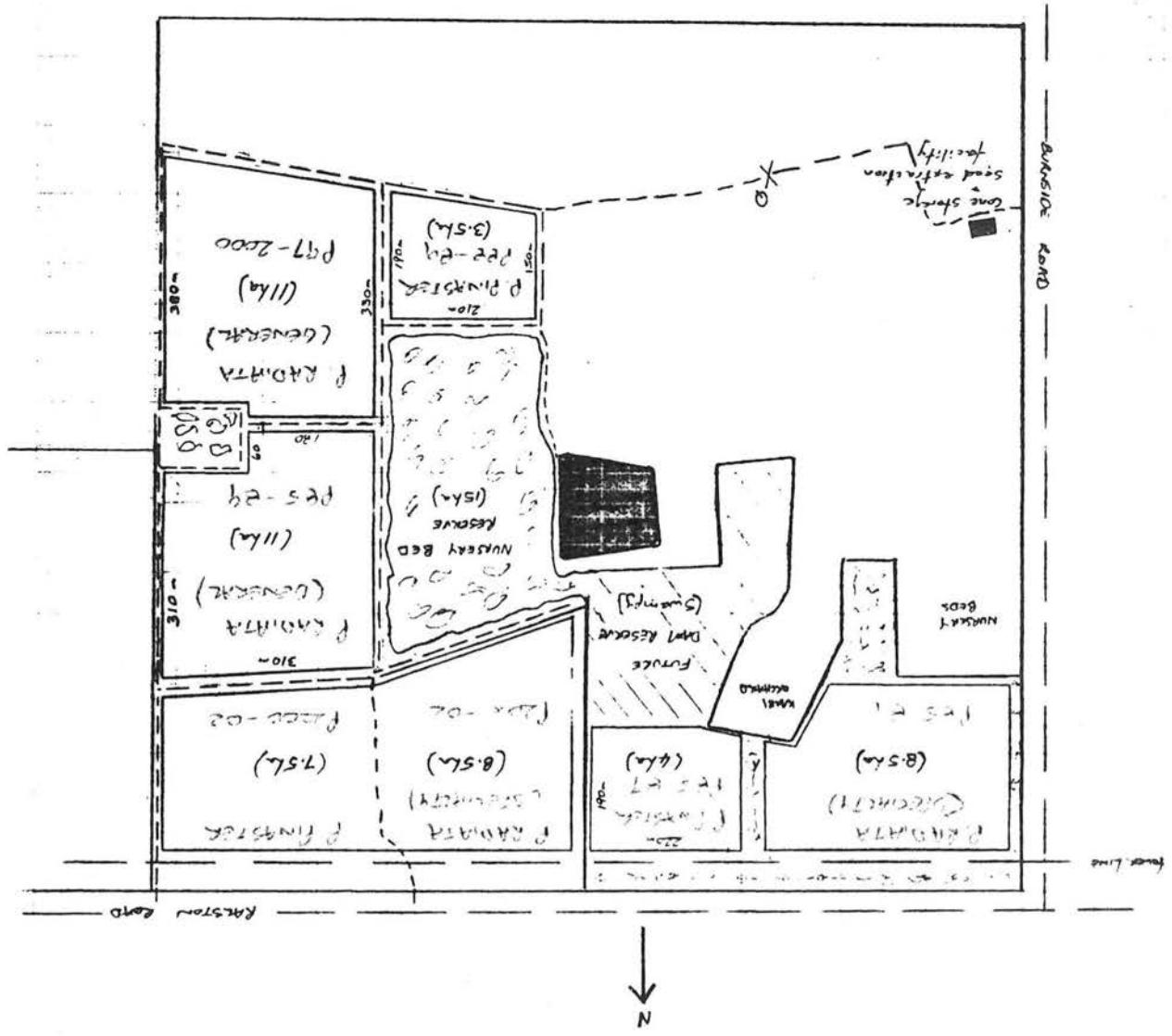
FINE SEED ORCHARDS

## WEST MOUNTAIN FOREST CENTRE



WIND DRAFTS AT MANGUM (1957-61)

SCALE (MM) 1 cm = 84.



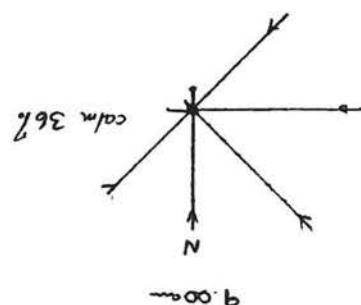
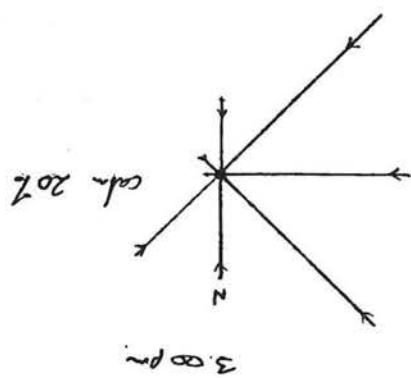
Przedmiescie (gmina) - 22 km  
Przedmiescie (szp) - 17 km  
Przedmiescie - 15 km

$$\text{Area of quadrilaterals} = 54 \text{ cm}^2$$

UV - as at 2015

## Fine Seeds & Chalks

## West Marine Forest Centre



WIND Distribution at Manzanares (1957-61) AUGUST PRECIPITATION PRECIPITATION

SCALE (MM) 1 cm = 84 m.

