

# Climate adaptation and provenance choice for revegetation: Insights from *Eucalyptus*

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# 1. Eucalypt populations exhibit a close adaptive response to their environment



LD Pryor

WD Jackson

- local-scale  
e.g. altitudinal or edaphic
- broad-scale  
e.g. macro-climate

Pryor LD (1957) Proc. Linn. Soc. NSW 81: 299-305

Barber & Jackson (1957) Nature 179: 1267-1269



# Numerous lines of evidence for climate adaptation

## 1. Correlative

- evidence of divergent selection  
e.g. divergence in functional trait ( $Q_{ST}$ ) exceeds neutral molecular expectations (drift) ( $F_{ST}$ )
- plus correlated with spatial variation in climate
- plus consistency with functional expectations

## 2. Direct (selection experiments)

- laboratory/glasshouse
- field trials:
  - *in situ* (e.g. reciprocal plantings in the wild)
  - *ex situ* (e.g. multi-site tree breeding trials)

# Range-wide seed collections and testing in breeding trials: the case of *E. globulus*

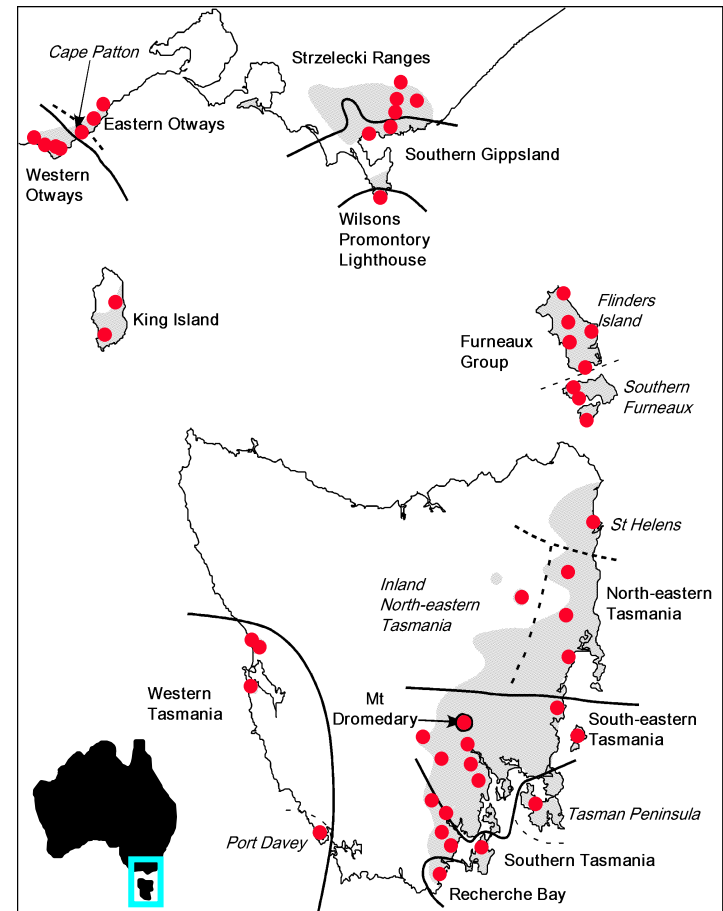
## Sampling localities

CSIRO 1987/88 base population seed collection

- 600+ open-pollinated families
- 46 localities



Trials designed with randomization at the family level (e.g. randomized incomplete block designs, 2 tree plots, 5 reps)



Dutkowski and Potts 1999

Lopez *et al.* 2001

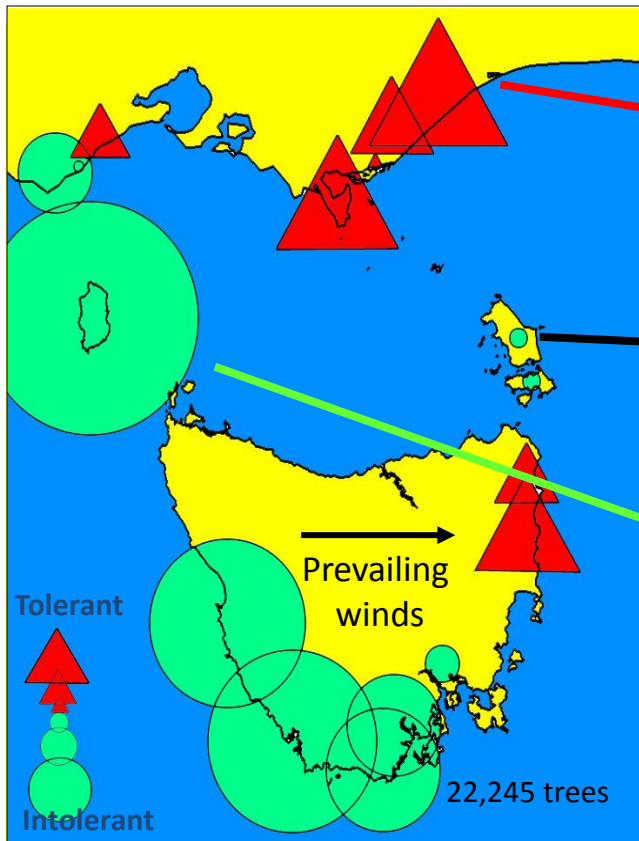
# Seed lots from the 87/88 CSIRO collection of *E. globulus* have been tested globally



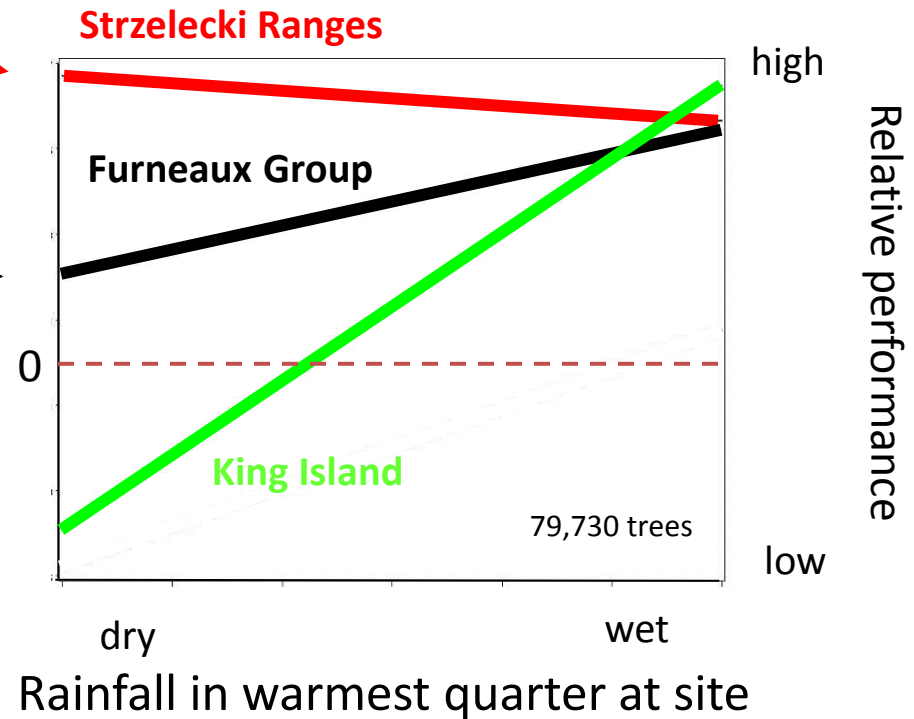


# Home-site climate predicts susceptibility to drought and growth on dry sites

## Drought tolerance



## Tree growth (dbh)



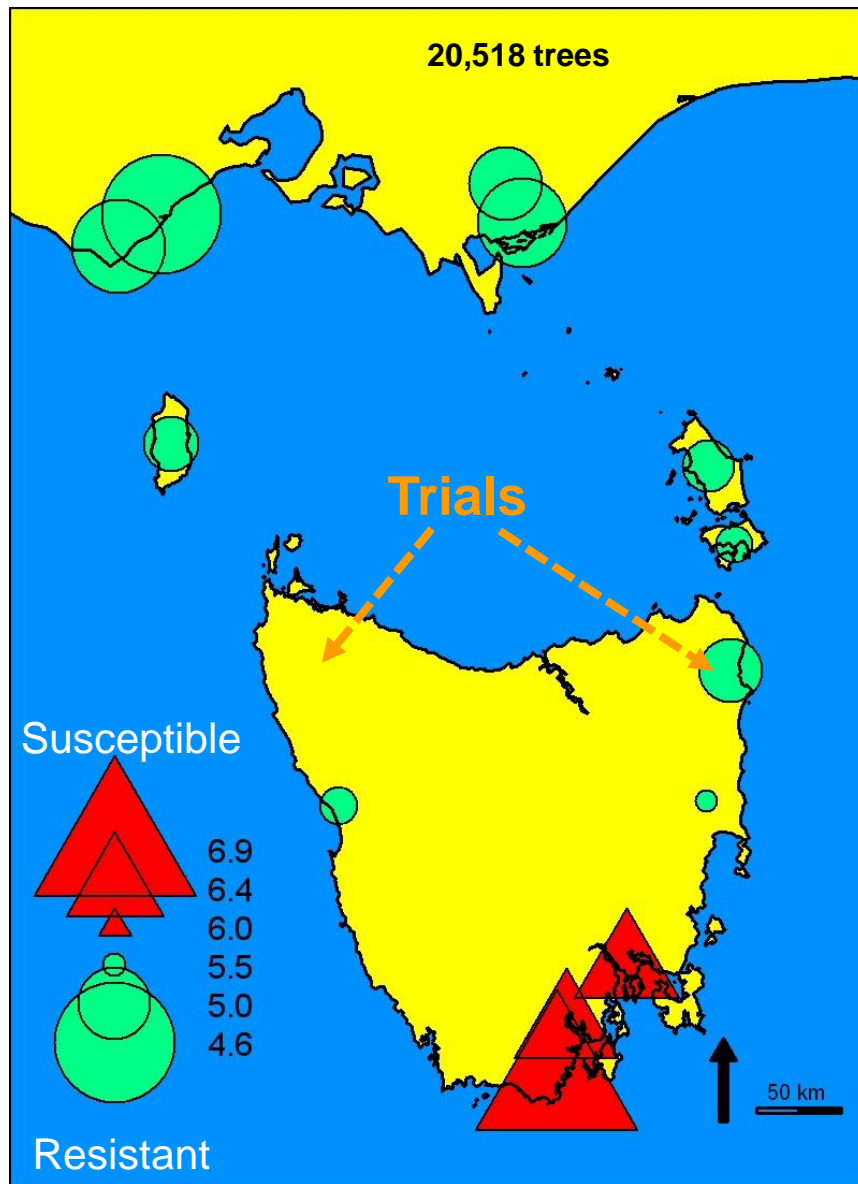
Costa e Silva *et al.* 2006 *Tree Genetics and Genomes* 2, 61-75.

1993/4 drought damage in 4 WA trials (5 year old)

Dutkowski and Potts 2012 *Tree Genetics & Genomes* 8, 757-773.

# Climate adaptation is multi-trait and involves selection by both abiotic and biotic factors

Genetic resistance to *Teratosphaeria* leaf disease increases in home-sites with high disease risk (i.e. high temperatures and high autumn rainfall)



% juvenile foliage damaged in 2007  
in 4 field trials in Tasmania

Hamilton *et al.* 2013, *Heredity* 110, 372–379







Climate adaptation is likely a  
genome-wide phenomenon

As sedentary, dominant terrestrial  
organisms, trees are likely to be  
strongly coupled, and exposed, to  
changes in the macro-climate

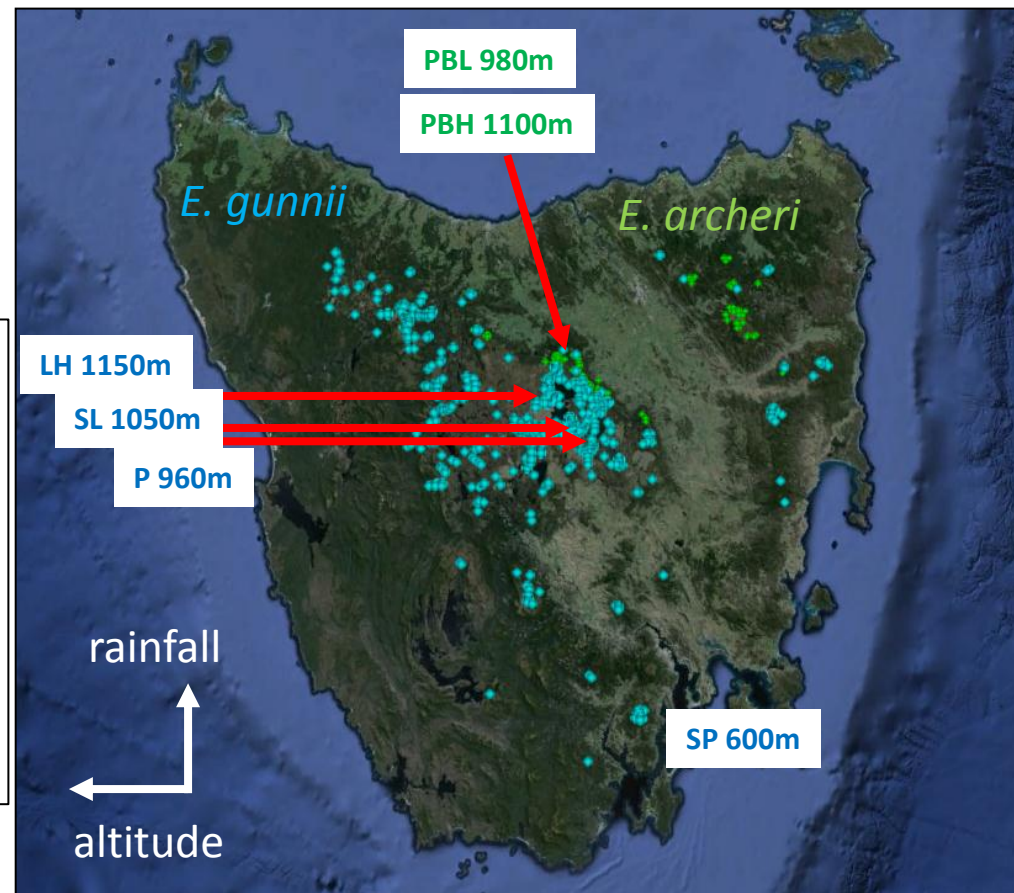


## 2. Climate change is impacting native gene pools and local provenance fitness

Clinal variation in the *E. gunnii*-*archeri* complex on the Central Plateau, Tasmania

Reciprocal transplant trials established in 1979 within wild populations:

- 4 sites
- 6 provenances
- 12 plants/provenance/site



Potts 1985, Aust J Bot 33: 687-704

Potts et al. 2001, Pap Proc R Soc Tasm 135: 57-61

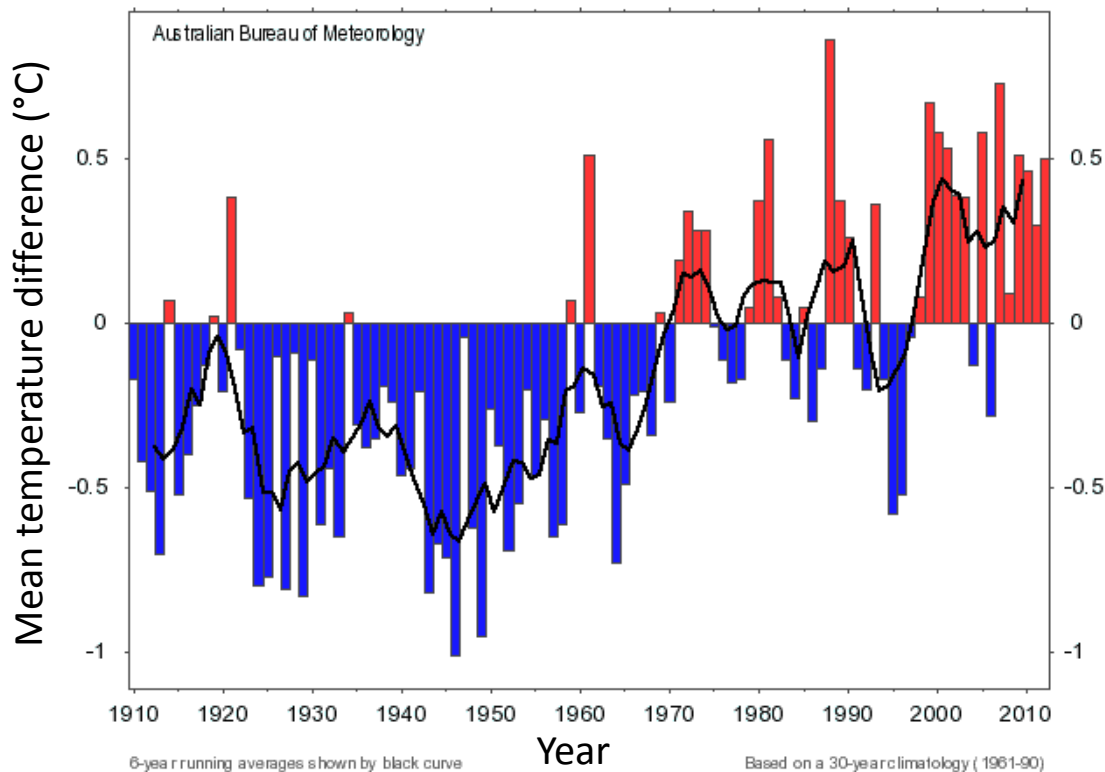
Calder & Kirkpatrick 2008, Aust J Bot 56: 684-692

Kremer et al. 2014, Functional Ecology 28: 22-36

# Local climates are changing

Central Plateau region has experienced declining autumn rainfall since 1970's and increasing temperatures

Deviation from long-term mean temperature in  
Tasmania (1910-2012)

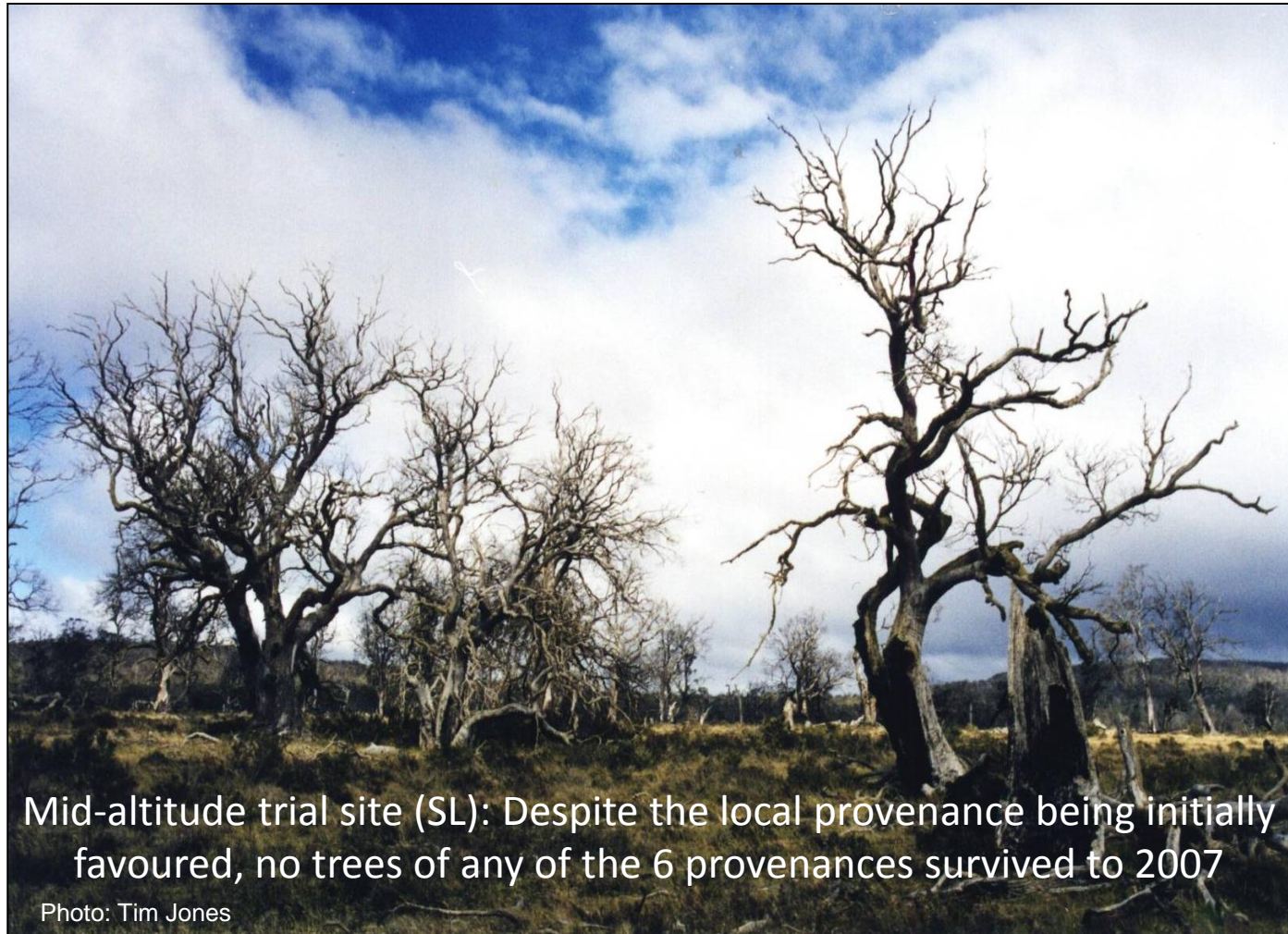


What was once 'locally' adapted may not be 'locally' adapted now or in the future



# ...and eucalypt populations are affected

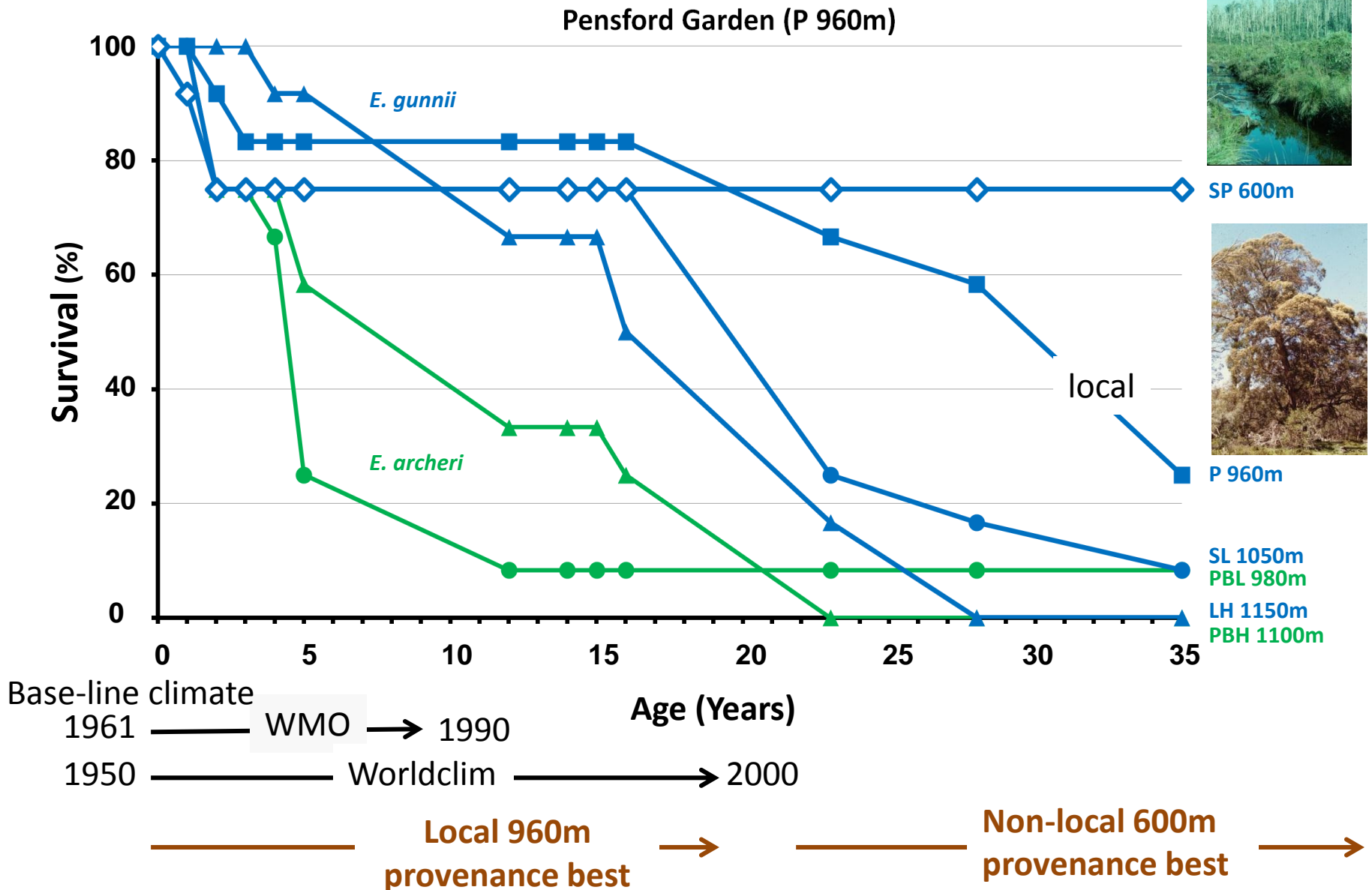
Major adult tree decline commenced in the early 1990's



Mid-altitude trial site (SL): Despite the local provenance being initially favoured, no trees of any of the 6 provenances survived to 2007

Photo: Tim Jones

# Local provenance no longer best at the benign low altitude site

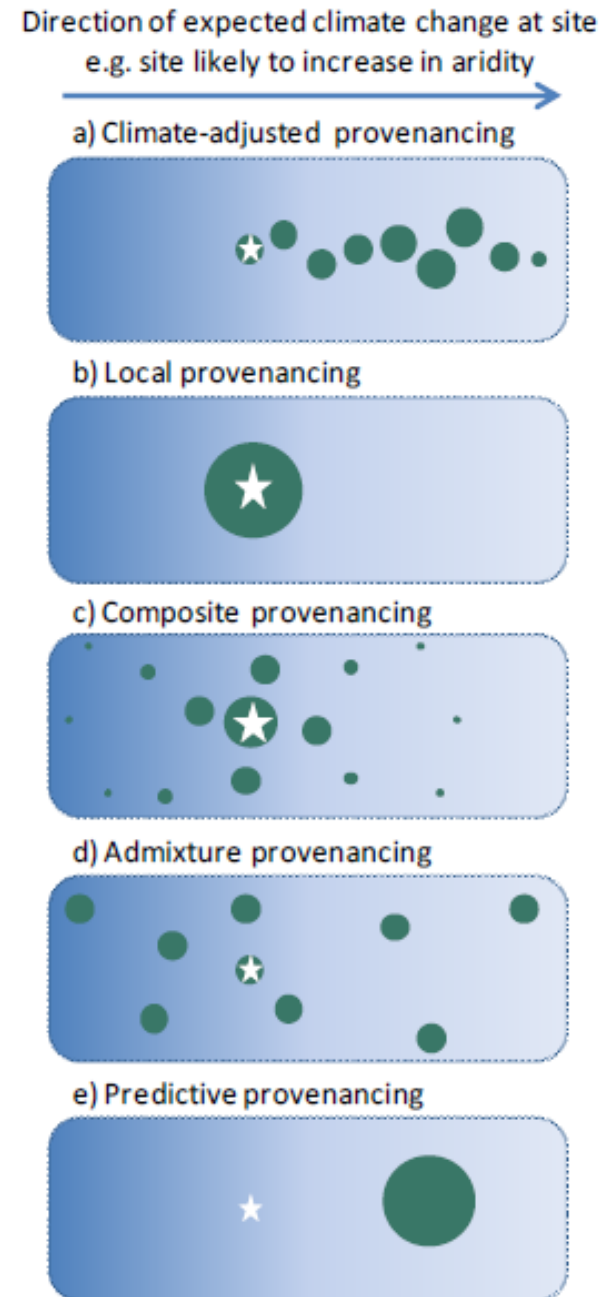




### 3. What are we doing?

## Testing provenance strategies for revegetation

Figure from Prober *et al.* 2015. *Frontiers in Ecology and Evolution* (doi: 10.3389/fevo.2015.00065)



# Building a long-term research infra-structure within GA's restoration plantings in Tasmania

c. 20% of restoration plantings are pedigreed local, multi-provenance and multi-species trials

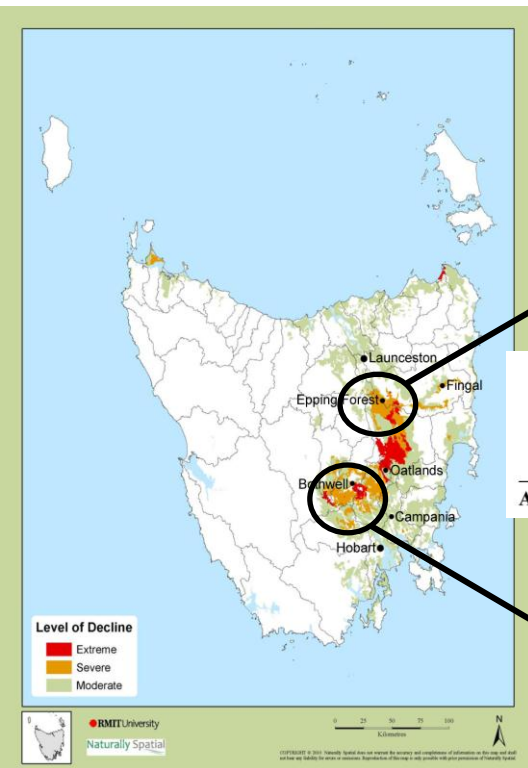
Connorville est. 2014



Dungrove est. 2010



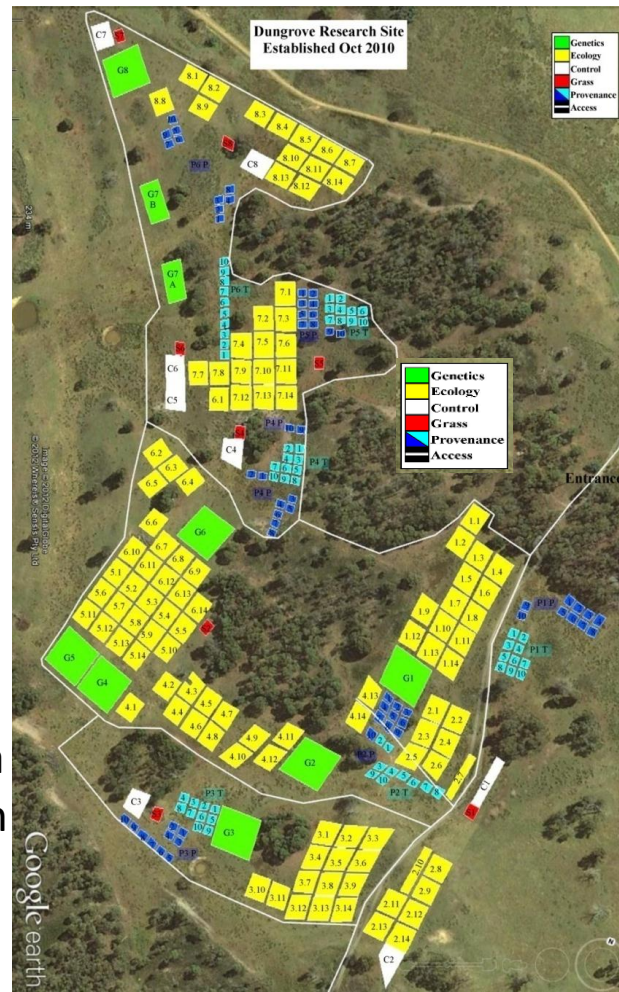
Bailey *et al.* 2013, 2015  
Gauli 2014



Two c. 20  
ha trial  
sites in  
2014

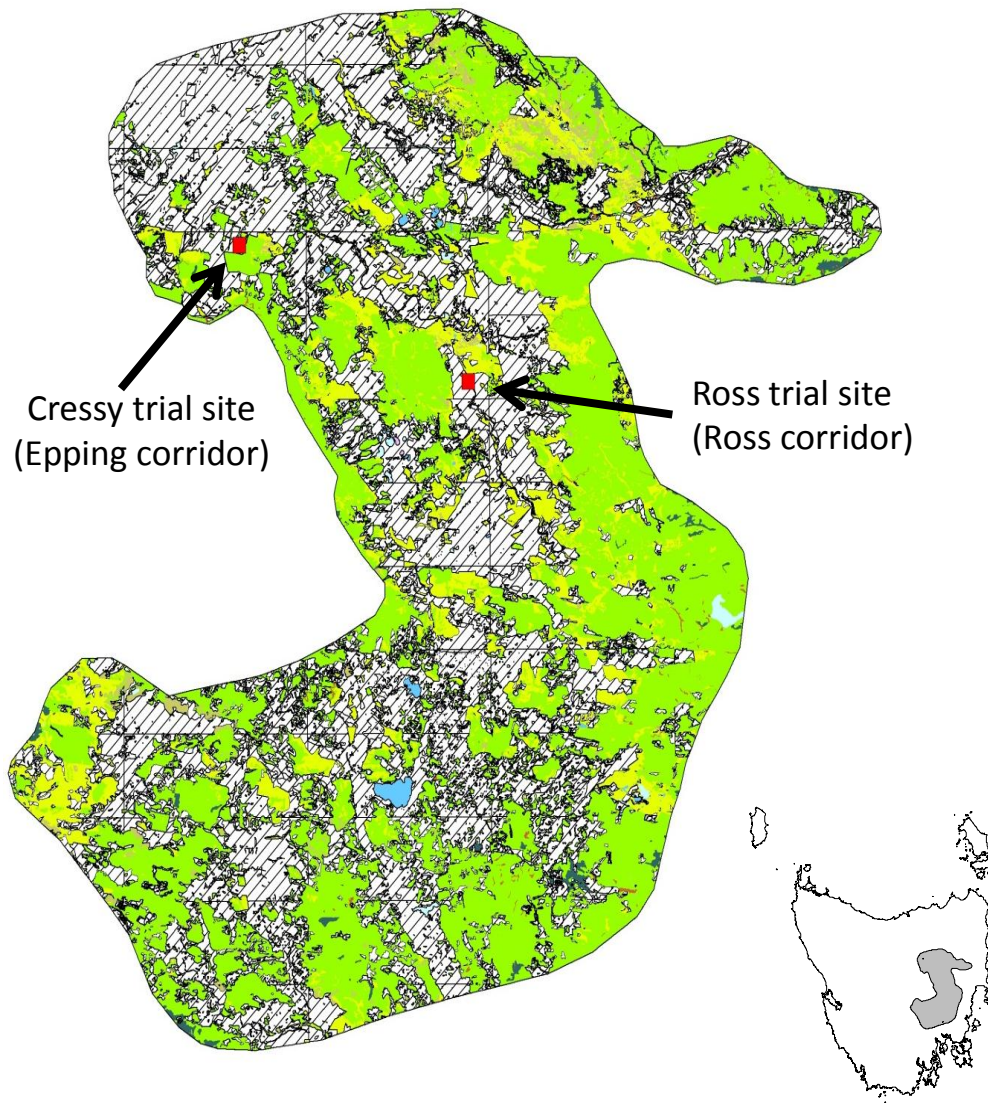


Three 30 ha  
trial sites in  
2010-11





# The Midlandscapes project



Consortium of stakeholders including GA, TLC, Bush Heritage, DPIPW and UTAS

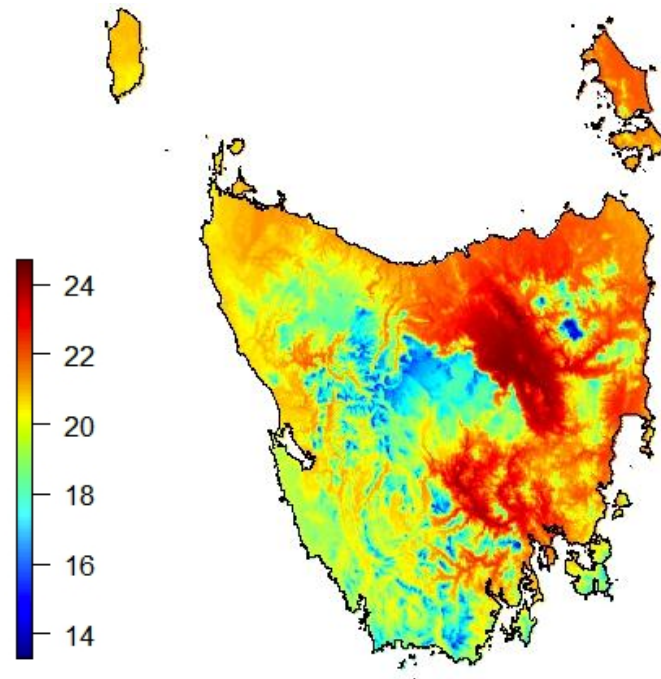
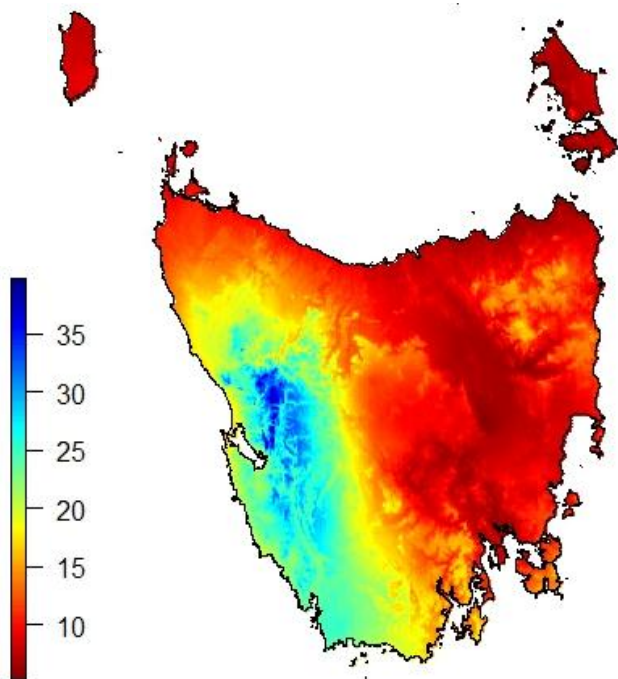
Midlandscape encompasses the Midlands Biodiversity Hotspot defined as the lowland plains and foothills between the Eastern Tiers and Western Tiers

Area of active research and habitat restoration projects



# A challenging climate – hot & dry (by Tasmanian standards)

- Lowest rainfall within the state
- Epicenter of high temperature extremes

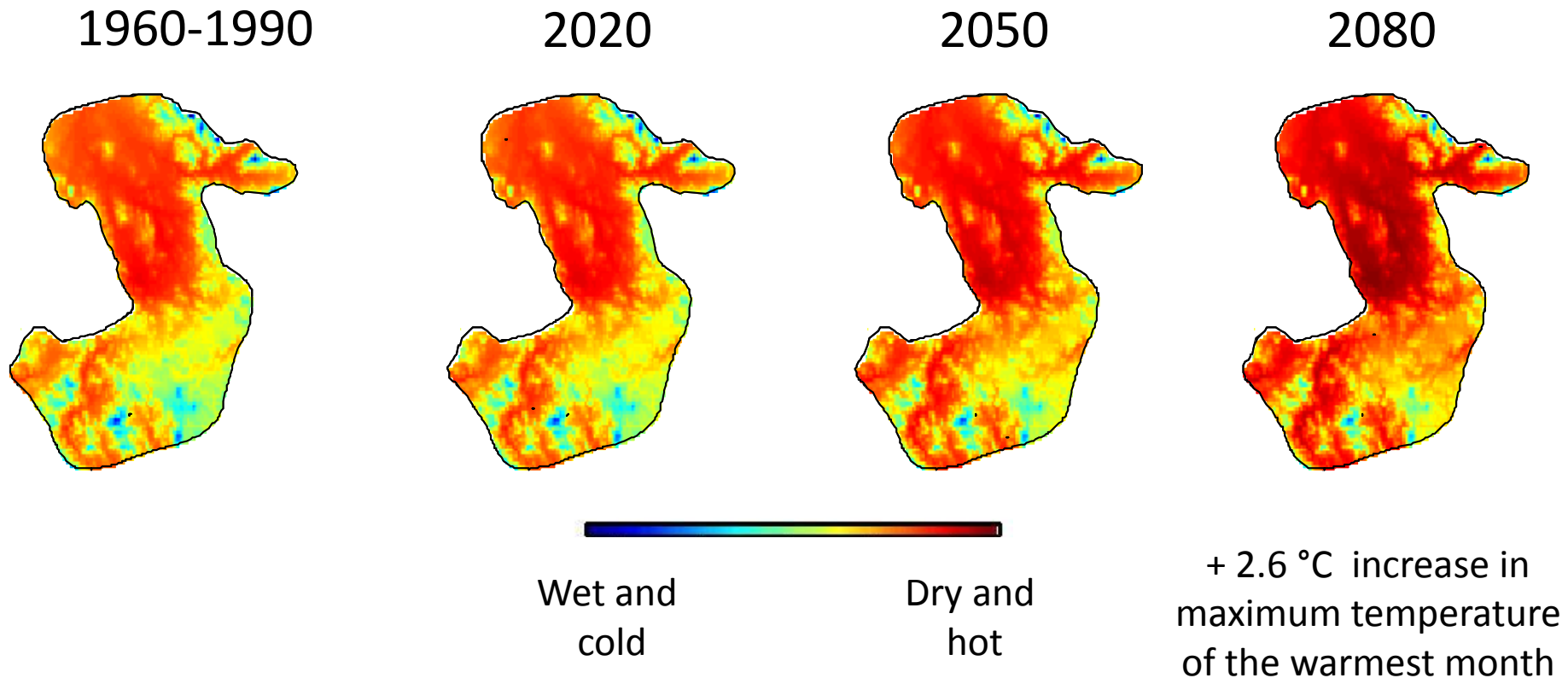


Precipitation of the driest month (mm)    Maximum temperature of the warmest month (°C)

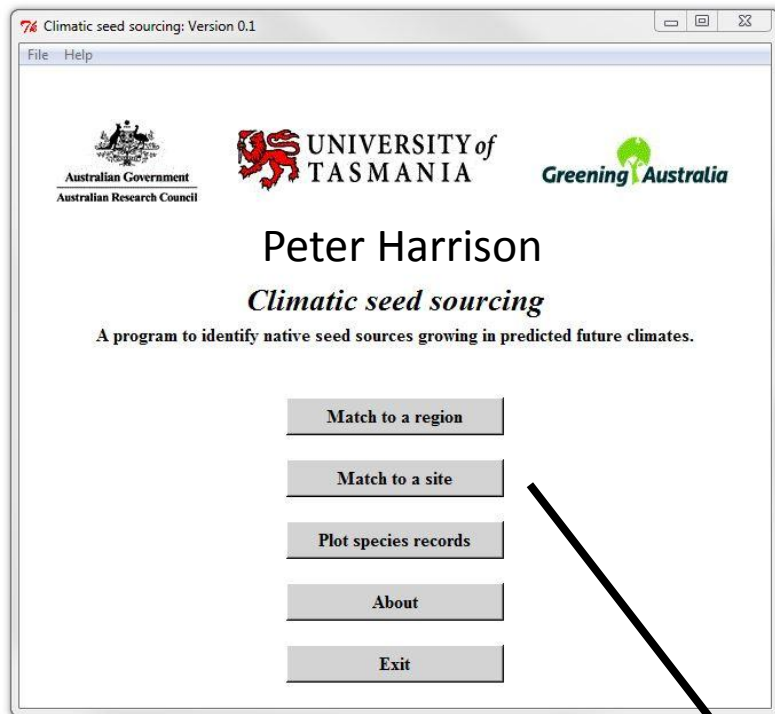


# Will get worse

Midlandscape predicted to become more arid during summer in the future



# Seed sourcing for future climates – the program



Options of the program

Climate-match a species to a region

Climate-match a species to a site

Set multiple parameters for the model and select the species you want to match

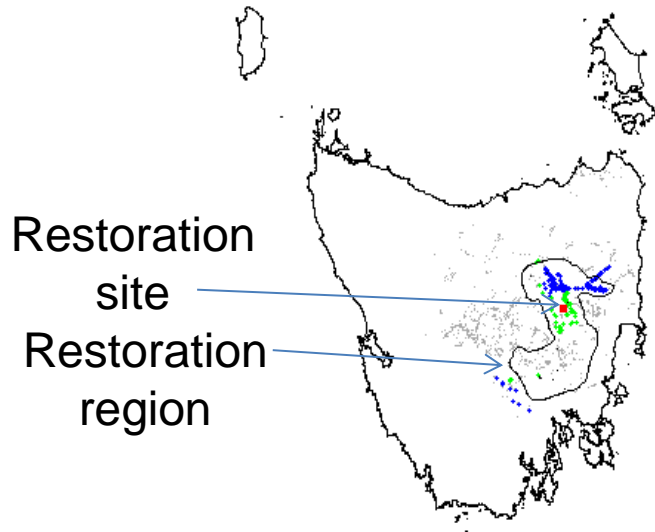
The 'Set Parameter Values' dialog box is titled 'Climatic matching to a site'. It contains the following fields and options:

- Set parameter values**
- Enter coordinates of the site(s)**
  - Latitude: (use a comma to separate multiple sites) -41.98078
  - Longitude: (use a comma to separate multiple sites) 147.4571
  - Select which region to restrict modelling to: [Dropdown menu]
- Select which species to climatically match to bioregion:**

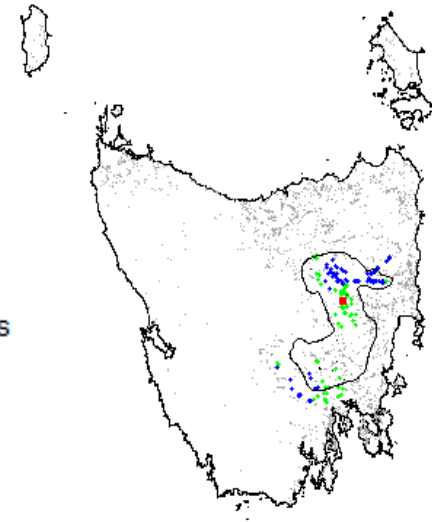
<i>Eucalyptus amygdalina</i>	<input type="checkbox"/>	<i>Eucalyptus pauciflora</i>	<input type="checkbox"/>
<i>Eucalyptus tenuiramis</i>	<input type="checkbox"/>	<i>Eucalyptus ovata</i>	<input type="checkbox"/>
<i>Eucalyptus rodwayi</i>	<input type="checkbox"/>	<i>Eucalyptus viminalis</i>	<input type="checkbox"/>
- Climatic threshold:** 2
- Keep results with a threshold less than:** 5
- Minimum number of points to return to screen:** 20
- What distance measure to use:** euclidean
- Plot output: (TRUE / FALSE)** TRUE
- How big is the subsampling radius around site(s): (km)** 5
- How many points to subsample within radius:** 20
- Buttons:** Run, Exit

# Seed sourcing for future climates – the output

*E. pauciflora* (2050s)



*E. ovata* (2050s)



pauciflora\_1\_2050

lon	lat	collect	ClimDist	GeoDist	location	frag
147.75	-41.757	1	0.94591	34.94	Avoca	1
147.8	-41.739	1	0.97314	39.28	Fingal	0
147.79	-41.746	1	0.98688	37.78	Fingal	1
147.79	-41.742	1	0.98688	38.2	Fingal	1
147.77	-41.752	1	1.04518	36.07	Avoca	1

ovata\_1\_2050

lon	lat	collect	ClimDist	GeoDist	location	frag
146.87	-42.687	1	0.82255	92.29	Glenora	1
146.87	-42.689	1	0.82255	92.4	Glenora	1
147.75	-41.757	1	0.94591	34.94	Avoca	1
147.8	-41.739	1	0.97314	39.28	Fingal	0
147.79	-41.746	1	0.98688	37.78	Fingal	1

Climate-adjusted provenancing





# Field testing seed sourcing strategies

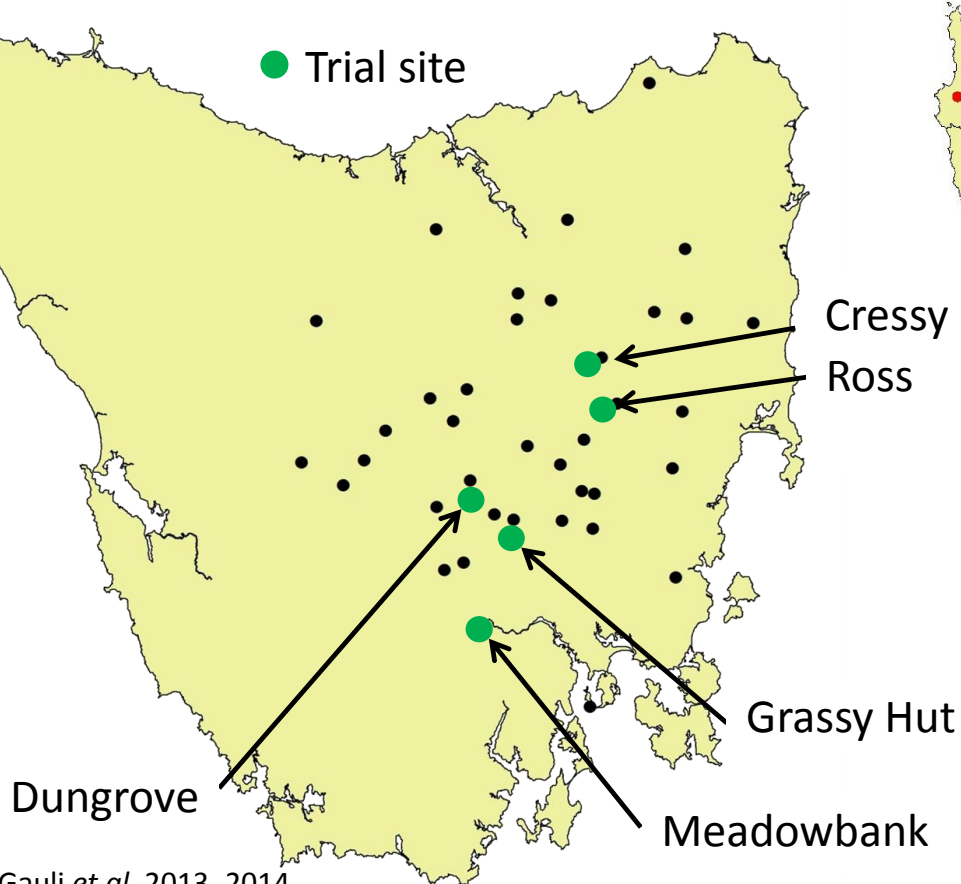
Seed collection localities in Tasmania (local + non local)

## *Eucalyptus pauciflora* [●]

280 Tasmanian trees (families)

37 Tasmanian provenances

13 Mainland provenances



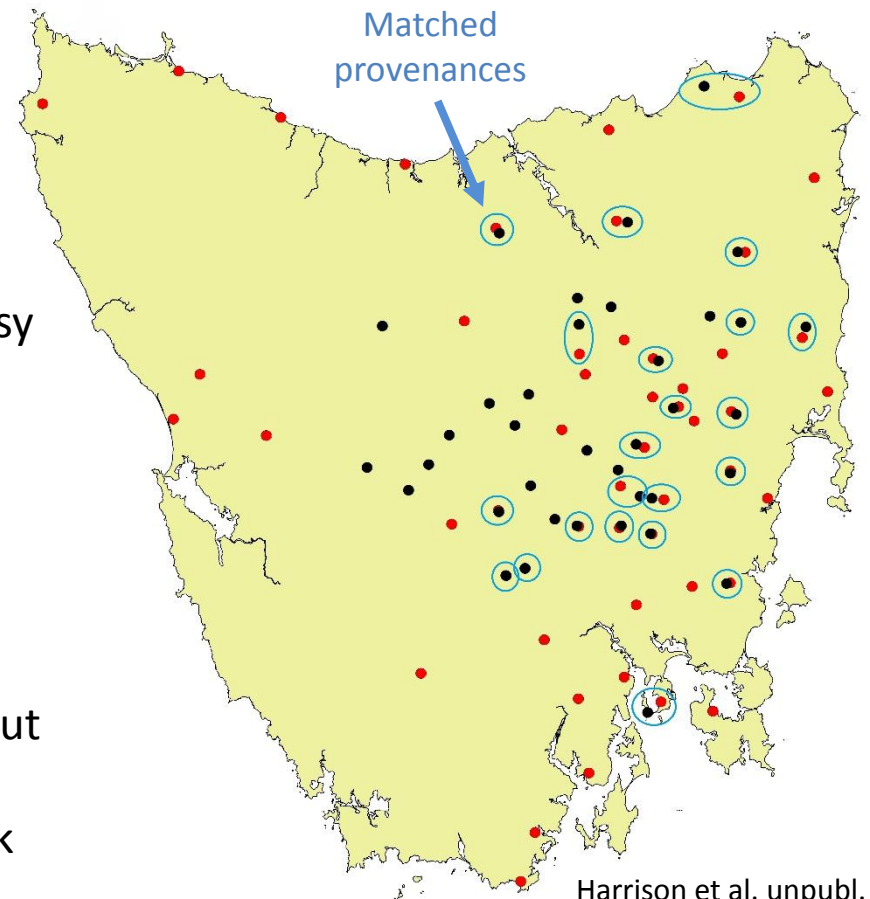
Gauli et al. 2013, 2014

## *Eucalyptus ovata* [●]

400 Tasmanian trees (families)

51 Tasmanian provenances

(22 paired with *E. pauciflora*)



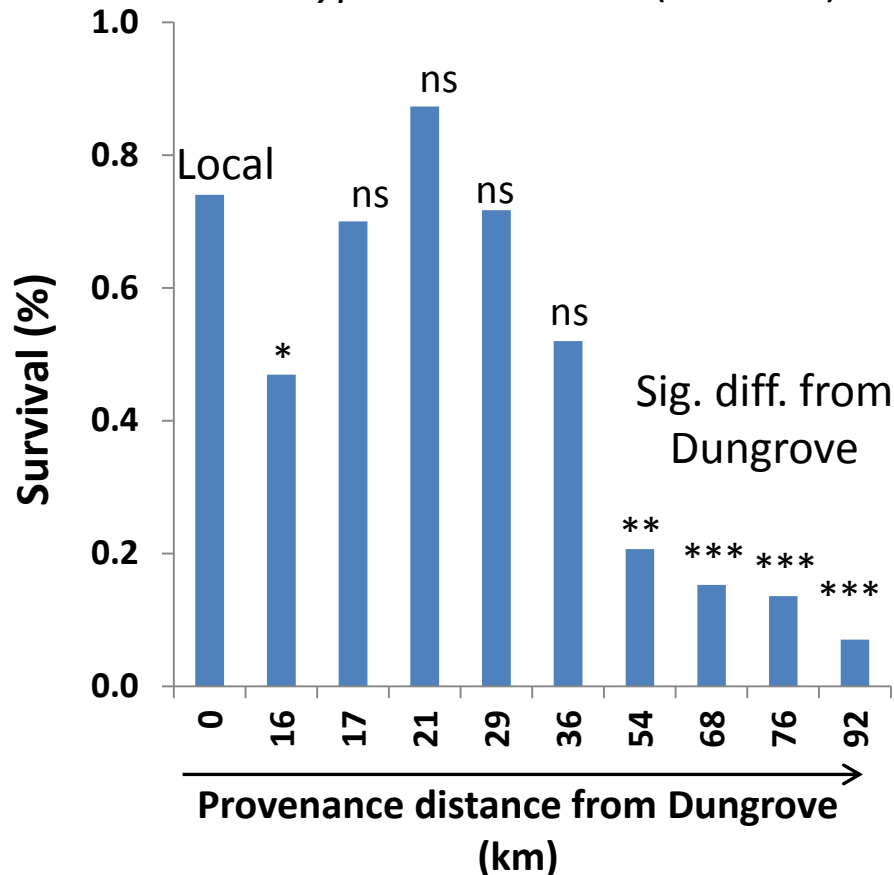
Harrison et al. unpubl.

# Climate change thresholds may be species-specific

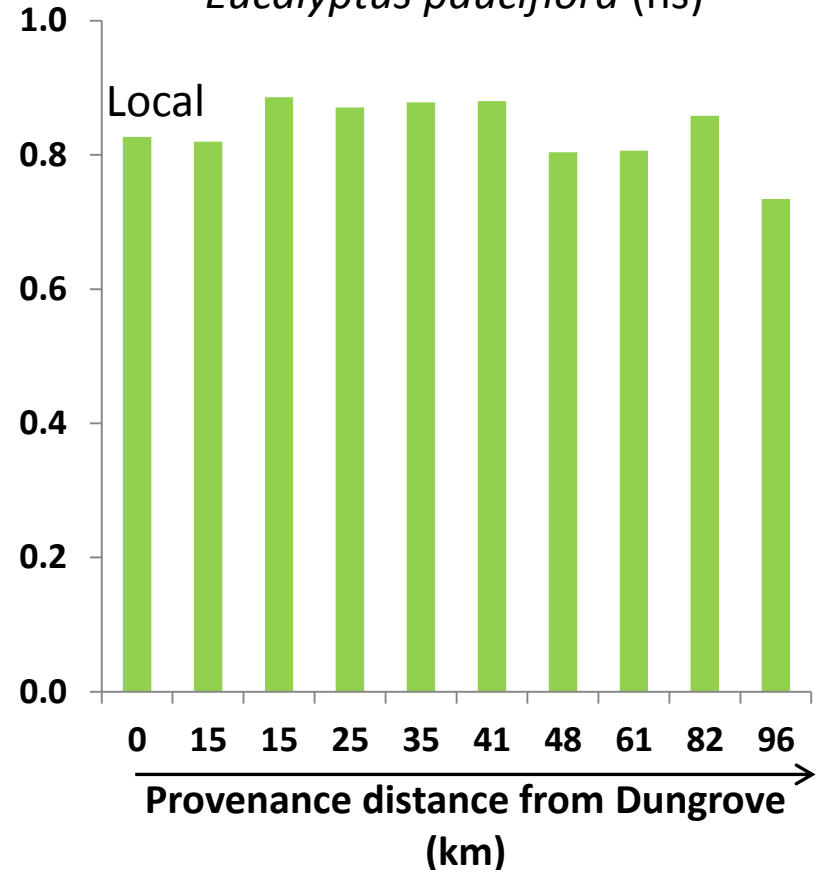
## Dungrove provenance trial

3 year survival in blocks of 25 trees from 10 provenances replicated 8 times

*Eucalyptus tenuiramis* ( $P < 0.001$ )



*Eucalyptus pauciflora* (ns)



# 4. There may be ways to short-cut field testing

## MOLECULAR ECOLOGY

Molecular Ecology (2014) 23, 2500–2513

doi: 10.1111/mec.12751

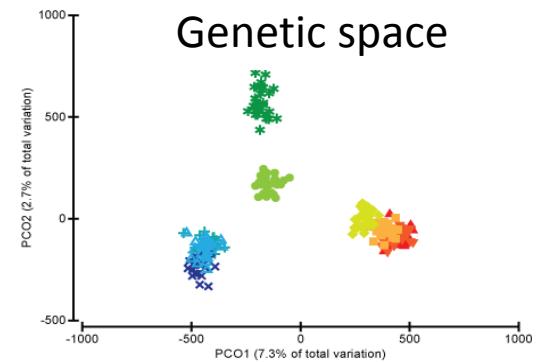
### Genome-wide scans detect adaptation to aridity in a widespread forest tree species

DOROTHY A. STEANE,\*†<sup>1</sup> BRAD M. POTTS,\*<sup>2</sup> ELIZABETH MCLEAN,‡ SUZANNE M. PROBER,§ WILLIAM D. STOCK,¶ RENÉ E. VAILLANCOURT\* and MARGARET BYRNE‡

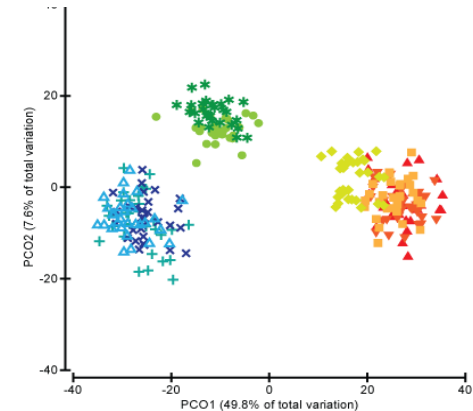
Genomic approaches to identify climatic drivers of adaptive variation

## The case of *E. tricarpa*

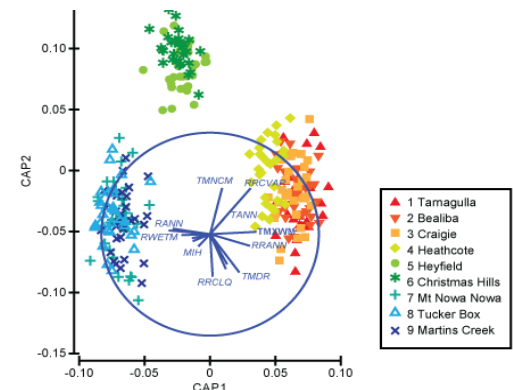
A combination of climate variables ‘aridity index’ maximally correlated with genome-wide changes in markers showing signatures of disruptive selection



## Adaptively enriched genetic space



## Climatically aligned adaptive genetic space



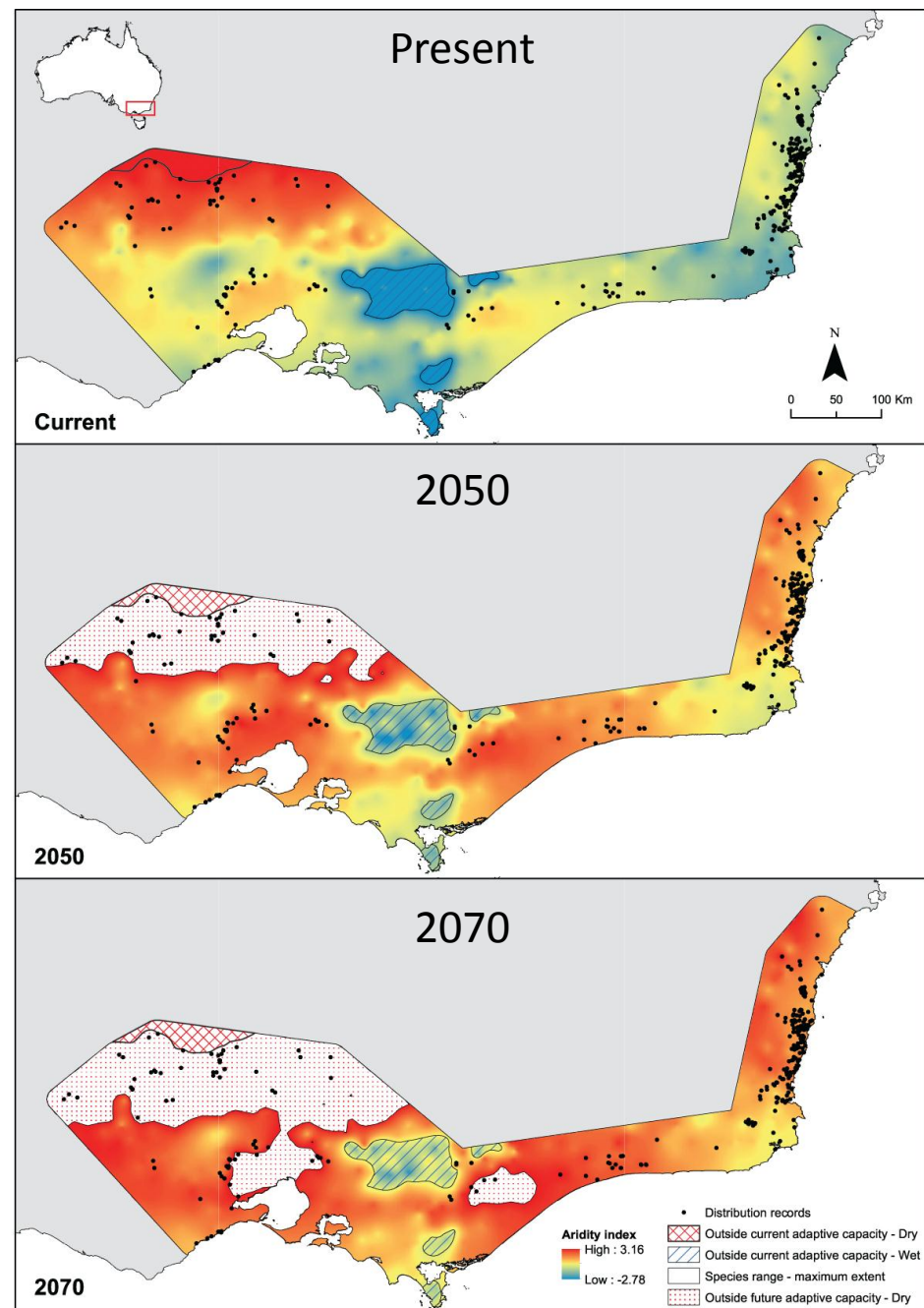


# Predicting adaptive surfaces for contemporary and future climates

Weighting climatic variables and their change according to likely selective impact

Can local populations adapt from the standing genetic variation and natural gene flow?

Harrison *et al.* unpubl data



CSIRO global climate model for 2050 and 2070



# 5. A wildcard: exotic pests and diseases

Exotic pests and diseases may impact the ecological and evolutionary trajectory of revegetation?

Feral fallow deer  
(*Dama dama*)

Deer damage at Dungrove was significantly greater on the faster growing, low altitude *E. pauciflora* provenances (Gauli 2014; Bailey *et al.* 2015)





# Thank you





# References

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