



Eocene to Oligocene vegetation and climate in the Tasmanian Gateway region controlled by changes in ocean currents and $p\text{CO}_2$

Michael Amoo¹, Ulrich Salzmann¹, Matthew J. Pound¹, Nick Thompson¹, and Peter K. Bijl²

¹ Department of Geography and Environmental Sciences, Northumbria University, Newcastle upon Tyne, UK

² Marine Palynology and Palaeoceanography, Utrecht University, Princetonlaan 8A, Utrecht, The Netherlands

Correspondence to: Michael Amoo (michael.amoo@northumbria.ac.uk)

Abstract. Considered as one of the most significant climate reorganisations of the Cenozoic period, the Eocene-Oligocene Transition (EOT; ca. 34.44–33.65) is characterised by global cooling and the first major glacial advance on Antarctica. While in the southern high-latitudes, the EOT cooling is primarily recorded in the marine realm, the extent and effect on terrestrial climate and vegetation is poorly documented. Here, we present a new, well-dated, continuous, high-resolution palynological (sporomorph) data and quantitative sporomorph-based climate estimates recovered from the East Tasman Plateau (ODP Site 1172) to reconstruct climate and vegetation dynamics from the late Eocene (37.97 Ma) to early Oligocene (33.06 Ma). Our results indicate three major climate transitions and four vegetation communities occupying Tasmania under different precipitation and temperature regimes: (i) a warm-temperate *Nothofagus*-Podocarpaceae dominated rainforest with paratropical elements from 37.97–37.52 Ma; (ii) cool-temperate *Nothofagus* dominated rainforest with secondary Podocarpaceae rapidly expanding and taking over regions previously occupied by the warmer taxa between 37.306–35.60 Ma; (iii) fluctuation between warm temperate - paratropical taxa and cool temperate forest from 35.50–34.49 Ma, followed by a cool phase across the EOT (34.30–33.82 Ma); (iv) post-EOT (earliest Oligocene) recovery characterised by a warm-temperate forest association from 33.55–33.06 Ma. Coincident with changes in stratification of water masses and sequestration of carbon from surface water in the Southern Ocean, our sporomorph-based temperature estimates between 37.52 Ma and 35.60 Ma (phase ii) showed 2–3 °C terrestrial cooling. The unusual fluctuation between warm and cold temperate forest between 35.50 to 34.59 Ma is suggested to be linked to the initial deepening of the Tasmanian Gateway allowing eastern Tasmania to come under the influence of warm water associated with the proto-Leeuwin Current (PLC). Further to the above, our terrestrial data show mean annual temperature declining by about 2 °C across the EOT before recovering in the earliest Oligocene. This phenomenon is synchronous with regional and global cooling during the EOT and linked to declining $p\text{CO}_2$. However, the earliest Oligocene climate rebound along eastern Tasmania is linked to transient recovery of atmospheric $p\text{CO}_2$ and sustained deepening of the Tasmanian Gateway, promoting PLC throughflow. The three main climate transitional events across the studied interval (late Eocene–earliest Oligocene) in the Tasmanian Gateway region suggest that changes in ocean circulation due to accelerated deepening of the Tasmanian Gateway may not have been solely responsible for the changes in terrestrial climate and vegetation dynamics, but a series of regional and global events, including a change in stratification of water masses, sequestration of carbon from surface waters, and changes in $p\text{CO}_2$ may have played vital roles.



35 1. Introduction

Palynological reconstruction demonstrate a high sensitivity of global vegetation to past changes in climate, leading to major shifts in biome distribution (Pound and Salzmann, 2017). The Eocene-Oligocene Transition (EOT; 34.44-33.65 Ma; Katz et al., 2008; Hutchinson et al., 2021) is one of the most important climate transitions of the Cenozoic and it is characterised by a shift from largely ice-free greenhouse conditions to an icehouse climate, involving the development of Antarctic cryosphere and global cooling (Liu et al., 2009; Pearson et al., 2009; Pagani et al., 2011; Hutchinson et al., 2021).

Tectonic opening of the southern gateways (Kennett, 1977), as well as a large and sharp drop in global atmospheric CO₂ (DeConto and Pollard, 2003; Huber et al., 2004; Zachos et al., 2008; Goldner et al., 2014; Ladant et al., 2014) have been proposed as possible drivers for this climate transition. The opening of the Australian-Antarctic Seaway (Tasmanian Gateway) and Drake Passage led to the strengthening of the Antarctic Circumpolar Current (ACC), which thermally isolated Antarctica (Kennett, 1977). However, marine geology, micropalaeontology and model simulation showed a potential time lag between the onset of the ACC and palaeogeographic changes, hence challenging a southern hemisphere tectonic driven global climate change at the EOT (Huber et al., 2004; Stickley et al., 2004; Goldner et al., 2014).

Although southern gateway opening and deepening have failed to fully explain Antarctic cooling at the EOT, the oceanographic changes following gateway opening and deepening have been reported to climatically impact Southern Ocean surface waters regionally (Stickley et al., 2004; Sijp et al., 2011; Houben et al., 2019; López-Quirós et al., 2021; Thompson et al., 2021). However, the extent and effect of the opening and deepening of the Tasmanian Gateway and its associated oceanographic changes on the coeval terrestrial climate and vegetation are not readily known. The lack of continuous and well-dated EOT terrestrial records place considerable limitations on detailed temporal and spatial reconstruction of vegetation and climate. These challenges are further compounded by the fact that the few late Eocene and early Oligocene terrestrial palynoflora records indicate a rather heterogeneous vegetation response at the EOT (Pound and Salzmann, 2017). For example, in southeastern Australia, the late Eocene to early Oligocene vegetation records indicate a shift from a warm-temperate to a cool-temperate rainforest (Korasidis et al., 2019; Lauretano et al., 2021) whereas in New Zealand, a warm humid rainforest persisted (Pocknall, 1989; Homes et al., 2015; Prebble et al., 2021). East Antarctica (Prydz Bay) saw the collapse of tall woody vegetation and their replacement by impoverished, taiga-like vegetation with dwarfed trees before the EOT during the late Eocene (Macphail and Truswell, 2004; Truswell and Macphail, 2009; Tibbett et al., 2021), whereas across the Drake Passage region major vegetation change did not take place until the early Oligocene, where there is a distinct expansion of gymnosperms and cryptogams indicating glacial expansion (Thompson et al., 2021).

To further our understanding of the timing and potential drivers of southern high-latitude terrestrial environment change at the EOT, this study presents a new sporomorph record recovered from ODP Site 1172 (Fig.1) on the East Tasman Plateau (ETP) spanning the late Eocene (37.97 Ma) to earliest Oligocene (33.06 Ma). The proximity of our study site to the Tasmanian Gateway places it in an excellent geographical position to identify potential climate or tectonic impacts on terrestrial vegetation of the Australo-Antarctica region. To further investigate potential links between the terrestrial and marine realm we also



compare our pollen-based quantitative climate estimates with newly published TEX₈₆-based sea-surface temperature (SST) and mean annual air temperature (MAAT_{soil}) reconstruction from the same site (Bijl et al., 2021). Our study reveals a significant
 70 terrestrial cooling ~3 Ma prior to the EOT, and a warming in the earliest Oligocene which is most likely controlled by transient rebound of atmospheric *p*CO₂ and sustained deepening of the Tasmanian Gateway.

2. Materials and methods

2.1. Tectonic evolution and depositional setting

Continental breakup and seafloor spreading between Australia and the continental blocks of Lord Howe Rise, Campbell
 75 Plateau, and New Zealand (LCNZ) started in the late Cretaceous (~75 Ma; Cande and Stock, 2004). Northward movement of Australia was propagated by rifting leading to the formation of the Tasman Sea and separation of northeastern Australia in the Paleocene (~60 Ma; Gaina et al., 1999). The series of tectonic events paved way for major ocean currents to flow along the coast of eastern Australia and Tasmania, the ETP, and South Tasman Rise (STR; Exon et al., 2004a). However, the Tasman promontory remained and separated the Australo-Antarctic gulf (AAG) from the Pacific Ocean until the late Eocene (~35.5
 80 Ma; Stickley et al., 2004). Our study site (ODP Site 1172 on the ETP; Fig.1) is located on one of the four continental blocks sampled during ODP Leg 189 (Exon et al., 2004b). Prior to the Tasman Sea break-up in the late Cretaceous (95 Ma), the ETP was part of Tasmania and the STR (Royer and Rollet, 1997; Exon et al., 2004b), subsiding slowly until the late Eocene. The ETP forms an oval platform presently located ~170 km southeast of Tasmania (43°57.6'S, 149° 55.7' E; Fig. 1a; Shipboard Scientific Party, 2001) at water depths of ~2620 m (Exon et al., 2004a) and enclosed by an 1800 m high seamount (Royer and
 85 Rollet, 1997). Bathymetric studies indicate that the ETP is connected to the east coast of Tasmania by the East Tasman Saddle (Royer and Rollet, 1997) which gives no indication of a deep basin in between (Hill and Exon, 2004). Dredging exercise confirms the continental origin of the of the plateau (Exon et al., 1997). However, the age of the guyot/seamount (dated as 36 Ma; Lanyon et al., 1993) disqualifies the ETP itself as the potential source of the terrestrial organic matter (Bijl et al., 2021). In addition, common Permo-Triassic reworked elements in our late Eocene–early Oligocene sporomorph assemblage likely
 90 indicate an eastern Tasmania sporomorph source, in line with the Permian–Triassic upper Parmeener Group containing terrestrial deposits and presently making up surface lithology across east Tasmania. Previous Paleocene-Eocene sporomorph assemblage presented from the ETP (ODP Site 1172) further supports an eastern Tasmania terrestrial palynomorph source (Contreras et al., 2014).

Lithologically, the marine sedimentary record is divided into three units: (i) shallow-marine, organic-rich middle Eocene to
 95 lower upper Eocene clay; (ii) a highly condensed middle upper Eocene to lowermost Oligocene glauconite-rich, shallow-marine silty-sandstone; (iii) lower Oligocene siliceous-rich, carbonate ooze (Stickley et al., 2004; Exon et al., 2001). Both Holes A and D of ODP Site 1172 on the East Tasman Plateau yielded EOT records and have been analysed for their pollen and spore content. The age model relies on magnetostratigraphy (which has particularly clear signal in the late Eocene; Stickley



et al., 2004; Fuller and Touchard, 2004) and biostratigraphy (dinoflagellate cyst, nannoplankton, and diatoms; Stickley et al., 2004; Bijl et al., 2013) as presented in Houben et al. (2019) and Bijl et al. (2021).

2.2. Study material

A total of 66 samples from the late Eocene to earliest Oligocene of ODP Site 1172 (37.97-33.06 Ma) were analysed for terrestrial palynomorphs to reconstruct palaeovegetation and palaeoclimate. These samples were prepared at the Laboratory of Palaeobotany and Palynology, Utrecht University following standard palynological processing techniques (Bijl et al., 2013). Sample processing involved treatment with 30% HCl and 38% HF and sieving residue through a 15 µm nylon mesh (Pross, 2001). The residues were mounted onto a microscope slide with glycerine gel as the mounting medium.

The Leica DM 500 and DM 2000 LED microscopes were used to analyse two slides for each sample at x400 or x1000 magnification. Where possible, 300 fossil spores and pollen grains (excluding reworked sporomorphs) were analysed for each sample, followed by further scanning of the entire microscope slide to record rare taxa. Aside from nine samples with counts below 50 grains, overall pollen preservation and counts were generally good. Reworked sporomorphs were identified based on the thermal maturation (colour) of their outer coat (exine) and occurrence outside their known stratigraphic range. Non-pollen palynomorphs were recorded but not added to the total pollen counts. Sporomorph percentages are calculated based on the total sum of pollen and spores, excluding reworked grains, and plotted using Tilia version 2.6.1 (Fig. 2; Grimm, 1990). Using Edward's and Cavalli-Sforza Chord Distance, we applied a stratigraphically constrained incremental sum-of-squares cluster analysis (CONISS, Grimm, 1987) to determine pollen assemblage zones (PZ; Fig.2). Sporomorph identification and botanical affinities (used for nearest living relative identification of fossil spores and pollen) were established using Macphail and Cantrill (2006); Macphail (2007); Truswell and Macphail (2009); Daly et al. (2011); Kumaran et al. (2011); Raine et al. (2011); Bowman et al. (2014); Stevens (2017); and Macphail and Hill (2018).

2.3. Bioclimatic analysis

The nearest-living relative (NLR) approach was used to estimate and reconstruct mean annual temperature (MAT), warm mean month temperature (WMMT), cold mean month temperature (CMMT) and mean annual precipitation (MAP). The bioclimatic analysis used in this study involved all pollen and spore taxa that could be related to an NLR and are listed in Table 1. The NLR is a uniformitarian approach based on the assumption that climate tolerance of extant taxa can be extended into the past. However, factors such as misidentification of fossil taxa and/or their NLRs, unresolved differences in climate tolerance between fossil taxa and their NLRs, climate tolerance of NLRs being potentially incomplete, and potential weakening in connection between fossil taxa and NLRs through deep time may pose some concerns and need to be considered prior to the application of the NLR-based climate reconstructions (Mosbrugger and Utescher, 1997; Mosbrugger, 1999; Pross, 2000; Utescher et al., 2000, 2014). Generally, these uncertainties and issues with the NLR approach increase when analysing plant remains or samples from deep-time geological records (Poole et al., 2005). To test the validity of our NLR-based climate



estimates, we compare them to previous published independent botanical or geochemical proxies in the southern high-latitude spanning the late Eocene to early Oligocene (e.g., Colwyn and Hren, 2019; Houben et al., 2019; Korasidis et al., 2019; Bijl et al., 2021; Lauretano et al., 2021; Tibbett et al., 2021). Overall, these are generally in agreement and provide a certain level of confidence in the utility of the NLR-based climate estimates.

The NLR analysis in this study is combined with the probability density function (PDF). The PDF works by statistically constraining the most likely climate co-occurrence envelope for an assemblage (Harbert and Nixon, 2015; Hollis et al., 2019). Bioclimatic analysis was performed using the dismo package in R (Hijmans et al., 2017) to cross-plot the modern distribution of the NLR from the Global Biodiversity Information Facility (GBIF; GBIF, 2021) with gridding from WorldCLIM climate surface (Fick and Hijmans, 2017). The datasets are then filtered to remove multiple entries per climate grid cell, plants whose botanical affinity are vague or doubtful, redundant, and occurrences termed exotic (e.g., garden plants). Filtering was performed to avoid bias in the probability function which may likely lead to results leaning towards a particular location (Reichgelt et al., 2018). To test the robustness of the dataset, bootstrapping was applied which was followed by calculating the likelihood of a taxon that occurs at a specific climate variable using the mean and standard deviation of modern range of each taxon (Kühl et al., 2002; Willard et al., 2019). For a more detailed explanation of this method see Willard et al. (2019) and Klages et al. (2020).

2.4. Quantitative and statistical analyses

Diversity indices (rarefaction, Shannon diversity index, equitability) were generated using PAST statistical software (Hammer et al., 2001) with sample counts of ≥ 75 individuals. Rarefaction is an interpolation technique used to compare taxonomic diversity in samples of different sizes (Birks and Line, 1992; Birks et al., 2016). Shannon diversity index (H) is a measure of diversity which considers number of individuals as well as number of taxa, and evenness of the species present (Shannon, 1948). It ranges from 0 involving vegetation communities with a single taxon to higher values where taxa are evenly distributed (Legendre and Legendre, 2012). Equitability (J) on the other hand, measures the level of abundance and how they are distributed in an assemblage. Low J values indicate the dominance of a few species in the population (Hayek and Buzas, 2010). Pollen Zones (PZ) have been defined following stratigraphically constrained analysis (CONISS; Grimm, 1987) in Tilia (Vers. 2.6.1) using total sum of squares with chord distance square root transformation (Cavalli-Sforza and Edwards, 1967). In addition, we used Detrended Correspondence Analysis (DCA; (Hill and Gauch, 1980) sample scores to measure sample-to-sample variance. DCA sample scores were generated using the Vegan package (Oksanen et al., 2019) of R statistical software (R Core Team, 2019).



3. Results

3.1. Palynological results from ODP Site 1172

The late Eocene-early Oligocene samples from the East Tasman Plateau (ODP Site 1172) yielded moderate to well-preserved sporomorphs. ~~Of the 66 samples analysed, nine do not contain sufficient pollen counts and were not used in our analyses.~~ Eighty-one (81) individual sporomorph taxa were recorded across the studied section. The sporomorph record is dominated by *Nothofagidites* spp., making between 38% to 48% of all non-reworked sporomorphs across the studied interval (Fig.2). *Podocarpidites* spp., *Myricipites harrisii*, *Cyathidites* spp., *Phyllocladidites mawsonii* and *Araucariacites australis* form significant components of the palynoflora and occur with varying frequency (Fig.2).

Based on results from rarefaction, the average diversity for the entire studied section was 21.0 ± 2.0 taxa/sample at 75 individuals. The sporomorph record, based on CONISS is grouped into four pollen zones (PZ; Fig. 2); PZ 1 (early late Eocene; 37.97–37.52 Ma), PZ 2 (late Eocene-latest Eocene; 37.30–35.60 Ma), PZ 3 (latest Eocene-earliest Oligocene 35.50–33.36 Ma), and PZ 4 (earliest Oligocene; 33.25–33.06 Ma). All the four zones consist of characteristic palynoflora assemblages that are described below. Taxa names in bracket refer to the NLR.

3.1.1. Pollen Zone 1 (37.97–37.52 Ma; 7 samples)

Pollen zone 1 is dominated by *Nothofagidites* spp. (*Nothofagus*), which accounts for ~48% of all non-reworked palynomorphs. Taxa belonging to the *Brassospora* (~28%) subgenus of *Nothofagus* make up the most abundant component, followed by *Fuscospora* (19%) and *Lophozonia* (1%), respectively. Other angiosperms (non-*Nothofagus*) on average account for 24% of all sporomorphs. These are represented, mainly in order of decreasing occurrence, by *Myricipites harrisii* (*Gymnostoma*), *Proteacidites pseudomoides* (*Carnarvonia*), *Proteacidites* spp., *Spinizonocolpites* spp. (*Arecaceae*), *Malvacearumpollis mannanensis* (*Malvaceae*), and *Malvacipollis* spp. (*Euphorbiaceae*). The abundance of gymnosperms is generally low throughout PZ 1 and accounts for about 16% of all non-reworked palynomorphs. These are also represented mainly, in order of decreasing occurrence by *Podocarpidites* spp. (*Podocarpaceae*), *Phyllocladidites mawsonii* (*Lagarostrobos*), *Dacrydiumites praecupressinoides* (*Dacrydium*) and *Araucariacites australis* (*Araucariaceae*). Ferns and mosses account for about 12% of the total sporomorphs and are represented by *Cyathidites* spp. (*Cyatheaceae*), *Dictyophyllidites* sp. (*Gleicheniaceae*), *Gleicheniidites* sp. (*Gleicheniaceae*), *Laevigatosporites* spp. (*Blechnaceae*) and *Stereisporites* sp. (*Sphagnum*).

Quantitatively, sporomorph diversity for this zone based on rarefied values is 21.61 ± 1.32 species per sample at 75 individuals. With respect to the diversity indices, the yields for Shannon diversity (H) are between 2.52 and 2.88, averaging at 2.75 ± 1.12 . Equitability (J) scores are set between 0.85 and 0.92, with an average of 0.89 ± 0.02 (Fig. 3; Table 2).

3.1.2. Pollen Zone 2 (37.30–35.60 Ma; 27 samples)

PZ 2 sees the decline of *Nothofagidites* spp., to about 42%. The *Brassospora*-type remains the dominant *Nothofagus* subgenus, but with a substantial decline in abundance from about 28% in PZ 1 to 22% in PZ 2. The *Fuscospora* and *Lophozonia* subgenus



190 however, accounted for 19% and 1%, respectively (Fig. 2). Other angiosperms (non-*Nothofagus*) in comparison to PZ 1 see a decline from about 24% to 17%. In order of decreasing abundance, the most significant taxa among non-*Nothofagus* angiosperms are *Myricipites harrisii* (*Gymnostoma*), *Proteacidites* spp. (*Proteaceae*), *Malvacearumpollis mannanensis* (*Malvaceae*) and *Malvacipollis* spp. (*Euphorbiaceae*). A sharp decline in *Proteacidites pseudomoides* (*Carnarvonina*) is coupled with the disappearance of *Spinizonocolpites* spp. (*Arecaceae*). Gymnosperms, on the other hand, almost doubled in relative abundance from about 16% in PZ 1 to over 29% in PZ 2. Gymnosperm taxa in order of decreasing abundance dominated by *Podocarpus* spp. *Araucariacites australis* (*Araucariaceae*), *Dacrydiumites praecupressinoides* (*Dacrydium*) and *Phyllocladidites mawsonii* (*Lagarostrobos*). *Microcachryidites antarcticus* (*Microcachrys*) is a taxon which first appears in this zone and forms an important component of the gymnospermous assemblage. In addition to the above, cryptogams decline slightly in this zone accounting for roughly 10% of the total sporomorphs. The main members of this group are *Cyathidites* spp. (*Cyatheaceae*), *Gleicheniidites* (*Gleicheniaceae*) and *Laevigatosporites* spp. (*Blechnaceae*).
 200 This zone has lower diversity than PZ 1. Based on rarefied values, the average diversity for PZ 2 is 20.52 ± 2.31 species per sample at 75 individuals. The results for Shannon diversity index (H) are between 2.40–2.99, averaging at 2.66 ± 0.16 . Equitability is set between 0.82 and 0.93, with an average of 0.88 ± 0.03 (Fig. 3; Table 2).

3.1.3. Pollen Zone 3 (35.50–33.36 Ma; 20 samples)

205 Zone 3 shows a further decline in *Nothofagidites* spp. to approximately 38%. However, the *Brassospora*-type *Nothofagus* sees a slight increase in abundance while the *Fuscospora*-type *Nothofagus* declines sharply from the peak 19% observed in PZ 2 to 12%. The *Lophozonia*-type remains stable (~1%). Other angiosperms (non-*Nothofagus*) see a slight decline and account for ~14% of all non-reworked sporomorphs. These are represented mainly by *Myricipites harrisii* (*Gymnostoma*) and *Proteacidites* spp. (*Proteaceae*), while *Malvacipollis* spp. (*Euphorbiaceae*), and *Malvacearumpollis mannanensis* (*Malvaceae*).
 210 Another important observation in this interval is the re-appearance of *Spinizonocolpites* spp. (*Arecaceae*) and *Proteacidites pseudomoides* (*Carnarvonina*). However, in contrast to PZ 1, *Spinizonocolpites* spp. are not consistently present. Gymnosperms increase slightly in this zone, accounting for about 31%. The gymnosperms remain dominant with *Podocarpidites* spp. (*Podocarpaceae*). However, other important taxa such as *Araucariacites australis* (*Araucariaceae*), *Phyllocladidites mawsonii* (*Lagarostrobos*) and *Microcachryidites* (*Microcachrys*), decline. *Dacrydiumites praecupressinoides* (*Dacrydium*) reaches its
 215 peak abundance in this zone. Cryptogams significantly increase in abundance and in order of abundance are represented by *Cyathidites* spp. (*Cyatheaceae*), *Laevigatosporites* spp. (*Blechnaceae*), *Osmundacidites* (*Osmundaceae*), *Polypodiisporites radiatus* (*Polypodiaceae*), and *Clavifera* spp. (*Gleicheniaceae*).
 Based on rarefied values, the average diversity for this PZ is 21.37 ± 1.81 species per sample. The results for Shannon diversity (H) are between 2.44–2.86, averaging at 2.66 ± 0.1 . Equitability (J) is set between 0.82–0.91, averaging at 0.87 ± 0.02 (Fig. 3; Table 2).
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3.1.4. Pollen Zone 4 (33.25–33.06 Ma; 3 samples)

The percentage abundance of *Nothofagidites* spp. (*Nothofagus*) including *Brassospora* (~23%), *Fuscospora* (12%) and *Lophozonia*-types remain unchanged, whereas other angiosperms percentages increase substantially from 14% in PZ 3 to ~20%. In order of decreasing abundance, these are represented by *Myricipites harrisii* (*Gymnostoma*) and *Proteacidites pseudomoides* (*Carnarvon*). PZ 4 also sees the emergence of new angiosperms such as *Sapotaceoidapollenites* cf. *latizonatus* (*Sapotaceae*) and *Parsonsidites psilatus* (*Parsonsia*). Gymnosperms, however, see a sharp decline in this interval accounting for about 10% of total sporomorph taxa with *Podocarpidites* spp. (*Podocarpaceae*) and *Dacrydium praecupressinoides* (*Dacrydium*) being the main components. *Microcachryidites antarcticus* (*Microcachrys*), *Araucariacites australis* (*Araucariaceae*), *Phyllocladidites mawsonii* (*Lagarostrobos*) showed significant decline whereas cryptogams increase to ~20%. The cryptogams are represented, in order of decreasing abundances by *Cyathidites* spp. (*Cyatheaceae*), *Laevigatosporites* spp. (*Blechnaceae*), *Dictyophyllidites* sp. (*Gleicheniaceae*) and *Cibotiidites tuberculiformis* (*Schizaeaceae*). Average diversity (21.16 ± 1.37 species per sample) is slightly lower than in PZ 3. The results for Shannon diversity (H) are between 2.42–2.72, averaging at 2.54 ± 0.15 . Equitability (J) is set between 0.80–0.87, averaging at 0.83 ± 0.03 (Fig. 3.; Table 2).

4. Discussion

4.1. Vegetation composition and altitudinal zonation

Throughout the studied section, abundant *Nothofagidites* spp. with common *Podocarpidites* spp. *Myricipites harrisii* and *Phyllocladidites mawsonii* indicate the presence of *Nothofagus*-dominated temperate rainforest (Truswell and Macphail, 2009; Bowman et al., 2014) that likely grew across lowland and mid-altitude elevations in eastern Tasmania. The occurrence of *Araucariacites australis*, *Microcachryidites antarcticus*, and *Proteacidites parvus* may also suggest a component of the sporomorph assemblage reflect higher altitudes with more open forest conditions (Macphail, 1999; Kershaw and Wagstaff, 2001). In addition, pollen taxa belonging to *Arecaceae*, *Gymnostoma*, and *Carnarvon*, indicate the existence of a paratropical vegetation community that grew in sheltered lowland and coastal areas (Huurdeeman et al., 2021). The paratropical rainforest likely occupied lowlands and coastal areas while temperate rainforest likely grew at higher elevation, similar to vegetation communities that prevailed on Wilkes Land and Tasmania during the early to mid-Eocene (Pross et al., 2012; Contreras et al., 2013, 2014). The existence of different vegetation communities, whose NLRs today grow under different temperatures and elevations, suggest that vegetation across eastern Tasmania were subject to climatic gradients related to differences in elevation and/or distance to the coastline. This is supported by reports of a topographic divide between sites facing the cool Tasman current (Gippsland basin, eastern Tasmania) and the westerly located south Australian basins (Holdgate et al., 2017) that may have served as the location for higher altitude temperate forest taxa. The following sub-sections further describe each of these vegetation communities in detail.



4.1.1. Lowland to mid-altitude *Nothofagus-Podocarpus* rainforest

Abundant *Nothofagidites* spp. with common *Podocarpidites* spp., *Myricipites harrisii*, and *Phyllocladidites mawsonii* and Cyatheaceae give an indication of a lowland to mid-altitude *Nothofagus-Podocarpus* dominated rainforest thriving under high precipitation regimes (MAP > 1300 mm/yr) in Tasmania during the late Eocene to the earliest Oligocene. The main canopy is primarily made up *Nothofagidites* spp. (*Nothofagus*/southern beech) and Podocarps (*Dacrydiumites*, *Podocarpidites*, *Dacrycarpites*) with rare Cupressaceae trees. Southern beech forests can either occur as pure stands or a mixed forest, making the definition and recognition of regional or local forest types from fossil pollen and spore challenging. Today, pure beech stands in New Zealand are mostly montane to subalpine, and lowland mixed beech forests are associated with diverse broadleaf angiosperms and canopy-emergent gymnosperms (Ogden et al., 1996). Following Dettmann et al. (1990), we categorise *Nothofagidites* pollen taxa into the *Brassospora*, *Fuscospora* and *Lophozonia* subgenera. Extant *Fuscospora* and *Lophozonia*-types thrive in cool temperate conditions in Tasmania, southeastern Australia, New Zealand, and southern South America (Hill, 1994, 2017; Veblen et al., 1996; Read et al., 2005) while the *Brassospora*-type are today restricted to warm temperate-subtropical conditions in New Guinea and New Caledonia (Hill and Dettmann, 1996; Veblen et al., 1996). These *Brassospora*-type *Nothofagus* grow today at lower to mid altitudes that receive high and consistent rainfall but, also in montane and subalpine areas (typically above 500m), pointing to their wide ecological and climate tolerance (MAT: 10.6 to 23.5 °C; Read et al., 2005).

Myricipites harrisii (Casuarinaceae) has two potential NLR, *Casuarina/Allocasuarina* and *Gymnostoma*. *Casuarina/Allocasuarina* have xeromorphic features indicating adaptation to arid climate with frequent fires (Hill, 2017; Lee et al., 2016). We selected the rainforest clade *Gymnostoma* as the most likely NLR of our fossil taxon *Myricipites harrisii* based on the subtropical affinities of the associated palynoflora. This is also supported by Paleogene vegetation reconstruction of southeastern Australia based on macrofossil remains which indicate rainforest communities (Christophel et al., 1987; Macphail et al., 1994; Hill, 2017) with *Gymnostoma* being common from the Paleocene to Oligocene and later being replaced by *Casuarina/Allocasuarina* (sclerophyll taxa) in the Miocene (Evi et al., 1995; Boland et al., 2006; Holdgate et al., 2015).

Dacrydium cupressinum is suggested as the most likely NLR of *Dacrydiumites praecupressinoides* (Rimu; Raine et al., 2011). Today, *Dacrydium cupressinum* occur as a minor component in the Kauri Forest of Northland, New Zealand and occur as emergent taxa commonly associated with *Agathis australis* (Araucariaceae) and *Podocarpus totara* (Farjon, 2010). The NLR of *Phyllocladidites mawsonii*, *Lagarostrobos franklinii* (Tasmanian Huon Pine; Raine et al., 2011) is very abundant at Site 1172. *Lagarostrobos* are evergreen cool temperate riparian trees that grow in Tasmania close to riverbanks (Farjon, 2010; Hill, 1994, 2017). Apart from forming groves that mark stream courses in low altitudes (Hill and Macphail, 1983; Farjon, 2010), they may also be found away from water courses on wet hillsides in temperate forest (Farjon, 2010; Bowman et al., 2014).

The two possible NLR relatives for *Proteacidites pseudomoides* are *Carnarvonina* and *Lomatia*. *Carnarvonina* thrives in warm temperate to paratropical areas such as wet northeastern Australia (Mabberley, 1997; Cooper and Cooper, 2004) and grows into large trees (Hyland, 1995). *Lomatia* grows as shrubs and small trees in remnant gallery warm temperate rainforests for



285 example, along creek lines on sandstones in Northern Sydney (Bowman et al., 2014; Myerscough et al., 2007). *Carnarvon*
 is selected as the likely NLR relative because of their significant increase in intervals where warmth-loving taxa such as
 Areaceae, *Brassospora*-type *Nothofagus*, Gleicheniaceae, and Cyatheaceae thrive.

4.1.2. High altitude temperate rainforest and shrubland

290 Components of the palynoflora that reflect higher altitude and more open vegetation on soils with low fertility are
Araucariacites australis, *Proteacidites parvus*, *Microcachrydites antarcticus* (Kershaw and Wagstaff, 2001; Macphail et al.,
 1999). Today, Araucariaceae trees grow in cool temperate forests in Chile and Argentina and extend to the tree line (Veblen
 et al., 1996; Sanguinetti and Kitzberger, 2008). In the Andes, trees belonging to Araucariaceae are found at altitudes of 600-
 800 m a.s.l and they receive high amounts of annual rainfall (2000-3000 mm/yr) as well as experiencing hot and dry spells in
 summer (Farjon, 2010). Araucariaceae build pure stands at higher altitude or mixed Valdivian rainforest at lower altitudes
 295 (Farjon, 2010). Increase in araucarian sporomorph taxa between 37.30–35.60 Ma in Tasmania give an indication of a dense,
 emergent cover of Araucariaceae thriving in relatively dry environments (Kershaw and Wagstaff, 2001). *Microcachrys* (Raine
 et al., 2011), the nearest living relative of *Microcachrydites antarcticus* is a creeping shrub that grows in alpine/subalpine
 areas and are today restricted to western Tasmania under boreal to cool temperate conditions (Truswell and Macphail, 2009;
 Biffin et al., 2012; Carpenter et al., 2011). Therefore, increase in this Tasmanian endemic alpine shrub (*Microcachrys*) from
 300 37.30-35.60 Ma together with *Bellendena* (low-growing protea shrub; NLR of *Proteacidites parvus*), and Araucariaceae
 (emergent canopy) suggest that the vegetation thriving in the higher altitudes in Tasmania preferred cool temperate conditions.

4.2. Subtropical vegetation and early-late Eocene cooling from 37.97-35.60 Ma

Throughout PZ 1 (37.97–37.52 Ma), abundant *Nothofagus* (especially *N. brassii*-type) with secondary Podocarpaceae,
Gymnostoma, along with minor Areaceae, *Carnarvon*, and cryptogams suggest the presence of a temperate *Nothofagus*-
 305 dominated rainforest with subtropical elements. Sporomorph-based climate estimates indicate these forests grew under MAT
 between 14.2–15.1 °C and a MAP of 1467–1681 mm/yr (Fig. 4). The vegetation-based summer temperature reconstructions
 of ca. 18.5 °C closely corroborate the brGDGT-biomarker reconstructions from the same site (Bijl et al., 2021), supporting the
 notion of a potential seasonal bias of this palaeothermometer (Contreras et al., 2014; Naafs et al., 2017). The warmth-loving
 taxa formed the main lowland forest components occupying sheltered areas and lowland subtropical coastal zones (Dowe,
 2010; Carpenter et al., 2012; Tripathi and Srivastava, 2012; Verma et al., 2020) and swamps (Kershaw, 1988). Sporomorph-
 310 based temperature estimates yield cold month mean temperature (CMMT) well above freezing (11.2–12.5 °C; Fig. 4). The
 decline and to a large extent, the absence of cool-temperate taxa coupled with persistent warm temperate (12–17 °C; Emanuel
 et al., 1985) to subtropical taxa (17–24 °C; Emanuel et al., 1985) taxa, further points to the expansion of warm temperate –
 paratropical rainforest up into the mid-altitudes and uplands.



315 The *Nothofagus*-dominated rainforest continued into PZ 2 (37.30–35.60 Ma). However, at ~37.30 Ma, a distinct environmental
 change occurred, leading to a drop and in some instances, the demise of warm-temperate and subtropical taxa (*Nothofagus*
 subgenus *Brassospora*, *Carnarvonina*, *Arecaceae*; Fig. 2). This vegetation change continued throughout PZ 2 with a
 concomitant rise in relative abundance of *Lagarostrobos*, *Microcachrys*, and decline in diversity (Table 2) ~3 Ma prior to the
 EOT. The increased occurrence of microthermal taxa points to a cool-temperate (southern beech) dominated rainforest with
 320 secondary Podocarpaceae expanding into lowland regions previously occupied by mesothermal taxa. The late Eocene cool
 temperate *Nothofagus*–Podocarpaceae dominated rainforest have been suggested to resemble modern Valdivian rainforest of
 Chile (Veblen, 1982; Cantrill and Poole, 2012; Bowman et al., 2014), cool temperate *Nothofagus* dominated rainforest with
 riparian *Lagarostrobos* restricted to river gullies in Victoria, Australia (Read and Hill, 1985) or on fertile soils in lowland
 Tasmania (Read and Hill, 1985; Macphail, 2007; Francis et al., 2008).

325 This interpretation is reflected in our sporomorph-based MAT estimates indicated by a 2–3 °C decline in MAT (Fig.4). Our
 findings also corroborate previous late Eocene studies throughout Australia indicating an increase in *Nothofagus* subgenus
Fuscospora with substantial decline in *Brassospora*-type *Nothofagus*, demise of most Proteaceae, *Arecaceae*, and most
 Australian angiosperm (Falkenberry et al., 1978; Kershaw, 1988; Christophel and Greenwood, 1989; Truswell, 1993; Martin, 1994, 2006;
 Macphail et al., 1994; Partridge and Dettmann, 2003; Korasidis et al., 2019). In line with the vegetation change, biomarker-
 330 based reconstruction from Site 1172 also indicates declining SSTs by ca. 2–3 °C starting around 37.5 Ma (Fig.4). However,
 the cooling indicated by both independent proxies is not reflected by the lipid biomarker-based terrestrial MAT estimates and
 the reason for this disparate trend remains unknown.

The transition from a warm-temperate rainforest with paratropical elements to cool temperate forests in the Tasmanian
 Gateway region also matches an early-late Eocene cooling (37.3 Ma) in the Southern Ocean (Kerguelen Plateau) ~3 Ma prior
 335 to the EOT (Villa et al., 2008, 2014; Scher et al., 2014). The 2–3 °C sporomorph-based MAT (100–200 kyr) cooling around
 37.3 Ma coincides with a regional transient (~140 kyr) cooling event at ODP Site 738 (Kerguelen Plateau) known as the
 Priabonian Oxygen Maximum (PrOM; Scher et al., 2014). The PrOM event, placed within magnetochron C17n.1n of the late
 Eocene, points to the temporary growth of ice sheets on East Antarctica based on positive excursion in benthic $\delta^{18}\text{O}$ (Scher et
 al., 2014). However, on the Kerguelen Plateau, differences in neodymium isotopic composition (ϵ_{Nd}) between bottom waters
 340 and terrigenous sediments point to changes in sediment provenance as opposed to changes in reorganisation of ocean currents
 (Scher et al., 2014). The transient 2–3 °C sporomorph-based MAT cooling phase is followed by a period of sustained cooler
 climate from 37.2 Ma to 35.6 Ma (Fig.4). This sustained cooler climate may have led to the climate threshold of the frost-
 sensitive (subtropical) taxa being exceeded, hence their continued decline and demise. In the marine realm, endemic-Antarctic
 dinoflagellate cysts become dominant at Site 1172 (Fig. 3; Stickley et al., 2004; Houben et al., 2019). The dominant endemic-
 345 Antarctic dinocyst in addition to general sea surface circulation models (Huber et al., 2004) point to the East Tasman Plateau
 and east Tasmania being bathed by relatively cool Antarctic-derived surface waters (Houben et al., 2019) which is consistent
 with TEX₈₆-based sea surface temperature records (~3–4 °C cooling; Houben et al., 2019; Bijl et al., 2021). Regionally, this
 sustained cool-temperate terrestrial MATs matches Oligotrophic conditions associated with low nutrients, stratification of



water mass, and increase in efficiency of ocean biologic pump, which favoured cooling as a result of carbon being sequestered from surface water in the Southern Ocean (Villa et al., 2008, 2014).

Close to the top of PZ 2 (35.7 Ma; Fig.4), branched GDGT-based MATs and SSTs show strong and rapid cooling, which is not mirrored by the pollen-based climate estimates. However, strong fluctuations of gymnosperms and an increase in cryptogams (Fig.2 and Fig.3) and diversity towards the top of PZ 2 indicate increasing environmental disturbance that might be linked to their recorded change in lipid biomarkers. The divergence between the different proxy signals could be related to their different origins and transport mechanisms. Whereas the lipid biomarkers are strongly controlled by the depositional settings, including river run-off, tectonic and geographic evolution (Bijl et al., 2021), the terrestrial palynological signal mainly consist of well dispersed pollen and spores.

4.3. Warm and cold temperate terrestrial climate fluctuation from 35.50-34.59 Ma

PZ 3 (35.50–33.36 Ma) is characterised by a major shift in sporomorph assemblages represented by increase in *Podocarpus*, decline in *Lagarostrobos*, *Microcachrys*, Araucariaceae and *Fusca*-type *Nothofagus*, with the re-emergence of subtropical and warm-temperate taxa. The peak in tree ferns, especially Cyatheaceae, indicate a period of disturbance (Majda et al., 2001) within this interval of shift in vegetation. Sporomorph-based temperature reconstructions indicate several fluctuations between warm and cool climate phase with MAT between 10.6–15.3 °C (Fig.4). In the regional Australo-Antarctic area, a similar phase of warming and cooling is observed in the late Eocene (35.8–34.7 Ma) climate records of Prydz Bay (Passchier et al., 2017; Tibbett et al., 2021) and Southern Australia (Benbow et al., 1995). Again, pollen-based WMMTs at Site 1172 closely match lipid biomarker derived MATs (Fig.4). Our sporomorph-based warm and cool climate fluctuation phase between 35.50 to 34.59 Ma in comparison, is recorded as a recovery phase in lipid biomarker-based MAT reconstruction. The fluctuations of pollen-based temperature estimates may be at least partly caused by the proxy method that relies on presence-absence data. However, a more detailed proxy comparison is hampered by the relative low resolution of the lipid biomarker in PZ 3.

Expansion and restriction of cool-temperate and warm-temperate forests which indicate cooling and warming phases are consistent with previous late Eocene geochemical, sedimentological, and palynological studies reporting an increase in sea surface temperature (TEX₈₆-based SST; Houben et al., 2019; Bijl et al., 2021), widespread deposition of glauconite (Stickley et al., 2004), and increase in cosmopolitan and protoperidinioid dinocyst (Fig.3; Stickley et al., 2004; Houben et al., 2019; Bijl et al., 2021). Though the glauconitic unit is interpreted to mark deepening and current inception due to widening of the Tasmanian Gateway (Stickley et al., 2004), a more recent counterargument links the deposition of the greensand to atmospheric-forced invigorated circulation in the Southern Ocean which helped to prepare Antarctica for rapid expansion of ice (Houben et al., 2019) and further circulation change ~2 Ma later (at the EOT). However, ocean model studies (Baatsen et al., 2016) in addressing the deposition of greensands along the south Australian margin, point to a further expansion in the eastward throughflow into the southwest Pacific Ocean. Our sporomorph-based MAT consequently showed an average 2 °C rise in temperature between 35.50–34.59 Ma coinciding with earlier reports of the initial deepening of the Tasmanian Gateway (Stickley et al., 2004). This is further supported by the common appearance of low-latitude cosmopolitan dinoflagellate cyst



taxa which rather than being supplied by the East Australian Current, are reported to have been sourced from the through-flow associated with the eastward proto-Leeuwin Current (Huber et al., 2004; Stickley et al., 2004; Houben et al., 2019). These events, coupled with ~ 2 °C recovery in SSTs (TEX₈₆-based; Houben et al., 2019; Bijl et al., 2021) between 35.7–34.59 Ma most likely point to warm surface waters associated with the Australo-Antarctic Gulf (AAG) influencing ODP Site 1172 (Houben et al., 2019), which at this time is reported to have been close to land (eastern Tasmania; Stickley et al., 2004), thereby affecting terrestrial climate and vegetation.

4.4. EOT cooling and climate rebound in earliest Oligocene from 34.30-33.06 Ma

At the onset of the EOT, our sporomorph record provides evidence for a return to a sustained cooler period on Tasmania spanning 34.30 to 33.82 Ma. This EOT cool phase coincides with the demise of *Spinizonocolpites* sp. (Arecaceae), a drop in Cyatheaceae and Gleicheniaceae, a slight increase in *Microcachrys* and *Lagarostrobos*. The palynoflora assemblage during the EOT is further characterized by a drop in overall angiosperm (non-*Nothofagus*) diversity with gymnosperms and *Nothofagus* (*Brassospora*-type) being common and co-dominating. Previous studies in southeast Australia (e.g., Macphail et al., 1994; Benbow et al., 1995; Holdgate et al., 2017; Lauretano et al., 2021) record a contemporaneous drop in angiosperm diversity and the final demise in Arecaceae (thermophilous elements) at the end of the Eocene (Pole and Macphail, 1996; Martin, 2006). Quantitatively, our sporomorph-based MAT reconstruction records a ~ 2 °C decline across the EOT (Fig.4) which coincides with ~ 2.4 °C and 5 °C cooling step in southeastern Australia (MBT'5me-based MAATsoil; Lauretano et al., 2021) and East Antarctica (Prydz Bay; MBT'5me-based MAATsoil; Tibbett et al., 2021) respectively. This cooling in our terrestrial record further matches the principal geochemical signature of EOT in the marine realm, which is $\sim +1.5$ ‰ excursion of oxygen isotope ratio of deep-sea benthic foraminifera (Zachos et al., 1996; Coxall et al., 2005; Pälike et al., 2006; De Vleeschouwer et al., 2017; Fig.5) associated with global cooling (Zanazzi et al., 2007; Pearson et al., 2009; Pagani et al., 2011; Hutchinson et al., 2021; Tibbett et al., 2021). This cooling at the EOT have been linked to global decline in atmospheric $p\text{CO}_2$ (Pearson et al., 2009; Lauretano et al., 2021)

Between $\sim 33.25 - 33.06$ Ma (PZ 4), the sporomorph-based climate estimates indicate a warming with MATs between 12.7-15.3 °C (Fig.4). In addition, the presence of warmth-loving taxa notably Sapotaceae, *Parsonsia* (Silkpod), and Polypodiaceae (subtropical epiphytes) further indicate a warming phase. The pollen flora resembles Oligocene warm-temperate *brassii* southern beech dominated forests of Karamu in the Waikato coal measures of New Zealand (Pocknall, 1985). The increase of *brassii*-type *Nothofagus* coupled with the appearance of Sapotaceae, and subtropical epiphytes suggests that, at least locally on lowlands, eastern Tasmania was warm enough to accommodate warm-temperate vegetation in the earliest Oligocene. Previous earliest Oligocene studies in Southeast Australia (Korasidis et al., 2019) show the presence of a cool temperate rainforest community. The palynoflora on east Antarctica (Askin, 2000; Askin and Raine, 2000; Prebble et al., 2006; Tibbett et al., 2021) and northeast Tasmania (Hill and Macphail, 1983) suggest an early Oligocene cold-temperate *Nothofagus* (subgenus *Lophozonia* or *Fuscospora*)-Podocarpaceae vegetation. These northern Tasmania and east Antarctica palynoflora are however reported to have most likely been made up of prostrate deciduous dwarf trees (Francis and Hill, 1996) or small



415 stature closed southern beech/podocarp refugia with a vegetation community that likely struggled to survive (Askin, 2000; Askin and Raine, 2000; Prebble et al., 2006; Francis et al., 2008; Tibbett et al., 2021). However, rather than a regional scrub (e.g., in Antarctica), the slight increase in angiosperms (other than *Nothofagus*) and cryptogams point to a local warm temperate forest growing along eastern Tasmania in the earliest Oligocene. Today, temperate forests in New Zealand and Tasmania host a diverse range of cryptogams as compared to scrub communities that do not offer other taxa to thrive under the low, closed canopies (Prebble et al., 2006).

420 Terrestrial cooling observed across the EOT followed by rapid recovery in the earliest Oligocene matches a partial return to warmer temperatures in previously reported terrestrial (Colwyn and Hren, 2019; Lauretano et al., 2021) and marine studies (Katz et al., 2008; Lear et al., 2008; Liu et al., 2009; Houben et al., 2012). The synchronicity between terrestrial and marine records suggest that, in addition to localised events (sustained Tasmanian Gateway deepening and widening; Stickley et al., 425 2004), the EOT and earliest Oligocene ETP record may also be responding to a much wider regional or global event. The most common explanation for global cooling (Zanazzi et al., 2007; Pagani et al., 2011; Hutchinson et al., 2021; Tibbett et al., 2021) across the EOT and transient warming in the earliest Oligocene have been ascribed to the decline in concentration of atmospheric carbon dioxide ($p\text{CO}_2$) and its recovery or rebound in the earliest Oligocene respectively (Pearson et al., 2009; Heures and Rickaby, 2015; Anagnostou et al., 2016; Fig. 5). Our results suggest that the warming, or at least the lack of 430 sustained cooling following the EOT in eastern Tasmania, may be related to a combination of $p\text{CO}_2$ recovery (Pearson et al., 2009) coupled with sustained Tasmanian gateway deepening and widening (Stickley et al., 2004, Houben et al., 2019) allowing the influx of more warm surface waters from AAG into the southwest Pacific thereby affecting terrestrial climate and vegetation along eastern Tasmania.

5. Conclusions

435 The late Eocene–early Oligocene vegetation reconstructed from terrestrial palynomorphs recovered from ODP Site 1172 (East Tasman Plateau) is characterised by three major climate transitions.

- 1) The early-late Eocene sporomorph record suggests a distinct 2–3 °C terrestrial cooling at 37.30 Ma coupled with a transition from a warm-temperate *Nothofagus*-Podocarpaceae dominated rainforest with paratropical elements to a cool-temperate *Nothofagus* dominated rainforest with secondary Podocarpaceae. This terrestrial cooling at 37.30 Ma 440 and sustained cool climate from 37.2–35.60 Ma coincides with long term SST decline from ~23 to 19 °C at ODP Site 1172, regional transient cooling event (PrOM) at ODP Site 738 (Kerguelen Plateau; Scher et al., 2014), and a relatively long-term regional Southern Ocean cooling due to carbon being sequestered from surface water (Villa et al., 2008, 2014).
- 2) Expansion and restriction of cool and warm temperate forests from 35.5–34.49 Ma, followed by a period of cooling across the EOT (34.30–33.82 Ma). This terrestrial climate fluctuation in this zone is consistent with latest Eocene 445 geochemical, sedimentological and palynological studies reporting an increase in SST, recovery in MBT’5me-based



MAATsoil (biomarker thermometry), widespread deposition of glauconite and common occurrence of low-latitude cosmopolitan and protoperidinioid dinocyst. These are interpreted to be linked to the initial deepening of the Tasmanian Gateway paving way for the warm water associated with the PLC to affect both terrestrial and marine climate in this region.

- 3) Post-EOT (earliest Oligocene) recovery characterised by a warm-temperate forest association from 33.55–33.06 Ma. This earliest Oligocene recovery in Tasmanian terrestrial temperatures following prior cooling across the EOT coincides with rebound of atmospheric $p\text{CO}_2$ at the earliest Oligocene glacial maximum (EOGM; Pearson et al., 2009) coupled with icesheet expansion in Antarctica (Galeotti et al., 2016), and sustained deepening of the Tasmanian Gateway (Stickley et al., 2004).

Our study shows that, against backdrop of global cooling in the late Eocene (sustained decline in $p\text{CO}_2$), a series of regional events in the marine realm, including a change in stratification of water masses, sequestration of carbon from surface water and, changes in ocean circulation due to Tasmanian Gateway accelerated deepening may have had a knock-on effect in driving terrestrial climate and vegetation change in the Tasmanian Gateway region.

6. Author contributions

MA and US conceived, designed and led this study. PKB supplied the palynological samples for this study and provided the biomarker thermometry data. MA and US undertook palynological analyses. MA interpreted palynological data and performed sporomorph-based bioclimatic analyses. MA prepared the manuscript with contributions from all co-authors.

7. Competing interests

The authors declare that they have no conflict of interest.

8. Acknowledgements

Samples for this study were supplied by the Ocean Drilling Program (ODP) sponsored by the US National Science Foundation under the management of the Joint Oceanographic Institutions (JOI). MA acknowledges funding from Northumbria University Research Development Fund (RDF). NT received funding from the Natural Environment Research Council (NERC)-funded Doctoral Training Partnership ONE Planet [NE/S007512/1]. PKB acknowledges funding from the European Research Council for starting grant #802835, OceaNice.



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Figure captions

Figure 1: (A) Location of East Tasman Plateau (ODP Site 1172; red star) and present-day Tasmania (Quilty, 2001). Tasmania landmass in green, and submerged ODP Site 1172 in grey with water depth of ~2620m. (B) Early Oligocene palaeogeography and palaeoceanography of the Tasmanian Gateway. ODP Site 1172 is marked by black five-pointed star. Surface currents are modified after reconstructions by Stickley et al. (2004). TC = Tasman current, PLC = proto-Leeuwin current, ACountC = Antarctic Counter Current AAG = Australo Antarctic Gulf. Solid red arrows indicate warmer ocean currents from the AAG, and solid blue arrows indicate cooler ocean currents. Arrow size also points to the relative strength of the current. Figure is modified after Hoem et al. (2021)

Figure 2: Sporomorph assemblages and relative abundance of major sporomorph taxa (Angiosperms, Gymnosperms, Cryptogams) recovered from the late Eocene - early Oligocene of ODP Site 1172. Angiosperms' relative abundance are marked by blue bars, Gymnosperms by red bars, and Cryptogams by green bars. In the Angiosperms group, *Nothofagidites* is further divided into subgenera. These are *Brassospora* (B), *Fuscospora* (F) and *Lophozonia* (L)-types. CONISS ordination constrains our late Eocene – early Oligocene sporomorph assemblages into four distinct pollen zones (PZ 1- PZ 4) or vegetation and climate phases. Age model is after Houben et al. (2019) and Bijl et al. (2021).

Figure 3: Sporomorph percentage abundance, diversity and Detrended Correspondence Analysis (DCA) results for ODP Site 1172. Percentage abundance for the major groups (Gymnosperms, Other Angiosperms, *Nothofagus* and Cryptogams) are presented for all samples with pollen counts ≥ 75 grains. The DCA results are derived from the sample scores of Axis-1 (measures sample-to sample variance) and shows four distinct compositional groupings as observed with CONISS for the late Eocene - early Oligocene Site 1172 samples. Diversity is calculated based on Sander's rarefaction analysis with samples rarefied at 75 grains/individuals. The Shannon diversity index (H) and evenness (J) are calculated for all samples with counts ≥ 75 grains. Relative percentage abundance of endemic-Antarctic and protoperidinioid dinoflagellate cyst, magnetostratigraphy and age model after Houben et al. (2019). Gippsland basin spore-pollen zonation after Holdgate et al. (2017).

Figure 4: Comparison of our sporomorph-based climate estimates, MAAT_{soil} values based on MBT'5me, TEX₈₆-based SST and sample score for DCA Axis 1 from the late Eocene – early Oligocene of ODP Site 1172. Sporomorph-based estimates are based on the use of the nearest living relative (NLR) and probability density function (PDF). The ranges of the climate estimates show the mathematical error and not the real range, which may have been a result of uncertainties associated with the use of the NLR approach. Green broken lines indicate average temperatures for sporomorph-based MATs. Biomarker thermometry data are from Bijl et al. (2021). The ~790 kyr interval corresponding to the EOT (34.44-33.65 Ma; Hutchinson et al., 2021) are marked with orange horizontal bar. Age model after Houben et al. (2019).

Figure 5. Comparison of the sporomorph-based MAT in the Tasmanian Gateway region across the EOT and earliest Oligocene to regional and global marine EOT and earliest Oligocene records. (A) Marine benthic foraminiferal calcite $\delta^{18}\text{O}$ record from ODP Site 1218 (Pälike et al., 2006). (B) Marine $\delta^{11}\text{B}$ -derived atmospheric $p\text{CO}_2$ record (Anagnostou et al., 2016). (C) Terrestrial temperature change across the EOT and earliest Oligocene based on our sporomorph-based MATs from ODP Site 1172.



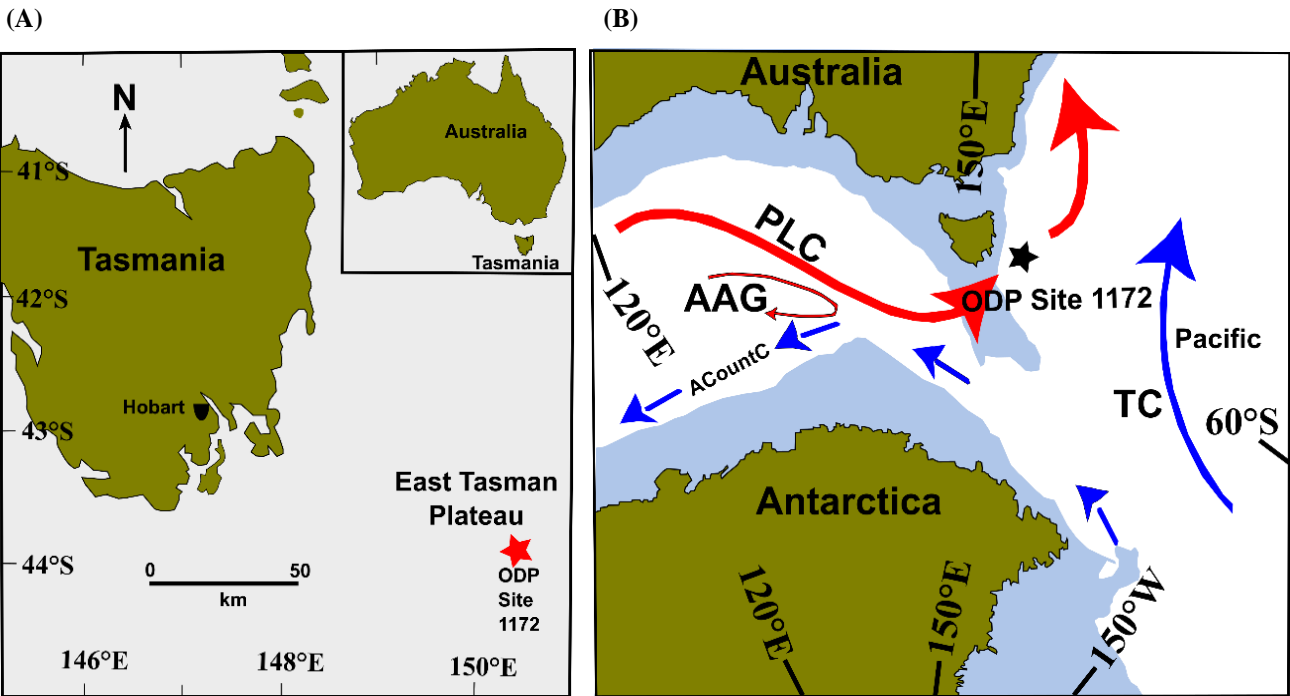
Table captions

Table 1: List of sporomorph taxa from the late Eocene to early Oligocene of ODP Site 1172 accompanied by botanical affinities, literature sources, nearest living relatives (NLR) selected for climatic reconstruction, and inferred climate range from (Macphail, 2007).

Table 2: Summary of quantitative species diversity and Axis 1, DCA sample score between the late Eocene to early Oligocene from ODP Site 1172.



890 **Figure 1.**



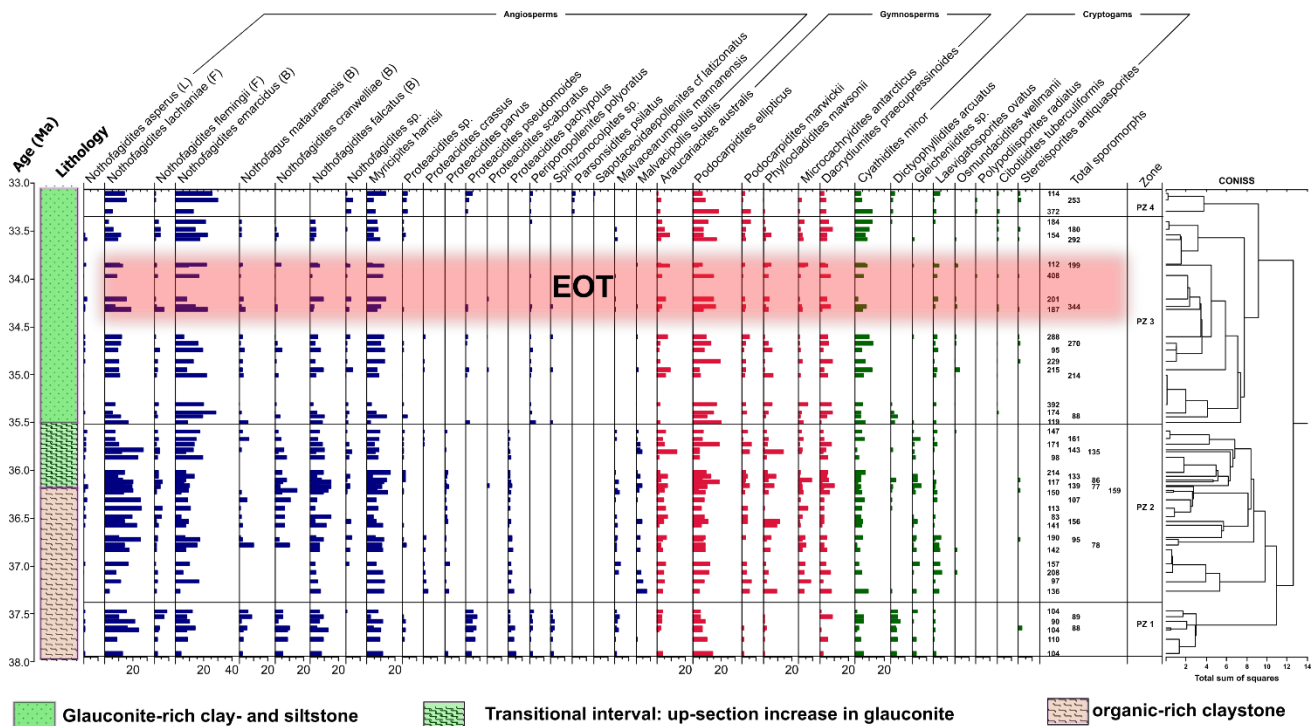
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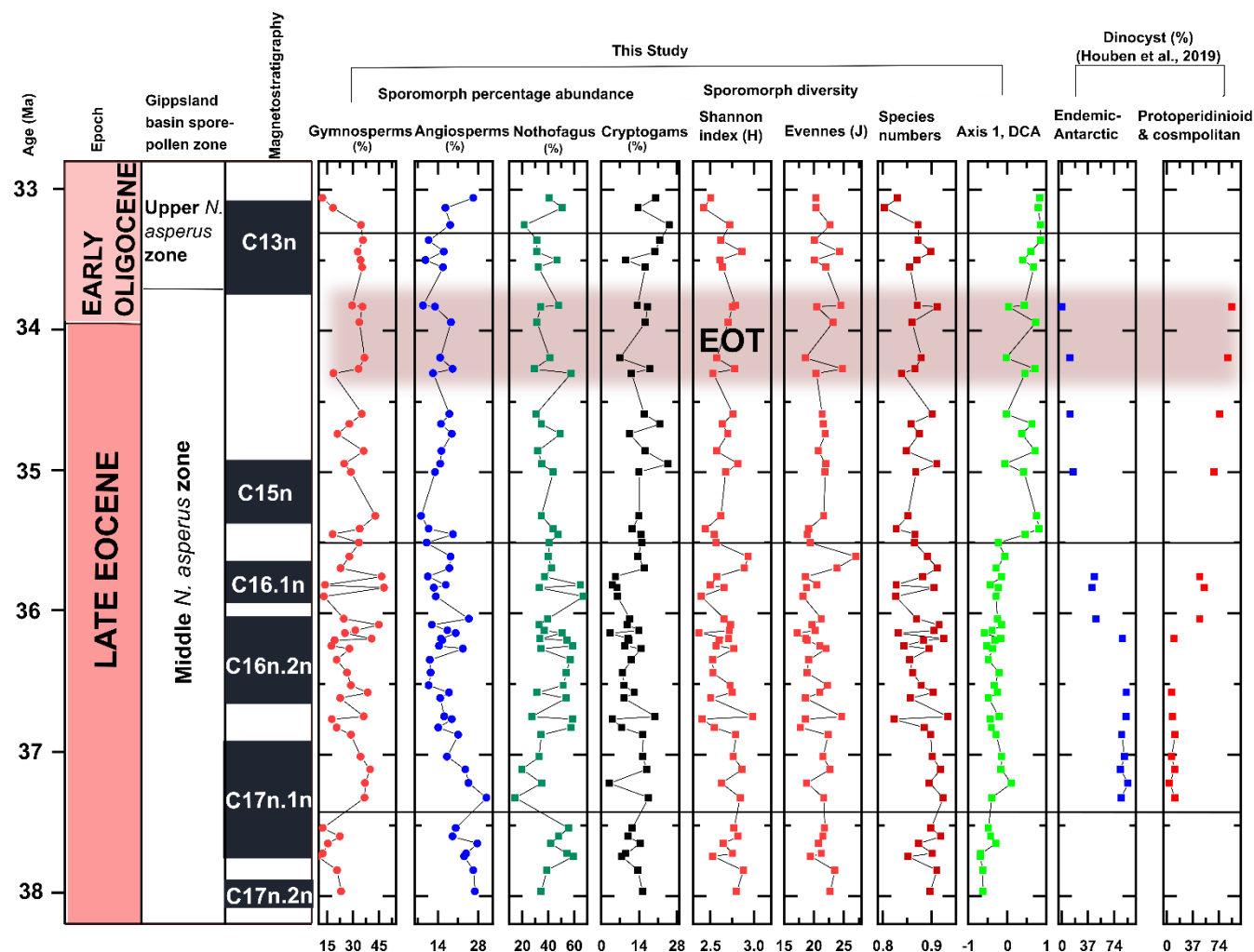


Figure 2.





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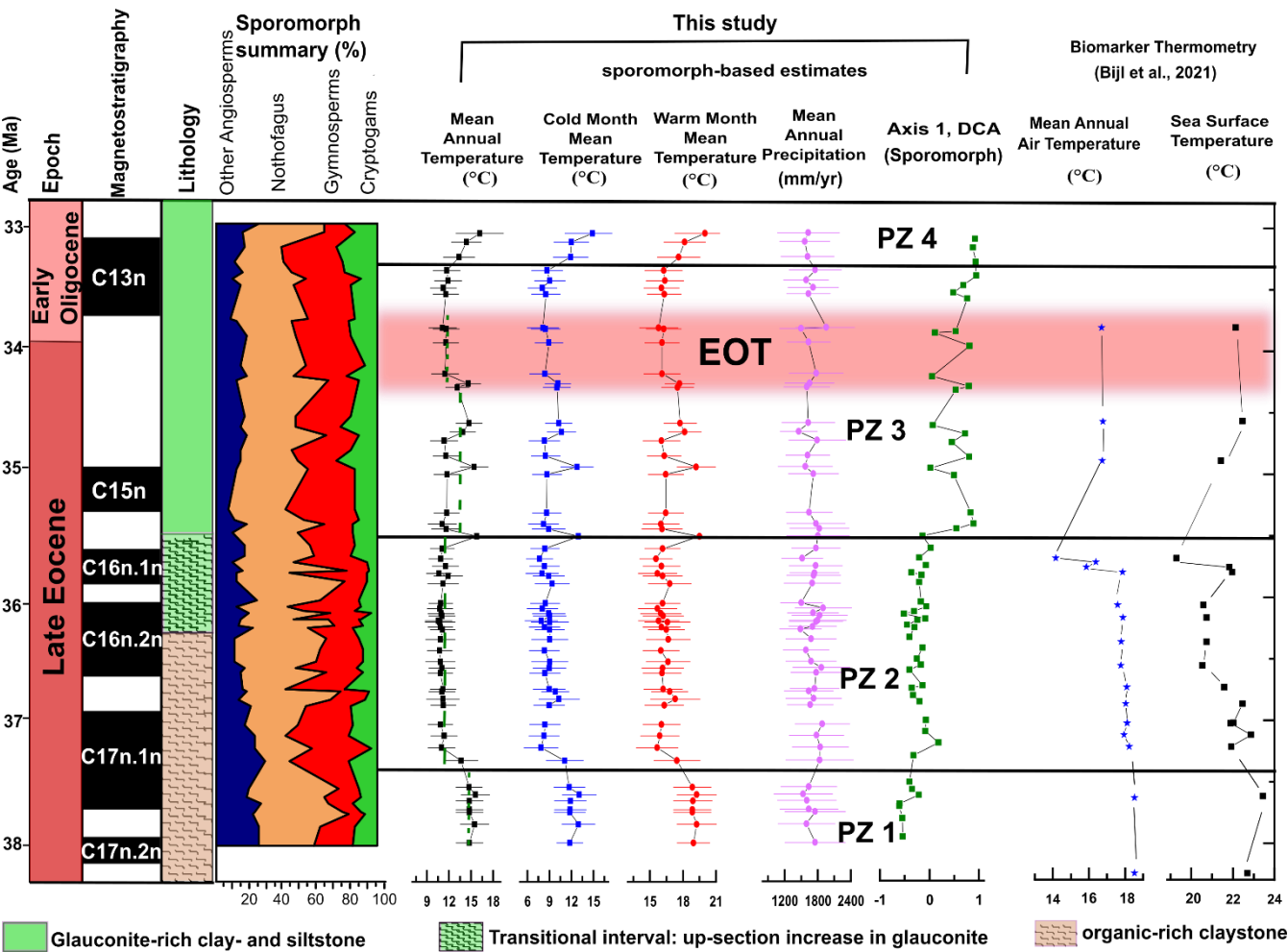


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Figure 4.

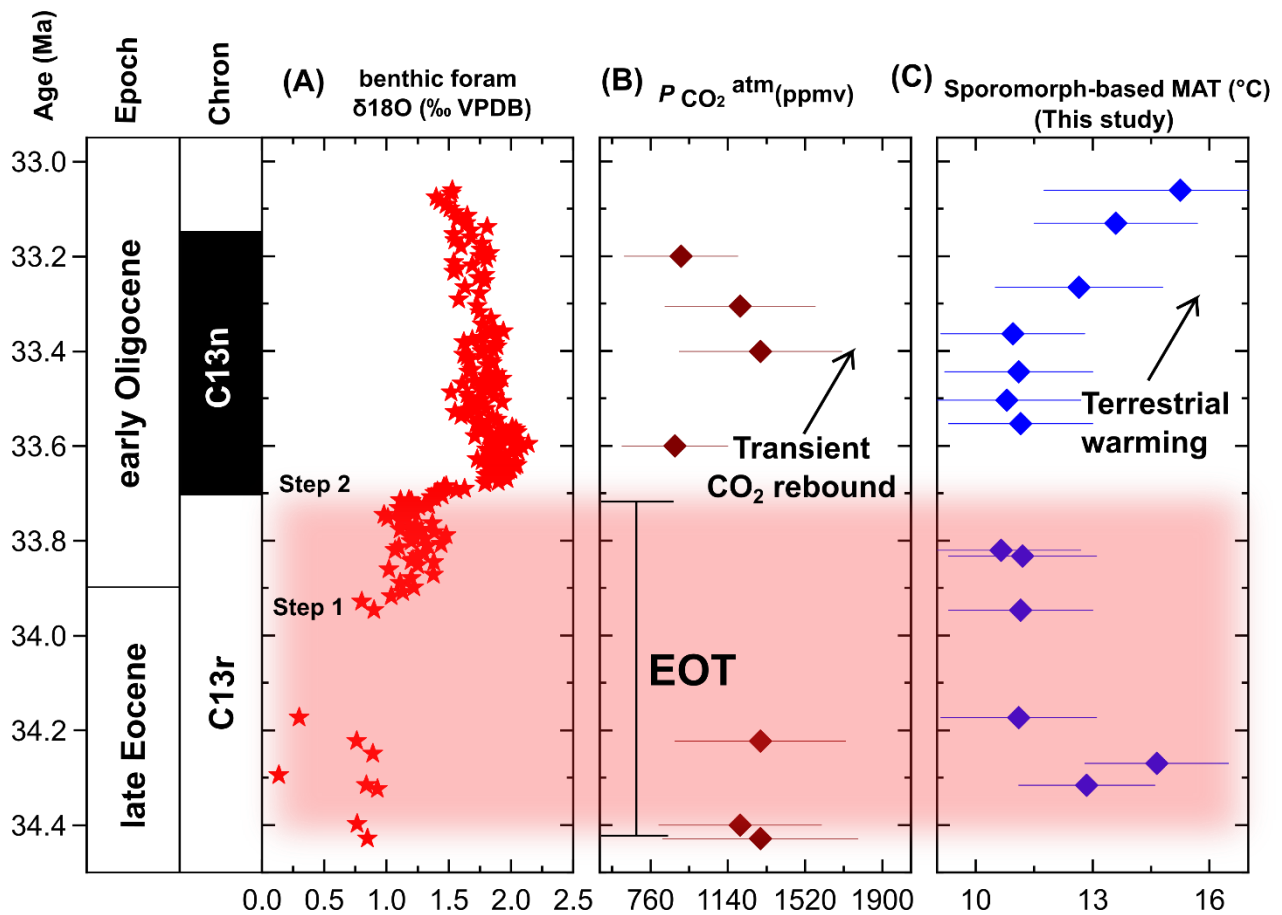


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955 **Figure 5.**



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970 **Table 1.**

Fossil taxon	Botanical Affinity	Source	Selected NLR for climate analysis	Inferred climate Range (Macphail, 2007)
Gymnosperms				
<i>Araucariacites australis</i>	Araucariaceae	Raine et al. (2011)	Araucariaceae	Lower to upper ?mesotherm
<i>Dilwynites granulatus</i>	Araucariaceae	Raine et al. (2011)	Araucariaceae	Lower to upper ?mesotherm
<i>Dacrydiumites preacupressinoides</i>	Podocarpaceae	Raine et al. (2011)	<i>Dacrydium cupressinum</i>	
<i>Podocarpidites ellipticus</i>	Podocarpaceae	Raine et al. (2011)	Podocarpaceae	Microtherm to? megatherm
<i>Podocarpidites</i> spp.	Podocarpaceae	Truswell & Macphail (2009)	Podocarpaceae	Microtherm to ?megatherm
<i>Dacrycarpites australiensis</i>	Podocarpaceae	Truswell & Macphail (2009)	Podocarpaceae	Upper microtherm to lower mesotherm
<i>Podocarpidites marwickii</i>	Podocarpaceae	Raine et al. (2011)	Podocarpaceae	Microtherm to ?megatherm
<i>Phyllocladidites mawsonii</i>	<i>Lagarostrobos</i>	Raine et al. (2011)	<i>Lagarostrobos</i>	Upper microtherm to lower mesotherm
<i>Phyllocladidites reticulasaccatus</i>	Podocarpaceae	Raine et al. (2011)	Podocarpaceae	
<i>Microcachrydites antarcticus</i>	Podocarpaceae	Raine et al. (2011)	<i>Microcachrys</i>	Upper microtherm to lower mesotherm
<i>Taxodiaceapollenites hiatus</i>	Cupressaceae	Raine et al. (2011)	Cupressaceae	
<i>Microaladites</i> sp.	Podocarpaceae	Raine et al. (2011)	Podocarpaceae	Upper microtherm to lower mesotherm
Angiosperms				
<i>Malvacipollis subtilis</i>	Euphorbiaceae	Raine et al. (2011)	Euphorbiaceae	
<i>Myricipites harrisii</i>	Casuarinaceae	Raine et al. (2011) Macphail (2007)	<i>Gymnostoma</i>	Lower mesotherm to megatherm
<i>Nothofagidites flemingii</i>	<i>Nothofagus</i> subg. <i>Fuscospora</i>	Raine et al. (2011)	<i>Nothofagus</i> subg. <i>Fuscospora</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites</i> spp.	<i>Nothofagus</i>	Raine et al. (2011)	<i>Nothofagus</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites emarcidus</i>	<i>Nothofagus</i> subg. <i>Brassospora</i>	Truswell & Macphail (2009)	<i>Nothofagus</i> subg. <i>Brassospora</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites falcatus</i>	<i>Nothofagus</i> subg. <i>Brassospora</i>	Raine et al. (2011)	<i>Nothofagus</i> subg. <i>Brassospora</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites lachlaniae</i>	<i>Nothofagus</i> subg. <i>Fuscospora</i>	Raine et al. (2011)	<i>Nothofagus</i> subg. <i>Fuscospora</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites matauraensis</i>	<i>Nothofagus</i> subg. <i>Brassospora</i>	Raine et al. (2011)	<i>Nothofagus</i> subg. <i>Brassospora</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites waipawaensis</i>	<i>Nothofagus</i> subg. <i>Fuscospora</i>	Raine et al. (2011)	<i>Nothofagus</i> subg. <i>Fuscospora</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites asperus</i>	<i>Nothofagus</i> subg. <i>Lophozonia</i>	Truswell & Macphail (2009)	<i>Nothofagus</i> subg. <i>Lophozonia</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites cranwelliae</i>	<i>Nothofagus</i> subg. <i>Brassospora</i>	Raine et al. (2011)	<i>Nothofagus</i> subg. <i>Brassospora</i>	Upper microtherm to lower mesotherm
<i>Nothofagidites senectus</i>	<i>Nothofagus</i>	Raine et al. (2011)	<i>Nothofagus</i>	
<i>Nothofagidites brachyspinulosus</i>	<i>Nothofagus</i> subg. <i>Fuscospora</i>	Raine et al. (2011)	<i>Nothofagus</i> subg. <i>Fuscospora</i>	
<i>Proteacidites crassus</i>	Proteaceae	Raine et al. (2011)	Proteaceae	Lower to upper mesotherm
<i>Proteacidites pachypolus</i>	Proteaceae	Macphail & Hill (2018)	Proteaceae	Lower to upper mesotherm
<i>Proteacidites pseudomoides</i>	Proteaceae	Raine et al. (2011)	<i>Carnarvonina</i>	Lower to upper mesotherm
<i>Proteacidites leightonii</i>	Proteaceae	Truswell & Macphail (2009)	Proteaceae	Lower to upper mesotherm
<i>Proteacidites reticulatus</i>	Proteaceae	Truswell & Macphail (2009) Raine et al. (2011)	Proteaceae	Lower to upper mesotherm



<i>Proteacidites scaboratus</i>	Proteaceae	Raine et al. (2011)	Proteaceae	Lower to upper mesotherm
<i>Proteacidites similis</i>	Proteaceae	Raine et al. (2011)	Proteaceae	Lower to upper mesotherm
<i>Proteacidites parvus</i>	Proteaceae	Bowman et al. (2014) Raine et al. (2011)	<i>Bellendena</i>	Lower to upper mesotherm
<i>Periporopollenites polyoratus</i>	Caryophyllaceae	Raine et al. (2011)	Caryophyllaceae	
<i>Parsonsidites psilatus</i>	Trimeniaceae <i>Parsonsia</i>	Raine et al. (2011)	<i>Parsonsia</i>	
<i>Spinizonocolpites</i> sp.	Areaceae	Raine et al. (2011) Kumaran et al. 2011	Areaceae	Upper mesotherm to megatherm
<i>Tricolpites trioblatus</i>	Scrophulariaceae Convolvulaceae	Raine et al. (2011)	<i>Hebe</i>	Lower to upper mesotherm
<i>Chenopodipollis chenopodioides</i>	Amaranthaceae	Stevens (2017)	Amaranthaceae subfamily Chenopodioideae	
<i>Malvacearumpollis mannanensis</i>	Malvaceae	Raine et al. (2011)	Malvaceae	
<i>Nupharipollis mortonensis</i>	Araceae Nymphaeaceae	Raine et al. (2011)	<i>Nuphar</i>	Upper mesotherm to megatherm
<i>Sapotaceoidaepollenites cf latizonatus</i>	Sapotaceae	Raine et al. (2011)	Sapotaceae	
Cryptogams				
<i>Cyathidites australis</i>	Cyatheaceae	Raine et al. (2011) Macphail (1994)	Cyatheaceae	Upper microtherm to lower mesotherm
<i>Cyathidites minor</i>	Cyatheaceae	Raine et al. (2011)	Cyatheaceae	Upper microtherm to lower mesotherm
<i>Cyathidites</i> sp.	Cyatheaceae	Raine et al. (2011)	Cyatheaceae	Upper microtherm to lower mesotherm
<i>Laevigatosporites ovatus</i>	Blechnaceae	Raine et al. (2011) Truswell & Macphail (2009)	Blechnaceae	
<i>Osmundacidites wellmanii</i>	Osmundaceae	Raine et al. (2011)	<i>Todea</i>	
<i>Osmundacidites</i> sp.	Osmundaceae	Raine et al. (2011) Raine et al. (2011) Truswell & Macphail (2009)	Osmundaceae	
<i>Baculatisporites comaumensis</i>	Osmundaceae, Hymenophyllaceae	Truswell & Macphail (2009) Macphail and Cantrill (2006)	<i>Hymenophyllum</i>	
<i>Gleicheniidites senonicus</i>	Gleicheniaceae	Truswell & Macphail (2009)	Gleicheniaceae	
<i>Gleicheniidites</i> spp.	Gleicheniaceae	Raine et al. (2011) Truswell & Macphail (2009)	Gleicheniaceae	
<i>Dictyophyllidites arcuatus</i>	Gleicheniaceae	Raine et al. (2011)	Gleicheniaceae	
<i>Kuylisporites waterbolkii</i>	Cyatheaceae	Raine et al. (2011)	Cyatheaceae	Upper microtherm to lower mesotherm
<i>Clavifera rudis</i>	Gleicheniaceae	Raine et al. (2011)	Gleicheniaceae	
<i>Clavifera triplex</i>	Gleicheniaceae	Raine et al. (2011) Truswell & Macphail (2009) Truswell & Macphail (2009)	Gleicheniaceae	
<i>Laevigatosporites major</i>	Blechnaceae	Raine et al. (2011) Macphail & Hill (2018)	Blechnaceae	
<i>Stereisporites antiquasporites</i>	Sphagnaceae	Truswell & Macphail (2009)	<i>Sphagnum</i>	± microtherm
<i>Ceratosporites equalis</i>	Selaginellaceae	Raine et al. (2011)	<i>Selaginella</i>	
<i>Cibotioidites tuberculiformis</i>	Schizaeaceae	Raine et al. (2011) Daly et al. (2011)	Schizaeaceae	
<i>Polypodiisporites radiatus</i>	Polypodiaceae	Raine et al. (2011)	Polypodiaceae	
<i>Retriletes austroclavatidites</i>	Lycopodiaceae	Raine et al. (2011)	<i>Lycopodium</i>	



Table 2.

Analysis	Pollen Zone 1		Pollen Zone 2		Pollen Zone 3		Pollen Zone 4	
	<i>Mean</i>	<i>(SD)</i>	<i>Mean</i>	<i>(SD)</i>	<i>Mean</i>	<i>(SD)</i>	<i>Mean</i>	<i>(SD)</i>
Rarefaction (75 individuals)	21.61	1.32	20.52	2.31	21.37	1.81	21.16	1.37
Shannon index (H)	2.75	1.12	2.66	0.16	2.66	0.10	2.54	0.15
Equitability (J)	0.89	0.02	0.88	0.03	0.87	0.02	0.83	0.03
DCA (Axis 1, sample scores)	-0.55	0.15	-0.29	0.15	0.44	0.33	0.83	0.03