Is succession-based management of coastal dune forest restoration valid?

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Abstract

Here we test the predictions of succession in a restoration context to ascertain the validity of

succession-based management. Our study took place in seven coastal dune forest sites of various

ages regenerating after mining disturbance. We used our 16 yr of data on 4 taxa to test the

predictions of succession theories. Patterns in turnover for all taxa showed a decelerating

decrease contradicting Clements's classical theory of succession. Changes in composition

followed patterns predicted by the individualistic model of succession. Trends in species

diversity measures did not always match predictions with bird species diversity declining in most

sites over time. Regional disturbances may explain this phenomenon. Succession-based

management is a valid approach to dune forest rehabilitation as long as restoration managers

recognise disturbance as an ecological reality.

Keywords: Coastal dune forest, Chronosequence, Mining, Rehabilitation, Succession,

### Introduction

The theory of succession has been a fundamental concept in ecology for over 100 yr (at least since Cowles' publication in 1901) and is perhaps the most enduring of all ecological theories (Walker and del Moral 2008). This theory describes the progressive and deterministic change in species composition and dynamics over time and forms the conceptual basis of restoration ecology (Young et al. 2005). Successional theory may be ideal as a basis for restoration as it is conceptually simple and encapsulates the ability of ecosystems to recover from disturbances. The manipulation of this restorative ability is a fundamental concern of restoration ecology (Walker et al. 2007).

The use of successional theory to frame restoration management is common, but testing the predictions of this theory in a restoration setting is not (Walker and del Moral 2008, but see Prach and Pyšek 2001 and Řehounková and Prach 2008). Restoration practice often relies more upon horticulture, agronomy, and engineering than on succession (Hodačova and Prach 2003, Young et al. 2005). Hobbs and colleagues (2007) posit that this separation stems from cultural and conceptual differences between restoration practitioners and those studying succession. Many aspects of succession make it unattractive to restoration practitioners. For example, successional studies are often on a time scale of hundreds or thousands of years, whereas a restoration project may only last 20 or 30 yr (Dobson et al. 1997). Importantly, there have been numerous examples of the failure of successional trajectories to achieve restoration targets (e.g., Zedler and Callaway 1999, Suding et al. 2004).

Many factors may render succession unpredictable, including priority effects (Connell and Slatyer 1977), edaphic and topographical heterogeneity (Cutler 2009), propagule

availability, species dispersal (Lanta and Lepš 2009), persistence of pioneers (Tsuyuzaki 2009), herbivory (Baniya et al. 2009), and the sporadic establishment of strong dominants (Walker and del Moral 2003). In addition, the structure, composition, and connectivity of the landscape may also have strong effects on species composition (Grainger et al. 2011). These and other factors may compromise the efficacy of succession-based restoration management (Suding et al. 2004). Succession theory, however, offers several predictions of the trends in species composition and other community properties that can be expected after a disturbance event. The outcomes of restoration actions should be predictable using successional theory (Van Andel and Aronson 2006). Temporal trends expected from succession include directional changes in species composition, increased species diversity (in early succession at least; Connell 1978) and increased stability (Dobson et al. 1997).

Perhaps the most alluring aspect of succession for restoration ecology is the concept that after a disturbance event, habitats will predictably recover their former structure and function (Walker & del Moral 2007). Accordingly, there will be a directional progression in species composition, which becomes more similar over time to an undisturbed community (Pickett et al. 1987; Wassenaar et al. 2005). The traits of species should also be predictable so that immediately after the disturbance, species with life histories that are adapted to survive in harsh conditions colonize the site (Dobson et al. 1997). Increasingly, competitive interactions will structure the community, whereby pioneer species are replaced by species that are superior competitors (Dobson 1997). This increased competition should eventually lead to community stability (Anderson 2007).

The definition of stability is ambiguous in the context of succession (Drury & Nisbet 1973); however, many authors have equated stability with compositional turnover (e.g. Anderson

2007). The rate of species turnover is predicted to be greatest at the earliest stages of succession (Drury & Nisbet 1973). This pattern occurs as the regional species pool becomes exhausted and the rate of competition increases as a community assembles, making it more difficult for new species to colonize (Tilman 1997). There are different responses of turnover expected for different theories of succession; the classical theory of Clements (1916) predicts spikes in turnover rate at each discrete community transition. However, if species are independent units (Gleason 1926) then turnover will be high initially and then decline to a continuous low level (a decelerating decrease). Anderson (2007) showed this response to be common in plant and arthropod successions.

If the reassembly of biotic communities after disturbance is predictable then species diversity may be expected to follow one of three possible trends. If succession leads to increased structural heterogeneity then species richness and diversity should increase (Odum 1969). If all the components of the species assemblage are present at the beginning of succession (Egler 1954) diversity and richness are maximized in the early stages of succession. The mid-stages of succession, however, may be the stages at which the species that are capable of establishing have done so, but competition has not yet filtered the species assemblage, leading to the highest levels of diversity (Howard & Lee 2003). Therefore, an initial increase in species richness and diversity will be followed in the later stages of succession by a decline, in the absence of further disturbance (Connell 1978).

Here we report on a restoration program that relies on successional processes to restore coastal dune forests destroyed by mining (see van Aarde et al. 1996a). The program began in 1977, and our research initiatives commenced during 1991. Several of our earlier papers (e.g., Ferreira and van Aarde 1996, van Aarde et al.1996a, 1996b, Kritzinger and van Aarde 1998,

Grainger et al. 2011) implied that the recovery of communities on these sub-tropical dune forests is consistent with models of succession i.e. deterministic. However, none focused on evaluating if successional patterns underlie forest regeneration following rehabilitation, as practiced here. In this paper, we wish to ascertain if the assumption that succession is a valid model for the restoration of coastal dune forest is correct. We assess 4 community level trends expected to result from ecological succession. Several of our previous papers have addressed the convergence of regenerating coastal dune forest with an undisturbed reference forest, so we will not address this aspect of succession herein (e.g.,van Aarde et al. 1996b, Davis et al. 2003, Redi et al. 2005, Wassenaar et al. 2005). We used data on trees and herbaceous plants from regenerating coastal dune forests to determine: 1) if the rate of species turnover decreases as coastal dune forest develops; 2) if there is a sequence of changing species "types" from pioneer species adapted to harsh conditions to species adapted to high levels of competition; 3) if this sequence of types is directional and consistent across all sites with similar climatic conditions; and 4) if species diversity increases, decreases, or fluctuates.

### Methods

## Study Sites and Restoration Process

The study area included circa 60 km of coastline between Richards Bay Town (28°43'S, 32°12'E) and the Sokhulu forest (28°27'S, 32°25'E) in KwaZulu-Natal, South Africa. The climate is humid and subtropical, with mean annual rainfall of  $1488 \pm 447.3$  mm [mean  $\pm$  SD, n = 34 yr between 1976 and 2009; data courtesy of Richards Bay Minerals (RBM),  $28^{\circ}$  41.153 S  $32^{\circ}$  8. 251 E]. Rainfall peaks in February, and the mean ( $\pm$  SD) temperature is  $23.8 \pm 3.50$  °C (monthly temperature between 2006 and 2009 (data courtesy of RBM)). These coastal dunes have been mined for minerals since 1977 (van Aarde et al. 1996a).

RBM aims to return indigenous coastal dune vegetation to one third of its mined area. The mining process (described in full in van Aarde et al. 1996a) destroys all vegetation in front of the mine-pond. Prior to mining, the topsoil is removed and stockpiled. Immediately post-mining, sand dunes are mechanically reshaped and topsoil replaced. The topsoil is then stabilized using drift-fencing and seeded with exotic annual plants (sunhemps, *Crotolaria sp.* and sunflowers, *Helianthus sp.*). These annuals do not successfully reproduce in regenerating sites and are absent past the third year of regeneration (Conservation Ecology Research Unit, University of Pretoria, unpublished data). We refer to this stabilization of dunes and return of topsoil as the "kick-start" to succession (van Aarde et al. 1996a, van Aarde et al. 1996c), after which there is minimal management intervention (removal of non-native plant species that successfully disperse to the sites and herbivores), and subsequently the restoration relies on natural successional processes (van Aarde et al. 1996a, van Aarde et al. 1996b).

## Data Collection

We conducted trees surveys in 1999, 2001 and 2005 in 7 (6 in 1999) restored sites, following the methods described by Wassenaar and others (2005). We identified self-supporting woody plants greater than 1.7 m in height in 7 16x16-m quadrats per site. In 2009, we used the Point-Centre Quarter method (PCQ) along randomly located transects that ran perpendicular to the sea (Cottam and Curtis 1956).

To survey the herbaceous plant community, we identified and counted all plant species below 1 m in height in 10 randomly located (5 in 2003) plots in each rehabilitating site and in the undisturbed reference forest (Wassenaar et al. 2005). A plot consisted of 10, 1-m<sup>2</sup> quadrats placed in a 2 x 5-m pattern with 5 m separating each quadrat. Data were collected in 1995, 1999, 2003, and 2005 (Table 1). Tree seedlings (< 1.7 m in height) were not included in the analyses.

# **Analysis**

Previous work in the study region has relied on the chronosequence approach (e.g., Ferreira and van Aarde 1997, Davis et al. 2003, Wassenaar et al. 2005), which has been criticised because it ignores site-specific effects (Chazdon et al. 2007, Johnson and Miyanishi 2008). To determine if trends in successional patterns were not just artefacts of the chronosequence, we identified trends (where data allowed) within individual sites across several survey events and compared them to trends across a chronosequence (substituting space for time). We used data from all of our 16 (at most) survey years to produce each chronosequence. We could not use data from the same site at different ages because this would have violated the assumptions of a chronosequence. Instead, we constructed simulated chronosequences by a stratified random sampling procedure, where a site was only included once within each chronosequence. The data could come from any one of the survey years. This procedure was repeated 1000 times, and the mean values were used in the chronosequence.

To test the assumption that sites of a similar age would have similar composition we used non-metric multidimensional scaling (NMDS) using the function 'metaMDS' of the package VEGAN (v.1.15-3; Oksanen et al. 2008) in the R statistical software (v. 2.8.1; R Core Team Development 2008). We used the Bray-Curtis dissimilarity index and set the 'zerodist' argument to add a small positive value to zero dissimilarities. We confirmed patterns of community composition shown in the NMDS using analysis of similarity (using the function 'anosim' in R's VEGAN package). Further, to test if changes in species composition were related to the age of sites we used a Mantel test using the function 'mantel' once again in the VEGAN package. This test measures concordance between two distance matrices - community similarity versus time in this instance. We determined species habitat associations from published sources for herbaceous plants (Pooley 1998), trees (Coates-Palgrave 2003), and birds (Gibbon 2006). No independent sources of information exist for millipede habitat associations.

To test the assumption that pioneers do not replace themselves, we used diameter at breast height (DBH) measurements from the 2009 tree survey to create size class distribution plots for all sites combined. We only assessed the most abundant tree species, including Sweet thorn (*Acacia karroo*; the most abundant tree in the rehabilitating sites), White stinkwood (*Celtis africana*), Turkey berry (*Canthium inerme*), Coast silver oak (*Brachlaena discolour*), Dune false currant (*Allophylus natalensis*), Pigeon wood (*Trema orientalis*), Coastal golden leaf (*Bridelia micrantha*), Tassel berry (*Antidesma venosum*) and Black monkey-orange (*Strychnous gerrardii*). We combined data for Coastal red milkwood (*Mimusops caffra*) and Bush red milkwood (*M. obovata*) as well as data for Natal mahogany (*Trichilia emetica*) and Forest mahogany (*T. dregeana*) as distinction between these related species is difficult.

We calculated species turnover as the average of species gains and losses between survey events, expressed as a proportion of the mean species richness during the survey period (Anderson 2007). Gains were defined as the number of new species added to the community, and losses were defined as the number of species lost from the community. We did not consider the reappearance of a previously present species in the community as ecologically significant; rather, we assumed that this was an artefact of sampling or a stochastic event. Therefore, these data were not counted in the calculation of turnover. We used regression analysis to assess how much variation in turnover could be attributed to regeneration age.

We calculated species richness as the number of species per plot. We calculated species diversity using the Shannon index of diversity and evenness using the Smith and Wilson evenness index (E<sub>var</sub>; Smith and Wilson 1996). We compared the regression slopes of individual sites to those predicted by randomized chronosequences using the method described by Zar (1984) and calculated in the program GraphPad Prism 3.0. Data were tested for departure from linearity using a runs test in GraphPad Prism 3.0.

## **Results**

Species turnover

Species turnover for both taxa followed the expected pattern of a decelerating decline in turnover rate with increased regeneration age (Figure 1). Regeneration age explained at least 30% of the variation in turnover rate (non-linear regression, trees:  $R^2 = 0.31$ , herbaceous plants:  $R^2 = 0.50$ ).

Change in life-history traits

The stress of the NMDS ordinations was relatively high (>20 %) with a two-dimensional ordination (i.e., k = 2); however, the results of the ANOSIM confirmed that community composition for both taxa differed between site ages (trees: R = 0.47, p<0.001; herbaceous plants: R = 0.31, p < 0.001, Figure 2). The tree community ordination showed clearer convergence between the plots in sites of a similar age (Figure 2a; NMDS, stress = 19.00, k = 3, non-metric fit  $r^2 = 0.98$ , linear fit  $r^2 = 0.93$ ). For the herbaceous plants, plots in sites aged between 1 and 5 yr old, and those between 6 and 10 yr old were separated from the plots >11 yr old, which overlapped considerably (Figure 2b; NMDS, stress = 17.25, k = 3, non-metric fit  $r^2 = 0.95$ , linear fit  $r^2 = 0.83$ ). The Mantel test results confirm these changes in species composition with regeneration age (Mantel test; trees: r = 0.48, p < 0.001; herbaceous plants: r = 0.38, p < 0.001). For both trees and herbaceous plants species identity appears to broadly shift from species adapted to harsh environments to those more typical of forests (Figure 2). Of the 11 tree species for which we assessed size class distributions only Acacia karroo and Antidesma venosum did not show the reverse-J pattern indicative of active recruitment (we only show results for four species in Figure 3).

Trends in species richness, diversity and evenness

Site and taxon specific chronosequential trajectories for richness, diversity, and evenness were idiosyncratic (Table 2). Herbaceous plant richness and diversity decreased in the youngest site with increased age and increased in older sites. This pattern was also evident in the youngest site for tree evenness (Table 2).

## **Discussion**

Our findings suggest that succession is a valid model of coastal dune forest restoration. The rate of succession (species turnover) exhibited by all the taxa showed a decelerating decrease; sites that have recently undergone disturbance had a greater compositional turnover. As communities aged, the number of available microsites apparently declined, and interspecific competition may have increased (Gross 1980, Tilman 1997). Compositional stability is a key concept in the theory of succession (McCook 1994, Anderson 2007), and the declining decrease in the rate of turnover fits with Egler's (1954) theory of initial floristics, Gleason's (1926) individualistic theory of succession, and Connell's (1978) intermediate disturbance hypothesis. This pattern does not support Clements's (1916) classical theory of succession.

The shift in species composition from species adapted to harsh environments to those species that are superior competitors appears to be ubiquitous in forest succession (example.g., Kardol et al. 2005, Cutler et al. 2008, Lebrija-Trejos et al. 2008). However, Chazdon and colleagues (2007) and Johnson and Miyanishi (2008) suggest that this trend in species compositional changes is an artefact of the chronosequence approach. Our results dispute this, as we have demonstrated that sites of a similar age share a similar species composition of coastal dune forest trees and herbaceous plants. This change in species composition may be driven by differences in species longevity, tolerance to shade, eventual size (a competitive advantage for forest trees), timing of colonization, and patterns of recruitment (Fajardo and González 2009). It is evident from the tree community in particular, that recruitment of *A. karroo*, the first tree to colonize regenerating sites, is limited under a canopy consisting of adults of the same species; whereas, other longer-lived pioneer and forest species were actively recruiting. This pattern fits the predictions of Connell and Slayter's (1977) facilitation mechanism of succession. Without empirical knowledge of species tolerance to shade, it is difficult to conclude that light intensity is

driving the replacement of plant species in coastal dune forest. However, this is a plausible driver of plant species replacement (Woods 2000, Fajardo and González 2009, but see Dietze and Clarke 2008 for counter arguments to this paradigm), and changes in the tree community may influence changes in the other taxa (Kritzinger and van Aarde 1998).

Trends in species richness and diversity for both trees and herbaceous plants appeared to match expectations overall. As sites aged, they increased in the number and diversity of species. For the herbaceous plants, the youngest site included in our survey (established in 2000, 3 to 5 yr old) decreased in species richness as it aged, which could be consistent with Egler's (1954) initial floristics model. However, this is unlikely, as older sites still gained species with time. The loss of species was more possibly the result of a change in habitat type from grassland to scrub and woodland.

The differences in chronosequence predictions and observed dynamics, although small, highlight an important proviso when using the chronosequence approach. Both regional scale climatic perturbations, such as drought, and site specific differences in the response to disturbance may obscure trends in diversity and richness (Foster and Tilman 2000, Svensson et al. 2009).

# Succession Drives Coastal Dune Forest Restoration

The patterns of age-related species turnover, trends in species type, richness, diversity, and evenness within regenerating coastal dune forest followed the trends expected from Gleason's (1926) individualistic model of successional theory. Gleason's model follows many of the same predictions of trends in communities over time as Clements's (1916) classical model. The difference is that in Gleason's (1926) model community properties are the sum of individual

species and these species will appear and disappear in a successional sere as independent units. Our observational study can only hint at the mechanisms of change in species composition over the course of succession. To identify the mechanism that is driving these changes, we may have to change the focus of our research from the observational to the experimental. Successional theory does provide hypotheses that can be tested experimentally; for example, Connell and Slayter's (1977) 3 pathways of community succession: facilitation, inhibition, or tolerance (although see McCook 1994, for a critique of these models) must be an avenue of future research at the study site. In addition, we must gather more information on the natural history of species in the regenerating coastal dune forest to fully understand the processes of forest regeneration.

Based on our assessment, succession (Gleason's 1926 individualistic model) is a valid model for the restoration of sub-tropical coastal dune forests. However, departures from the expected patterns do occur, which are likely the result of global, regional, or local scale disruptions and disturbance (Trimble and van Aarde 2011) and landscape composition (Grainger et al. 2011). It is imperative that any restoration project that relies on successional-based management allows for and expects external disruptions to the pattern of succession (Walker and del Moral 2003). In fact, continued disruption and disturbance should be embraced as a natural part of ecosystem dynamics.

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Table 1. Survey years and site ages for each rehabilitating coastal dune forest site in KwaZulu-Natal, South Africa, indicated by the year of establishment 1977 to 2003. Surveys for herbaceous plants, and trees took place in different survey years. Highlighted in grey are the survey years for each.

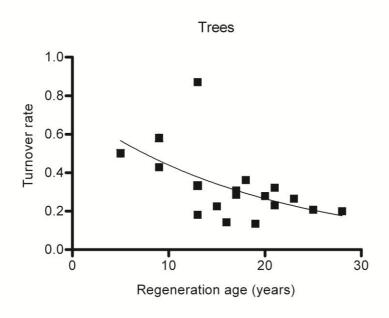
Survey year	1993	1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009
Survey year	1993	1774	1993	1990	1991	1990	1999	2000	2001	2002	2003	2004	2003	2000	2007	2000	2009
Herbs																	
Trees																	
1977	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32
1980	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29
1984	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25
1988	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21
1992	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17
1996	X	X	X	X	1	2	3	4	5	6	7	8	9	10	11	12	13
2000	X	X	X	X	X	X	X	X	1	2	3	4	5	6	7	8	9
2003	X	X	X	X	X	X	X	X	X	X	X	1	2	3	4	5	6

Table 2. The slope of the regression lines for species richness, diversity, and evenness for herbaceous plants, and trees occurring in regenerating sites in KwaZulu-Natal, South Africa. The randomized chronosequence is the expected trend, and the individual sites is the observed trend. Asterisks (\*) indicate a significant ( $\alpha = 0.05$ ) difference between the slope of the expected and observed regression line.

	Expected		Observed			
	Slope of the regression line	Significantly non-zero (alpha = 0.05)	Slope of the regression line	Significantly non-zero (alpha = 0.05)		
Herbaceous plants – species per transect						
Randomised chronosequence	-0.06 ± 0.20	No				
18 to 28 years of regeneration			0.34 ± 0.14	Yes		
15 to 25 years of regeneration			0.61 ± 0.16*	Yes		
11 to 21 years of regeneration			0.50 ± 0.12*	Yes		
7 to 17 years of regeneration			0.99 ± 0.12*	Yes		
3 to 13 years of regeneration			0.17 ± 0.17	No		
3 to 9 years of regeneration			1.25 ± 0.34*	Yes		
3 to 5 years of regeneration			-2.82 ± 0.94*	Yes		
Trees – species per transect						
Randomised chronosequence	$0.42 \pm 0.07$	Yes				
14 to 28 years of regeneration			0.64 ± 0.15	Yes		
11 to 25 years of regeneration			0.058 ± 0.15*	No		
8 to 21 years of regeneration			$0.64 \pm 0.14$	Yes		
4 to 17 years of regeneration			0.58 ± 0.12	Yes		
9 to 13 years of regeneration			$0.81 \pm 0.40$	Yes		
5 to 9 years of regeneration			1.02 ± 0.30	Yes		
1 to 5 years of regeneration			0.71 ± 0.19	Yes		

Randomised chronosequence	$0.02 \pm 0.00$	Yes			
18 to 28 years of regeneration			$0.03 \pm 0.01$	Yes	
15 to 25 years of regeneration			$0.04 \pm 0.01$	Yes	
11 to 21 years of regeneration			$0.03 \pm 0.01$	Yes	
7 to 17 years of regeneration			$0.06 \pm 0.01^*$	Yes	
3 to 13 years of regeneration			$0.02 \pm 0.02^*$	No	
3 to 9 years of regeneration			$0.13 \pm 0.03$	Yes	
3 to 5 years of regeneration			$-0.13 \pm 0.05$	Yes	
Trees – diversity					
Randomised chronosequence	$0.08 \pm 0.02$	Yes			
14 to 28 years of regeneration			$0.03 \pm 0.02^*$	No	
11 to 25 years of regeneration			$0.04 \pm 0.04^*$	No	
8 to 21 years of regeneration			$0.21 \pm 0.02$	Yes	
4 to 17 years of regeneration			$0.12 \pm 0.03$	Yes	
9 to 13 years of regeneration			$0.17 \pm 0.05$	Yes	
5 to 9 years of regeneration			$0.05 \pm 0.05^*$	No	
4 to Fiveers of responsible			$0.06 \pm 0.02$	Yes	
1 to 5 years of regeneration			0.00 ± 0.02	100	
Herbaceous plants – evenness			0.00 1 0.02	100	
Herbaceous plants –	0.00 ±0.01	No	0.00 ± 0.02		
Herbaceous plants – evenness	0.00 ±0.01	No	-0.00 ± 0.00	No	
Herbaceous plants – evenness Randomised chronosequence	0.00 ±0.01	No			
Herbaceous plants – evenness  Randomised chronosequence 18 to 28 years of regeneration	0.00 ±0.01	No	-0.00 ± 0.00	No	
Herbaceous plants – evenness  Randomised chronosequence 18 to 28 years of regeneration 15 to 25 years of regeneration	0.00 ±0.01	No	-0.00 ± 0.00 -0.01 ± 0.00	No No	
Herbaceous plants – evenness  Randomised chronosequence 18 to 28 years of regeneration 15 to 25 years of regeneration 11 to 21 years of regeneration	0.00 ±0.01	No	-0.00 ± 0.00 -0.01 ± 0.00 -0.01 ± 0.00*	No No Yes	
Herbaceous plants – evenness  Randomised chronosequence 18 to 28 years of regeneration 15 to 25 years of regeneration 11 to 21 years of regeneration 7 to 17 years of regeneration	0.00 ±0.01	No	$-0.00 \pm 0.00$ $-0.01 \pm 0.00$ $-0.01 \pm 0.00^*$ $-0.00 \pm 0.00$	No No Yes No	
Herbaceous plants – evenness  Randomised chronosequence 18 to 28 years of regeneration 15 to 25 years of regeneration 11 to 21 years of regeneration 7 to 17 years of regeneration 3 to 13 years of regeneration	0.00 ±0.01	No	$-0.00 \pm 0.00$ $-0.01 \pm 0.00$ $-0.01 \pm 0.00^*$ $-0.00 \pm 0.00$ $0.00 \pm 0.01$	No No Yes No	
Herbaceous plants – evenness  Randomised chronosequence 18 to 28 years of regeneration 15 to 25 years of regeneration 11 to 21 years of regeneration 7 to 17 years of regeneration 3 to 13 years of regeneration 3 to 9 years of regeneration	0.00 ±0.01	No	$-0.00 \pm 0.00$ $-0.01 \pm 0.00$ $-0.01 \pm 0.00^{*}$ $-0.00 \pm 0.00$ $0.00 \pm 0.01$ $-0.00 \pm 0.01$	No No Yes No No	
Herbaceous plants – evenness  Randomised chronosequence 18 to 28 years of regeneration 15 to 25 years of regeneration 11 to 21 years of regeneration 7 to 17 years of regeneration 3 to 13 years of regeneration 3 to 9 years of regeneration 3 to 5 years of regeneration	0.00 ±0.01 0.03 ± 0.01	No	$-0.00 \pm 0.00$ $-0.01 \pm 0.00$ $-0.01 \pm 0.00^{*}$ $-0.00 \pm 0.00$ $0.00 \pm 0.01$ $-0.00 \pm 0.01$	No No Yes No No	

11 to 25 years of regeneration	-0.01 ± 0.01*	No
8 to 21 years of regeneration	-0.00 ± 0.01*	No
4 to 17 years of regeneration	$0.00 \pm 0.01^*$	No
9 to 13 years of regeneration	$0.00 \pm 0.00^*$	No
5 to 9 years of regeneration	-0.00 ± 0.01*	No
1 to 5 years of regeneration	-0.01 ± 0.00*	Yes



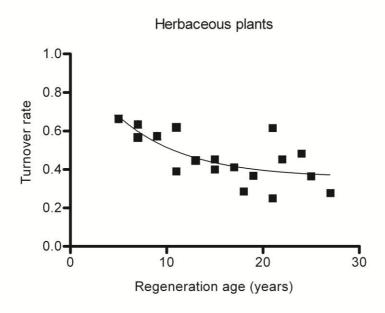
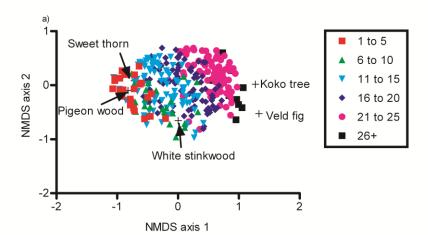


Figure 1. The rate of herbaceuous plant, tree, bird, and millipede species turnover in regenerating coastal dune forests in KwaZulu-Natal, South Africa. .



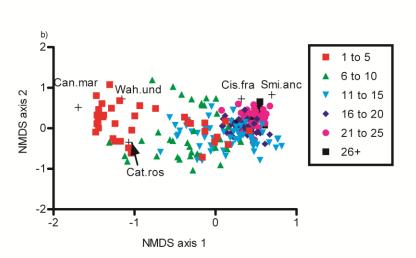


Figure 2. Ordination of regenerating coastal dune forest communities of known age in KwaZulu-Natal, South Africa using non-metric multidimensional scaling (NMDS). Sites of similar age (grouped together in 5-yr categories) are indicated by the same symbol. Species are indicated by small black crosses (+) for the: a) trees; b) herbaceous plants. Labels for the herbaceous plants (b) refer to "Can.mar" = Bay bean (*Canavalia maritima*), "Cat.ros" = Madagascar periwinkle (*Catharanthus roseus*), "Cis.fra" = Forest grape vine (*Cissus fragilis*), "Smi.anc" = Wild sarsaparilla (*Smilax anceps*) and "Wah.und" = Giant bell flower (*Wahlenbergia undulata*).

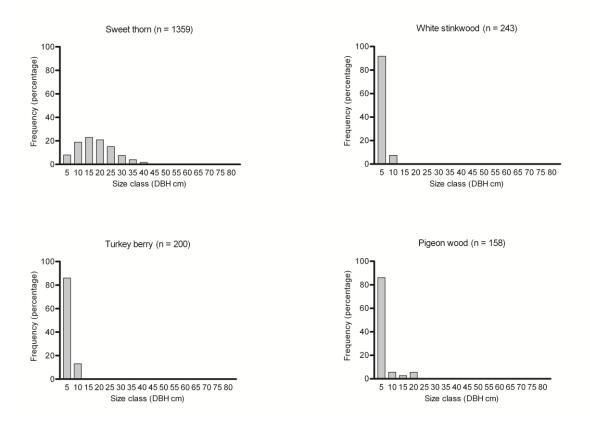


Figure 3. Size class distributions of the dominant pioneer tree Sweet thorn (*Acacia karroo*), and the most abundant three other tree species (White stinkwood, *Celtis africana*; Turkey berry, *Canthium inerme*; Pigeon wood, *Trema orientalis*) within the regenerating coastal dune forests in KwaZulu-Natal, South Africa. The only tree that did not show a regenerating population is the pioneer *Acacia karroo*.