# Weed analysis

### Craig Simpkins

#### Impact of weeds

Ran model for 1000 steps for 15 repeats with ground weeds on and off. Weed parameters were randomly selected based on initial settings and only testing the clustered scenario. The parameter values were:

- trad-init-cover = 0.150
- trad-spread-local = 0.20
- trad-spread-long = 0.025
- trad-growth = 1.0
- smother-f = 0.20

#### Impact of weed parameters (i.e. sensitivity)

Sensitivity analysis model runs were run for 1000 steps, repeated six times (due to time constraints) for every parameter value combination. Note that parameter values were altered by 100% above and below baseline levels, e.g., if baseline was 0.5 runs were conducted at 0 and 1. This high degree of change was done to allow for analysis of impact of extreme changes in addition to sensitivity.

#### **Abundance**

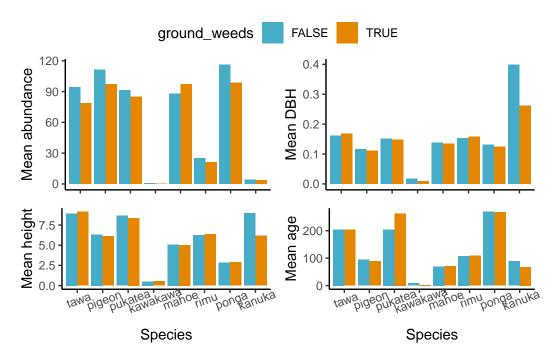
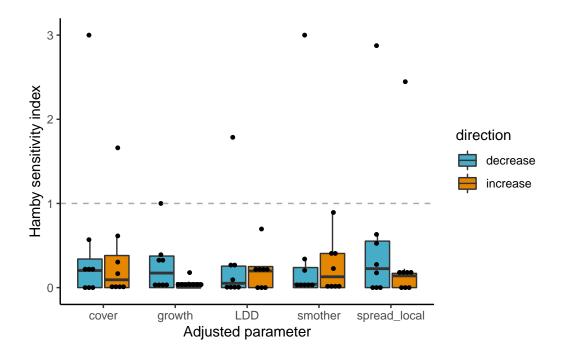
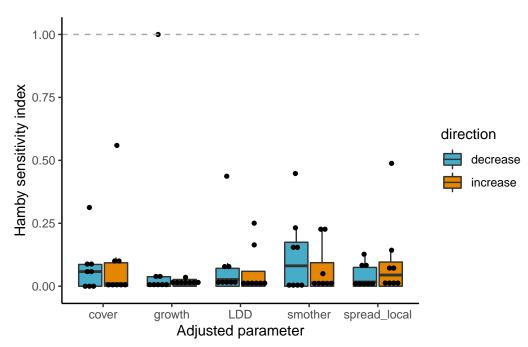


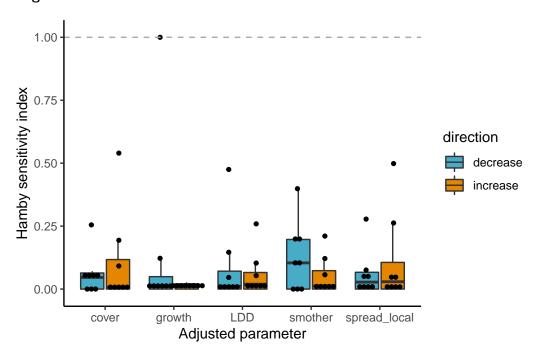
Figure 1: Composition measures for each species taken at 1000 time steps (years) with and with out ground weed cover, presented as the mean of the 15 model repeats.



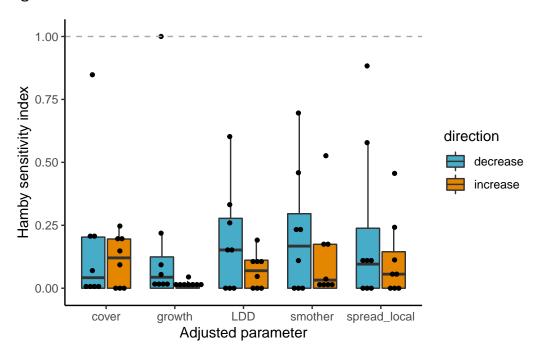
### DBH



### Height



### Age



### Resilience to extreme parameter values

Selected the parameter settings which had the largest percentage change compared to the no ground weed runs.

#### **Abundance**

$\operatorname{trad}_{\underline{\cdot}}$	_init_cove	mother_	_frad_spread_lotaald_	_spread_	_l <b>oræ</b> dgro	wtspecies	$abun\_diff$	abun_per
	0.0	0.0	0.0	0	0.00	kanuka	0.000000	NaN
	0.0	0.0	0.0	0	0.00	kawakay	v <b>a</b> 0.000000	NaN
	1.0	1.0	0.0	0	1.25	mahoe	45.600000	130.03802
	0.0	0.5	0.5	1	0.00	pigeon	204.400000	127.00911
	0.5	0.0	1.0	1	2.50	ponga	63.900000	86.42922
	1.0	0.5	1.0	1	1.25	pukatea	-	74.87036
							125.133333	}
	1.0	1.0	1.0	1	0.00	$\operatorname{rimu}$	2.066667	775.00000
	0.0	0.0	0.0	1	0.00	tawa	74.733333	81.58661

DBH

$\overline{\mathrm{trad}}$	_init	_cov <b>er</b> mother_	_frad_spread_	_lo <b>ta</b> ald	spread_	_lonagd_	_grow	tspecies	$\mathrm{dbh}_{\underline{\ }}$	_diff	dbh_	_per
	0.	0.0	0.0		0.0		0.00	kanuka	0.00	00000	N	aN
	0.	0.0	0.0		0.0	(	0.00	kawakaw	<b>10.00</b>	00000	N	aN
	0.	$5 \qquad 0.5$	0.5		0.5	(	0.00	mahoe		-	100.0	00000
									0.11	35333		
	0.	$5 \qquad 0.0$	0.5		0.5		1.25	pigeon		-	44.36	3851
									0.05	41000		
	0.	0.0	0.5		0.0		2.50	ponga		-	42.84	1038
									0.06	08333		
	0.	0.0	0.5		1.0		2.50	pukatea		-	29.59	9950
									0.04	73000		
	1.	0.0	1.0		0.0		1.25	rimu	0.28	67667	727.8	33418
	1.	0.0	0.5		1.0	(	0.00	tawa		-	39.72	2544
									0.06	17333		

# Height

trad_init	_covemoth	er_frad_s	pread_lo <b>tra</b> ld_	_spread_	_lotnæd_gro	wtspecies	hgt_diff	hgt_per
(	0.0	.0	0.0	0.0	0.00	kanuka	0.000000	NaN
(	0.0	.0	0.0	0.0	0.00	kawakav	v <b>a</b> 0.000000	NaN
(	0.5	.5	0.5	0.5	0.00	mahoe	-	100.00000
							4.287933	
(	0.5	.0	0.5	0.5	1.25	pigeon	-	40.07619
							2.601800	
(	0.5	.0	1.0	0.0	0.00	ponga	-	38.50556
							1.116867	
(	0.0	.0	0.5	1.0	2.50	pukatea	-	30.48589
							2.767367	
(	0.0	.0	0.0	1.0	1.25	rimu	6.348333	676.31392
]	.0 0.	.0	0.5	1.0	0.00	tawa	-	37.83667
							3.352833	

# Age

trad_init_cove	mother_fr	ad_spread_lo <b>tra</b> ld_	_spread_	_l <b>ora</b> gdgrov	vtspecies	$age\_diff$	age_per
0.0	0.0	0.0	0.0	0.00	kanuka	0.00000	NaN
0.0	0.0	0.0	0.0	0.00	kawakay	va0.00000	NaN

$\operatorname{trad}_{\_}$	_initc	cov <b>em</b> other_	_frad_spread_l	otrald_spread_	_l <b>orag</b> dgrov	wtspecies	age_diff	age_per
	0.5	0.5	0.5	0.5	0.00	mahoe	-	100.00000
							41.00000	
	0.5	1.0	1.0	0.5	0.00	pigeon	66.76667	76.97925
	0.5	0.5	0.0	0.5	1.25	ponga	-	47.20057
							88.80000	
	0.5	0.5	0.0	0.0	0.00	pukatea	106.40000	42.57135
	1.0	0.0	1.0	0.0	1.25	rimu	121.03333	264.26492
	0.0	1.0	0.5	0.5	0.00	tawa	-	46.72602
							108.46667	