

Supplementary file for: Techow N.M.S.M., O’Ryan C., Robertson C.J.R. & Ryan P.G. 2016. The origins of white-chinned petrels killed by long-line fisheries off South Africa and New Zealand. *Polar Research* 35. Correspondence: Peter Ryan, Percy FitzPatrick Institute of African Ornithology, Department of Science and Technology-National Research Foundation Centre of Excellence, University of Cape Town, Rondebosch 7701, South Africa. E-mail pryan31@gmail.com

Supplementary Table S1. Variable sites in the cytochrome b fragment for the white-chinned petrel (GenBank accession numbers: EU053404-EU053425). Nucleotide positions are relative to the published white-chinned petrel sequence (Genbank Accession number U74350). Haplotypes specific to a sampling location were indicated as follows: South Georgia (SG), Marion Island (M) in the Prince Edward Islands, New Zealand regional population (NZL), Antipodes (An), Auckland Island (Au). Haplotypes found in bycatch samples were indicated by a B (wcpB18-22).

Haplotype	Nucleotide position																											Sample size ^a				
	1	1	1	1	1	1	1	1	2	2	2	2	3	3	3	3	3	3	3	3	3	4	4	4	4	4	5	5	5	6		
	0	1	3	5	6	6	7	8	0	2	8	9	0	3	3	5	7	8	9	0	5	6	6	6	8	4	4	7	9	3		
	8	7	5	8	2	4	7	7	7	5	5	1	0	3	6	1	6	4	3	0	3	2	5	9	1	0	3	6	5	7		
Atlantic/ Indian Ocean (two Csp61 fragments)																											36					
wcp1	T	C	G	C	A	A	T	G	T	A	T	T	A	G	G	A	G	C	A	T	T	T	C	A	C	T	C	A	A	G	36	
wcpSG2	T	2
wcpSG3	T	3		
wcpSG4	G	1		
wcpM5	C	1		
wcpM6	A	1		
wcpB20	A	1		
wcpB21	G	1		
wcpB22	A	1		
New Zealand Islands (three Csp61 fragments, apart from wcpB19)																												22				
wcpNZL9	C	G	.	C	.	.	T	.	.	.	G	22			
wcpNZL7	C	G	.	C	.	.	T	13			
wcpNZL8	C	G	.	.	.	G	T	5			
wcpNZL14	C	G	.	.	.	T	4			
wcpNZL12	C	.	.	.	A	.	.	G	.	C	.	.	T	3			
wcpAn13	C	.	C	G	.	C	.	.	T	2			
wcpAu16	C	G	.	.	.	A	.	.	G	.	C	.	.	T	2			

wcpAn10 C A	. . . G	. . G	T	1
wcpAn11 C	G	C	. . T	C	. . G
wcpAu15 C	G	C	C	. . T	. . G
wcpAu17	. . . T C	G	. . . G	T
wcpB18 C	. . C	G	C	. . T
wcpB19 C C	. T	. T	1

^aThe number of individuals sharing a particular haplotype.