Biochemical and molecular biomarkers and their association with anthropogenic chemicals in wintering Manx shearwaters (*Puffinus puffinus*)

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Supplementary files

Table S1

Primers used for qPCR biomarkers genes of Manx shearwaters (*Puffinus puffinus*) listed in Table 1.

Gene identification and symbol	Primer sequence	Tm*	%GC**
	F: GTCCGAATGATGAAGACAGGTGAGAGTG	60	50
Aryl hydrocarbon receptor (AhR)	R: GGTCTTTGCGTGGCAATGATGTAATCC	60	48.1
Cutochromo DAEO 1A E (CVD1AE)	F: ACAGTTTGGTGATGTGGCTGCTG	59.9	52.2
Cytochrome P450 1A-5 (CYP1A5) —	R: TCTGTAGGAAATGGAGGAATCGCTTGTTG	59.9	44.8
Estrogon recentor sinks 1 (ECD1)	F: GCAGAGGTGGGCGAATGATGAAAC	60.2	54.2
Estrogen receptor alpha 1 (ESR1)	R: ACAGGGCTGGACTGTTCTTCTTGTTATG	60.1	46.2
	F: GTACGGCTTGTGAGAGGGCAAAG	59.9	56.5
Heat shock protein 70 kDa (HSP70) —	R: CGGGCACGAGTAATGGAAGTGTAGAA	59.9	50
	F: CCACAGAATGATCTTCTAGCTCACCCTAAG	59.3	46.7
UDP-glucuronosyl-transferase (UGT1)	R: GCGTTGTCCATCTGGTCTCCAAATAAAG	59.3	46.4

* Tm = melting temperature, ** %GC = GC content.

Matrix of association among variables related to Manx shearwaters (*Puffinus puffinus*) stranded along the southern and southeastern coast of Brazil from 2016 to 2020. Linear regression was used to evaluate association between two quantitative variables (Day of the semester and Body mass). Kruskal-Wallis tests were (H) used to evaluate association between quantitative and categorical variables. Chi-Square tests and Cramer's V statistic (V) were used to evaluate association between two categorical variables.

Variable	Body mass	Region	Status	Sex	Age class	Gastrointestinal parasites	Kidney parasites	Plastics
	R = -0.263	H = 5.634	H = 1.257	H = 0.367	H = 0.124	H = 0.124	H = 2.256	H = 3.555
Day of the semester	<i>p</i> < 0.001***	df = 2	df = 1	df = 1	df = 1	df = 1	df = 1	df = 1
	p < 0.001	<i>p</i> = 0.060	<i>p</i> = 0.262	<i>p</i> = 0.545	<i>p</i> = 0.211	<i>p</i> = 0.725	<i>p</i> = 0.133	<i>p</i> = 0.059
		H = 2.001	H = 16.275	H = 1.757	H = 4.094	H = 0.241	H = 3.967	H = 0.033
Body mass		df = 2	df = 1	df = 1	df = 1	df = 1	df = 1	df = 1
		<i>p</i> = 0.368	<i>p</i> < 0.001	<i>p</i> = 0.185	<i>p</i> = 0.043*	<i>p</i> = 0.624	<i>p</i> = 0.046*	<i>p</i> = 0.857
Desien			V = 0.261	V = 0.035	V = 0.186	V = 0.316	V = 0.129	V = 0.212
Region			<i>p</i> = 0.005**	<i>p</i> = 0.909	<i>p</i> = 0.068	p < 0.001*	p = 0.275	<i>p</i> = 0.031*
Status				V < 0.001	V = 0.028	V = 0.090	V = 0.010	V = 0.098
Status				p > 0.999	p = 0.731	<i>p</i> = 0.264	<i>p</i> = 0.903	<i>p</i> = 0.223
Cov					V = 0.092	V = 0.013	V = 0.029	V = 0.126
Sex					p = 0.253	<i>p</i> = 0.870	p = 0.718	<i>p</i> = 0.116
						V = 0.195	V = 0.135	V = 0.264
Age class						p = 0.015*	<i>p</i> = 0.092	$p = 0.001^*$
Gastrointestinal parasites							V = 0.031	V = 0.222
							<i>p</i> = 0.698	<i>p</i> = 0.006**
Kidnov parasitas								V < 0.001
Kidney parasites								p > 0.999

Significance codes: *** *p* < 0.001, ** *p* < 0.01, * *p* < 0.05.

Summary of General Additive Models (GAM) for the liver enzymatic activity of glutathione S-transferases (GST) and ethoxy-resorufin O-deethylase (EROD) in Manx shearwaters (*Puffinus puffinus*) stranded along the southern and southeastern coast of Brazil. Coefficient estimates are provided for qualitative and non-smoothed quantitative variables. P-values (*p*) and significance codes are provided for all variables. The percentage of deviance explained are provided for each model.

		GST		EROD			
	Estimate	р	Signif.	Estimate	р	Signif.	
Non-smoothed variables							
Region ("PR")	-0.076	0.325		0.046	0.813		
Region ("SP-RJ")	-0.001	0.989		0.124	0.449		
Status ("Beach carcass")	0.041	0.529		-0.283	0.092	(*)	
Sex ("Female")	0.032	0.527		0.050	0.707		
Age class ("Adult")	-0.098	0.117		0.070	0.667		
Gastrointestinal parasites ("True")	0.033	0.576		-0.077	0.619		
Kidney parasites ("True")	0.028	0.597		0.151	0.271		
Plastics ("True")	-0.013	0.834		-0.200	0.216		
Year (integer)	0.052	0.046	*	0.242	0.001	***	
Smoothed variables							
Day of the semester	1.505	0.196		0.802	0.001	**	
Body mass	0.226	0.231		0.392	0.103		
PAHs	0.260	0.206		<0.001	0.918		
PCBs	0.483	0.149		<0.001	0.710		
НСВ	<0.001	0.623		0.604	0.083	(*)	
Drins	0.786	0.037	*	< 0.001	0.410		
Mirex	1.898	0.027	*	<0001	0.627		
Intercept	-105.320	0.044	*	-486.023	0.001	***	
Deviance explained		14.0%	•		15.7%		

Significance codes: *** *p* < 0.001, ** *p* < 0.01, * *p* < 0.05, (*) *p* < 0.1.

Summary of General Additive Models (GAM) for the liver immunodetection level of the cytochrome P450 1A proteins (CYP1A) and the transcription levels of the *cytochrome P450 1A-5 (CYP1A5)* and *aryl hydrocarbon receptor (AhR)* genes in Manx shearwaters (*Puffinus puffinus*) stranded along the southern and southeastern coast of Brazil. Coefficient estimates are provided for qualitative and non-smoothed quantitative variables. P-values (*p*) and significance codes are provided for all variables. The percentage of deviance explained are provided for each model.

	CYP1A			CYP1A5			AhR		
	Estimate	р	Signif.	Estimate	р	Signif.	Estimate	р	Signif.
Non-smoothed variables									
Region ("PR")	0.022	0.026	*	-0.497	0.021	*	-0.472	0.035	*
Region ("SP-RJ")	0.016	0.051	(*)	-0.348	0.054	(*)	-0.075	0.681	
Status ("Beach carcass")	0.009	0.312		0.308	0.096	(*)	-0.274	0.156	
Sex ("Female")	0.009	0.157		-0.196	0.172		0.064	0.665	
Age class ("Adult")	-0.002	0.800		-0.281	0.111		-0.217	0.229	
Gastrointestinal parasites ("True")	0.001	0.845		-0.108	0.519		-0.092	0.597	
Kidney parasites ("True")	0.007	0.319		-0.018	0.904		-0.293	0.056	(*)
Plastics ("True")	0.005	0.536		-0.111	0.523		0.296	0.099	(*)
Year (integer)	-0.003	0.351		-0.042	0.578		0.073	0.356	
Smoothed variables							· · · · · ·		
Day of the semester		0.011	*		0.001	**		0.198	
Body mass		0.151			0.136			0.124	
PAHs		0.310			0.149			0.082	(*)
PCBs		0.982			0.034	*		0.987	
НСВ		0.243			0.001	**		0.257	
Drins		0.086	(*)		0.402			0.545	
Mirex		0.038	*		0.060	(*)		0.995	
Intercept	6.433	0.350		91.638	0.550		-141.750	0.377	
Deviance explained		18.2%			22.2%			17.4%	

Significance codes: *** *p* < 0.001, ** *p* < 0.01, * *p* < 0.05, (*) *p* < 0.1.

Summary of General Additive Models (GAM) for the liver transcription levels of the *estrogen receptor alpha 1* (*ESR1*), *heat shock protein 70* (*HSP70*) and *UDP-glucuronosyl-transferase* (*UGT1*) genes in Manx shearwaters (*Puffinus puffinus*) stranded along the southern and southeastern coast of Brazil. Coefficient estimates are provided for qualitative and non-smoothed quantitative variables. P-values (*p*) and significance codes are provided for all variables. The percentage of deviance explained are provided for each model.

	ESR1			HSP70			UGT1		
	Estimate	р	Signif.	Estimate	р	Signif.	Estimate	р	Signif.
Non-smoothed variables									
Region ("PR")	-0.295	0.156		-0.169	0.643		-0.730	<0.001	***
Region ("SP-RJ")	-0.212	0.221		-0.147	0.627		-0.507	0.001	***
Status ("Beach carcass")	-0.012	0.945		0.239	0.453		0.224	0.147	
Sex ("Female")	-0.109	0.435		-0.324	0.184		-0.088	0.467	
Age class ("Adult")	-0.142	0.405		0.366	0.217		-0.375	0.012	*
Gastrointestinal parasites ("True")	-0.337	0.041	*	0.103	0.714		-0.225	0.114	
Kidney parasites ("True")	-0.317	0.031	*	0.032	0.899		-0.306	0.015	*
Plastics ("True")	0.089	0.599		-0.061	0.831		-0.189	0.193	
Year (integer)	-0.009	0.904		0.356	0.004	**	0.147	0.022	*
Smoothed variables							· · · ·		
Day of the semester		0.237			0.124			0.674	
Body mass		0.108			0.002	**		0.084	(*)
PAHs		0.831			0.157			0.953	
PCBs		0.225			0.156			0.003	**
НСВ		0.048	*		0.944			0.001	***
Drins		0.404			0.502			0.517	
Mirex		0.037	*		0.876			0.669	
Intercept	21.245	0.885		-714.771	0.004	**	-289.254	0.025	*
Deviance explained		14.4%			36.7%			25.7%	

Significance codes: *** *p* < 0.001, ** *p* < 0.01, * *p* < 0.05, (*) *p* < 0.1.

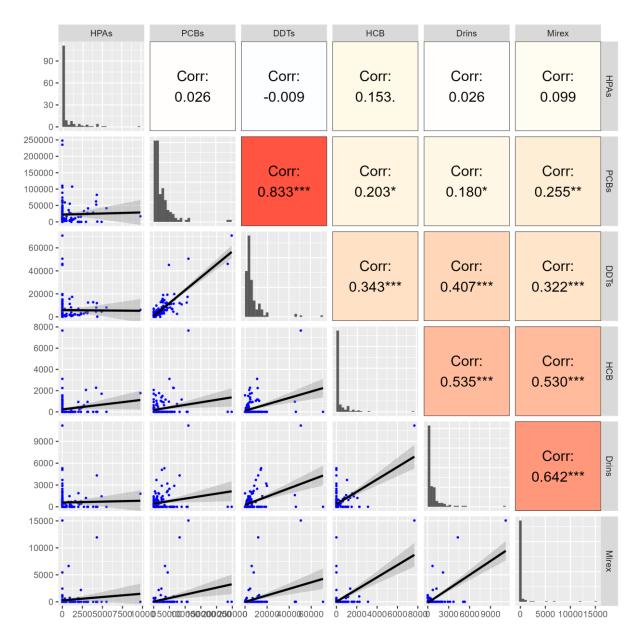


Fig. S1. Correlogram among pollutant concentrations in Manx shearwaters (*Puffinus puffinus*) stranded along the southern and southeastern coast of Brazil. Diagonal plots represent the histogram of each variable. Bottom left plots represent scatter plots of each pair of variables (blue dots) as well as the corresponding linear regression (black line) and its 95% confidence interval (gray shaded area). Top right squares represent the Spearman's correlation coefficient, with background color indicating the strength of the correlation (white = 0, red = \pm 1). Significance of the correlations: * *p* < 0.05, ** *p* < 0.01, *** *p* < 0.001.

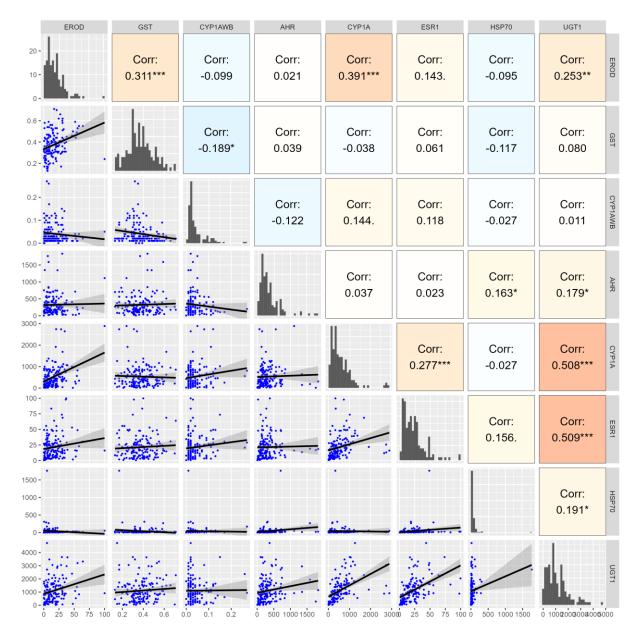


Fig. S2. Correlogram among biomarker results in Manx shearwaters (*Puffinus puffinus*) stranded along the southern and southeastern coast of Brazil. Diagonal plots represent the histogram of each variable. Bottom left plots represent scatter plots of each pair of variables (blue dots) as well as the corresponding linear regression (black line) and its 95% confidence interval (gray shaded area). Top right squares represent the Spearman's correlation coefficient, with background color indicating the strength of the correlation (white = 0, red = \pm 1). Significance of the correlations: * *p* < 0.05, ** *p* < 0.01, *** *p* < 0.001.