

Biomathematics and Statistics Scotland

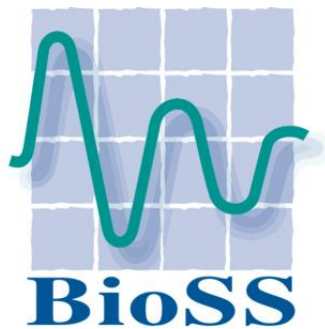
**Additional Work Coquet Colony – Seabird
Tracking Data (Under Agreement C10-0206-0387)**

CONTRACT No: C10-0206-0387

Report submitted to:

Joint Nature Conservation Committee

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In addition to this report, there are ancillary files associated with this project:

(i) Spreadsheet files of grid predictions for the Coquet Colony

5.

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1. Introduction

The Joint Nature Conservation Committee (JNCC) is working on the identification of important marine areas around the UK that are used by five species of tern during the breeding season. For the four larger tern species (Arctic, common, roseate and Sandwich terns), visual tracking data are available from boat surveys.

A Phase I project completed by BioSS earlier in 2012 used these data to learn about important associations between terns usage/preference and environmental covariates, and to predict usage/preference. During subsequent Phase II work it became apparent that for some environmental covariates there were some issues with outliers and skewed distributions. BioSS was therefore asked to use the methodology developed from the Phase I project to re-run the models for the Coquet colony after removing outliers and transforming variables where necessary.

2. Methods

We followed the methodology described in the report for Phase I (Brewer et al., 2012a) with modifications described below.

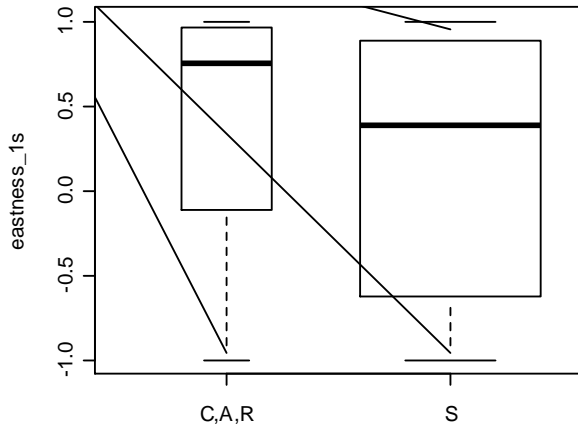
The boxplots in the Phase II report (Brewer et al., 2012b) show a negatively skewed distribution for sea surface temperature, with low values occurring near the shore. The extent to which these data are reliable is uncertain. Removal of values considered unreliable would have resulted in considerable loss of data, resulting in a possible bias of the predictive model when applied to grid cells close to shore; for this reason, sea surface temperature (SST) was excluded from the Phase II analysis. For the current work we fitted models both excluding data points with extreme values of SST (for April this is interpreted as less than 6°C, for May as less than 8°C, and for June as less than 9.3°C) and models that excluded the SST variables entirely.

Chlorophyll concentrations and wave and current shear stresses had positively skewed distributions and were therefore log-transformed as in the Phase II report. The Phase II report did not include boxplots for eastness, northness, slope, `sum_front_sd`, or `spring_front_sd`, as the decision had been taken to remove them after consideration of biological realism. For consistency with the Phase I report, these are presented as part of this report. The boxplots show that slope has a skewed distribution and hence this was also log-transformed prior to analysis. Where necessary, a small constant was added to variables prior to log transformation to avoid taking logs of zero values.

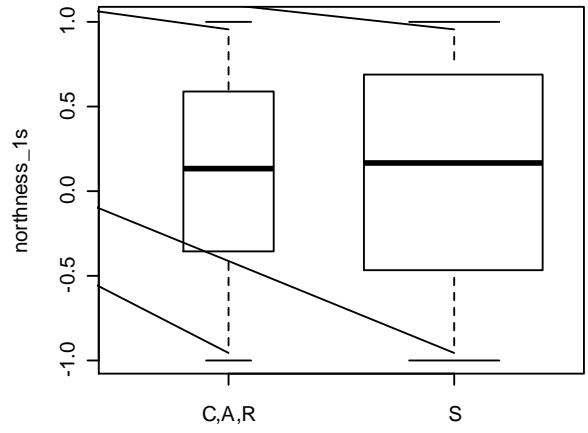
It was apparent that a model selected for roseate terns (including SST with outliers removed) by the stepwise AIC procedure did not in fact have minimum AIC, as the model selected using LRT had a lower AIC. This is to be expected, as the stepwise procedure is never guaranteed to find an optimal model; note that running an exhaustive search would be prohibitive given the size of the data set and the number of variables, and at least in a predictive sense the model is still likely to be a “good” model, if not the “best” (a concept whose legitimacy is doubtful in any case). Instead, some additional code has now been written which starts with a model having the distance to colony variable as sole explanatory variable; the original code for performing stepwise selection started with all the potential covariates in the model. Where the models selected by the two methods are the same, we can be reasonably confident that we have found a minimum AIC model. However, where they differ, the two models should be examined to see which one is better. Code was also added for calculating correlations between variables in the selected model and variance inflation factors (VIFs).

It was suggested by JNCC that grid cells within 500m of the colony should be removed. However, as there were four grid cells at distances of approximately 500m (489, 494, 507 and 512 metres, respectively), a cut-off of 500m would be rather arbitrary and all four cells were therefore removed.

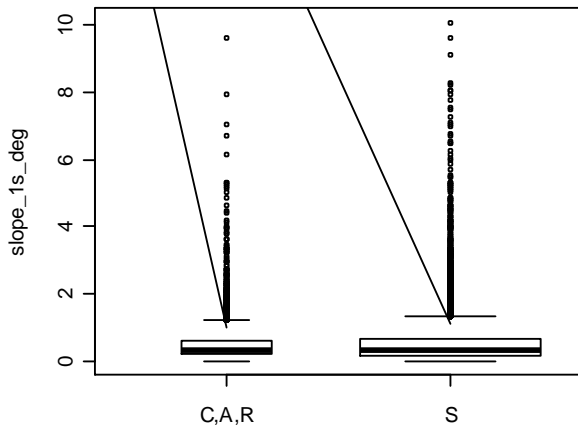
Coquet: eastness_1s



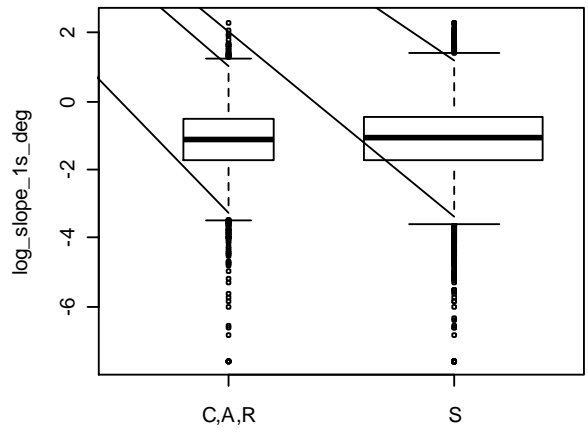
Coquet: northness_1s



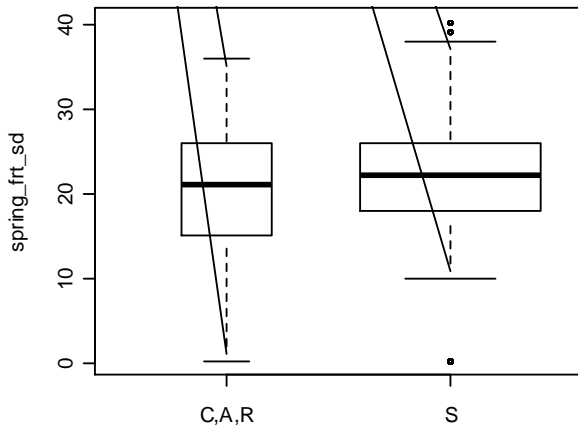
Coquet: slope_1s_deg



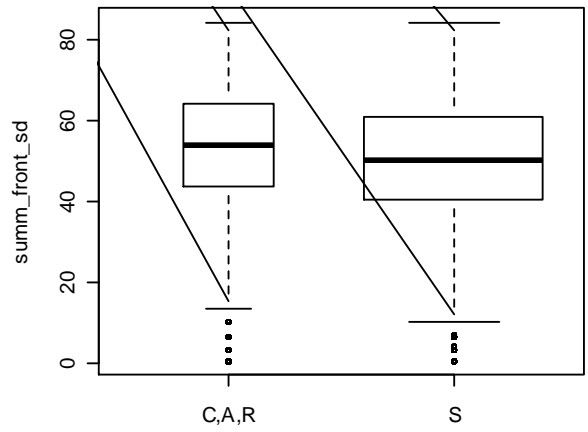
Coquet: log_slope_1s_deg



Coquet: spring_ft_sd



Coquet: summ_front_sd



3. Results

The results are presented in the Appendix.

For Arctic terns, models both with and without SST involve `dist_col`, `chl_june` and `sum_front_sd`, `bathy_1sec`. The models including SST have two additional variables: `sst_may` and `ss_current`. Fitted values from the two models are very similar, so we use the model without SST for prediction, as this allows us to make predictions for all grid points.

For common terns the SST variables are not selected in the LRT or GAM models. We therefore base predictions on a model that excludes SST and use the GAM model which has a nonlinear term in `dist_col` and linear terms in `chl_june` and `bathy_1sec`.

For roseate terns, the model excluding SST is more parsimonious, involving just three variables (`dist_col`, `chl_june` and `sst_may`) instead of seven, some of which have large VIFs.

For Sandwich terns SST variables were not selected in the AIC, BIC, LRT or GAM models; we did not therefore consider models involving SST any further. The GAM selection procedure selects a model involving `dist_col`, `chl_may` and `bathy_1sec`. There is a very high correlation (0.96) between `ss_wave` and `bathy_1sec`, resulting in VIFs of 7.1 and 9.5, respectively, but this should not be an issue if LRT or GAM models are used for prediction as `ss_wave` is not selected.

References

Brewer M.J., Potts J.M., Duff E.I. & Elston D.A. (2012a). To carry out tern modelling under the Framework Agreement C10-0206-0387. Report submitted to: Joint Nature Conservation Committee.

Brewer M.J., Potts J.M., Duff E.I. & Elston D.A. (2012b). Prediction of New Colonies – Seabird Tracking Data (Under Agreement C10-0206-0387). Report submitted to: Joint Nature Conservation Committee.

Appendix

Arctic terns – SST included with outliers removed

AIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + chl_june + sst_may +  
     summ_front_sd + ss_current + bathy_1sec, family = "binomial",  
     data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.96827	-0.03506	-0.01563	-0.00825	1.96597

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-16.18994	7.11271	-2.276	0.02283	*
dist_col	-0.22366	0.04388	-5.097	3.45e-07	***
chl_june	1.77415	0.70781	2.507	0.01219	*
sst_may	1.33067	0.75854	1.754	0.07939	.
summ_front_sd	-0.02412	0.01366	-1.765	0.07750	.
ss_current	0.74834	0.42481	1.762	0.07814	.
bathy_1sec	-0.05801	0.02218	-2.616	0.00891	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 524.57 on 47322 degrees of freedom
Residual deviance: 401.17 on 47316 degrees of freedom
AIC: 57.086

Number of Fisher Scoring iterations: 8

BIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col, family = "binomial",  
     data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.79847	-0.03649	-0.01705	-0.00697	1.92602

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.88128	0.18899	-4.663	3.11e-06 ***
dist_col	-0.21249	0.02855	-7.443	9.86e-14 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 524.57 on 47322 degrees of freedom
Residual deviance: 414.38 on 47321 degrees of freedom
AIC: 49.348

Number of Fisher Scoring iterations: 8

LRT Selected Model:

Call:

```
glm(formula = formula.glm, family = "binomial", data =  
complete.data.to.analyse,  
weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.96827	-0.03506	-0.01563	-0.00825	1.96597

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-16.18994	7.11271	-2.276	0.02283	*
dist_col	-0.22366	0.04388	-5.097	3.45e-07	***
chl_june	1.77415	0.70781	2.507	0.01219	*
sst_may	1.33067	0.75854	1.754	0.07939	.
summ_front_sd	-0.02412	0.01366	-1.765	0.07750	.
ss_current	0.74834	0.42481	1.762	0.07814	.
bathy_1sec	-0.05801	0.02218	-2.616	0.00891	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 524.57 on 47322 degrees of freedom

Residual deviance: 401.17 on 47316 degrees of freedom

AIC: 57.086

Number of Fisher Scoring iterations: 8

Single term deletions

Model:

```
SEARCH_FORAGE ~ dist_col + chl_june + sst_may + summ_front_sd +  
  ss_current + bathy_lsec
```

	Df	Deviance	AIC	LRT	Pr(>Chi)	
<none>		401.17	57.086			
dist_col	1	450.62	104.536	49.450	2.034e-12	***
chl_june	1	407.32	61.234	6.148	0.013156	*
sst_may	1	404.32	58.235	3.149	0.075975	.
summ_front_sd	1	404.47	58.387	3.301	0.069249	.
ss_current	1	404.30	58.214	3.128	0.076969	.
bathy_lsec	1	408.56	62.477	7.391	0.006556	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Now checking for interactions with Year.

Single term deletions

Model:

```
SEARCH_FORAGE ~ Year * (dist_col + chl_june + sst_may +  
  summ_front_sd +  
  ss_current + bathy_lsec)
```

	Df	Deviance	AIC	LRT	Pr(>Chi)
--	----	----------	-----	-----	----------

<none>		390.56	84.201		
Year:dist_col	2	393.09	82.731	2.53009	0.2822
Year:chl_june	2	390.78	80.419	0.21754	0.8969
Year:sst_may	2	391.32	80.955	0.75403	0.6859
Year:summ_front_sd	2	391.23	80.868	0.66693	0.7164
Year:ss_current	2	390.80	80.434	0.23329	0.8899
Year:bathy_lsec	2	391.30	80.941	0.73940	0.6909

No significant Year interactions.

Running GAM.

GAM Model selected (ML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_june, k = 3) + s(sst_may,
  k = 3) + s(summ_front_sd, k = 3) + s(ss_current, k = 3) +
  s(bathy_lsec, k = 3)
```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-4.0518	0.3258	-12.44	<2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value
--	-----	--------	--------	---------

```

s(dist_col)      1      1 25.982 3.45e-07 ***
s(chl_june)     1      1  6.283 0.01219 *
s(sst_may)      1      1  3.077 0.07939 .
s(summ_front_sd) 1      1  3.117 0.07750 .
s(ss_current)   1      1  3.103 0.07814 .
s(bathy_1sec)   1      1  6.841 0.00891 **

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq. (adj) = 0.232 Deviance explained = 23.5%

ML score = 200.59 Scale est. = 1 n = 47323

GAM Model selected (REML output):

Family: binomial

Link function: logit

Formula:

```

SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_june, k = 3) + s(sst_may,
  k = 3) + s(summ_front_sd, k = 3) + s(ss_current, k = 3) +
  s(bathy_1sec, k = 3)

```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-3.9105	0.3054	-12.8	<2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(dist_col)	1.625	1.859	32.810	5.87e-08	***
s(chl_june)	1.000	1.000	6.321	0.0119	*
s(sst_may)	1.000	1.000	2.701	0.1003	
s(summ_front_sd)	1.000	1.000	2.033	0.1539	
s(ss_current)	1.000	1.000	2.159	0.1417	
s(bathy_1sec)	1.000	1.000	7.055	0.0079	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq. (adj) = 0.235 Deviance explained = 24.1%

REML score = 204.51 Scale est. = 1 n = 47323

Running INLA.

Call:

```
c("inla(formula = formula.inla, family = \"binomial\", data =
complete.data.to.analyse, ", " weights = weights, verbose =
TRUE)")
```

Time used:

Pre-processing	Running inla	Post-processing	Total
8.252414	1014.204581	441.823976	1464.280971

Fixed effects:

	mean	sd	0.025quant	0.5quant	0.975quant
(Intercept)	-15.80343264	6.96513547	-29.71785598	-15.71790518	-2.381378967
chl_june	1.84899787	0.72981499	0.41120145	1.85139713	3.273482817

sst_may	1.28141835	0.74649658	-0.16186267	1.27373868
2.768318318				
summ_front_sd	-0.01794380	0.01365707	-0.04562755	-0.01764065
0.008024216				
ss_current	0.58411433	0.43613776	-0.26880017	0.58324510
1.442128817				
bathy_1sec	-0.05970494	0.02132887	-0.10275585	-0.05929434
0.018983481				
dist_coll	-10.55911704	2.32672743	-15.58000967	-10.39287378
6.445102056				
dist_col2	-2.12409860	3.83919303	-10.68494460	-1.71775894
4.315279905				

kld

(Intercept)	0.0068190888
chl_june	0.0000995303
sst_may	0.0063532090
summ_front_sd	0.0047399743
ss_current	0.0005071765
bathy_1sec	0.0056925455
dist_coll	0.0684144995
dist_col2	0.0092390962

Random effects:

Name	Model	Max KLD
UserFunction0	NoModelName	
UserFunction1	NoModelName	
mesh.points	SPDE model	

Model hyperparameters:

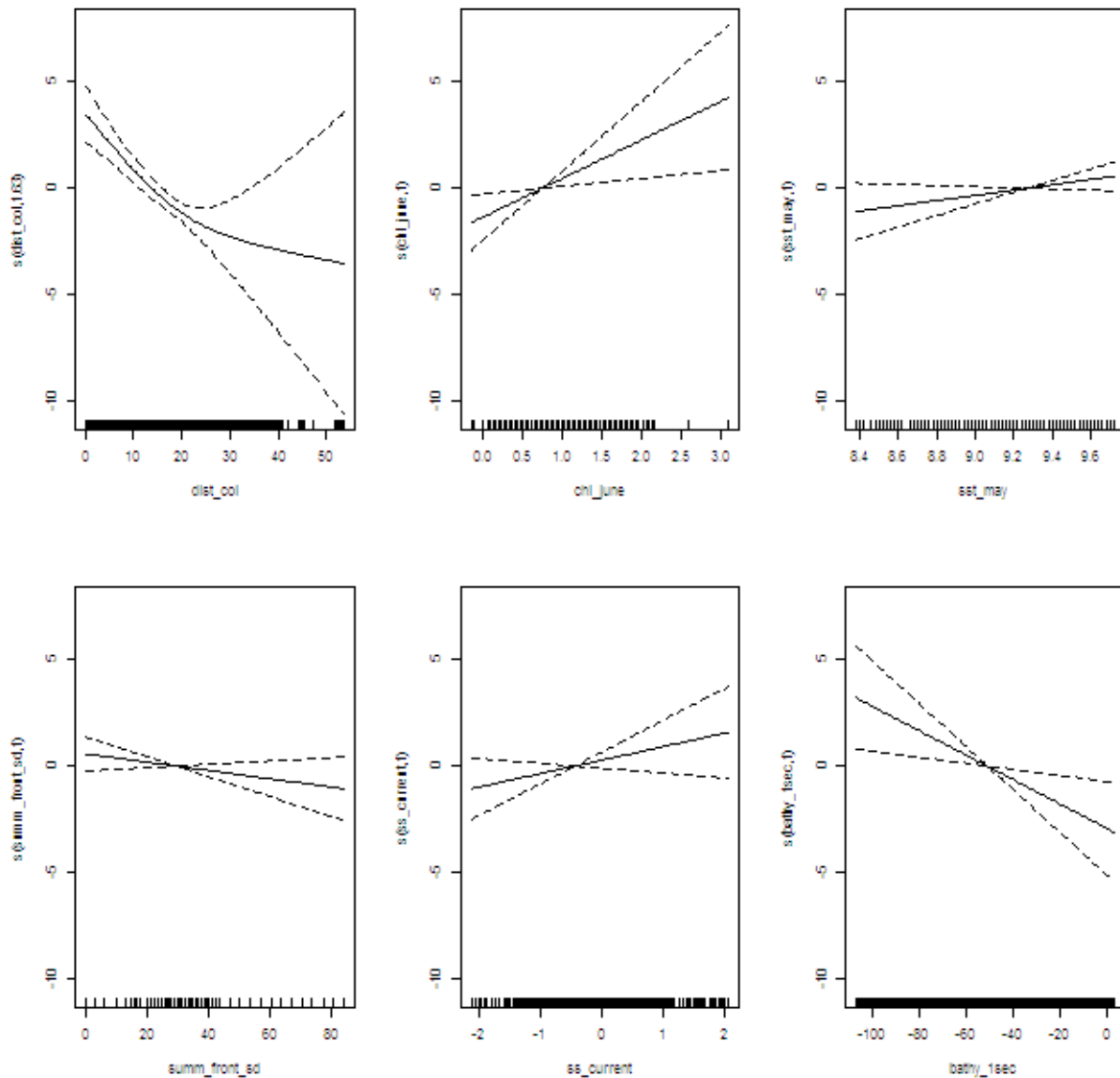
	mean	sd	0.025quant	0.5quant	0.975quant
T.0 for mesh.points-basisT	7.521	3.185	1.250	7.530	13.754

Expected number of effective parameters(std dev): 7.993(0.001666)

Number of equivalent replicates : 5920.87

Marginal Likelihood: -233.79

	mean	sd	0.025quant	0.5quant	0.975quant	kld
(Intercept)	-15.8034	6.9651	-29.7179	-15.7179	-2.3814	0.0068
chl_june	1.8490	0.7298	0.4112	1.8514	3.2735	0.0001
sst_may	1.2814	0.7465	-0.1619	1.2737	2.7683	0.0064
summ_front_sd	-0.0179	0.0137	-0.0456	-0.0176	0.0080	0.0047
ss_current	0.5841	0.4361	-0.2688	0.5832	1.4421	0.0005
bathy_1sec	-0.0597	0.0213	-0.1028	-0.0593	-0.0190	0.0057
dist_col1	-10.5591	2.3267	-15.5800	-10.3929	-6.4451	0.0684
dist_col2	-2.1241	3.8392	-10.6849	-1.7178	4.3153	0.0092



Arctic terns – SST excluded

AIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + chl_june + summ_front_sd +
     bathy_1sec, family = "binomial", data = complete.data.to.analyse,
     weights = weights)
```

Deviance Residuals:

Min 1Q Median 3Q Max

-0.91533 -0.03595 -0.01519 -0.00745 1.93331

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-3.80859	1.29761	-2.935	0.00333	**
dist_col	-0.22048	0.03941	-5.595	2.2e-08	***
chl_june	1.44940	0.68547	2.114	0.03447	*
summ_front_sd	-0.02344	0.01349	-1.738	0.08212	.
bathy_1sec	-0.05269	0.02061	-2.556	0.01058	*

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 542.78 on 47731 degrees of freedom
Residual deviance: 420.58 on 47727 degrees of freedom
AIC: 50.803

Number of Fisher Scoring iterations: 8

BIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col, family = "binomial",  
     data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.79352	-0.03705	-0.01715	-0.00702	1.93099

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-0.89725	0.18258	-4.914	8.91e-07	***
dist_col	-0.21127	0.02803	-7.537	4.81e-14	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 542.78 on 47731 degrees of freedom
Residual deviance: 429.21 on 47730 degrees of freedom
AIC: 46.903

Number of Fisher Scoring iterations: 8

LRT Selected Model:

Call:

```
glm(formula = formula.glm, family = "binomial", data =  
complete.data.to.analyse,  
weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.91533	-0.03595	-0.01519	-0.00745	1.93331

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-3.80859	1.29761	-2.935	0.00333	**
dist_col	-0.22048	0.03941	-5.595	2.2e-08	***
chl_june	1.44940	0.68547	2.114	0.03447	*

```

summ_front_sd -0.02344    0.01349   -1.738   0.08212 .
bathy_1sec    -0.05269    0.02061   -2.556   0.01058 *

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

```

Null deviance: 542.78  on 47731  degrees of freedom
Residual deviance: 420.58  on 47727  degrees of freedom
AIC: 50.803

```

Number of Fisher Scoring iterations: 8

Single term deletions

Model:

```
SEARCH_FORAGE ~ dist_col + chl_june + summ_front_sd + bathy_1sec
```

	Df	Deviance	AIC	LRT	Pr(>Chi)	
<none>		420.58	50.803			
dist_col	1	471.76	99.983	51.180	8.427e-13	***
chl_june	1	424.94	53.168	4.365	0.036688	*
summ_front_sd	1	423.79	52.014	3.210	0.073170	.
bathy_1sec	1	427.43	55.656	6.853	0.008851	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Now checking for interactions with Year.

Single term deletions

Model:

SEARCH_FORAGE ~ Year * (dist_col + chl_june + summ_front_sd +
bathy_lsec)

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		410.93	70.784		
Year:dist_col	2	414.05	69.900	3.11630	0.2105
Year:chl_june	2	411.18	67.029	0.24483	0.8848
Year:summ_front_sd	2	411.58	67.431	0.64715	0.7236
Year:bathy_lsec	2	411.61	67.464	0.68062	0.7116

No significant Year interactions.

Running GAM.

GAM Model selected (ML output):

Family: binomial

Link function: logit

Formula:

SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_june, k = 3) + s(summ_front_sd,
k = 3) + s(bathy_lsec, k = 3)

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-4.1100	0.3328	-12.35	<2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value
s(dist_col)	1	1	31.307	2.2e-08 ***

```

s(chl_june)      1      1  4.471  0.0345 *
s(summ_front_sd) 1      1  3.022  0.0821 .
s(bathy_1sec)    1      1  6.534  0.0106 *

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.202 Deviance explained = 22.5%

ML score = 210.29 Scale est. = 1 n = 47732

GAM Model selected (REML output):

Family: binomial

Link function: logit

Formula:

```

SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_june, k = 3) + s(summ_front_sd,
      k = 3) + s(bathy_1sec, k = 3)

```

Parametric coefficients:

```

      Estimate Std. Error z value Pr(>|z|)
(Intercept) -3.9027      0.2984  -13.08  <2e-16 ***

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

```

      edf Ref.df Chi.sq p-value
s(dist_col)      1.744  1.934 29.469 3.57e-07 ***
s(chl_june)      1.000  1.000  4.856  0.02756 *
s(summ_front_sd) 1.000  1.000  1.748  0.18614
s(bathy_1sec)    1.000  1.000  6.835  0.00894 **

```

Signif. codes: 0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.203 Deviance explained = 23.3%

REML score = 212.19 Scale est. = 1 n = 47732

Running INLA.

Call:

```
c("inla(formula = formula.inla, family = \"binomial\", data =
complete.data.to.analyse, \" \" weights = weights, verbose = TRUE)")
```

Time used:

Pre-processing	Running inla	Post-processing	Total
7.628414	1008.947372	79.825340	1096.401126

Fixed effects:

	mean	sd	0.025quant	0.5quant	0.975quant
(Intercept)	-3.90900218	1.31281510	-6.49076802	-3.90633291	-1.34186621
chl_june	1.58965424	0.70842570	0.19273306	1.59243788	2.97108358
summ_front_sd	-0.01658983	0.01340367	-0.04372188	-0.01630558	0.00893142
bathy_1sec	-0.05373979	0.01994945	-0.09356625	-0.05350362	-0.01527430
dist_col1	-9.68644917	1.96757797	-13.91786931	-9.55175439	-6.19284249
dist_col2	-0.86978841	3.37145595	-8.40630496	-0.50347308	4.76031320
		kld			
(Intercept)	4.286010e-04				
chl_june	1.351205e-05				
summ_front_sd	2.382156e-03				
bathy_1sec	1.230062e-03				
dist_col1	5.396084e-02				
dist_col2	3.868163e-03				

Random effects:

Name	Model	Max KLD
UserFunction0	NoModelName	
UserFunction1	NoModelName	
mesh.points	SPDE model	

Model hyperparameters:

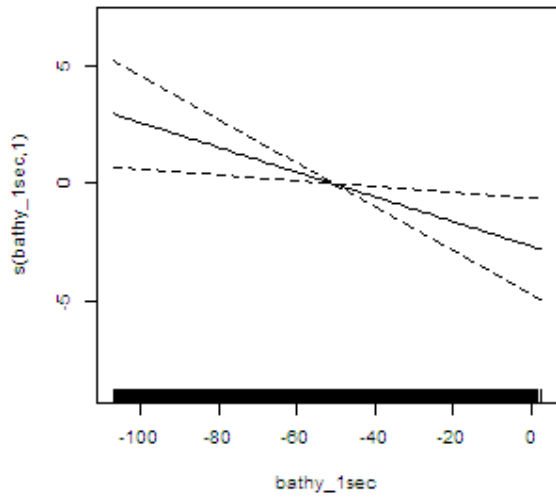
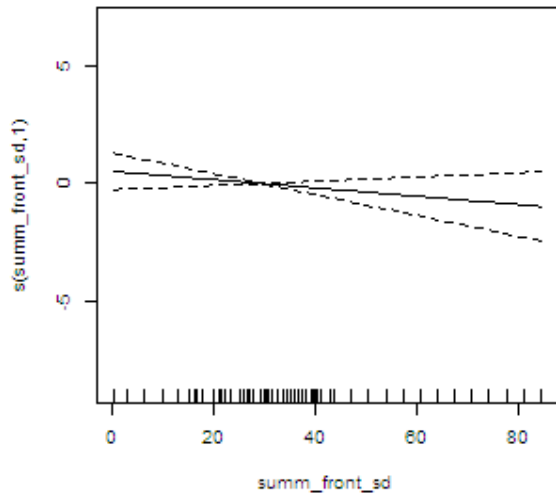
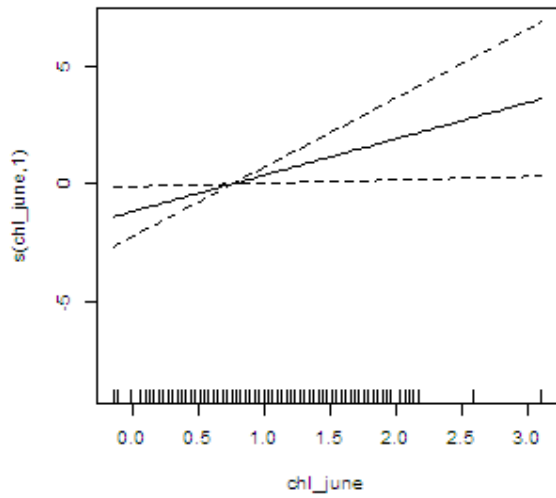
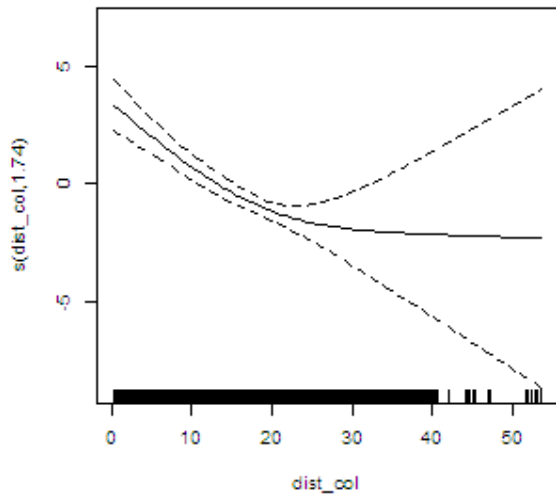
	mean	sd	0.025quant	0.5quant	0.975quant
T.0 for mesh.points-basisT	7.556	3.163	1.357	7.549	13.775

Expected number of effective parameters (std dev): 5.985 (0.0002149)

Number of equivalent replicates : 7975.61

Marginal Likelihood: -234.66

	mean	sd	0.025quant	0.5quant	0.975quant	kld
(Intercept)	-3.9090	1.3128	-6.4908	-3.9063	-1.3419	0.0004
chl_june	1.5897	0.7084	0.1927	1.5924	2.9711	0.0000
summ_front_sd	-0.0166	0.0134	-0.0437	-0.0163	0.0089	0.0024
bathy_1sec	-0.0537	0.0199	-0.0936	-0.0535	-0.0153	0.0012
dist_col1	-9.6864	1.9676	-13.9179	-9.5518	-6.1928	0.0540
dist_col2	-0.8698	3.3715	-8.4063	-0.5035	4.7603	0.0039



Common terns – SST included with outliers removed

AIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + chl_june + sst_april +
    bathy_1sec + sst_june, family = "binomial", data =
    complete.data.to.analyse,
    weights = weights)
```


Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.82985	-0.04135	-0.01208	-0.00441	2.26769

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	1.54375	14.04657	0.110	0.9125
dist_col	-0.26060	0.05551	-4.695	2.67e-06 ***
chl_june	1.96148	0.90724	2.162	0.0306 *
sst_april	1.76848	1.18084	1.498	0.1342
bathy_1sec	-0.04720	0.02424	-1.947	0.0515 .
sst_june	-1.58961	1.01222	-1.570	0.1163

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 406.92 on 29483 degrees of freedom
Residual deviance: 305.90 on 29478 degrees of freedom
AIC: 26.276

Number of Fisher Scoring iterations: 8

BIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col, family = "binomial",  
     data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.75057	-0.04414	-0.01844	-0.00799	2.25600

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-0.95103	0.21796	-4.363	1.28e-05	***
dist_col	-0.22265	0.03482	-6.395	1.61e-10	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 406.92 on 29483 degrees of freedom
Residual deviance: 322.84 on 29482 degrees of freedom
AIC: 17.005

Number of Fisher Scoring iterations: 8

LRT Selected Model:

Call:

```
glm(formula = formula.glm, family = "binomial", data =  
complete.data.to.analyse,  
weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.79572	-0.04154	-0.01434	-0.00651	2.26004

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-5.70345	1.54060	-3.702	0.000214	***
dist_col	-0.23234	0.04846	-4.795	1.63e-06	***
chl_june	2.62596	0.78605	3.341	0.000836	***
bathy_1sec	-0.05203	0.02343	-2.221	0.026370	*

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 406.92 on 29483 degrees of freedom
Residual deviance: 310.54 on 29480 degrees of freedom
AIC: 20.981

Number of Fisher Scoring iterations: 8

Single term deletions

Model:

SEARCH_FORAGE ~ dist_col + chl_june + bathy_1sec

	Df	Deviance	AIC	LRT	Pr(>Chi)	
<none>		310.54	20.981			
dist_col	1	347.84	56.278	37.296	1.015e-09	***
chl_june	1	322.20	30.635	11.653	0.0006409	***
bathy_1sec	1	315.88	24.320	5.338	0.0208620	*

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Now checking for interactions with Year.

Single term deletions

Model:

SEARCH_FORAGE ~ Year * (dist_col + chl_june + bathy_1sec)

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		303.73	35.990		
Year:dist_col	2	307.42	35.683	3.6924	0.1578
Year:chl_june	2	305.27	33.534	1.5437	0.4622
Year:bathy_1sec	2	304.32	32.589	0.5990	0.7412

No significant Year interactions.

Running GAM.

GAM Model selected (ML output):

Family: binomial

Link function: logit

Formula:

SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_june, k = 3) +
s(bathy_1sec,
k = 3)

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-4.3787	0.4312	-10.15	<2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(dist_col)	1	1	22.987	1.63e-06	***
s(chl_june)	1	1	11.160	0.000836	***
s(bathy_1sec)	1	1	4.931	0.026373	*

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq. (adj) = 0.0893 Deviance explained = 23.7%

ML score = 155.27 Scale est. = 1 n = 29484

GAM Model selected (REML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_june, k = 3) +  
s(bathy_1sec,  
  k = 3)
```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-4.8100	0.7859	-6.121	9.32e-10	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(dist_col)	1.526	1.776	11.403	0.00251	**
s(chl_june)	1.000	1.000	9.662	0.00188	**
s(bathy_1sec)	1.000	1.000	4.140	0.04189	*

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.0909 Deviance explained = 23.9%

REML score = 156.74 Scale est. = 1 n = 29484

Common terms – SST excluded

AIC Selected Model (forwards):

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + chl_may + strat_temp,
     family = "binomial", data = complete.data.to.analyse, weights =
weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.81262	-0.04578	-0.01442	-0.00435	2.27538

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	0.76896	2.25121	0.342	0.73267	
dist_col	-0.14588	0.04925	-2.962	0.00306	**
chl_may	0.51844	0.27789	1.866	0.06209	.
strat_temp	-0.93045	0.62090	-1.499	0.13399	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 430.38 on 29956 degrees of freedom
Residual deviance: 329.69 on 29953 degrees of freedom
AIC: 20.545

Number of Fisher Scoring iterations: 8

AIC Selected Model (backwards):

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + chl_apr + summ_front_sd +  
      strat_temp + sal_summ + sal_spring + ss_current + bathy_lsec,  
      family = "binomial", data = complete.data.to.analyse, weights =  
weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.85772	-0.04245	-0.01314	-0.00344	2.35404

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	2280.56304	6876.91998	0.332	0.74017
dist_col	-0.17217	0.06194	-2.780	0.00544 **
chl_apr	1.52198	0.64787	2.349	0.01881 *
summ_front_sd	-0.03109	0.01534	-2.026	0.04273 *

strat_temp	-2.28255	0.85454	-2.671	0.00756	**
sal_summ	310.64817	292.10740	1.063	0.28757	
sal_spring	-375.56977	170.74309	-2.200	0.02783	*
ss_current	-0.78831	0.40811	-1.932	0.05341	.
bathy_1sec	-0.06836	0.02577	-2.652	0.00799	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 430.38 on 29956 degrees of freedom
 Residual deviance: 319.88 on 29948 degrees of freedom
 AIC: 29.887

Number of Fisher Scoring iterations: 9

BIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col, family = "binomial",
     data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

	Min	1Q	Median	3Q	Max
	-0.75936	-0.04579	-0.01874	-0.00808	2.24320

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.92484	0.20783	-4.450	8.59e-06 ***

dist_col -0.22190 0.03366 -6.592 4.33e-11 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 430.38 on 29956 degrees of freedom
Residual deviance: 340.30 on 29955 degrees of freedom
AIC: 16.789

Number of Fisher Scoring iterations: 8

LRT Selected Model:

Call:

```
glm(formula = formula.glm, family = "binomial", data =  
complete.data.to.analyse,
```

```
weights = weights)
```

Deviance Residuals:

	Min	1Q	Median	3Q	Max
	-0.85772	-0.04245	-0.01314	-0.00344	2.35404

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	2280.56304	6876.91998	0.332	0.74017	
dist_col	-0.17217	0.06194	-2.780	0.00544	**
chl_apr	1.52198	0.64787	2.349	0.01881	*
summ_front_sd	-0.03109	0.01534	-2.026	0.04273	*

strat_temp	-2.28255	0.85454	-2.671	0.00756	**
sal_summ	310.64817	292.10740	1.063	0.28757	
sal_spring	-375.56977	170.74309	-2.200	0.02783	*
ss_current	-0.78831	0.40811	-1.932	0.05341	.
bathy_1sec	-0.06836	0.02577	-2.652	0.00799	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 430.38 on 29956 degrees of freedom
 Residual deviance: 319.88 on 29948 degrees of freedom
 AIC: 29.887

Number of Fisher Scoring iterations: 9

Single term deletions

Model:

SEARCH_FORAGE ~ dist_col + chl_apr + summ_front_sd + strat_temp +
 sal_summ + sal_spring + ss_current + bathy_1sec

	Df	Deviance	AIC	LRT	Pr(>Chi)	
<none>		319.88	29.887			
dist_col	1	330.35	38.352	10.4649	0.001217	**
chl_apr	1	325.45	33.452	5.5644	0.018329	*
summ_front_sd	1	324.26	32.265	4.3773	0.036420	*
strat_temp	1	328.55	36.549	8.6620	0.003249	**
sal_summ	1	322.66	30.664	2.7764	0.095661	.
sal_spring	1	325.95	33.956	6.0682	0.013764	*

```

ss_current      1    323.72 31.725   3.8375 0.050120 .
bathy_1sec     1    327.48 35.488   7.6006 0.005835 **

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Now checking for interactions with Year.

Single term deletions

Model:

```

SEARCH_FORAGE ~ Year * (dist_col + chl_apr + summ_front_sd +
    strat_temp + sal_summ + sal_spring + ss_current + bathy_1sec)

```

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		304.00	63.070		
Year:dist_col	2	305.45	60.520	1.4493	0.4845
Year:chl_apr	2	307.73	62.790	3.7194	0.1557
Year:summ_front_sd	2	305.77	60.834	1.7633	0.4141
Year:strat_temp	2	305.23	60.293	1.2224	0.5427
Year:sal_summ	2	304.06	59.128	0.0577	0.9716
Year:sal_spring	2	304.32	59.385	0.3142	0.8546
Year:ss_current	2	305.66	60.725	1.6548	0.4372
Year:bathy_1sec	2	307.56	62.623	3.5522	0.1693

No significant Year interactions.

Running GAM.

GAM Model selected (ML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(dist_col, k = 3) + s(summ_front_sd, k = 3) +  
s(strat_temp, k = 3) + s(bathy_1sec, k = 3)
```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-4.7091	0.5915	-7.961	1.71e-15	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(dist_col)	1	1	6.438	0.0112	*
s(summ_front_sd)	1	1	2.947	0.0860	.
s(strat_temp)	1	1	4.944	0.0262	*
s(bathy_1sec)	1	1	3.040	0.0812	.

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.154 Deviance explained = 23.5%

ML score = 164.64 Scale est. = 1 n = 29957

GAM Model selected (REML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(dist_col, k = 3) + s(summ_front_sd, k = 3) +
  s(strat_temp, k = 3) + s(bathy_1sec, k = 3)
```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-5.683	1.195	-4.756	1.98e-06 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value
s(dist_col)	1.698	1.909	3.931	0.1299
s(summ_front_sd)	1.000	1.000	3.306	0.0690 .
s(strat_temp)	1.000	1.000	3.027	0.0819 .
s(bathy_1sec)	1.000	1.000	1.977	0.1597

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.16 Deviance explained = 24%

REML score = 165.06 Scale est. = 1 n = 29957

Running INLA.

Call:

```
c("inla(formula = formula.inla, family = \"binomial\", data =
complete.data.to.analyse, \"\", \" weights = weights, verbose =
TRUE) ")
```

Time used:

Pre-processing	Running inla	Post-processing	Total
----------------	--------------	-----------------	-------

5.288409 366.195043 37.627266 409.110718

Fixed effects:

	mean	sd	0.025quant	0.5quant
0.975quant				
(Intercept)	2.88575072	2.54362974	-1.79018884	2.76958373
8.2069291341				
summ_front_sd	-0.02846866	0.01523990	-0.05919033	-0.02818933
0.0006633967				
strat_temp	-1.46885089	0.87216742	-3.29022247	-1.43056737
0.1382222306				
bathy_1sec	-0.03038028	0.02172686	-0.07360470	-0.03017156
0.0116340869				
dist_col1	-11.93446328	6.65679445	-26.73892393	-11.24937056
0.7195157446				
dist_col2	-18.66787888	12.57327954	-46.54660279	-17.41457803
2.6157828993				

kld

(Intercept)	4.410405e-04
summ_front_sd	6.402627e-04
strat_temp	2.082387e-05
bathy_1sec	9.155687e-04
dist_col1	1.193949e-03
dist_col2	2.080870e-03

Random effects:

Name	Model	Max KLD
UserFunction0	NoModelName	
UserFunction1	NoModelName	
mesh.points	SPDE model	

Model hyperparameters:

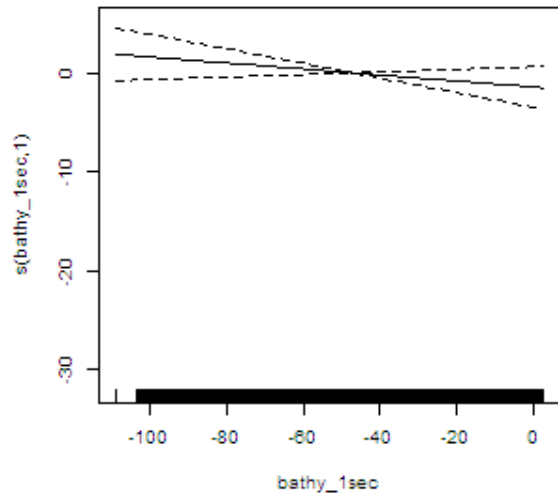
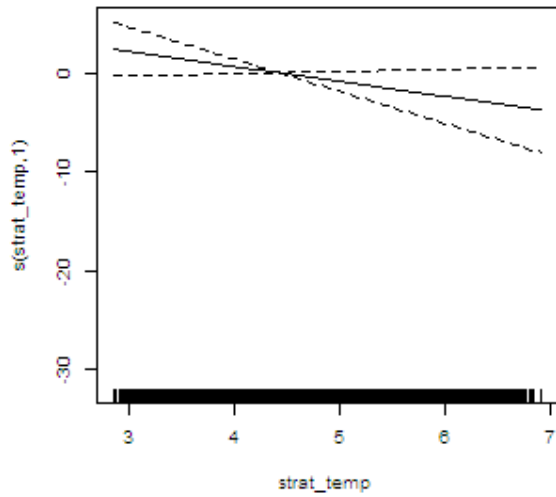
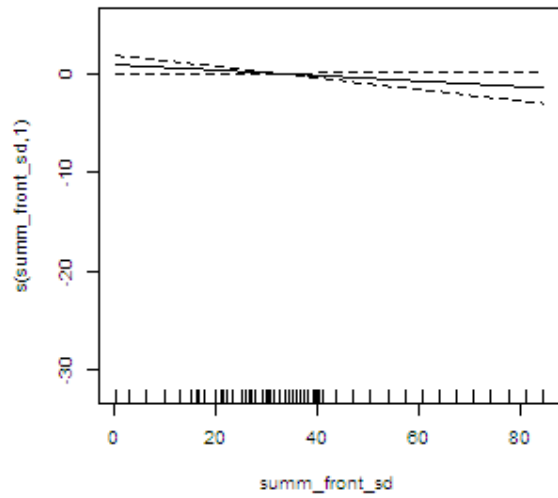
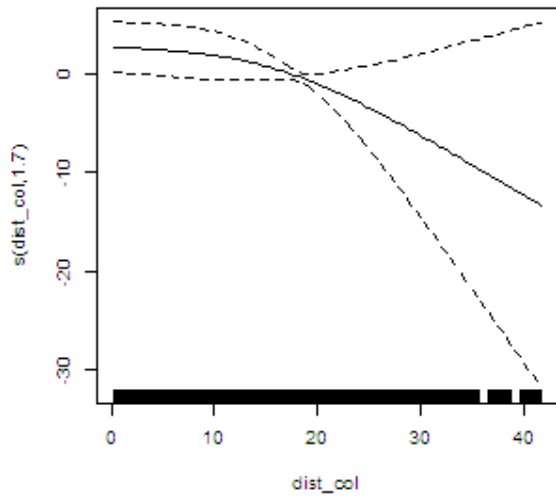
	mean	sd	0.025quant	0.5quant	0.975quant
T.0 for mesh.points-basisT	7.351	3.160	1.148	7.353	13.562

Expected number of effective parameters(std dev): 5.799(0.0001968)

Number of equivalent replicates : 5165.93

Marginal Likelihood: -188.06

	mean	sd	0.025quant	0.5quant	0.975quant	kld
(Intercept)	2.8858	2.5436	-1.7902	2.7696	8.2069	0.0004
summ_front_sd	-0.0285	0.0152	-0.0592	-0.0282	0.0007	0.0006
strat_temp	-1.4689	0.8722	-3.2902	-1.4306	0.1382	0.0000
bathy_1sec	-0.0304	0.0217	-0.0736	-0.0302	0.0116	0.0009
dist_col1	-11.9345	6.6568	-26.7389	-11.2494	-0.7195	0.0012
dist_col2	-18.6679	12.5733	-46.5466	-17.4146	2.6158	0.0021



Roseate terns – SST included with outliers removed

AIC Selected Model (forwards):

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + sst_may + chl_june,
     family = "binomial", data = complete.data.to.analyse, weights =
weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.26451	-0.01579	-0.00840	-0.00354	1.13378

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-61.4293	16.2506	-3.780	0.000157	***
dist_col	-0.1389	0.0762	-1.823	0.068265	.
sst_may	6.0750	1.7496	3.472	0.000516	***
chl_june	2.3922	0.9148	2.615	0.008922	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 152.037 on 23776 degrees of freedom
Residual deviance: 86.027 on 23773 degrees of freedom
AIC: 16.473

Number of Fisher Scoring iterations: 8

AIC Selected Model (backwards):

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + dist_shore + chl_may +  
    chl_june + sst_april + sst_may + sal_summ + sal_spring +  
    ss_wave, family = "binomial", data = complete.data.to.analyse,  
    weights = weights)
```

Deviance Residuals:

	Min	1Q	Median	3Q	Max
	-1.19098	-0.01262	-0.00396	-0.00090	1.31249

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-1189.3904	26136.1760	-0.046	0.963703
dist_col	-0.1561	0.1091	-1.432	0.152248
dist_shore	-0.9450	0.4381	-2.157	0.031021 *
chl_may	-2.1428	1.0046	-2.133	0.032928 *
chl_june	3.4515	1.8494	1.866	0.061996 .
sst_april	3.3277	2.3368	1.424	0.154433
sst_may	10.1408	2.9705	3.414	0.000641 ***
sal_summ	779.8877	1130.8640	0.690	0.490421
sal_spring	-749.1439	510.2635	-1.468	0.142063
ss_wave	-1.8782	1.0351	-1.815	0.069600 .

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 152.037 on 23776 degrees of freedom
 Residual deviance: 76.022 on 23767 degrees of freedom
 AIC: 27.365

Number of Fisher Scoring iterations: 10

BIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ chl_june + sst_may, family =
"binomial",
    data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.05757	-0.01643	-0.00858	-0.00473	1.30217

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-66.3809	16.2546	-4.084	4.43e-05	***
chl_june	3.6421	0.6319	5.764	8.23e-09	***
sst_may	6.3127	1.7134	3.684	0.000229	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 152.037 on 23776 degrees of freedom
Residual deviance: 89.986 on 23774 degrees of freedom
AIC: 15.403

Number of Fisher Scoring iterations: 8

LRT Selected Model:

Call:

```
glm(formula = formula.glm, family = "binomial", data =
complete.data.to.analyse,
```

weights = weights)

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.26451	-0.01579	-0.00840	-0.00354	1.13378

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-61.4293	16.2506	-3.780	0.000157	***
dist_col	-0.1389	0.0762	-1.823	0.068265	.
chl_june	2.3922	0.9148	2.615	0.008922	**
sst_may	6.0750	1.7496	3.472	0.000516	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 152.037 on 23776 degrees of freedom
Residual deviance: 86.027 on 23773 degrees of freedom
AIC: 16.473

Number of Fisher Scoring iterations: 8

Single term deletions

Model:

SEARCH_FORAGE ~ dist_col + chl_june + sst_may

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		86.027	16.473		

```

dist_col 1 89.986 18.432 3.9594 0.046610 *
chl_june 1 92.933 21.379 6.9063 0.008589 **
sst_may 1 102.194 30.639 16.1667 5.8e-05 ***

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Now checking for interactions with Year.

Single term deletions

Model:

```
SEARCH_FORAGE ~ Year * (dist_col + chl_june + sst_may)
```

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		77.284	31.878		
Year:dist_col	2	77.348	27.943	0.06452	0.9683
Year:chl_june	2	78.093	28.688	0.80975	0.6671
Year:sst_may	2	78.838	29.433	1.55476	0.4596

No significant Year interactions.

Running GAM.

GAM Model selected (ML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_june, k = 3) + s(sst_may,
k = 3)
```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-5.3353	0.7244	-7.366	1.76e-13	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(dist_col)	1	1	3.324	0.068290	.
s(chl_june)	1	1	6.838	0.008926	**
s(sst_may)	1	1	12.056	0.000516	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.547 Deviance explained = 43.4%

ML score = 43.014 Scale est. = 1 n = 23777

GAM Model selected (REML output):

Family: binomial

Link function: logit

Formula:

SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_june, k = 3) + s(sst_may,
k = 3)

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-5.3353	0.7244	-7.366	1.76e-13	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value
s(dist_col)	1	1	3.324	0.068293 .
s(chl_june)	1	1	6.837	0.008932 **
s(sst_may)	1	1	12.055	0.000517 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq. (adj) = 0.547 Deviance explained = 43.4%

REML score = 42.804 Scale est. = 1 n = 23777

Running INLA.

Call:

```
c("inla(formula = formula.inla, family = \"binomial\", data =
complete.data.to.analyse, \"\", \" weights = weights, verbose =
TRUE)\"")
```

Time used:

Pre-processing	Running inla	Post-processing	Total
4.009207	214.328776	28.969251	247.307234

Fixed effects:

	mean	sd	0.025quant	0.5quant
0.975quant				
(Intercept)	-58.3199202	15.24188825	-90.2301034	-57.615178
	30.30411312			
dist_col	-0.1490866	0.07567855	-0.3085743	-0.145206
	0.01101368			

chl_june 2.5220610 0.90212343 0.7957233 2.506268
4.33722865

sst_may 5.7345578 1.64441725 2.6936579 5.665215
9.15841756

kld

(Intercept) 0.0006666827

dist_col 0.0090943008

chl_june 0.0078334218

sst_may 0.0005512711

Random effects:

Name	Model	Max KLD
------	-------	---------

UserFunction0	NoModelName	
---------------	-------------	--

UserFunction1	NoModelName	
---------------	-------------	--

mesh.points	SPDE model	
-------------	------------	--

Model hyperparameters:

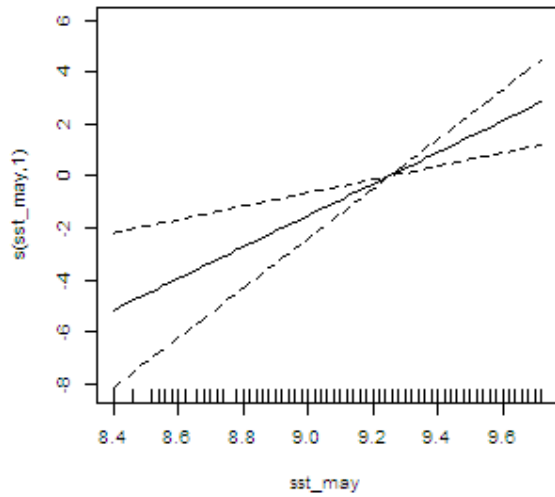
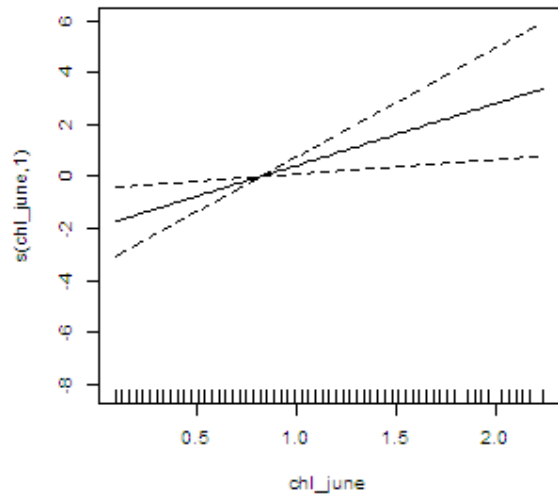
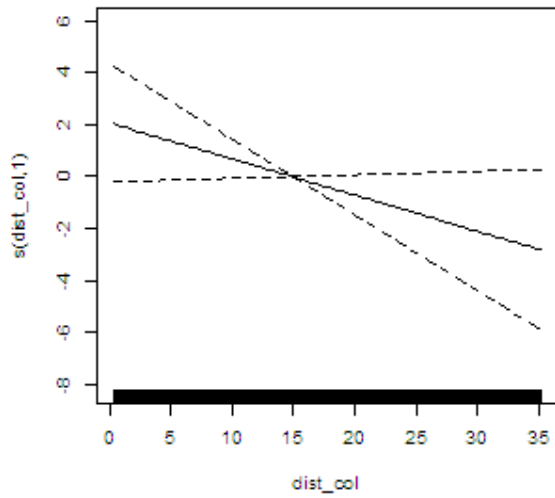
	mean	sd	0.025quant	0.5quant	0.975quant
T.0 for mesh.points-basisT	7.659	3.029	1.867	7.597	13.737

Expected number of effective parameters(std dev): 3.934(0.00264)

Number of equivalent replicates : 6043.23

Marginal Likelihood: -57.63

	mean	sd	0.025quant	0.5quant	0.975quant	kld
(Intercept)	-58.3199	15.2419	-90.2301	-57.6152	-30.3041	0.0007
dist_col	-0.1491	0.0757	-0.3086	-0.1452	-0.0110	0.0091
chl_june	2.5221	0.9021	0.7957	2.5063	4.3372	0.0078
sst_may	5.7346	1.6444	2.6937	5.6652	9.1584	0.0006



Roseate terns – SST excluded

AIC Selected Model (forwards):

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + spring_frt_sd + chl_apr +
     summ_front_sd + ss_current + dist_shore + spring_front, family =
     "binomial",
     data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.88454	-0.01414	-0.00228	-0.00007	1.46200

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-1.76959	1.69987	-1.041	0.29787
dist_col	-0.22721	0.08313	-2.733	0.00627 **
spring_frt_sd	-0.24877	0.07882	-3.156	0.00160 **
chl_apr	1.47884	0.81862	1.807	0.07084 .
summ_front_sd	0.07729	0.03733	2.071	0.03839 *
ss_current	-1.17329	0.60579	-1.937	0.05277 .
dist_shore	-0.74037	0.35381	-2.093	0.03639 *
spring_front	0.16653	0.08128	2.049	0.04048 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 207.04 on 24099 degrees of freedom
Residual deviance: 104.28 on 24092 degrees of freedom
AIC: 27.317

Number of Fisher Scoring iterations: 10

AIC Selected Model (backwards):

Call:

glm(formula = SEARCH_FORAGE ~ dist_col + dist_shore + chl_apr +

```
chl_may + chl_june + summ_front_sd + spring_front +
spring_frt_sd +
```

```
ss_current, family = "binomial", data =
complete.data.to.analyse,
```

```
weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.98022	-0.01264	-0.00213	-0.00013	1.32210

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-5.03524	3.04441	-1.654	0.0981 .
dist_col	-0.21226	0.08740	-2.429	0.0152 *
dist_shore	-0.62434	0.35334	-1.767	0.0772 .
chl_apr	2.33921	0.98792	2.368	0.0179 *
chl_may	-1.16329	0.69506	-1.674	0.0942 .
chl_june	2.45022	1.61685	1.515	0.1297
summ_front_sd	0.08477	0.03863	2.194	0.0282 *
spring_front	0.16968	0.08544	1.986	0.0470 *
spring_frt_sd	-0.20549	0.08453	-2.431	0.0151 *
ss_current	-1.41175	0.65603	-2.152	0.0314 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 207.04 on 24099 degrees of freedom
Residual deviance: 100.55 on 24090 degrees of freedom

AIC: 30.464

Number of Fisher Scoring iterations: 10

BIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + spring_frt_sd, family =  
"binomial",  
     data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

	Min	1Q	Median	3Q	Max
	-1.05505	-0.01404	-0.00571	-0.00208	1.32104

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.01991	0.33981	-0.059	0.95328
dist_col	-0.16688	0.05840	-2.858	0.00427 **
spring_frt_sd	-0.17563	0.04014	-4.376	1.21e-05 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 207.04 on 24099 degrees of freedom
Residual deviance: 123.49 on 24097 degrees of freedom
AIC: 19.175

Number of Fisher Scoring iterations: 8

LRT Selected Model:

Call:

```
glm(formula = formula.glm, family = "binomial", data =  
complete.data.to.analyse,  
weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.88454	-0.01414	-0.00228	-0.00007	1.46200

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-1.76959	1.69987	-1.041	0.29787
dist_col	-0.22721	0.08313	-2.733	0.00627 **
dist_shore	-0.74037	0.35381	-2.093	0.03639 *
chl_apr	1.47884	0.81862	1.807	0.07084 .
summ_front_sd	0.07729	0.03733	2.071	0.03839 *
spring_front	0.16653	0.08128	2.049	0.04048 *
spring_frt_sd	-0.24877	0.07882	-3.156	0.00160 **
ss_current	-1.17329	0.60579	-1.937	0.05277 .

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 207.04 on 24099 degrees of freedom

Residual deviance: 104.28 on 24092 degrees of freedom

AIC: 27.317

Number of Fisher Scoring iterations: 10

Single term deletions

Model:

SEARCH_FORAGE ~ dist_col + dist_shore + chl_apr + summ_front_sd +
spring_front + spring_frt_sd + ss_current

	Df	Deviance	AIC	LRT	Pr(>Chi)	
<none>		104.28	27.317			
dist_col	1	119.83	40.868	15.5510	8.031e-05	***
dist_shore	1	109.55	30.583	5.2658	0.0217480	*
chl_apr	1	107.64	28.676	3.3589	0.0668421	.
summ_front_sd	1	108.90	29.938	4.6204	0.0315946	*
spring_front	1	108.36	29.392	4.0741	0.0435460	*
spring_frt_sd	1	116.28	37.321	12.0038	0.0005309	***
ss_current	1	108.60	29.635	4.3176	0.0377206	*

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Now checking for interactions with Year.

Single term deletions

Model:

SEARCH_FORAGE ~ Year * (dist_col + dist_shore + chl_apr +
summ_front_sd +
spring_front + spring_frt_sd + ss_current)

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		90.621	57.791		
Year:dist_col	2	91.556	54.725	0.9349	0.6266
Year:dist_shore	2	92.143	55.312	1.5217	0.4673
Year:chl_apr	2	94.291	57.461	3.6701	0.1596
Year:summ_front_sd	2	91.370	54.539	0.7487	0.6877
Year:spring_front	2	90.625	53.795	0.0041	0.9980
Year:spring_frt_sd	2	90.663	53.832	0.0417	0.9794
Year:ss_current	2	91.068	54.237	0.4469	0.7997

No significant Year interactions.

Running GAM.

GAM Model selected (ML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(dist_col, k = 3) + s(dist_shore, k = 3) +
s(chl_apr,
  k = 3) + s(summ_front_sd, k = 3) + s(spring_front, k = 3) +
s(spring_frt_sd, k = 3) + s(ss_current, k = 3)
```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-8.973	2.066	-4.342	1.41e-05 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(dist_col)	1	1	7.470	0.00627	**
s(dist_shore)	1	1	4.377	0.03642	*
s(chl_apr)	1	1	3.263	0.07085	.
s(summ_front_sd)	1	1	4.286	0.03842	*
s(spring_front)	1	1	4.197	0.04050	*
s(spring_frt_sd)	1	1	9.960	0.00160	**
s(ss_current)	1	1	3.751	0.05278	.

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq. (adj) = 0.67 Deviance explained = 49.6%

ML score = 52.14 Scale est. = 1 n = 24100

GAM Model selected (REML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(dist_col, k = 3) + s(dist_shore, k = 3) +  
s(chl_apr,  
  k = 3) + s(summ_front_sd, k = 3) + s(spring_front, k = 3) +  
s(spring_frt_sd, k = 3) + s(ss_current, k = 3)
```

Parametric coefficients:

Estimate	Std. Error	z value	Pr(> z)
----------	------------	---------	----------

(Intercept) -9.837 2.253 -4.367 1.26e-05 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(dist_col)	1.000	1.000	7.633	0.00573	**
s(dist_shore)	1.000	1.000	5.105	0.02386	*
s(chl_apr)	1.000	1.000	4.588	0.03222	*
s(summ_front_sd)	1.000	1.000	3.995	0.04563	*
s(spring_front)	1.000	1.000	4.724	0.02974	*
s(spring_frt_sd)	1.000	1.000	8.515	0.00352	**
s(ss_current)	1.714	1.918	4.260	0.11081	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.686 Deviance explained = 50.8%

REML score = 49.085 Scale est. = 1 n = 24100

Running INLA.

Call:

```
c("inla(formula = formula.inla, family = \"binomial\", data =
complete.data.to.analyse, ", " weights = weights, verbose =
TRUE)")
```

Time used:

Pre-processing	Running inla	Post-processing	Total
4.274408	260.598457	134.519037	399.391902

Fixed effects:

	mean	sd	0.025quant	0.5quant	0.975quant
(Intercept)	-3.89146324	4.93212613	-14.58219272	-3.51669287	4.77858073
dist_col	-0.27480379	0.09203716	-0.47515565	-0.26738643	0.11388958
dist_shore	-0.94319517	0.36927706	-1.71605636	-0.92631322	0.26419945
chl_apr	2.14580048	0.90349962	0.42444768	2.12735928	3.97178222
summ_front_sd	0.08307844	0.03775422	0.01231117	0.08185252	0.16070465
spring_front	0.19264712	0.08358823	0.03141790	0.19163898	0.35954604
spring_frt_sd	-0.24121878	0.08033603	-0.40997595	-0.23731110	0.09410133
ss_current1	-0.53076514	6.66212229	-12.27682067	-1.02692769	13.88251817
ss_current2	-14.81097398	6.49787405	-28.64374683	-14.41859618	3.10700804

kld

(Intercept)	1.286369e-03
dist_col	2.146974e-02
dist_shore	3.360716e-02
chl_apr	2.002575e-02
summ_front_sd	1.647998e-02
spring_front	6.244368e-03
spring_frt_sd	4.518262e-03
ss_current1	7.561957e-07
ss_current2	4.181191e-02

Random effects:

Name	Model	Max KLD
UserFunction0	NoModelName	
UserFunction1	NoModelName	
mesh.points	SPDE model	

Model hyperparameters:

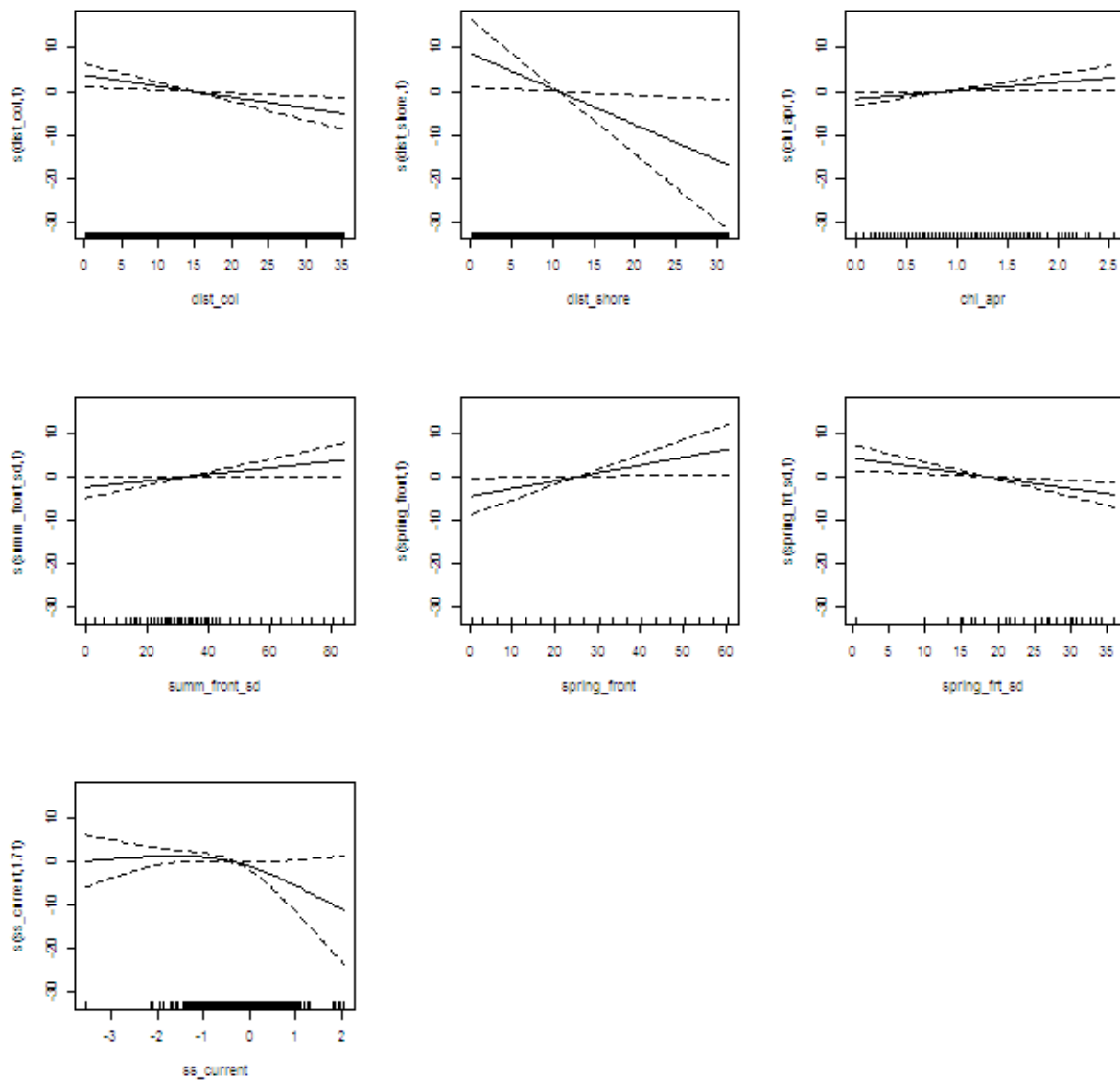
	mean	sd	0.025quant	0.5quant	0.975quant
T.0 for mesh.points-basisT	7.321	3.159	1.145	7.313	13.556

Expected number of effective parameters(std dev): 8.913(0.0002458)

Number of equivalent replicates : 2703.78

Marginal Likelihood: -88.70

	mean	sd	0.025quant	0.5quant	0.975quant	kld
(Intercept)	-3.8915	4.9321	-14.5822	-3.5167	4.7786	0.0013
dist_col	-0.2748	0.0920	-0.4752	-0.2674	-0.1139	0.0215
dist_shore	-0.9432	0.3693	-1.7161	-0.9263	-0.2642	0.0336
chl_apr	2.1458	0.9035	0.4244	2.1274	3.9718	0.0200
summ_front_sd	0.0831	0.0378	0.0123	0.0819	0.1607	0.0165
spring_front	0.1926	0.0836	0.0314	0.1916	0.3595	0.0062
spring_frt_sd	-0.2412	0.0803	-0.4100	-0.2373	-0.0941	0.0045
ss_current1	-0.5308	6.6621	-12.2768	-1.0269	13.8825	0.0000
ss_current2	-14.8110	6.4979	-28.6437	-14.4186	-3.1070	0.0418



Sandwich terns – SST included with outliers removed

AIC Selected Model (forwards):

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + dist_shore + chl_apr +
     chl_june, family = "binomial", data = complete.data.to.analyse,
     weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.11619	-0.01143	-0.00112	-0.00009	2.14536

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.13942	1.45755	-0.096	0.92380
dist_col	-0.07377	0.02565	-2.877	0.00402 **
dist_shore	-0.34391	0.14097	-2.440	0.01470 *
chl_apr	-1.46258	0.68294	-2.142	0.03223 *
chl_june	1.20165	0.56285	2.135	0.03277 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 373.27 on 55201 degrees of freedom
Residual deviance: 237.28 on 55197 degrees of freedom
AIC: 50.169

Number of Fisher Scoring iterations: 10

AIC Selected Model (backwards):

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + dist_shore + chl_apr +  
     chl_june + summ_front_sd + ss_wave, family = "binomial",  
     data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.96057	-0.00981	-0.00135	-0.00010	2.02261

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	1.67114	1.74933	0.955	0.3394
dist_col	-0.06693	0.02711	-2.468	0.0136 *
dist_shore	-0.36899	0.17130	-2.154	0.0312 *
chl_apr	-1.58321	0.72402	-2.187	0.0288 *
chl_june	1.30794	0.58894	2.221	0.0264 *
summ_front_sd	-0.03224	0.02332	-1.382	0.1669
ss_wave	-0.60335	0.38387	-1.572	0.1160

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 373.27 on 55201 degrees of freedom
Residual deviance: 233.55 on 55195 degrees of freedom
AIC: 52.258

Number of Fisher Scoring iterations: 11

BIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + dist_shore, family =  
"binomial",
```

```
data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.94821	-0.01150	-0.00123	-0.00010	2.15189

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.23169	0.30767	-0.753	0.45141
dist_col	-0.07501	0.02330	-3.220	0.00128 **
dist_shore	-0.35555	0.08826	-4.029	5.61e-05 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 373.27 on 55201 degrees of freedom
Residual deviance: 245.78 on 55199 degrees of freedom
AIC: 48.431

Number of Fisher Scoring iterations: 10

LRT Selected Model:

Call:

```
glm(formula = formula.glm, family = "binomial", data =  
complete.data.to.analyse,  
weights = weights)
```

Deviance Residuals:

	Min	1Q	Median	3Q	Max
	-1.11619	-0.01143	-0.00112	-0.00009	2.14536

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.13942	1.45755	-0.096	0.92380
dist_col	-0.07377	0.02565	-2.877	0.00402 **
dist_shore	-0.34391	0.14097	-2.440	0.01470 *
chl_apr	-1.46258	0.68294	-2.142	0.03223 *
chl_june	1.20165	0.56285	2.135	0.03277 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 373.27 on 55201 degrees of freedom
 Residual deviance: 237.28 on 55197 degrees of freedom
 AIC: 50.169

Number of Fisher Scoring iterations: 10

Single term deletions

Model:

SEARCH_FORAGE ~ dist_col + dist_shore + chl_apr + chl_june

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		237.28	50.169		
dist_col	1	250.48	61.370	13.2012	0.0002798 ***
dist_shore	1	245.09	55.981	7.8119	0.0051904 **


```
chl_apr      1    242.09 52.976  4.8068 0.0283469 *
chl_june     1    241.83 52.722  4.5525 0.0328715 *
```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Now checking for interactions with Year.

Single term deletions

Model:

```
SEARCH_FORAGE ~ Year * (dist_col + dist_shore + chl_apr + chl_june)
```

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		223.89	67.969		
Year:dist_col	2	225.15	65.231	1.2622	0.5320
Year:dist_shore	2	230.90	70.982	7.0132	0.0300 *
Year:chl_apr	2	223.96	64.040	0.0715	0.9649
Year:chl_june	2	225.26	65.340	1.3712	0.5038

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Year effects not consistent - run one year at a time.

Running GAM.

GAM Model selected (ML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(chl_apr, k = 3) + s(chl_june, k = 3) +  
s(strat_temp,  
  k = 3)
```

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-7.766	1.168	-6.651	2.92e-11	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(chl_apr)	1	1	3.759	0.05254	.
s(chl_june)	1	1	6.793	0.00915	**
s(strat_temp)	1	1	16.309	5.38e-05	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.261 Deviance explained = 35.7%

ML score = 119.96 Scale est. = 1 n = 55202

GAM Model selected (REML output):

Family: binomial

Link function: logit

Formula:

```
SEARCH_FORAGE ~ s(chl_apr, k = 3) + s(chl_june, k = 3) +  
s(strat_temp,
```

k = 3)

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-7.766	1.168	-6.65	2.93e-11	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(chl_apr)	1	1	3.756	0.05262	.
s(chl_june)	1	1	6.788	0.00918	**
s(strat_temp)	1	1	16.306	5.39e-05	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.261 Deviance explained = 35.7%

REML score = 121.18 Scale est. = 1 n = 55202

Sandwich terns – SST excluded

AIC Selected Model (forwards):

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + dist_shore + chl_may,  
     family = "binomial", data = complete.data.to.analyse, weights =  
weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.05311	-0.00859	-0.00059	-0.00003	2.22445

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-0.67811	0.66862	-1.014	0.310491	
dist_col	-0.08874	0.02378	-3.731	0.000191	***
dist_shore	-0.40978	0.09831	-4.168	3.07e-05	***
chl_may	0.36957	0.25108	1.472	0.141045	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 447.26 on 55926 degrees of freedom
Residual deviance: 272.26 on 55923 degrees of freedom
AIC: 58.191

Number of Fisher Scoring iterations: 10

AIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + dist_shore + chl_may +  
     ss_wave + bathy_1sec, family = "binomial", data =  
complete.data.to.analyse,  
     weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.16017	-0.00901	-0.00109	-0.00013	2.09111

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	1.92164	1.99426	0.964	0.3353
dist_col	-0.07300	0.02543	-2.870	0.0041 **
dist_shore	-0.23885	0.17394	-1.373	0.1697
chl_may	0.42894	0.25193	1.703	0.0886 .
ss_wave	-0.72665	0.52229	-1.391	0.1641
bathy_1sec	0.07223	0.03917	1.844	0.0652 .

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 447.26 on 55926 degrees of freedom
Residual deviance: 268.72 on 55921 degrees of freedom
AIC: 61.428

Number of Fisher Scoring iterations: 11

BIC Selected Model:

Call:

```
glm(formula = SEARCH_FORAGE ~ dist_col + dist_shore, family =  
"binomial",  
data = complete.data.to.analyse, weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-----	----	--------	----	-----

-0.93590 -0.00920 -0.00064 -0.00003 2.15522

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	0.19595	0.29277	0.669	0.503311	
dist_col	-0.08246	0.02221	-3.712	0.000206	***
dist_shore	-0.44069	0.09469	-4.654	3.26e-06	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 447.26 on 55926 degrees of freedom
Residual deviance: 274.44 on 55924 degrees of freedom
AIC: 57.509

Number of Fisher Scoring iterations: 10

LRT Selected Model:

Call:

```
glm(formula = formula.glm, family = "binomial", data =  
complete.data.to.analyse,  
weights = weights)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.93590	-0.00920	-0.00064	-0.00003	2.15522

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	0.19595	0.29277	0.669	0.503311
dist_col	-0.08246	0.02221	-3.712	0.000206 ***
dist_shore	-0.44069	0.09469	-4.654	3.26e-06 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 447.26 on 55926 degrees of freedom
Residual deviance: 274.44 on 55924 degrees of freedom
AIC: 57.509

Number of Fisher Scoring iterations: 10

Single term deletions

Model:

SEARCH_FORAGE ~ dist_col + dist_shore

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		274.44	57.509		
dist_col	1	297.06	78.134	22.624	1.970e-06 ***
dist_shore	1	315.30	96.372	40.863	1.633e-10 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Now checking for interactions with Year.

Single term deletions

Model:

SEARCH_FORAGE ~ Year * (dist_col + dist_shore)

	Df	Deviance	AIC	LRT	Pr(>Chi)
<none>		256.69	62.876		
Year:dist_col	2	257.58	59.772	0.8961	0.6388641
Year:dist_shore	2	271.50	73.684	14.8083	0.0006087 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Year effects not consistent - run one year at a time.

Running GAM.

GAM Model selected (ML output):

Family: binomial

Link function: logit

Formula:

SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_may, k = 3) +
s(bathy_1sec,
k = 3)

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-6.4775	0.6966	-9.298	<2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value	
s(dist_col)	1	1	12.382	0.000434	***
s(chl_may)	1	1	4.289	0.038356	*
s(bathy_1sec)	1	1	22.594	2e-06	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.397 Deviance explained = 39%

ML score = 136.38 Scale est. = 1 n = 55927

GAM Model selected (REML output):

Family: binomial

Link function: logit

Formula:

SEARCH_FORAGE ~ s(dist_col, k = 3) + s(chl_may, k = 3) +
s(bathy_1sec,
k = 3)

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-12.177	3.776	-3.225	0.00126	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value
--	-----	--------	--------	---------

```

s(dist_col) 1.782 1.952 6.33 0.0402 *
s(chl_may) 1.739 1.932 3.94 0.1318
s(bathy_1sec) 1.000 1.000 25.50 4.42e-07 ***

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq. (adj) = 0.392 Deviance explained = 40.6%

REML score = 136.39 Scale est. = 1 n = 55927

Running INLA.

Call:

```

c("inla(formula = formula.inla, family = \"binomial\", data =
complete.data.to.analyse, \"\", \" weights = weights, verbose =
TRUE) ")

```

Time used:

Pre-processing	Running inla	Post-processing	Total
9.500417	1320.886321	185.343926	1515.730664

Fixed effects:

	mean	sd	0.025quant	0.5quant	0.975quant
(Intercept)	-5.43840679	2.64987462	-10.94920787	-5.32945102	-0.5341445
bathy_1sec	0.06747274	0.01330845	0.04227278	0.06714279	0.0945406
dist_col1	-17.79413016	8.66266498	-36.63624746	-17.10742092	2.6315940
dist_col2	-28.58541689	16.41798545	-63.89194512	-27.45796809	0.5826683
chl_may1	10.03138044	4.89989045	0.98684934	9.81857212	20.2492269

chl_may2 2.32065696 1.10282970 0.23980301 2.29046736
4.5725689

kld

(Intercept) 0.0030535172

bathy_1sec 0.0041734944

dist_coll 0.0025193175

dist_coll2 0.0039443896

chl_may1 0.0026498437

chl_may2 0.0007893911

Random effects:

Name Model Max KLD

UserFunction0 NoModelName

UserFunction1 NoModelName

mesh.points SPDE model

Model hyperparameters:

	mean	sd	0.025quant	0.5quant	0.975quant
T.0 for mesh.points-basisT	8.013	3.160	1.763	8.033	14.162

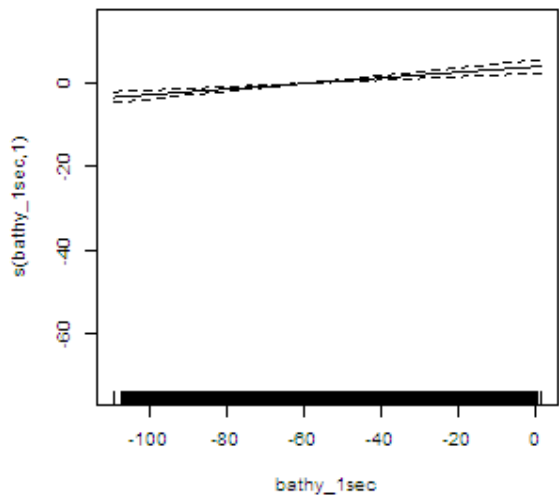
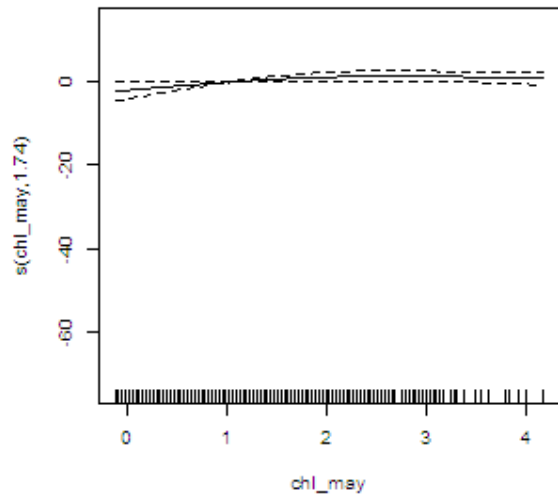
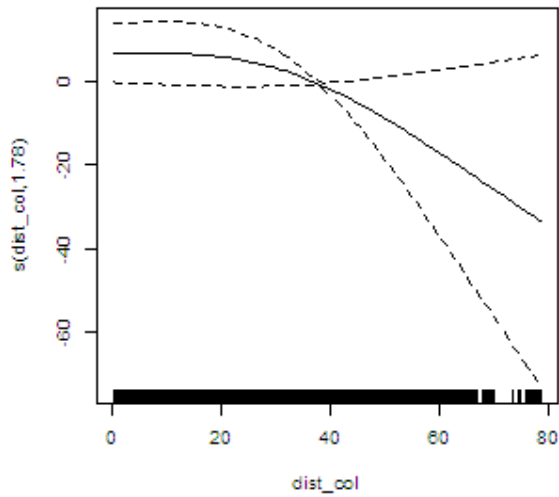
Expected number of effective parameters(std dev): 5.628(0.0004809)

Number of equivalent replicates : 9937.15

Marginal Likelihood: -152.14

	mean	sd	0.025quant	0.5quant	0.975quant	kld
(Intercept)	-5.4384	2.6499	-10.9492	-5.3295	-0.5341	0.0031
bathy_1sec	0.0675	0.0133	0.0423	0.0671	0.0945	0.0042
dist_coll	-17.7941	8.6627	-36.6362	-17.1074	-2.6316	0.0025

dist_col2	-28.5854	16.4180	-63.8919	-27.4580	0.5827	0.0039
chl_may1	10.0314	4.8999	0.9868	9.8186	20.2492	0.0026
chl_may2	2.3207	1.1028	0.2398	2.2905	4.5726	0.0008



Variance Inflation Factors

Arctic tern, AIC selected model, including SST

dist_col	chl_june	sst_may	summ_front_sd	ss_current
2.590696	4.579082	1.540922	2.526619	1.558960
bathy_1sec				
7.486712				

Arctic tern, AIC selected model, excluding SST

dist_col	chl_june	summ_front_sd	bathy_1sec
1.965858	4.317997	2.526375	6.674326

Common tern, AIC selected model, including SST

dist_col	chl_june	sst_april	bathy_1sec	sst_june
1.720706	4.623803	1.449510	4.555302	2.950160

Common tern, AIC selected model, excluding SST

dist_col	chl_apr	summ_front_sd	strat_temp	sal_summ
2.692582	2.557135	2.551321	4.646595	2.946827
sal_spring	ss_current	bathy_1sec		
5.469419	1.602503	6.241017		

Roseate tern, LRT selected model, including SST

dist_col	chl_june	sst_may
1.517489	1.539523	1.017911

Roseate tern, AIC selected model, excluding SST

dist_col	dist_shore	chl_apr	chl_may	chl_june
1.588470	5.518533	2.414380	2.174784	3.230091
summ_front_sd	spring_front	spring_frt_sd	ss_current	
5.786227	8.709124	5.425485	1.703359	

Sandwich tern, AIC selected model, including SST

dist_col	dist_shore	chl_apr	chl_june	summ_front_sd
1.124730	3.652339	1.817593	2.020926	2.729867
ss_wave				
2.996139				

Sandwich tern, AIC selected model, excluding SST

dist_col	dist_shore	chl_may	ss_wave	bathy_1sec
1.150402	3.750730	1.112642	7.076829	9.538748

	dist_col	dist_shore	chl_may	ss_wave	bathy_1sec
dist_col	1.0000000	0.7006419	-0.5509828	-0.5065455	-0.5723878
dist_shore	0.7006419	1.0000000	-0.7563967	-0.7518555	-0.8016200
chl_may	-0.5509828	-0.7563967	1.0000000	0.7533570	0.7775044

ss_wave	-0.5065455	-0.7518555	0.7533570	1.0000000	0.9612668
bathy_1sec	-0.5723878	-0.8016200	0.7775044	0.9612668	1.0000000