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**esem**  
*Release v0.2.0*

**D Watson-Parris**

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## **INSTALLING ESEM**

### **1.1 Using PyPi**

It is straightforward to install esem using pip, this will automatically include tensorflow (with GPU support):

```
$ pip install esem
```

Optionally also install GPFlow, keras or scikit-learn

```
$ pip install esem[gpflow]
```

Or

```
$ pip install esem[gpflow,keras,scikit-learn]
```

### **1.2 Using conda**

In order to make the most of the support for Iris and CIS creating a specific conda environment is recommended. If you don't already have conda, you must first download and install it. Anaconda is a free conda package that includes Python and many common scientific and data analysis libraries, and is available [here](#). Further documentation on using Anaconda and the features it provides can be found at <http://docs.continuum.io/anaconda/index.html>.

Having installed (mini-) conda - and ideally within a fresh environment - you can easily install CIS (and Iris) with the following command:

```
$ conda install -c conda-forge cis
```

It is then straightforward to install esem in to this environment using pip as above.

### **1.3 Dependencies**

If you choose to install the dependencies yourself, use the following command to check the required dependencies are present:

```
$ python setup.py checkdep
```



## WHAT'S NEW IN ESEM

### 2.1 What's new in ESEM 1.1

This page documents the new features added, and bugs fixed in ESEM since version 1.0. For more detail see all changes here: <https://github.com/duncanwp/ESEM/compare/1.0.0...1.1.0>

#### 2.1.1 ESEM 1.1 features

- We have added this What's New page for tracking the latest developments in ESEM!
- We have dropped the mandatory requirement of Iris to make installation of ESEM easier. We have also added support for xarray DataArrays so that users can use their preferred library for data processing.
- The `esem.emulator.Emulator.predict()` and `esem.emulator.Emulator.batch_stats()` methods can now accept pd.DataFrames to match the training interface. The associated doc-strings and signatures have been extended to reflect this.

#### 2.1.2 Bugs fixed

- Use tqdm.auto to automatically choose the appropriate progress bar for the context
- Fix `plot_validation` handling of masked data



## EMULATING WITH ESEM

ESEm provides a simple and streamlined interface to emulate earth system datasets, denoted  $Y$  in the following documentation. These datasets can be provided as iris Cubes, xarray DataArrays or numpy arrays and the resulting emulation results will preserve any associated metadata. The corresponding predictors ( $X$ ) can be provided as a numpy array or pandas *DataFrame*. This emulation is essentially just a regression estimating the functional form:

$$Y \approx f(X)$$

and can be performed using a variety of techniques using the same API.

### 3.1 Gaussian processes emulation

Gaussian processes (GPs) are a popular choice for model emulation due to their simple formulation and robust uncertainty estimates, particularly in cases where there is limited training data. Many excellent texts are available to describe their implementation and use (Rasmussen and Williams, 2005) and we only provide a short description here. Briefly, a GP is a stochastic process (a distribution of continuous functions) and can be thought of as an infinite dimensional normal distribution (hence the name).

The ESEm GP emulation module provides a thin wrapper around the [GPFlow](#) implementation. Please see their documentation for a detailed description of the implementation.

An important consideration when using GP regression is the form of the covariance matrix, or kernel. Typical kernels include: constant; linear; radial basis function (RBF; or squared exponential); and Matérn 3/2 and 5/2 which are only once and twice differentiable respectively. Kernels can also be designed to represent any aspect of the functions of interest such as non-stationarity or periodicity. This choice can often be informed by the physical setting and provides greater control and interpretability of the resulting model compared to e.g., Neural Networks.

---

**Note:** By default, ESEm uses a combination of linear, RBF and polynomial kernels which are suitable for the smooth and continuous parameter response expected for the examples used in this paper and related problems. However, given the importance of the kernel for determining the form of the functions generated by the GP we have also included the ability for users to specify combinations of other common kernels. See e.g., [Duvenaud, 2011](#) for a clear description of some common kernels and their combinations, as well as work towards automated methods for choosing them.

---

The framework provided by GPFlow also allows for multi-output GP regression and ESEm takes advantage of this to automatically provide regression over each of the output features provided in the training data. E.g.  $Y$  can be of arbitrary dimensionality. It will be automatically flattened and reshaped before being passed to GPFlow.

The most convenient way to setup a GPFlow emulator is using the `esem_gp_model()` function which can be imported directly:

```
from esem import gp_model
```

This creates a regression model with a default kernel as described above but provides a convenient interface for defining arbitrary kernels through addition and multiplication. For example, to initialize a model with a `Linear+Cosine()` kernel:

```
from esem import gp_model

# X_train and Y_train are our predictors and outputs, respectively.
model = gp_model(X_train, Y_train, kernel=['Linear', 'Cosine'], kernel_op='add')
```

Further details are described in the function description `esem.gp_model()`.

Examples of emulation using Gaussian processes can be found in [Emulating\\_using\\_GPs.ipynb](#) and [CMIP6\\_emulator.ipynb](#).

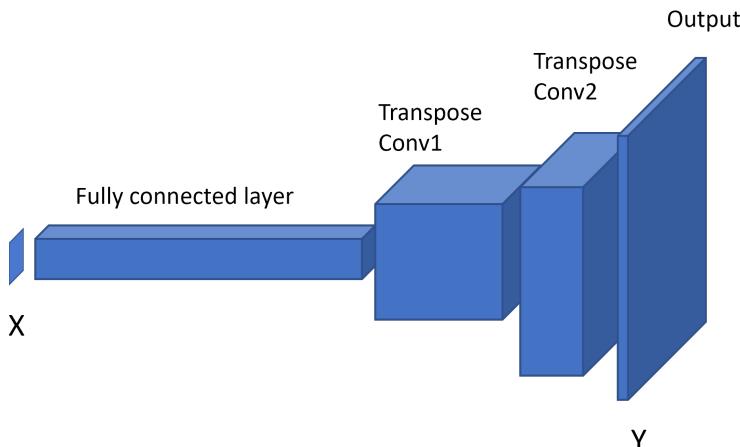
## 3.2 Neural network emulation

While fully connected neural networks have been used for many years, even in climate science, the recent surge in popularity has been powered by the increases in expressibility provided by deep, convolutional neural networks (CNNs) and the regularisation techniques which prevent these huge models from over-fitting the large amounts of training data required to train them.

Many excellent introductions can be found elsewhere but, briefly, a neural network consists of a network of nodes connecting (through a variety of architectures) the inputs to the target outputs via a series of weighted activation functions. The network architecture and activation functions are typically chosen a-priori and then the model weights are determined through a combination of back-propagation and (batch) gradient descent until the outputs match (defined by a given loss function) the provided training data. As previously discussed, the random dropping of nodes (by setting the weights to zero), termed dropout, can provide estimates of the prediction uncertainty of such networks.

The computational efficiency of such networks and the rich variety of architectures available have made them the tool of choice in many machine learning settings, and they are starting to be used in climate sciences for emulation, although the large amounts of training data required (especially compared to GPs) have so far limited their use somewhat.

ESEm provides a baseline CNN architecture based on the [Keras](#) library which essentially acts as a decoder - transforming input parameters into predicted 2(or 3) dimensional output fields:



This model can be easily constructed using the `esem.cnn_model()` function. It is possible to use any Keras model in this way though and there are many potential ways of improving / developing this simple model.

An example of emulation using this convolution neural network can be found in [Emulating\\_using\\_ConvNets.ipynb](#).

### 3.3 Random forest emulation

ESEm also provides the option for emulation with Random Forests using the open-source implementation provided by scikit-learn. Random Forest estimators are comprised of an ensemble of decision trees; each decision tree is a recursive binary partition over the training data and the predictions are an average over the predictions of the decision trees.

As a result of this architecture, Random Forests (along with other algorithms built on decision trees) have two main attractions. Firstly, they require very little pre-processing of the inputs as the binary partitions are invariant to monotonic rescaling of the training data. Secondly, and of particular importance for climate problems, they are unable to extrapolate outside of their training data because the predictions are averages over subsets of the training dataset.

This model can be constructed using the `esem.rf_model()` function. All of the relevant scikit-learn arguments and keyword-arguments can be provided through this interface.

An example of emulation using the random forest can be found in [CRM\\_Emotion\\_with\\_RandomForest.ipynb](#).

### 3.4 Data processing

Many of the above approaches make assumptions about, or simply perform better when, the training data is structured or distributed in a certain way. These transformations are purely to help the emulator fit the training data, and can complicate comparison with e.g. observations during calibration. ESEm provides a simple and transparent way of transforming the datasets for training, and this automatically un-transforms the model predictions to aid in observational comparison.

Where these transformations are strictly necessary for a given model then it will be included in the wrapper function. Other choices are left to the user to apply as required.

For example, to ‘whiten’ the data (that is, remove the mean and normalise by the standard deviation):

```
import esem
from esem import gp_model

# X_train and Y_train are our predictors and outputs, respectively.
model = gp_model(X_train, Y_train, data_processors=[esem.data_processors.Whiten()])
```

A full list of the data processors can be found in the [API documentation](#).

### 3.5 Feature selection

ESEm includes a simple utility function that wraps the scikit-learn LassoLarsIC regression tool in order to enable an initial feature (parameter) selection. This can be useful to reduce the dimensionality of the input space. Either the Akaike information criterion (AIC) or the Bayes Information criterion (BIC) can be used, although BIC is the default.

For example,

```
from esem import gp_model
from esem.utils import get_param_mask

# X and Y are our model parameters and outputs respectively.
```

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```
active_params = get_param_mask(X, y)

# The model parameters can then be subsampled either directly
X_sub = X[:, active_params]

# Or by specifying the GP active_dims
active_dims, = np.where(active_params)
model = gp_model(X, y, active_dims=active_dims)
```

Note, this estimate only applies to one-dimensional outputs. Feature selection for higher dimension outputs is a much harder task beyond the scope of this package.

## CALIBRATING WITH ESEM

Having trained a fast, robust emulator this can be used to calibrate our model against available observations. The following description closely follows that in our model description paper but with explicit links to the ESEm algorithms to aid clarity. Generally, this problem involves estimating the model parameters which could give rise to, or best match, the available observations.

In other words, we would like to know the posterior probability distribution of the input parameters:  $p(\theta|Y^0)$ .

Using Bayes' theorem, we can write this as:

$$p(\theta|Y^0) = p(Y^0|\theta) p(\theta) p(Y^0) \quad (4.1)$$

Where the probability of an output given the input parameters,  $p(Y^0|\theta)$ , is referred to as the likelihood. While the model is capable of sampling this distribution, generally the full distribution is unknown and intractable, and we must approximate this likelihood. Depending on the purpose of the calibration and assumptions about the form of  $p(Y^0|Y)$ , different techniques can be used. In order to determine a (conservative) estimate of the parametric uncertainty in the model for example, we can use approximate Bayesian computation (ABC) to determine those parameters which are plausible given a set of observations. Alternatively, we may wish to know the optimal parameters to best match a set of observations and Markov-Chain Monte-Carlo based techniques might be more appropriate. Both of these sampling strategies are available in ESEm and we describe each of them here.

### 4.1 Using approximate Bayesian computation (ABC)

The simplest ABC approach seeks to approximate the likelihood using only samples from the simulator and a discrepancy function  $\rho$ :

$$p(\theta|Y^0) \propto p(Y^0|Y) p(Y|\theta) p(\theta) \approx \int \mathbb{I}(\rho(Y^0, Y) \leq \epsilon) p(Y|\theta) p(\theta) dY \quad (4.2)$$

where the indicator function  $\mathbb{I}(x) = 1$  if  $x$  is true and  $\mathbb{I}(x) = 0$  if  $x$  is false, and  $\epsilon$  is a small discrepancy. This can then be integrated numerically using e.g., Monte-Carlo sampling of  $p(\theta)$ . Any of those parameters for which  $\rho(Y^0, Y) \leq \epsilon$  are accepted and those which do not are rejected. As  $\epsilon \rightarrow \infty$  therefore, all parameters are accepted and we recover  $p(\theta)$ . For  $\epsilon = 0$ , it can be shown that we generate samples from the posterior  $p(\theta|Y^0)$  exactly.

In practice however the simulator proposals will never exactly match the observations and we must make a pragmatic choice for both  $\rho$  and  $\epsilon$ . ESEm includes an implementation of the ‘implausibility metric’ which defines the discrepancy in terms of the standardized Cartesian distance:

$$\rho(Y^0, Y(\theta)) = \frac{|Y^0 - Y(\theta)|}{\sqrt{\sigma_E^2 + \sigma_Y^2 + \sigma_R^2 + \sigma_S^2}} = \rho(Y^0, \theta) \quad (4.3)$$

where the total standard deviation is taken to be the squared sum of the emulator variance ( $\sigma_E^2$ ) and the uncertainty in the observations ( $\sigma_Y^2$ ) and due to representation ( $\sigma_R^2$ ) and structural model uncertainties ( $\sigma_S^2$ ) as described in the paper.

Framed in this way,  $\epsilon$ , can be thought of as representing the number of standard deviations the (emulated) model value is from the observations. While this can be treated as a free parameter and may be specified in ESEm, it is common to choose  $\epsilon = 3$  since it can be shown that for unimodal distributions values of  $3\sigma$  correspond to a greater than 95% confidence bound. This approach is implemented in the `esem.abc_sampler.ABCSampler` class where  $\epsilon$  is referred to as a threshold since it defines the cut-off for acceptance.

In the general case, multiple ( $\mathcal{N}$ ) observations can be used and  $\rho$  can be written as a vector of implausibilities,  $\rho(Y_i^O, \theta)$  or simply  $\rho_i(\theta)$ , and a modified method of rejection or acceptance must be used. A simple choice is to require  $\rho_i < \epsilon \forall i \in \mathcal{N}$ , however this can become restrictive for large  $\mathcal{N}$  due to the curse of dimensionality. The first step should be to reduce  $\mathcal{N}$  through the use of summary statistics, such as averaging over regions, or stations, or by performing an e.g. Principle Component Analysis (PCA) decomposition.

An alternative is to introduce a tolerance ( $T$ ) such that only some proportion of  $\rho_i$  need to be smaller than  $\epsilon$  :  $\sum_{i=0}^{\mathcal{N}} H(\rho_i - \epsilon) < T$ , where  $H$  is the Heaviside function. This tolerance can be specified when sampling using the `esem.abc_sampler.ABCSampler`. An efficient implementation of this approach whereby the acceptance is calculated in batches on the GPU can be particularly useful when dealing with high-dimensional outputs `esem.abc_sampler.ABCSampler`. It is recommended however, to choose  $T = 0$  as a first approximation and then identify any particular observations which generate a very large implausibilities, since this provides a mechanism for identifying potential structural (or observational) errors.

A useful way of identifying such observations is using the `esem.abc_sampler.ABCSampler.get_implausibility()` method which returns the full implausibility matrix  $\rho_i$ . Note this may be very large (N-samples x N-observations) so it is recommended that only a subset of the full sample space be requested. The offending observations can then be removed and noted for further investigation.

Examples of the ABC sampling can be found in [Calibrating\\_GPs\\_using\\_ABC.ipynb](#).

## 4.2 Using Markov-chain Monte-Carlo

The ABC method described above is simple and powerful, but somewhat inefficient as it repeatedly samples from the same prior. In reality each rejection or acceptance of a set of parameters provides us with extra information about the ‘true’ form of  $p(\theta|Y^0)$  so that the sampler could spend more time in plausible regions of the parameter space. This can then allow us to use smaller values of  $\epsilon$  and hence find better approximations of  $p(\theta|Y^0)$ .

Given the joint probability distribution described by Eq. 2 and an initial choice of parameters  $\theta'$  and (emulated) output  $Y'$ , the acceptance probability  $r$  of a new set of parameters ( $\theta$ ) is given by:

$$r = \frac{p(Y^0|Y') p(\theta'|\theta) p(\theta)}{p(Y^0|Y) p(\theta|\theta') p(\theta)} \quad (4.4)$$

The `esem.sampler.MCMCSampler` class uses the TensorFlow-probability implementation of Hamiltonian Monte-Carlo (HMC) which uses the gradient information automatically calculated by TensorFlow to inform the proposed new parameters  $\theta$ . For simplicity, we assume that the proposal distribution is symmetric:  $p(\theta'|\theta) = p(\theta|\theta')$ , which is implemented as a zero log-acceptance correction in the initialisation of the TensorFlow target distribution. The target log probability provided to the TensorFlow HMC algorithm is then:

$$\log(r) = \log(p(Y^0|Y')) + \log(p(\theta')) - \log(p(Y^0|Y)) - \log(p(\theta)) \quad (4.5)$$

Note, that for this implementation the distance metric  $\rho$  must be cast as a probability distribution with values [0, 1]. We therefore assume that this discrepancy can be approximated as a normal distribution centred about zero, with standard deviation equal to the sum of the squares of the variances as described in Eq. 3:

$$p(Y^0|Y) \approx \frac{1}{\sigma_t \sqrt{2\pi}} e^{-\frac{1}{2} \left( \frac{Y^0 - Y}{\sigma_t} \right)^2}, \quad \sigma_t = \sqrt{\sigma_E^2 + \sigma_Y^2 + \sigma_R^2 + \sigma_S^2} \quad (4.6)$$

The `esem.sampler.MCMCSampler.sample()` method will then return the requested number of accepted samples as well as reporting the acceptance rate, which provides a useful metric for tuning the algorithm. It should be noted that MCMC algorithms can be sensitive to a number of key parameters, including the number of burn-in steps used (and discarded) before sampling occurs and the step size. Each of these can be controlled via keyword arguments to the `esem.sampler.MCMCSampler.sample()` method.

This approach can provide much more efficient sampling of the emulator and provide improved parameter estimates, especially when used with informative priors which can guide the sampler.

Examples of the MCMC sampling can be found in [Calibrating\\_GPs\\_using\\_MCMC.ipynb](#) and [CMIP6\\_emulator.ipynb](#).



## EXAMPLES

## 5.1 Emulating using GPs

```
[1]: import os
## Ignore my broken HDF5 install...
os.putenv("HDF5_DISABLE_VERSION_CHECK", '1')
```

```
[2]: import iris
from utils import get_bc_ppe_data

from esem import gp_model
from esem.utils import get_random_params

import iris.quickplot as qplt
import matplotlib.pyplot as plt
%matplotlib inline

C:\Users\duncan\miniconda3\envs\gcem_dev\lib\site-packages\h5py\__init__.py:40: UserWarning: h5py is running against HDF5 1.10.6 when it was built against 1.10.5, this may cause problems
'{0}.{1}.{2}'.format(*version.hdf5_builtin_version_tuple)
```

### 5.1.1 Read in the parameters and observables

```
[4]: ppe_params, ppe_aaod = get_bc_ppe_data()

C:\Users\duncan\miniconda3\envs\gcem_dev\lib\site-packages\iris\__init__.py:249: IrisDeprecation: setting the 'Future' property 'netcdf_promote' is deprecated and will be removed in a future release. Please remove code that sets this property.
warn_deprecated(msg.format(name))
C:\Users\duncan\miniconda3\envs\gcem_dev\lib\site-packages\iris\__init__.py:249: IrisDeprecation: setting the 'Future' property 'netcdf_promote' is deprecated and will be removed in a future release. Please remove code that sets this property.
warn_deprecated(msg.format(name))
```

```
[5]: n_test = 5

X_test, X_train = ppe_params[:n_test], ppe_params[n_test:]
Y_test, Y_train = ppe_aaod[:n_test, 0], ppe_aaod[n_test:, 0]
```

## 5.1.2 Setup and run the models

```
[9]: model = gp_model(X_train, Y_train)

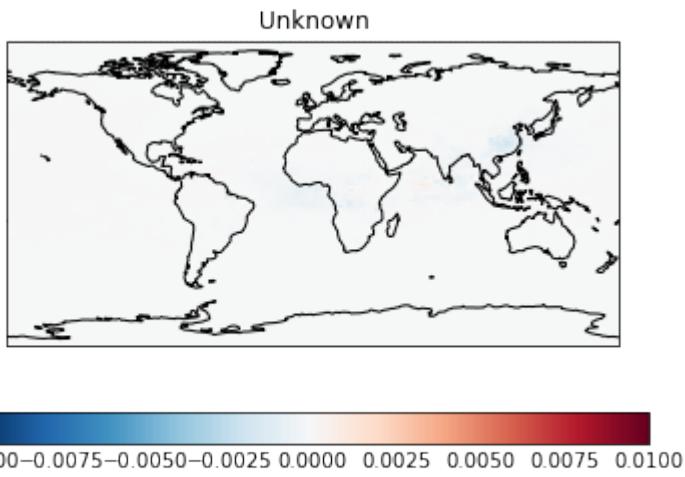
[15]: model.train()

[16]: m, v = model.predict(X_test.values)

[17]: ## validation_plot(Y_test.data.flatten(), m.data.flatten(), v.data.flatten())

[18]: qplt.pcolormesh((m.collapsed('sample', iris.analysis.MEAN)-Y_test.collapsed('job', iris.
   ↴analysis.MEAN)), cmap='RdBu_r', vmin=-0.01, vmax=0.01)
    plt.gca().coastlines()
C:\Users\duncan\miniconda3\envs\gcm_dev\lib\site-packages\iris\coords.py:1410: UserWarning: Collapsing a non-contiguous coordinate. Metadata may not be fully descriptive for 'sample'.
    warnings.warn(msg.format(self.name()))
C:\Users\duncan\miniconda3\envs\gcm_dev\lib\site-packages\iris\coords.py:1410: UserWarning: Collapsing a non-contiguous coordinate. Metadata may not be fully descriptive for 'job'.
    warnings.warn(msg.format(self.name()))
C:\Users\duncan\miniconda3\envs\gcm_dev\lib\site-packages\iris\coords.py:1193: UserWarning: Coordinate 'longitude' is not bounded, guessing contiguous bounds.
    'contiguous bounds.'.format(self.name()))
C:\Users\duncan\miniconda3\envs\gcm_dev\lib\site-packages\iris\coords.py:1193: UserWarning: Coordinate 'latitude' is not bounded, guessing contiguous bounds.
    'contiguous bounds.'.format(self.name()))

[18]: <cartopy.mpl.feature_artist.FeatureArtist at 0x19ea2c99388>
```



```
[19]: ## Note the model variance is constant across the outputs
v.data.max()

[19]: 1.175408691451335e-06
```

```
[15]: get_random_params(3, int(1e5)).shape
[15]: (100000, 3)

[16]: m, sd = model.batch_stats(get_random_params(3, int(1e3)))
100%|#####| 1000/1000 [00:09<00:00, 149.00sample/s]

[17]: m, sd = model.batch_stats(get_random_params(3, int(1e4)), batch_size=10)
100%|#####| 10000/10000 [00:07<00:00, 1459.07sample/s]

[19]: m, sd = model.batch_stats(get_random_params(3, int(1e6)), batch_size=10000)
100%|#####| 1000000/1000000 [00:09<00:00, 122569.61sample/s]

[ ]:
```

## 5.2 Emulating using CNNs

```
[1]: import iris

from utils import get_bc_ppe_data

from esem import cnn_model
from esem.utils import get_random_params

import iris.quickplot as qplt
import matplotlib.pyplot as plt
%matplotlib inline
```

### 5.2.1 Read in the parameters and data

```
[2]: ppe_params, ppe_aaod = get_bc_ppe_data()

[3]: ## Ensure the time dimension is last - this is treated as the color 'channel'
ppe_aaod.transpose((0,2,3,1))

[4]: n_test = 5

X_test, X_train = ppe_params[:n_test], ppe_params[n_test:]
Y_test, Y_train = ppe_aaod[:n_test], ppe_aaod[n_test:]

[5]: Y_train
[5]: <iris 'Cube' of Absorption optical thickness - total 550nm / (1) (job: 34; latitude: 96;
longitude: 192; time: 12)>
```

## 5.2.2 Setup and run the models

[6]: model = cnn\_model(X\_train, Y\_train)

[7]: model.train()

```
Epoch 1/100
4/4 [=====] - 3s 71ms/step - loss: 1.1407 - val_loss: 0.4622
Epoch 2/100
4/4 [=====] - 0s 19ms/step - loss: 1.1396 - val_loss: 0.4620
Epoch 3/100
4/4 [=====] - 0s 17ms/step - loss: 1.1391 - val_loss: 0.4618
Epoch 4/100
4/4 [=====] - 0s 19ms/step - loss: 1.1382 - val_loss: 0.4613
Epoch 5/100
4/4 [=====] - ETA: 0s - loss: 1.048 - 0s 18ms/step - loss: 1.1360 - val_loss: 0.4577
Epoch 6/100
4/4 [=====] - 0s 17ms/step - loss: 1.1256 - val_loss: 0.4536
Epoch 7/100
4/4 [=====] - 0s 18ms/step - loss: 1.1114 - val_loss: 0.4446
Epoch 8/100
4/4 [=====] - 0s 19ms/step - loss: 1.0957 - val_loss: 0.4372
Epoch 9/100
4/4 [=====] - 0s 20ms/step - loss: 1.0746 - val_loss: 0.4246
Epoch 10/100
4/4 [=====] - 0s 18ms/step - loss: 1.0569 - val_loss: 0.4152
Epoch 11/100
4/4 [=====] - 0s 20ms/step - loss: 1.0326 - val_loss: 0.4073
Epoch 12/100
4/4 [=====] - 0s 19ms/step - loss: 1.0055 - val_loss: 0.4009
Epoch 13/100
4/4 [=====] - 0s 20ms/step - loss: 0.9815 - val_loss: 0.3892
Epoch 14/100
4/4 [=====] - 0s 21ms/step - loss: 0.9567 - val_loss: 0.3797
Epoch 15/100
4/4 [=====] - 0s 22ms/step - loss: 0.9285 - val_loss: 0.3679
Epoch 16/100
4/4 [=====] - 0s 22ms/step - loss: 0.9015 - val_loss: 0.3472
Epoch 17/100
4/4 [=====] - 0s 21ms/step - loss: 0.8915 - val_loss: 0.3343
Epoch 18/100
4/4 [=====] - 0s 20ms/step - loss: 0.8643 - val_loss: 0.3255
Epoch 19/100
4/4 [=====] - 0s 19ms/step - loss: 0.8362 - val_loss: 0.3176
Epoch 20/100
4/4 [=====] - 0s 19ms/step - loss: 0.8201 - val_loss: 0.3009
Epoch 21/100
4/4 [=====] - 0s 19ms/step - loss: 0.7953 - val_loss: 0.2862
Epoch 22/100
4/4 [=====] - 0s 19ms/step - loss: 0.7777 - val_loss: 0.2778
Epoch 23/100
4/4 [=====] - 0s 18ms/step - loss: 0.7578 - val_loss: 0.2747
```

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Epoch 24/100
4/4 [=====] - 0s 19ms/step - loss: 0.7465 - val_loss: 0.2592
Epoch 25/100
4/4 [=====] - 0s 22ms/step - loss: 0.7223 - val_loss: 0.2570
Epoch 26/100
4/4 [=====] - 0s 19ms/step - loss: 0.7036 - val_loss: 0.2337
Epoch 27/100
4/4 [=====] - 0s 20ms/step - loss: 0.6804 - val_loss: 0.2253
Epoch 28/100
4/4 [=====] - 0s 18ms/step - loss: 0.6666 - val_loss: 0.2308
Epoch 29/100
4/4 [=====] - 0s 18ms/step - loss: 0.6502 - val_loss: 0.2109
Epoch 30/100
4/4 [=====] - 0s 19ms/step - loss: 0.6315 - val_loss: 0.1978
Epoch 31/100
4/4 [=====] - 0s 18ms/step - loss: 0.6138 - val_loss: 0.1819
Epoch 32/100
4/4 [=====] - 0s 19ms/step - loss: 0.5970 - val_loss: 0.1738
Epoch 33/100
4/4 [=====] - 0s 21ms/step - loss: 0.5801 - val_loss: 0.1640
Epoch 34/100
4/4 [=====] - 0s 22ms/step - loss: 0.5622 - val_loss: 0.1643
Epoch 35/100
4/4 [=====] - 0s 18ms/step - loss: 0.5502 - val_loss: 0.1508
Epoch 36/100
4/4 [=====] - 0s 18ms/step - loss: 0.5405 - val_loss: 0.1625
Epoch 37/100
4/4 [=====] - 0s 18ms/step - loss: 0.5311 - val_loss: 0.1517
Epoch 38/100
4/4 [=====] - 0s 18ms/step - loss: 0.5191 - val_loss: 0.1331
Epoch 39/100
4/4 [=====] - 0s 19ms/step - loss: 0.5063 - val_loss: 0.1264
Epoch 40/100
4/4 [=====] - 0s 18ms/step - loss: 0.4976 - val_loss: 0.1217
Epoch 41/100
4/4 [=====] - 0s 18ms/step - loss: 0.4865 - val_loss: 0.1159
Epoch 42/100
4/4 [=====] - 0s 18ms/step - loss: 0.4715 - val_loss: 0.1112
Epoch 43/100
4/4 [=====] - 0s 19ms/step - loss: 0.4604 - val_loss: 0.1131
Epoch 44/100
4/4 [=====] - 0s 19ms/step - loss: 0.4498 - val_loss: 0.1002
Epoch 45/100
4/4 [=====] - 0s 21ms/step - loss: 0.4404 - val_loss: 0.0969
Epoch 46/100
4/4 [=====] - 0s 20ms/step - loss: 0.4320 - val_loss: 0.0938
Epoch 47/100
4/4 [=====] - 0s 18ms/step - loss: 0.4242 - val_loss: 0.0905
Epoch 48/100
4/4 [=====] - 0s 18ms/step - loss: 0.4173 - val_loss: 0.0874
Epoch 49/100
4/4 [=====] - 0s 19ms/step - loss: 0.4105 - val_loss: 0.0903

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Epoch 50/100
4/4 [=====] - 0s 18ms/step - loss: 0.4032 - val_loss: 0.0825
Epoch 51/100
4/4 [=====] - 0s 21ms/step - loss: 0.3931 - val_loss: 0.0872
Epoch 52/100
4/4 [=====] - 0s 22ms/step - loss: 0.3891 - val_loss: 0.0839
Epoch 53/100
4/4 [=====] - 0s 22ms/step - loss: 0.3813 - val_loss: 0.0768
Epoch 54/100
4/4 [=====] - 0s 22ms/step - loss: 0.3767 - val_loss: 0.0764
Epoch 55/100
4/4 [=====] - 0s 21ms/step - loss: 0.3720 - val_loss: 0.0776
Epoch 56/100
4/4 [=====] - 0s 21ms/step - loss: 0.3671 - val_loss: 0.0723
Epoch 57/100
4/4 [=====] - 0s 20ms/step - loss: 0.3601 - val_loss: 0.0719
Epoch 58/100
4/4 [=====] - 0s 19ms/step - loss: 0.3570 - val_loss: 0.0708
Epoch 59/100
4/4 [=====] - 0s 20ms/step - loss: 0.3562 - val_loss: 0.0704
Epoch 60/100
4/4 [=====] - 0s 20ms/step - loss: 0.3519 - val_loss: 0.0699
Epoch 61/100
4/4 [=====] - 0s 19ms/step - loss: 0.3488 - val_loss: 0.0689
Epoch 62/100
4/4 [=====] - 0s 20ms/step - loss: 0.3444 - val_loss: 0.0727
Epoch 63/100
4/4 [=====] - 0s 19ms/step - loss: 0.3427 - val_loss: 0.0795
Epoch 64/100
4/4 [=====] - 0s 20ms/step - loss: 0.3420 - val_loss: 0.0691
Epoch 65/100
4/4 [=====] - 0s 19ms/step - loss: 0.3360 - val_loss: 0.0678
Epoch 66/100
4/4 [=====] - 0s 21ms/step - loss: 0.3336 - val_loss: 0.0664
Epoch 67/100
4/4 [=====] - 0s 19ms/step - loss: 0.3303 - val_loss: 0.0689
Epoch 68/100
4/4 [=====] - 0s 19ms/step - loss: 0.3297 - val_loss: 0.0669
Epoch 69/100
4/4 [=====] - 0s 19ms/step - loss: 0.3246 - val_loss: 0.0659
Epoch 70/100
4/4 [=====] - 0s 19ms/step - loss: 0.3246 - val_loss: 0.0682
Epoch 71/100
4/4 [=====] - 0s 20ms/step - loss: 0.3217 - val_loss: 0.0660
Epoch 72/100
4/4 [=====] - 0s 18ms/step - loss: 0.3196 - val_loss: 0.0656
Epoch 73/100
4/4 [=====] - 0s 19ms/step - loss: 0.3201 - val_loss: 0.0667
Epoch 74/100
4/4 [=====] - 0s 18ms/step - loss: 0.3166 - val_loss: 0.0653
Epoch 75/100
4/4 [=====] - 0s 18ms/step - loss: 0.3151 - val_loss: 0.0666

```

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```

Epoch 76/100
4/4 [=====] - 0s 19ms/step - loss: 0.3137 - val_loss: 0.0657
Epoch 77/100
4/4 [=====] - 0s 18ms/step - loss: 0.3143 - val_loss: 0.0648
Epoch 78/100
4/4 [=====] - 0s 18ms/step - loss: 0.3112 - val_loss: 0.0647
Epoch 79/100
4/4 [=====] - 0s 18ms/step - loss: 0.3084 - val_loss: 0.0674
Epoch 80/100
4/4 [=====] - 0s 19ms/step - loss: 0.3087 - val_loss: 0.0650
Epoch 81/100
4/4 [=====] - 0s 19ms/step - loss: 0.3056 - val_loss: 0.0709
Epoch 82/100
4/4 [=====] - 0s 18ms/step - loss: 0.3082 - val_loss: 0.0651
Epoch 83/100
4/4 [=====] - 0s 18ms/step - loss: 0.3048 - val_loss: 0.0644
Epoch 84/100
4/4 [=====] - 0s 19ms/step - loss: 0.3036 - val_loss: 0.0669
Epoch 85/100
4/4 [=====] - 0s 18ms/step - loss: 0.3032 - val_loss: 0.0662
Epoch 86/100
4/4 [=====] - 0s 18ms/step - loss: 0.3030 - val_loss: 0.0672
Epoch 87/100
4/4 [=====] - 0s 18ms/step - loss: 0.3016 - val_loss: 0.0691
Epoch 88/100
4/4 [=====] - 0s 18ms/step - loss: 0.3014 - val_loss: 0.0706
Epoch 89/100
4/4 [=====] - 0s 18ms/step - loss: 0.3006 - val_loss: 0.0648
Epoch 90/100
4/4 [=====] - 0s 18ms/step - loss: 0.2988 - val_loss: 0.0670
Epoch 91/100
4/4 [=====] - 0s 17ms/step - loss: 0.2985 - val_loss: 0.0670
Epoch 92/100
4/4 [=====] - 0s 21ms/step - loss: 0.2977 - val_loss: 0.0635
Epoch 93/100
4/4 [=====] - 0s 20ms/step - loss: 0.2967 - val_loss: 0.0638
Epoch 94/100
4/4 [=====] - 0s 23ms/step - loss: 0.2969 - val_loss: 0.0654
Epoch 95/100
4/4 [=====] - 0s 19ms/step - loss: 0.2954 - val_loss: 0.0648
Epoch 96/100
4/4 [=====] - 0s 21ms/step - loss: 0.2973 - val_loss: 0.0638
Epoch 97/100
4/4 [=====] - 0s 23ms/step - loss: 0.2949 - val_loss: 0.0668
Epoch 98/100
4/4 [=====] - 0s 24ms/step - loss: 0.2945 - val_loss: 0.0641
Epoch 99/100
4/4 [=====] - 0s 22ms/step - loss: 0.2936 - val_loss: 0.0648
Epoch 100/100
4/4 [=====] - 0s 23ms/step - loss: 0.2932 - val_loss: 0.0651

```

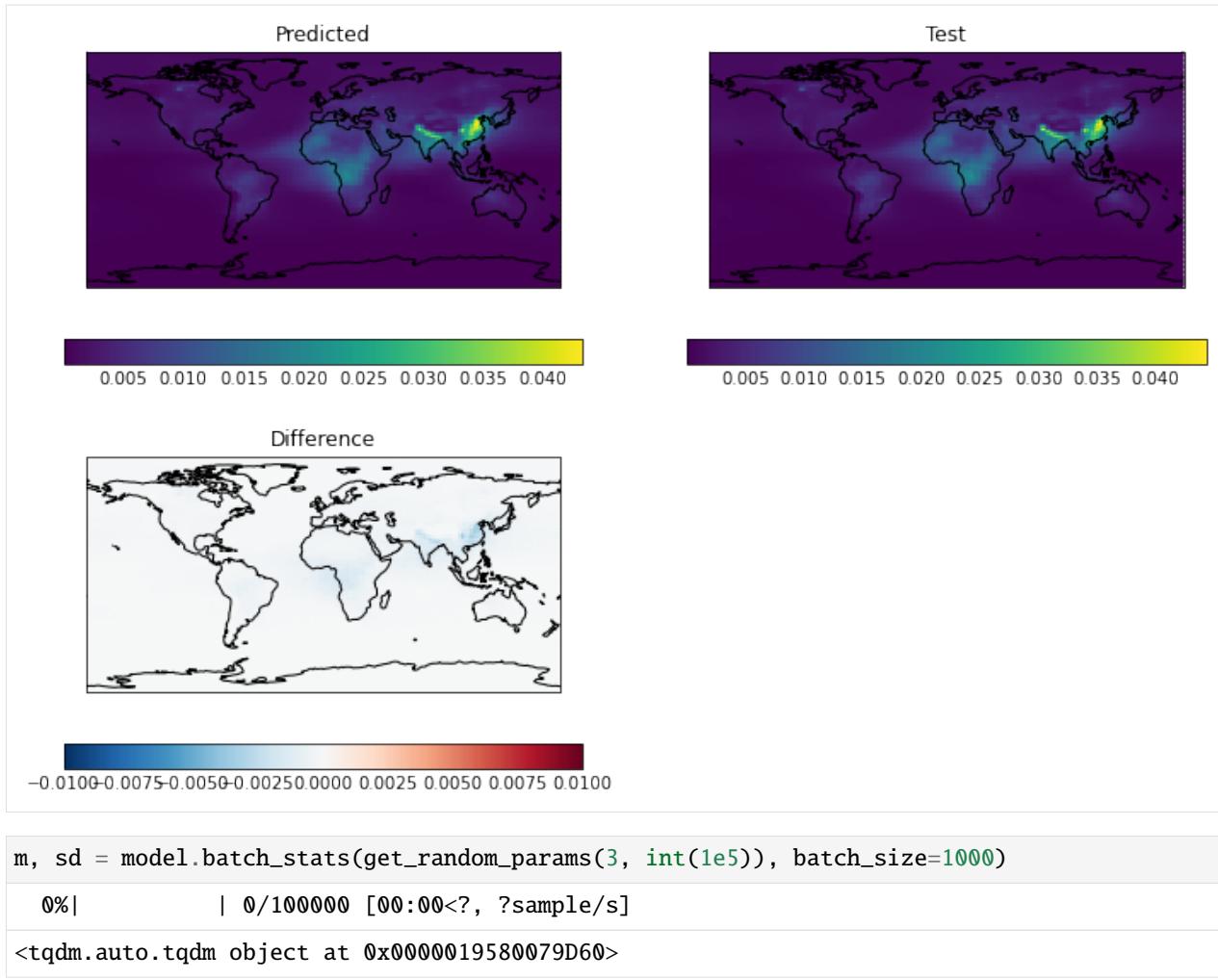
[8]: `m, v = model.predict(X_test.to_numpy())`

```
[9]: ## TODO: Tidy this up a bit
plt.figure(figsize=(12, 8))
plt.subplot(2,2,1)
qplt.pcolormesh(m[0].collapsed('time', iris.analysis.MEAN))
plt.gca().set_title('Predicted')
plt.gca().coastlines()

plt.subplot(2,2,2)
qplt.pcolormesh(Y_test[0].collapsed('time', iris.analysis.MEAN))
plt.gca().set_title('Test')
plt.gca().coastlines()

plt.subplot(2,2,3)
qplt.pcolormesh((m.collapsed(['sample', 'time'], iris.analysis.MEAN)-Y_test.collapsed([
    'job', 'time'], iris.analysis.MEAN)), cmap='RdBu_r', vmin=-0.01, vmax=0.01)
plt.gca().coastlines()
plt.gca().set_title('Difference')

C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\coords.py:1803:_
    ↵UserWarning: Coordinate 'longitude' is not bounded, guessing contiguous bounds.
    warnings.warn(
C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\coords.py:1803:_
    ↵UserWarning: Coordinate 'latitude' is not bounded, guessing contiguous bounds.
    warnings.warn(
C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\coords.py:1803:_
    ↵UserWarning: Coordinate 'longitude' is not bounded, guessing contiguous bounds.
    warnings.warn(
C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\coords.py:1803:_
    ↵UserWarning: Coordinate 'latitude' is not bounded, guessing contiguous bounds.
    warnings.warn(
C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\coords.py:1979:_
    ↵UserWarning: Collapsing a non-contiguous coordinate. Metadata may not be fully_
    ↵descriptive for 'sample'.
    warnings.warn(msg.format(self.name()))
C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\coords.py:1979:_
    ↵UserWarning: Collapsing a non-contiguous coordinate. Metadata may not be fully_
    ↵descriptive for 'job'.
    warnings.warn(msg.format(self.name()))
C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\coords.py:1803:_
    ↵UserWarning: Coordinate 'longitude' is not bounded, guessing contiguous bounds.
    warnings.warn(
C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\coords.py:1803:_
    ↵UserWarning: Coordinate 'latitude' is not bounded, guessing contiguous bounds.
    warnings.warn(
[9]: Text(0.5, 1.0, 'Difference')
```



## 5.3 Random Forest Example: Cloud-resolving model sensitivity

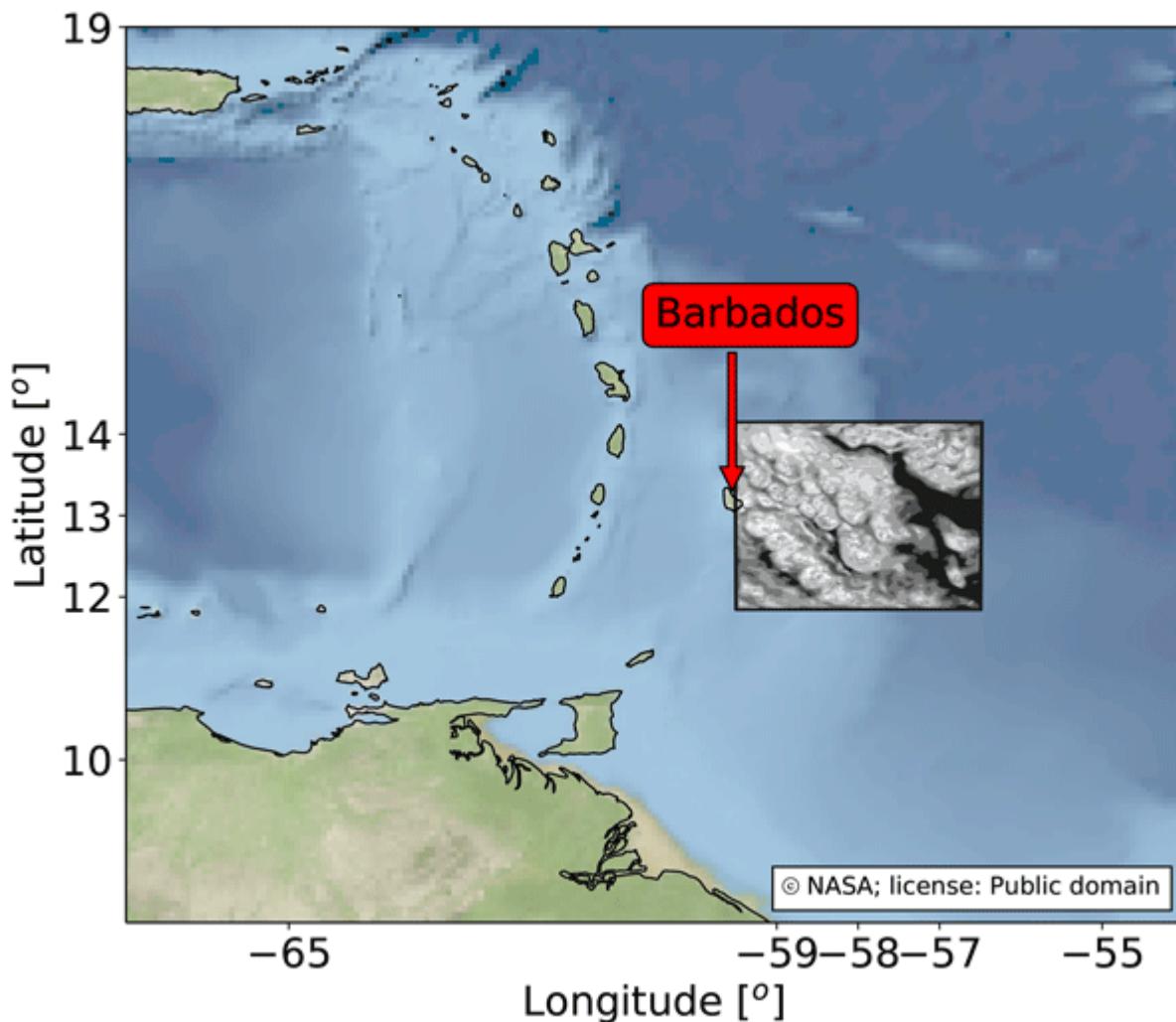
In this example, we will use an ensemble of large-domain simulations of realistic shallow cloud fields to explore the sensitivity of shallow precipitation to local changes in the environment.

The simulation data we use for training the emulator is taken from a recent study Dagan and Stier (2020), where they performed ensemble daily simulations for one month-long period during December 2013 over the ocean to the East of Barbados, such that they sampled variability associated with shallow convection. Each day of the month consisted of two runs, both forced by realistic boundary conditions taken from reanalysis, but with different cloud droplet number concentrations (CDNC) to represent clean and polluted conditions. The altered CDNC was found to have little impact on the precipitation rate in the simulations, and so we simply treat the CDNC change as a perturbation to the initial conditions, and combine the two CDNC runs from each day together to increase the amount of data available for training the emulator. At hourly resolution, this provides us with 1488 data points.

However, given that the amount of precipitation is strongly tied to the local cloud regime, not fully controlling for cloud regime can introduce spurious correlations when training the emulator. As such we also filter out all hours which are not associated with shallow convective clouds. To do this, we consider domain-mean vertical profiles of total cloud water content (liquid + ice),  $q_t$ , and filter out all hours where the vertical sum of  $q_t$  below 600hPa exceeds  $10^{-6}$  kg/kg. This

condition allows us to filter out hours associated with the onset and development of deep convection in the domain, as well as masking out hours with high cirrus layers or hours dominated by transient mesoscale convective activity which is advected in by the boundary conditions. After this, we are left with 850 hourly data point which meet our criteria and can be used to train the emulator.

In this example we emulate hourly precip using domain-mean features: liquid water path (LWP), geopotential height at 700hPa ( $z:\text{math:}_700$ ), Estimated Inversion Strength (EIS), sea-surface temperature (SST) and the vertical pressure velocity at 700hPa ( $w700$ ).



#### References:

Dagan, G. and Stier, P.: Ensemble daily simulations for elucidating cloud–aerosol interactions under a large spread of realistic environmental conditions, *Atmos. Chem. Phys.*, 20, 6291–6303, <https://doi.org/10.5194/acp-20-6291-2020>, 2020.

```
[1]: import numpy as np
import pandas as pd
import iris

from utils import get_crm_data
from esem.utils import plot_results, prettyfify_plot, add_121_line, leave_one_out
```

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```
from esem import rf_model

from matplotlib import pyplot as plt
plt.style.use('default')
%matplotlib inline
```

Concatenate 20cdnc and 200cdnc runs into one dataframe

```
[2]: df_crm = get.crm_data()
df_crm
```

	precip	pres_msl	LWP	WS10	lhfl_s	shfl_s	\
0	0.004593	101407.410	0.035898	6.639860	-167.53857	5.745860	
1	0.006900	101356.266	0.044468	6.822748	-176.93939	4.438721	
2	0.008916	101316.420	0.051559	6.798490	-182.61697	3.649221	
3	0.008932	101270.490	0.057509	6.756970	-188.87599	3.033055	
4	0.016204	101256.270	0.064226	6.763690	-194.85498	2.826119	
..	...	...	...	...	...	...	...
845	0.063121	101309.750	0.064794	8.253145	-191.23718	12.219704	
846	0.064601	101303.110	0.063914	8.326073	-192.57118	11.947702	
847	0.046773	101332.234	0.059974	8.404624	-193.80084	12.372276	
848	0.056623	101394.280	0.062895	8.385845	-192.18195	13.336615	
849	0.064975	101438.690	0.069100	8.429897	-192.28928	13.679647	
							\
	LTS	w500	w600	w700	wmax850	wstd850	zg500
0	13.180252	-0.014463	-0.012311	-0.010275	-0.000024	0.000947	56627.516
1	13.279678	-0.015064	-0.012710	-0.008676	0.000030	0.000382	56572.645
2	13.333527	-0.014811	-0.012014	-0.006025	0.000642	0.000511	56525.613
3	13.328018	-0.013470	-0.012141	-0.004758	0.001519	0.000476	56471.332
4	13.317032	-0.010917	-0.011119	-0.003158	0.003252	0.000958	56443.758
..	...	...	...	...	...	...	...
845	10.142059	-0.024480	-0.006400	-0.007968	-0.000044	0.001105	56084.273
846	10.162674	-0.019426	0.000300	-0.003904	-0.000034	0.000588	56071.547
847	10.166580	-0.014384	0.004355	-0.000284	-0.000251	0.000650	56079.492
848	10.149658	-0.016936	0.002702	0.000667	0.000013	0.000509	56117.140
849	10.164475	-0.021005	0.000413	0.000406	-0.000102	0.000838	56146.297
							\
	zg600	zg700	rh850	rh700	u_shear	EIS	\
0	42694.220	30541.566	67.243774	60.067740	-4.662799	0.989443	
1	42640.473	30488.172	69.299180	58.453730	-4.322696	1.130803	
2	42593.594	30442.703	71.522900	56.912193	-3.925541	1.242463	
3	42539.062	30390.662	74.115690	55.652990	-3.556973	1.304206	
4	42510.805	30364.852	77.510765	54.434470	-3.319007	1.362710	
..	...	...	...	...	...	...	...
845	42214.140	30222.945	83.696740	77.278465	-5.993636	-2.696190	
846	42206.734	30214.715	84.196236	77.536760	-5.848422	-2.673406	
847	42225.984	30236.312	84.394960	77.754560	-5.663757	-2.643809	
848	42272.740	30286.500	84.437530	78.009740	-5.427930	-2.635981	
849	42305.730	30322.684	84.389620	78.030650	-5.215088	-2.612224	
							\
	SST						
0	301.173248						
1	301.173248						

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```

2    301.173248
3    301.173248
4    301.173248
...
845  300.126465
846  300.126465
847  300.126465
848  300.126465
849  300.126465

[850 rows x 20 columns]

```

Extract the precipitation timeseries as target data

```
[3]: precip = df_crm['precip'].to_numpy().reshape(-1, 1)
```

### 5.3.1 Visualize the precipitation landscape

In the ensemble, shallow precipitation is highly correlated with many different physical features. Most obviously there is a high correlation with liquid water path (LWP), 10-metre windspeed (WS10) and geopotential height at 700hPa ( $z_{700}$ ).

We can use these correlations to create “collapsing spaces” for investigating the relationships between shallow precipitation and the local meteorological environment.

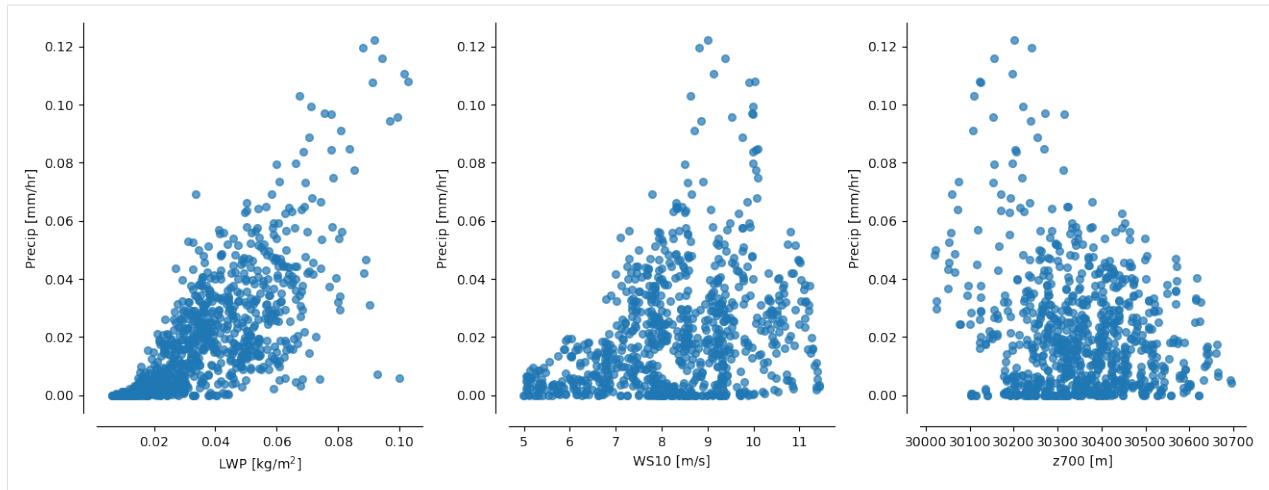
```
[4]: fig, axs = plt.subplots(ncols=3, figsize=(13,5), dpi=100, gridspec_kw={'width_ratios':[1, 1, 1]})

axs[0].scatter(df_crm['LWP'], df_crm['precip'], s=30, marker='o', alpha=0.7, label=r'CDNC  
↔$_{20}$ and CDNC$_{200}$')
axs[0].set_xlabel('LWP [kg/m$^2$]')
axs[0].set_ylabel('Precip [mm/hr]')
prettify_plot(axs[0])

axs[1].scatter(df_crm['WS10'], df_crm['precip'], s=30, marker='o', alpha=0.7, label=r'  
↔'CDNC$_{20}$ and CDNC$_{200}$')
axs[1].set_xlabel('WS10 [m/s]')
axs[1].set_ylabel('Precip [mm/hr]')
prettify_plot(axs[1])

axs[2].scatter(df_crm['zg700'], df_crm['precip'], s=30, marker='o', alpha=0.7, label=r'  
↔'CDNC$_{20}$ and CDNC$_{200}$')
axs[2].set_xlabel('z700 [m]')
axs[2].set_ylabel('Precip [mm/hr]')
prettify_plot(axs[2])

fig.tight_layout()
plt.savefig("Figs/1hr_D13shal_lwp-ws-pr.png", dpi=200)
```



Also, good to note that each of these predictors ``(LWP, WS10, z700)`` are mutually uncorrelated (see plots below)

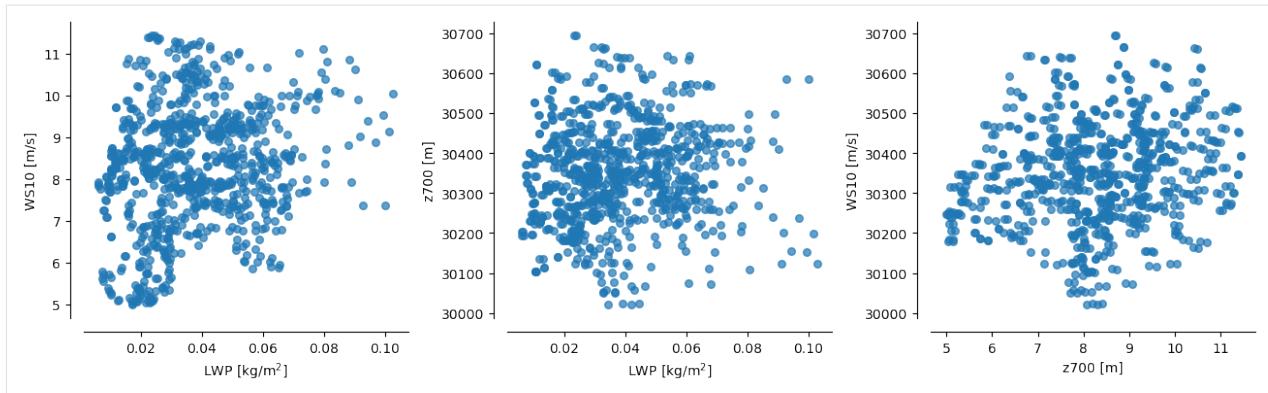
```
[5]: fig, axs = plt.subplots(ncols=3, figsize=(13,4), dpi=100, gridspec_kw={'width_ratios':[1, 1, 1]})

axs[0].scatter(df_crm['LWP'], df_crm['WS10'], s=30, marker='o', alpha=0.7, label=r'CDNC$_{20}$ and CDNC$_{200}$')
axs[0].set_xlabel('LWP [kg/m$^{2}$]')
axs[0].set_ylabel('WS10 [m/s]')
prettify_plot(axs[0])

axs[1].scatter(df_crm['LWP'], df_crm['zg700'], s=30, marker='o', alpha=0.7, label=r'CDNC$_{20}$ and CDNC$_{200}$')
axs[1].set_xlabel('LWP [kg/m$^{2}$]')
axs[1].set_ylabel('z700 [m]')
prettify_plot(axs[1])

axs[2].scatter(df_crm['WS10'], df_crm['zg700'], s=30, marker='o', alpha=0.7, label=r'CDNC$_{20}$ and CDNC$_{200}$')
axs[2].set_xlabel('z700 [m]')
axs[2].set_ylabel('WS10 [m/s]')
prettify_plot(axs[2])

fig.tight_layout()
plt.savefig("Figs/1hr_D13shal_lwp-ws-pr.png", dpi=200)
```



**Stratifying precip by pairs of these predictors creates nice “collapsing spaces”**

Nice for illustrating how the emulated surface compares to the raw data

```
[6]: fig, axs = plt.subplots(ncols=3, figsize=(13,5), dpi=100, gridspec_kw={'width_ratios':[1, 1, 0.05]})

sc1 = axs[0].scatter(df_crm['LWP'], df_crm['zg700'], c=df_crm['precip'],
                     vmin=0, vmax=0.12, s=30, marker='o', alpha=0.7, label=r'CDNC_{20} and CDNC_{200}')
axs[0].set_xlabel(r'LWP [kg m^{-2}]')
axs[0].set_ylabel(r'z_{700} [m]')

axs[0].set_title("Hourly output: Dec2013, shallow clouds")
axs[0].legend(loc='lower right')

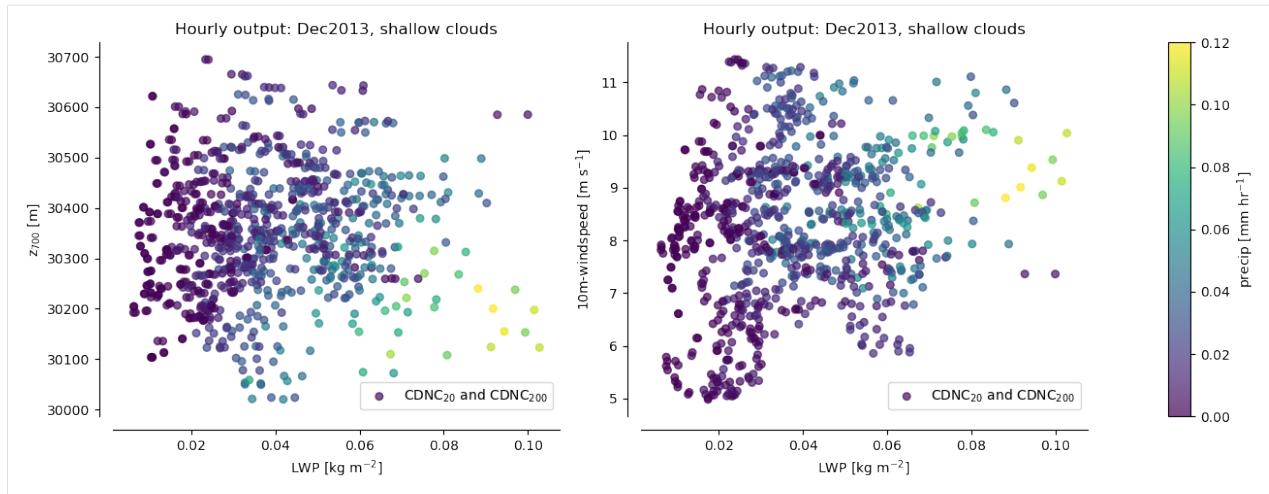
prettify_plot(axs[0])

sc2 = axs[1].scatter(df_crm['LWP'], df_crm['WS10'], c=df_crm['precip'],
                     vmin=0, vmax=0.12, s=30, marker='o', alpha=0.7, label=r'CDNC_{20} and CDNC_{200}')
axs[1].set_xlabel(r'LWP [kg m^{-2}]')
axs[1].set_ylabel(r'10m-windspeed [m s^{-1}]')

axs[1].set_title("Hourly output: Dec2013, shallow clouds")
axs[1].legend(loc='lower right')

prettify_plot(axs[1])

fig.colorbar(sc1, cax=axs[2], label=r'precip [mm hr^{-1}]')
fig.tight_layout()
# plt.savefig("Figs/1hr_D13shal_lwp-ws-pr.png", dpi=200)
```



### 5.3.2 Emulation

Our aim is to emulate shallow precipitation as a function of the environmental conditions, and then plot the predictions in LWP-z700 space to compare with the scatter points above.

To do this we choose a set of predictors which are typical “cloud-controlling factors” such as SST, Estimated Inversion Strength, vertical velocity at 700 hPa, LWP and z700. Other variables could also be chosen and it’s worth exploring this just to get a sense for how the model behaves.

After validating the model using Leave-One-Out cross-validation, we then retrain the model using the full dataset, and use this model to predict the precipitation across a wide range of values. Finally, for the purpose of plotting in LWP-z700 space, we reduce the dimensionality of our final prediction by averaging over all features with aren’t LWP or z700. This gives us a smooth field to compare with the scatter points.

```
[7]: params = df_crm.loc[:, ['LWP', 'zg700', 'EIS', 'SST', 'w700']]

print("The input params are: \n", params, "\n")
The input params are:
      LWP      zg700       EIS       SST      w700
0  0.035898  30541.566  0.989443  301.173248 -0.010275
1  0.044468  30488.172  1.130803  301.173248 -0.008676
2  0.051559  30442.703  1.242463  301.173248 -0.006025
3  0.057509  30390.662  1.304206  301.173248 -0.004758
4  0.064226  30364.852  1.362710  301.173248 -0.003158
..    ...
845 0.064794  30222.945 -2.696190  300.126465 -0.007968
846 0.063914  30214.715 -2.673406  300.126465 -0.003904
847 0.059974  30236.312 -2.643809  300.126465 -0.000284
848 0.062895  30286.500 -2.635981  300.126465  0.000667
849 0.069100  30322.684 -2.612224  300.126465  0.000406
[850 rows x 5 columns]
```

### LeaveOneOut cross-validation and plotting

```
[8]: %time

# Ignore the mountain of warnings
import warnings
from sklearn.exceptions import DataConversionWarning
warnings.filterwarnings(action='ignore', category=DataConversionWarning)

outp = np.asarray(leave_one_out(Xdata=params, Ydata=precip, model='RandomForest', n_
estimators=50, random_state=0))
```

Wall time: 2min 12s

```
<timed exec>:6: VisibleDeprecationWarning: Creating an ndarray from ragged nested_
sequences (which is a list-or-tuple of lists-or-tuples-or ndarrays with different_
lengths or shapes) is deprecated. If you meant to do this, you must specify
'dtype=object' when creating the ndarray.
```

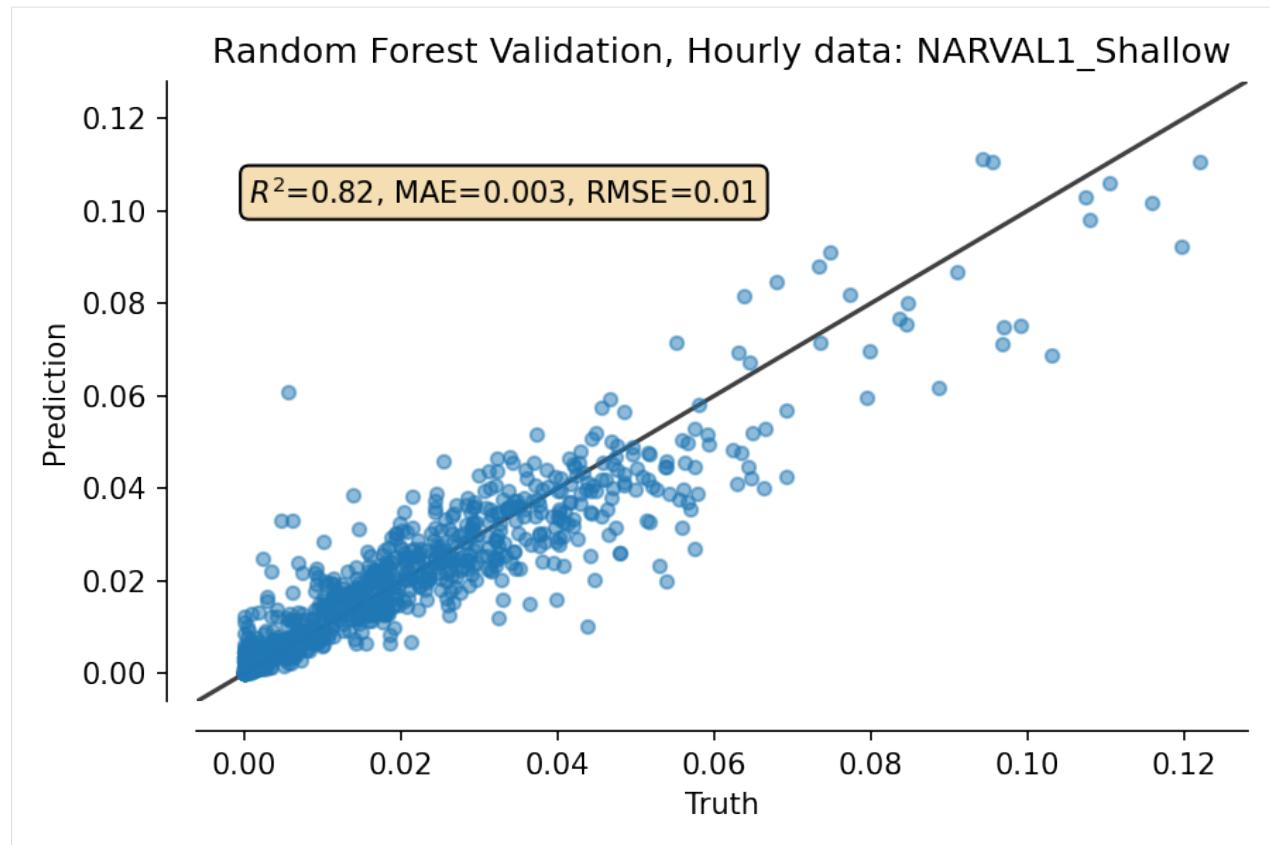
```
[9]: truth_RF, pred_RF = outp[:,0], outp[:,1]
```

```
[10]: from sklearn.metrics import mean_squared_error

""" Validation plot """
fig, ax = plt.subplots(dpi=150)

plot_results(ax, truth_RF, pred_RF, title="Random Forest Validation, Hourly data:_"
_NARVAL1_Shallow")

fig.tight_layout()
```



Now, retrain model on all data, and extrapolate over whole space

```
[11]: X_train = params.to_numpy()
Y_train = precip.ravel()
model = rf_model(X_train, Y_train)
```

```
[12]: model.train()
```

```
[13]: %time

# Now, make grid for plotting RF predictions
# more n_points means higher resolution, but takes exponentially longer
n_points = 30

min_vals = params.min()
max_vals = params.max()

# For uniform prediction over full params space
space=np.linspace(min_vals, max_vals, n_points)

# Reshape to (N,D)
reshape_to_ND = np.transpose(space)
Xs_uniform = np.meshgrid(*reshape_to_ND)
test = np.array([_.flatten() for _ in Xs_uniform]).T
```

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```
# Predict
predictions, _ = model.predict(test)
predictions = predictions.reshape(Xs_uniform[0].shape)

# Now, take mean over all parameters except [LWP, zg700], assumed to be first 2 indices
predictions_reduced = np.mean(predictions, axis=tuple(range(2, predictions.ndim)))

Wall time: 1min 35s
```

[14]: # Now, make grid for plotting RF predictions

```
LWP_grid = np.linspace(min_vals['LWP'], max_vals['LWP'], num=n_points)
zg700_grid = np.linspace(min_vals['zg700'], max_vals['zg700'], num=n_points)

lwp, zg = np.meshgrid(LWP_grid, zg700_grid)
```

### 5.3.3 Plot!

[15]:

```
fig, ax = plt.subplots(ncols=3, figsize=(7,4), dpi=250, gridspec_kw={'width_ratios':[1, 0.05, 0.05]})

fig.suptitle("Hourly output: Dec2013, shallow clouds")

cp = ax[0].pcolormesh(LWP_grid, zg700_grid,
                      predictions_reduced, vmin=0, vmax=0.12, alpha=1)

fig.colorbar(cp, cax=ax[1], orientation='vertical', shrink=0.05, label=r'Precip [mm hr$^{-1}$]')

"""Overlap errors"""
ax[0].scatter(df_crm['LWP'], df_crm['zg700'], c=df_crm['precip'],
              vmin=0, vmax=0.12, s=30, marker='o', edgecolors="None")

ers = ax[0].scatter(df_crm['LWP'], df_crm['zg700'], c=(truth_RF-pred_RF)/(truth_RF+pred_RF),
                     facecolors="None",
                     vmin=-1, vmax=1, s=30, marker='o', lw=0.7, alpha=0.5, cmap='seismic')
ers.set_facecolors("None")
fig.colorbar(ers, cax=ax[2], label=r'$\frac{\text{Truth-Prediction}}{\text{Truth+Prediction}}$')

for idx, _ in enumerate(ax[:1]):
    _.set_xlabel(r'LWP [kg m$^{-2}$]')
    _.set_ylabel(r'z$_{700}$ [m]')

    if idx==0:
        _.set_xlim(min_vals['LWP'], max_vals['LWP'])
        _.set_ylim(min_vals['zg700'], max_vals['zg700'])

fig.tight_layout()
```

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```

fig.subplots_adjust(top=0.85) # Put this AFTER tight_layout() call !

"""\Add validation plot as inset to first axis"""
axins = ax[0].inset_axes([0.79, 0.79, 0.2, 0.2], # x0, y0, width, height
                       transform=ax[0].transAxes)
axins.scatter(truth_RF, pred_RF, s=2, alpha=0.3)
add_121_line(axins)
axins.set_xlabel('Model', position=(0.5,0), fontsize=8, labelpad=-0.01)
axins.set_xticks([0, 0.05, 0.1])
axins.set_xticklabels(labels=[0, 0.05, 0.1], fontdict={'fontsize':6})

axins.set_ylabel('Emulator', position=(0,0.5), fontsize=8, labelpad=-0.01)
axins.set_yticks([0, 0.05, 0.1])
axins.set_yticklabels(labels=[0, 0.05, 0.1], fontdict={'fontsize':6})

axins.tick_params(axis='both', which='major', pad=0.01)

# plt.savefig("./Figs/1hr_D13shal_lwp-zg700-pr_w-errorpoints.png", dpi=300)

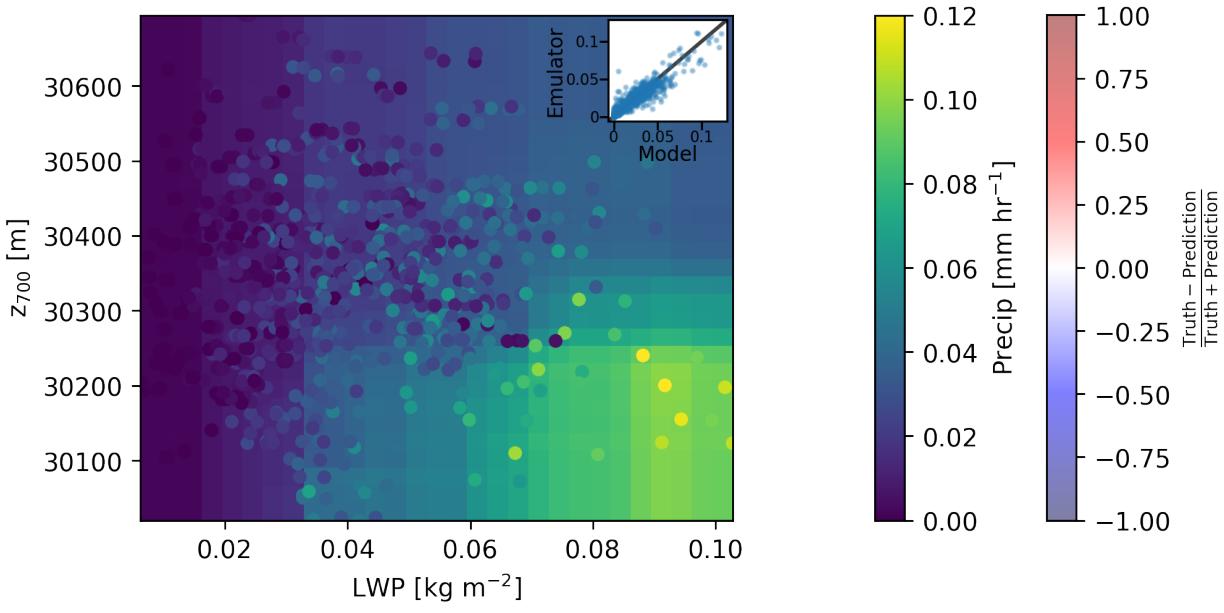
```

```

C:\Users\duncan\AppData\Local\Temp\ipykernel_21624\1073200818.py:4: UserWarning: MatplotlibDeprecationWarning: shading='flat' when X and Y have the same dimensions as C is deprecated since 3.3. Either specify the corners of the quadrilaterals with X and Y, or pass shading='auto', 'nearest' or 'gouraud', or set rcParams['pcolor.shading']'. This will become an error two minor releases later.
cp = ax[0].pcolormesh(LWP_grid, zg700_grid,

```

Hourly output: Dec2013, shallow clouds



[ ]:

## 5.4 Calibrating GPs using ABC

```
[1]: import warnings
warnings.filterwarnings('ignore') # Ignore all the iris warnings...

[2]: import pandas as pd
import cis
import iris

from utils import get_aeronet_data, get_bc_ppe_data

from esem.utils import validation_plot, plot_parameter_space, get_random_params,_
    ensemble_collocate
from esem import gp_model
from esem.abc_sampler import ABCSampler, constrain

import iris.quickplot as qplt
import matplotlib.pyplot as plt
%matplotlib inline
```

### 5.4.1 Read in the parameters and observables

```
[3]: aaod = get_aeronet_data()
print(aaod)

Ungridded data: Absorption_AOD440nm / (1)
Shape = (10098,)

Total number of points = 10098
Number of non-masked points = 10098
Long name = Absorption_AOD440nm
Standard name = None
Units = 1
Missing value = -999.0
Range = (5.1e-05, 0.47236)
History =
Coordinates:
longitude
    Long name =
    Standard name = longitude
    Units = degrees_east
    Missing value = None
    Range = (-155.576755, 141.3407)
    History =
latitude
    Long name =
    Standard name = latitude
    Units = degrees_north
    Missing value = None
    Range = (-35.495807, 79.990278)
```

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```

History =
time
  Long name =
  Standard name = time
  Units = days since 1600-01-01 00:00:00
  Missing value = None
  Range = (cftime.DatetimeGregorian(2017, 1, 1, 12, 0, 0, has_year_
zero=False), cftime.DatetimeGregorian(2018, 1, 2, 12, 0, 0, has_year_zero=False))
History =

```

[4]: # Read in the PPE parameters, AAOD and DRE  
`ppe_params, ppe_aaod, ppe_dre = get_bc_ppe_data(dre=True)`

[5]: # Take the annual mean of the DRE  
`ppe_dre, = ppe_dre.collapsed('time', iris.analysis.MEAN)`

WARNING:root:Creating guessed bounds as none exist in file  
WARNING:root:Creating guessed bounds as none exist in file  
WARNING:root:Creating guessed bounds as none exist in file  
WARNING:root:Creating guessed bounds as none exist in file

## 5.4.2 Collocate the model on to the observations

[6]: `col_ppe_aaod = ensemble_collocate(ppe_aaod, aaod)`

[7]: `n_test = 8`  
`X_test, X_train = ppe_params[:n_test], ppe_params[n_test:]`  
`Y_test, Y_train = col_ppe_aaod[:n_test], col_ppe_aaod[n_test:]`

[8]: `Y_train`  
[8]: <iris 'Cube' of Absorption optical thickness - total 550nm / (1) (job: 31; obs: 10098)>

## 5.4.3 Setup and run the models

### Explore different model choices

[9]: `from esem.utils import leave_one_out, prediction_within_ci`  
`from scipy import stats`  
`import numpy as np`  
`from esem.data_processors import Log`  
`res_l = leave_one_out(X_train, Y_train, model='GaussianProcess', data_`  
`_processors=[Log(constant=0.1)], kernel=['Linear', 'Exponential', 'Bias'])`  
`r2_values_l = [stats.linregress(x.data.compressed(), y.data[:, ~x.data.mask]).`  
`_flatten()**2 for x,y,_ in res_l]`

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```

ci95_values_l = [prediction_within_ci(x.data.flatten(), y.data.flatten(), v.data.
    ↪flatten())[2].sum()/x.data.count() for x,y,v in res_l]
print("Mean R^2: {:.2f}".format(np.asarray(r2_values_l).mean()))
print("Mean proportion within 95% CI: {:.2f}".format(np.asarray(ci95_values_l).mean()))

res = leave_one_out(X_train, Y_train, model='GaussianProcess', kernel=['Linear', 'Bias'])
r2_values = [stats.linregress(x.data.flatten(), y.data.flatten())[2]**2 for x,y,v in res]
ci95_values = [prediction_within_ci(x.data.flatten(), y.data.flatten(), v.data.
    ↪flatten())[2].sum()/x.data.count() for x,y,v in res]
print("Mean R^2: {:.2f}".format(np.asarray(r2_values).mean()))
print("Mean proportion within 95% CI: {:.2f}".format(np.asarray(ci95_values).mean()))

# Note that while the Log pre-processing leads to slightly better R^2, the model is_
    ↪under-confident and_
# has too large uncertainties which would adversely affect our implausibility metric.

Mean R^2: 1.00
Mean proportion within 95% CI: 1.00
Mean R^2: 0.99
Mean proportion within 95% CI: 0.94

```

## Build final model

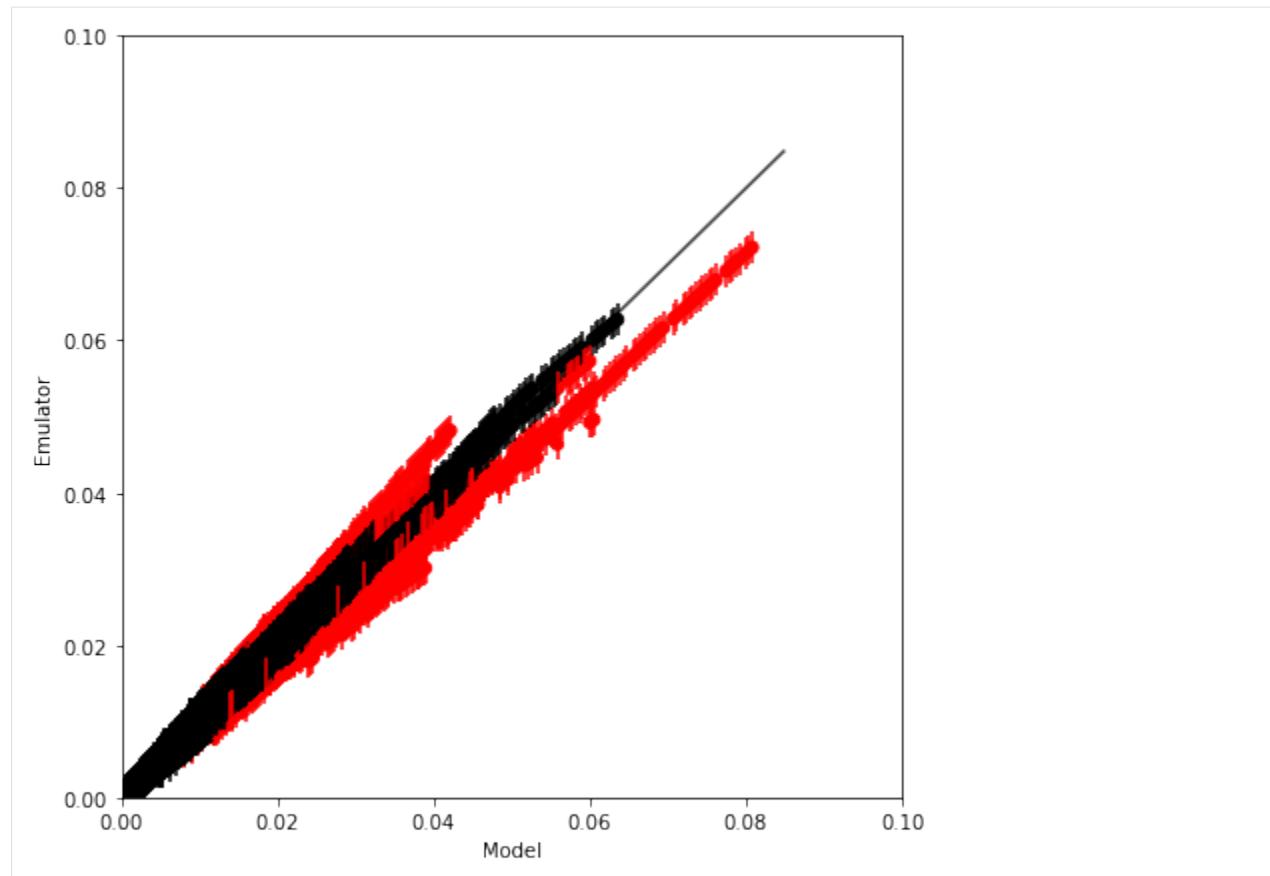
[10]: model = gp\_model(X\_train, Y\_train, kernel=['Linear', 'Bias'])

[11]: model.train()

[12]: m, v = model.predict(X\_test.values)

[13]: validation\_plot(Y\_test.data.flatten(), m.data.flatten(), v.data.flatten(),
 minx=0, maxx=0.1, miny=0., maxy=0.1)

Proportion of 'Bad' estimates : 5.52%



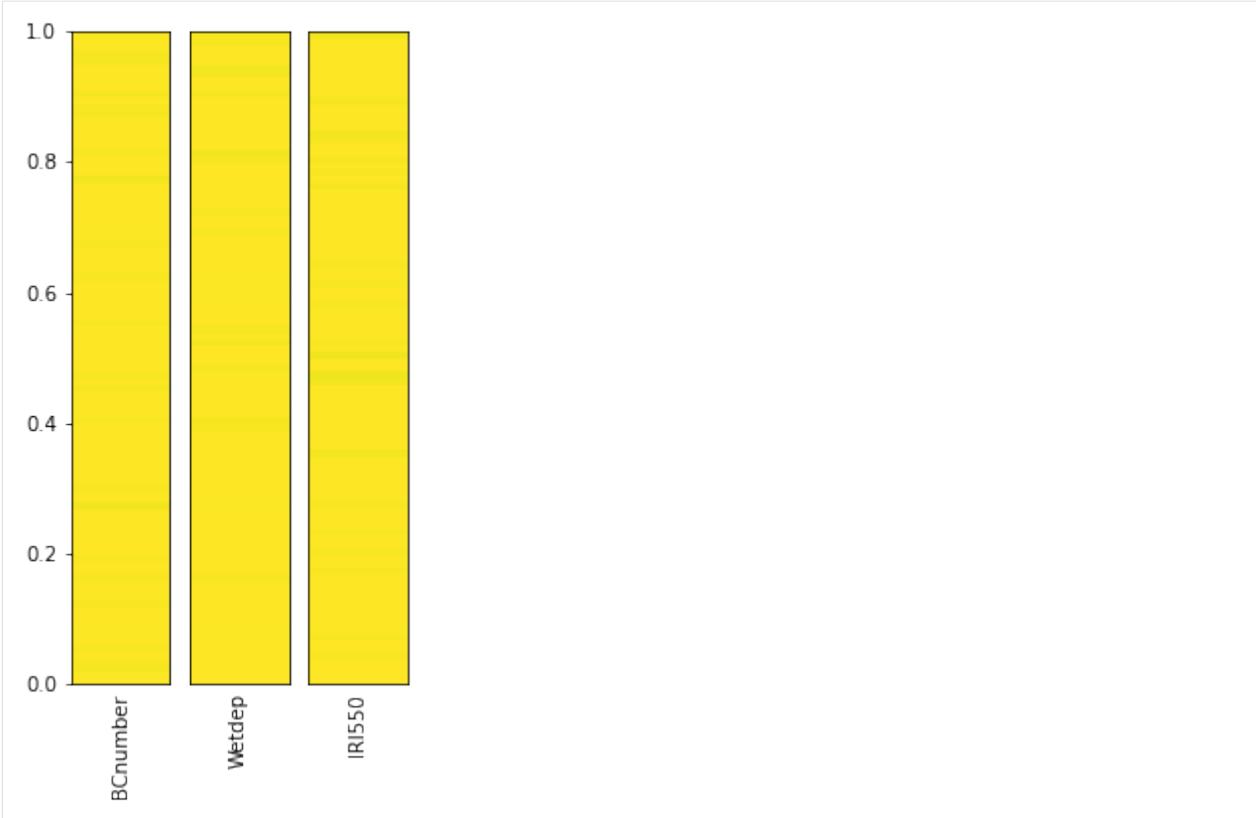
#### 5.4.4 Sample and constrain the models

Emulating 1e6 sample points directly would require 673 Gb of memory so we can either run 1e6 samples for each point, or run the constraint everywhere, but in batches. Here we do the latter, optioanlly on the GPU, using the ‘naive’ algorithm for calculating the running mean and variance of the various properties.

The rejection sampling happens in a similar manner so that only as much memory as is used for one batch is ever used.

```
[14]: # In this case
sample_points = pd.DataFrame(data=get_random_params(3, int(1e6)), columns=X_train.columns)
```

```
[15]: # Note that smoothing the parameter distribution can be slow for large numbers of points
plot_parameter_space(sample_points, fig_size=(3,6), smooth=False)
```



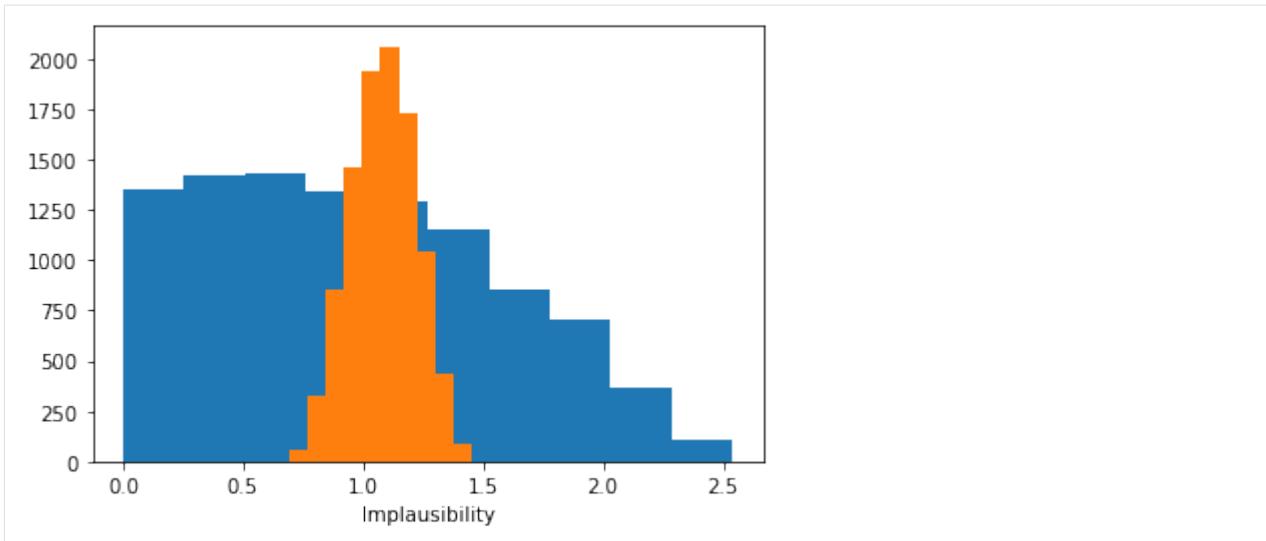
```
[16]: # Setup the sampler to compare against our AeroNet data
sampler = ABCSampler(model, aaod, obs_uncertainty=0.5, repres_uncertainty=0.5)

[17]: # Calculate the implausibility for each sample against each observation - note this can
      # be very large so we only sample a fraction!
implaus = sampler.get_implausibility(sample_points[::-100], batch_size=1000)

# The implausibility distributions for different observations can be very different.
_ = plt.hist(implaus.data[:, 1400])
_ = plt.hist(implaus.data[:, 14])
plt.gca().set(xlabel='Implausibility')

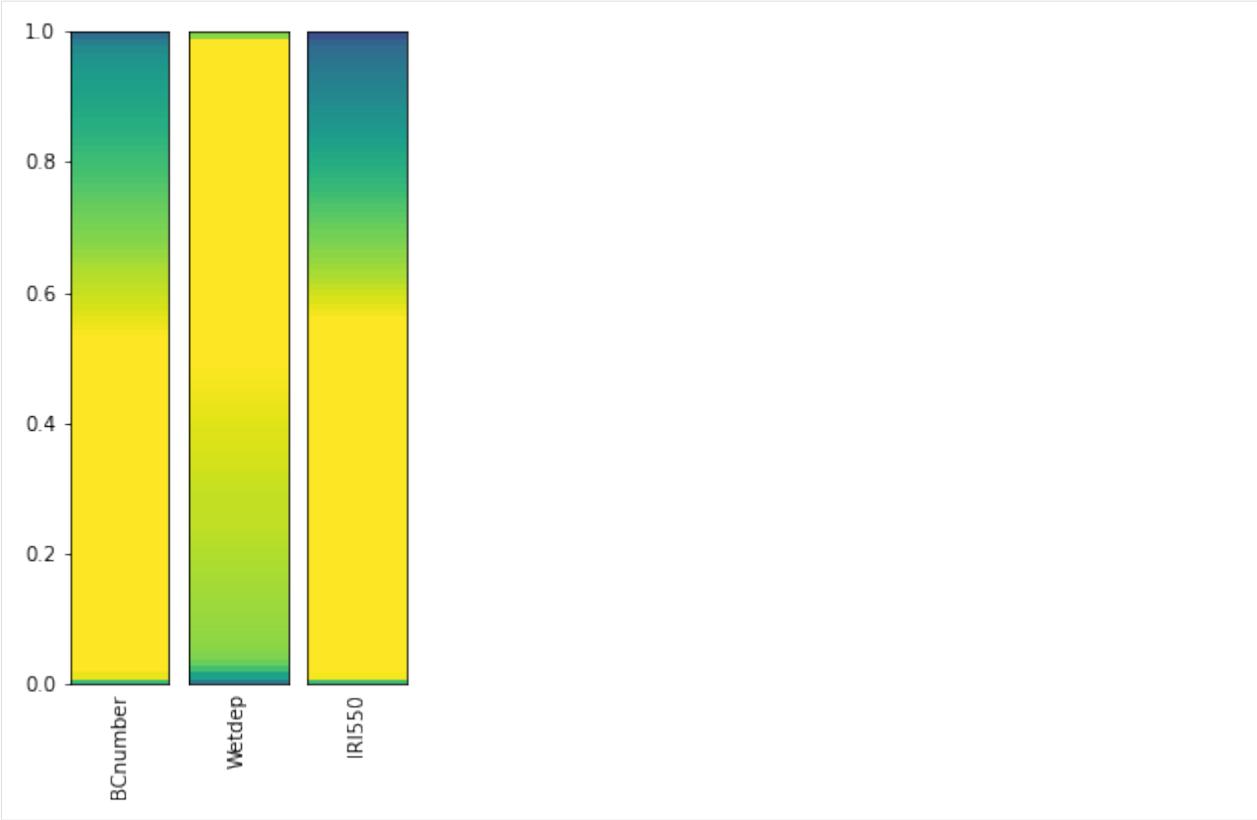
0%| 0/10000 [00:00<?, ?sample/s]
<tqdm.auto.tqdm object at 0x00000236A10BA5E0>

[17]: [Text(0.5, 0, 'Implausibility')]
```



```
[18]: # Find the valid samples in our full 1million samples by comparing against a given
      ↪ tolerance and threshold
valid_samples = sampler.batch_constrain(sample_points, batch_size=10000, tolerance=.1)
print("Remaining points: {}".format(valid_samples.sum()))
0%|          | 0/1000000 [00:00<?, ?sample/s]
<tqdm.auto.tqdm object at 0x000002398006DC70>Remaining points: 729474
```

```
[19]: # Plot the reduced parameter distribution
constrained_sample = sample_points[valid_samples]
plot_parameter_space(constrained_sample, fig_size=(3, 6))
```



```
[20]: # We can also easily plot the joint distributions
# Only plot every one in 100 points as scatter plots with large numbers of points are
# slow...
import matplotlib

# Mimic Seaborn scaling without requiring the whole package
scale = 1.5
matplotlib.rcParams['font.size'] = 12 * scale
matplotlib.rcParams['axes.labelsize'] = 12 * scale
matplotlib.rcParams['axes.titlesize'] = 12 * scale
matplotlib.rcParams['xtick.labelsize'] = 11 * scale
matplotlib.rcParams['ytick.labelsize'] = 11 * scale
matplotlib.rcParams['lines.linewidth'] = 1.5 * scale
matplotlib.rcParams['lines.markersize'] = 6 * scale
#
m, _ = model.predict(constrained_sample[::100].values)
Zs = m.data
# Plot the emulated AAOD value (averaged over observation locations) for each point
grr = pd.plotting.scatter_matrix(constrained_sample[::100], c=Zs.mean(axis=1),
                                 figsize=(12, 10), marker='o',
                                 hist_kwds={'bins': 20,}, s=20, alpha=.8, vmin=1e-3,
                                 vmax=1e-2, range_padding=0.,
                                 density_kwds={'range': [[0., 1.], [0., 1.]], 'colormap':
                                 'viridis'},
                                 )

```

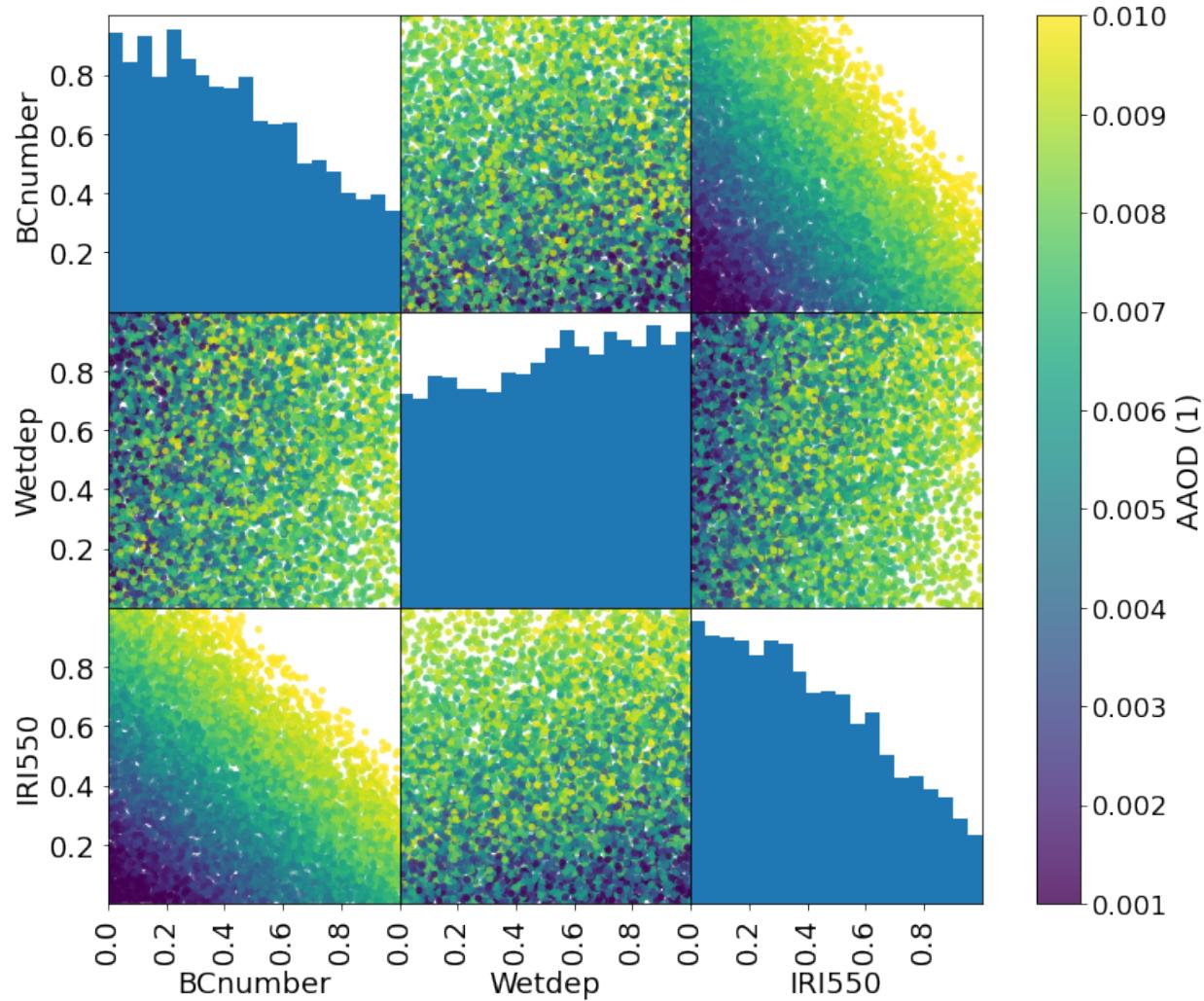
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```
# Matplotlib dragons...
grr[0][0].set_yticklabels([0.2, 0.4, 0.6, 0.8], fontsize=12 * scale)
for i in range(2):
    grr[i+1][0].set_yticklabels([0.0, 0.2, 0.4, 0.6, 0.8], fontsize=12 * scale)
for i in range(3):
    grr[2][i].set_xticks([0.0, 0.2, 0.4, 0.6, 0.8])
    grr[2][i].set_xticklabels([0.0, 0.2, 0.4, 0.6, 0.8], fontsize=12 * scale)

plt.colorbar(grr[0][1].collections[0], ax=grr, use_gridspec=True, label='AAOD (1)')

plt.savefig('BCPPE_constrained_params_paper.png', transparent=True)
```



## 5.4.5 Explore the uncertainty in Direct Radiative Effect of Aerosol in constrained sample-space

```
[21]: dre_test, dre_train = ppe_dre[:n_test], ppe_dre[n_test:]

ari_model = gp_model(X_train, dre_train, name="ARI", kernel=['Linear', 'Bias'])
ari_model.train()
```

```
[22]: # Calculate the mean and std-dev DRE over each set of sample points

unconstrained_mean_ari, unconstrained_sd_ari = ari_model.batch_stats(sample_points, ↴
    ↴batch_size=10000)
constrained_mean_ari, constrained_sd_ari = ari_model.batch_stats(constrained_sample, ↴
    ↴batch_size=10000)

0%|          | 0/1000000 [00:00<?, ?sample/s]
<tqdm.auto.tqdm object at 0x00000239A2B2D310>
0%|          | 0/729474 [00:00<?, ?sample/s]
<tqdm.auto.tqdm object at 0x00000239ACFEFDF0>
```

```
[23]: # The original (unconstrained DRE)
qplt.pcolormesh(unconstrained_sd_ari, vmin=0., vmax=1)
plt.gca().coastlines()
```

```
[23]: <cartopy.mpl.feature_artist.FeatureArtist at 0x239ad149610>

Ensemble standard deviation in mean instantaneous sw forcing all sky top of the atmosphere



A world map showing the ensemble standard deviation in mean instantaneous shortwave forcing (W m-2) for the entire sky at the top of the atmosphere. The map uses a color scale from dark purple (0.0) to bright yellow (1.0). The highest uncertainty is concentrated over the Sahara Desert and parts of East Asia, where values exceed 0.8 W m-2.



0.0 0.2 0.4 0.6 0.8 1.0  
W m-2



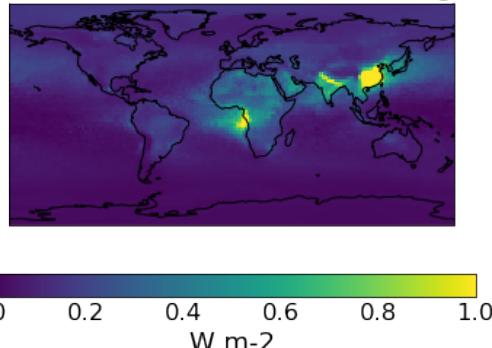
```
[24]: # The constrained DRE
qplt.pcolormesh(constrained_sd_ari, vmin=0., vmax=1)
plt.gca().coastlines()
```



```
[24]: <cartopy.mpl.feature_artist.FeatureArtist at 0x239af7e3700>
```


```

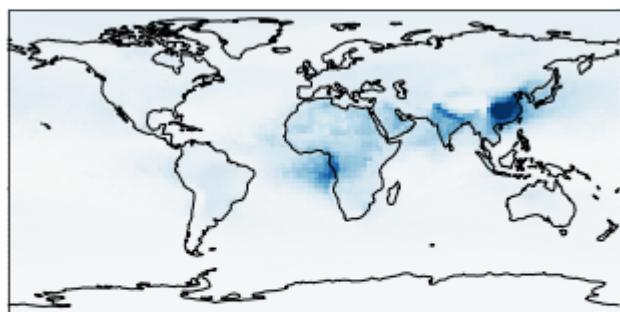
Ensemble standard deviation in mean instantaneous sw forcing all sky top of the atmosphere



```
[25]: # The change in spread after the constraint is applied
qplt.pcolormesh((constrained_sd_ari-unconstrained_sd_ari), cmap='RdBu_r', vmin=-5e-1,
                  vmax=5e-1)
plt.gca().coastlines()
```

```
[25]: <cartopy.mpl.feature_artist.FeatureArtist at 0x239af86c310>
```

Unknown



-0.4    -0.2    0.0    0.2    0.4  
W m<sup>-2</sup>

```
[ ]:
```

## 5.5 Calibrating GPs using MCMC

```
[1]: import os
# Ignore my broken HDF5 install...
os.putenv("HDF5_DISABLE_VERSION_CHECK", '1')
```

```
[2]: import pandas as pd
import numpy as np
import iris
```

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```
from utils import get_aeronet_data, get_bc_ppe_data
from esem import gp_model
from esem.sampler import MCMCSampler

import os

import matplotlib.pyplot as plt
%matplotlib inline

# GPU = "1"

# os.environ["CUDA_VISIBLE_DEVICES"] = GPU
```

### 5.5.1 Read in the parameters and observables

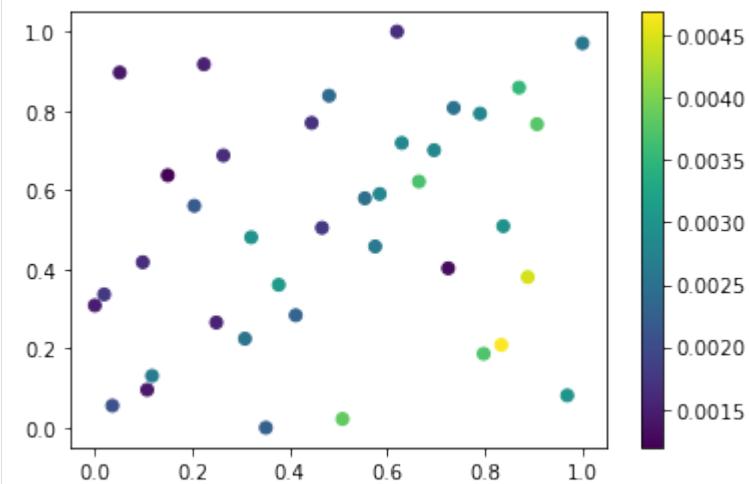
[3]: ppe\_params, ppe\_aaod = get\_bc\_ppe\_data()

[4]: # Calculate the global, annual mean AAOD (CIS will automatically apply the weights)
mean\_aaod, = ppe\_aaod.collapsed(['latitude', 'longitude', 'time'], 'mean')

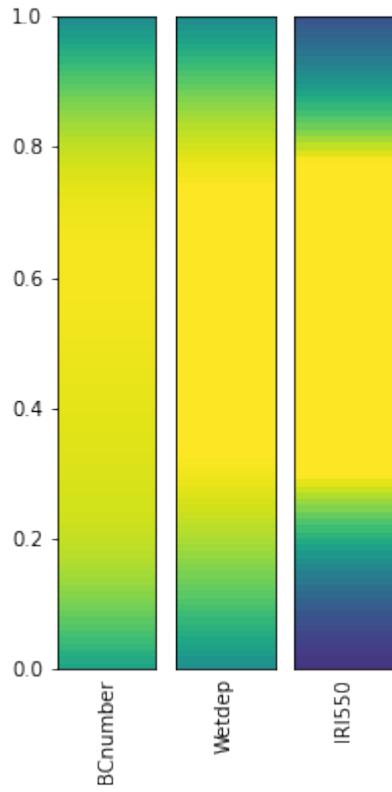
WARNING:root:Creating guessed bounds as none exist in file  
 WARNING:root:Creating guessed bounds as none exist in file  
 WARNING:root:Creating guessed bounds as none exist in file  
 C:\Users\duncan\miniconda3\envs\climatebench\lib\site-packages\iris\analysis\cartography.  
 ↵py:394: UserWarning: Using DEFAULT\_SPHERICAL\_EARTH\_RADIUS.  
 warnings.warn("Using DEFAULT\_SPHERICAL\_EARTH\_RADIUS.")

[5]: plt.scatter(ppe\_params.BCnumber, ppe\_params.Wetdep, c=mean\_aaod.data)  
 plt.colorbar()

[5]: <matplotlib.colorbar.Colorbar at 0x1807b701fd0>



```
[6]: from esem.utils import plot_parameter_space
plot_parameter_space(ppe_params, fig_size=(3,6))
c:\users\duncan\pycharmprojects\gcem\esem\utils.py:119: MatplotlibDeprecationWarning:
shading='flat' when X and Y have the same dimensions as C is deprecated since 3.3. 
Either specify the corners of the quadrilaterals with X and Y, or pass shading='auto',
'nearest' or 'gouraud', or set rcParams['pcolor.shading']. This will become an error
two minor releases later.
ax.pcolor(X, Y, vals[:, np.newaxis], vmin=0, vmax=1)
```



```
[7]: n_test = 8
X_test, X_train = ppe_params[:n_test], ppe_params[n_test:]
Y_test, Y_train = mean_aaod[:n_test], mean_aaod[n_test:]
```

## 5.5.2 Setup and run the models

```
[8]: model = gp_model(X_train, Y_train)
WARNING: Using default kernel - be sure you understand the assumptions this implies.
Consult e.g. http://www.cs.toronto.edu/~duvenaud/cookbook/ for an excellent
description of different kernel choices.
```

```
[9]: model.train()
```

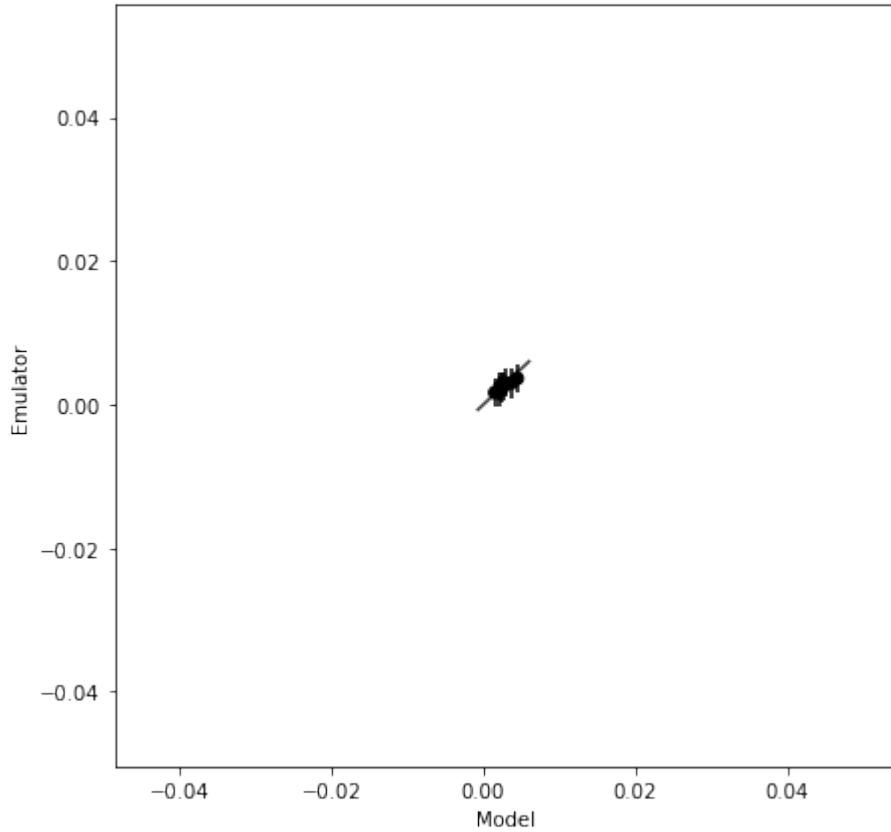
```
[10]: m, v = model.predict(X_test.values)
```

```
[11]: Y_test.data
```

```
[11]: masked_array(data=[0.002822510314728863, 0.002615034735649063,
                      0.002186747212596059, 0.0015112621889264352,
                      0.004456776854431466, 0.0025203727123839117,
                      0.0022378865435589133, 0.003730134076583199],
                  mask=[False, False, False, False, False, False, False, False],
                  fill_value=1e+20)
```

```
[12]: from esem.utils import validation_plot
```

```
validation_plot(Y_test.data.flatten(), m.data.flatten(), v.data.flatten())
Proportion of 'Bad' estimates : 0.00%
```



```
[13]: # Set the objective as one of the test datasets
sampler = MCMCSampler(model, Y_test[0])
```

```
[14]: samples = sampler.sample(n_samples=800, mcmc_kwargs=dict(num_burnin_steps=100) )
Acceptance rate: 0.9262387969566703
```

```
[15]: new_samples = pd.DataFrame(data=samples, columns=ppe_params.columns)
m, _ = model.predict(new_samples.values)
```

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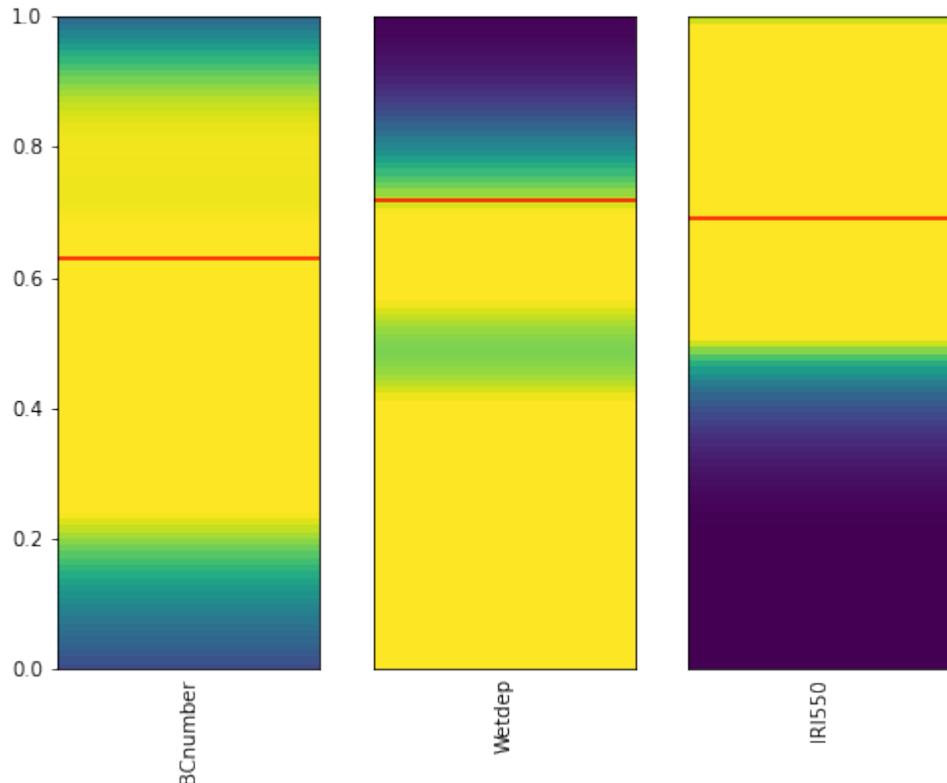
Zs = m.data

```
[16]: print("Sample mean: {}".format(Zs.mean()))
print("Sample std dev: {}".format(Zs.std()))
```

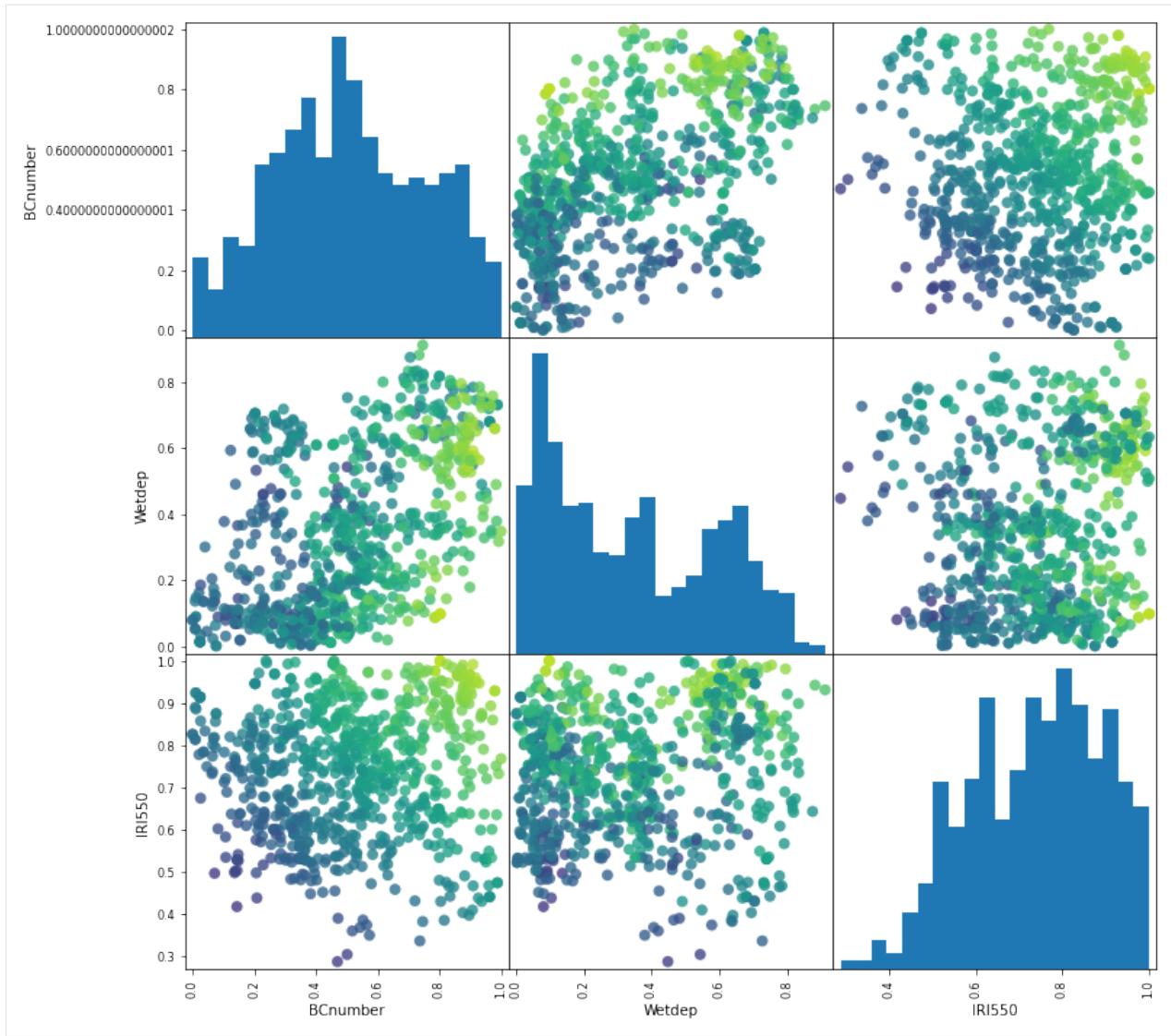
Sample mean: 0.003150297212253644  
 Sample std dev: 0.0006305448129278108

```
[17]: plot_parameter_space(new_samples, target_df=X_test.iloc[0])
```

```
c:\users\duncan\pycharmprojects\gcm\esem\utils.py:119: MatplotlibDeprecationWarning:
  ~shading='flat' when X and Y have the same dimensions as C is deprecated since 3.3. ~
  ~Either specify the corners of the quadrilaterals with X and Y, or pass shading='auto', ~
  ~'nearest' or 'gouraud', or set rcParams['pcolor.shading']. This will become an error~
  ~two minor releases later.
  ax.pcolor(X, Y, vals[:, np.newaxis], vmin=0, vmax=1)
```



```
[18]: grr = pd.plotting.scatter_matrix(new_samples, c=Zs, figsize=(12, 12), marker='o',
                                     hist_kwds={'bins': 20}, s=60, alpha=.8, vmin=1e-3, ~
                                     ~vmax=5e-3)
```



```
[19]: from esem.abc_sampler import ABCSampler, constrain
from esem.utils import get_random_params
```

```
[20]: sampler = ABCSampler(model, Y_test[0])

samples = sampler.sample(n_samples=2000, threshold=0.5)
valid_points = pd.DataFrame(data=samples, columns=ppe_params.columns)
Acceptance rate: 0.33641715727502103
```

```
[21]: m, _ = model.predict(valid_points.values)
Zs = m.data
```

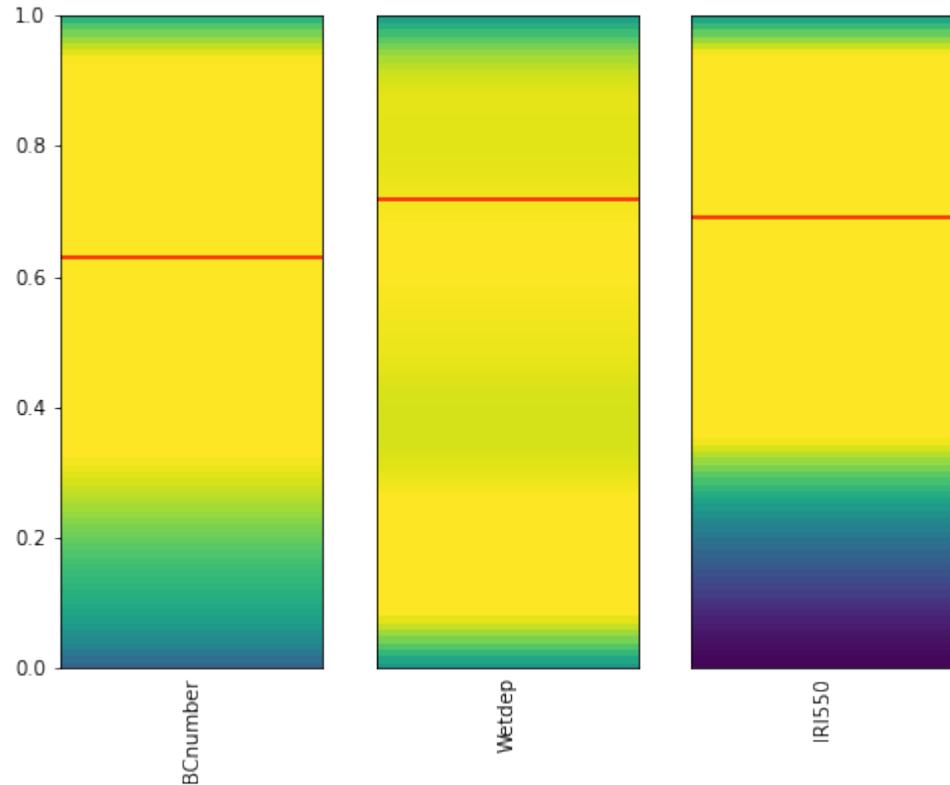
```
[22]: print("Sample mean: {}".format(Zs.mean()))
print("Sample std dev: {}".format(Zs.std()))
Sample mean: 0.002788395703569595
```

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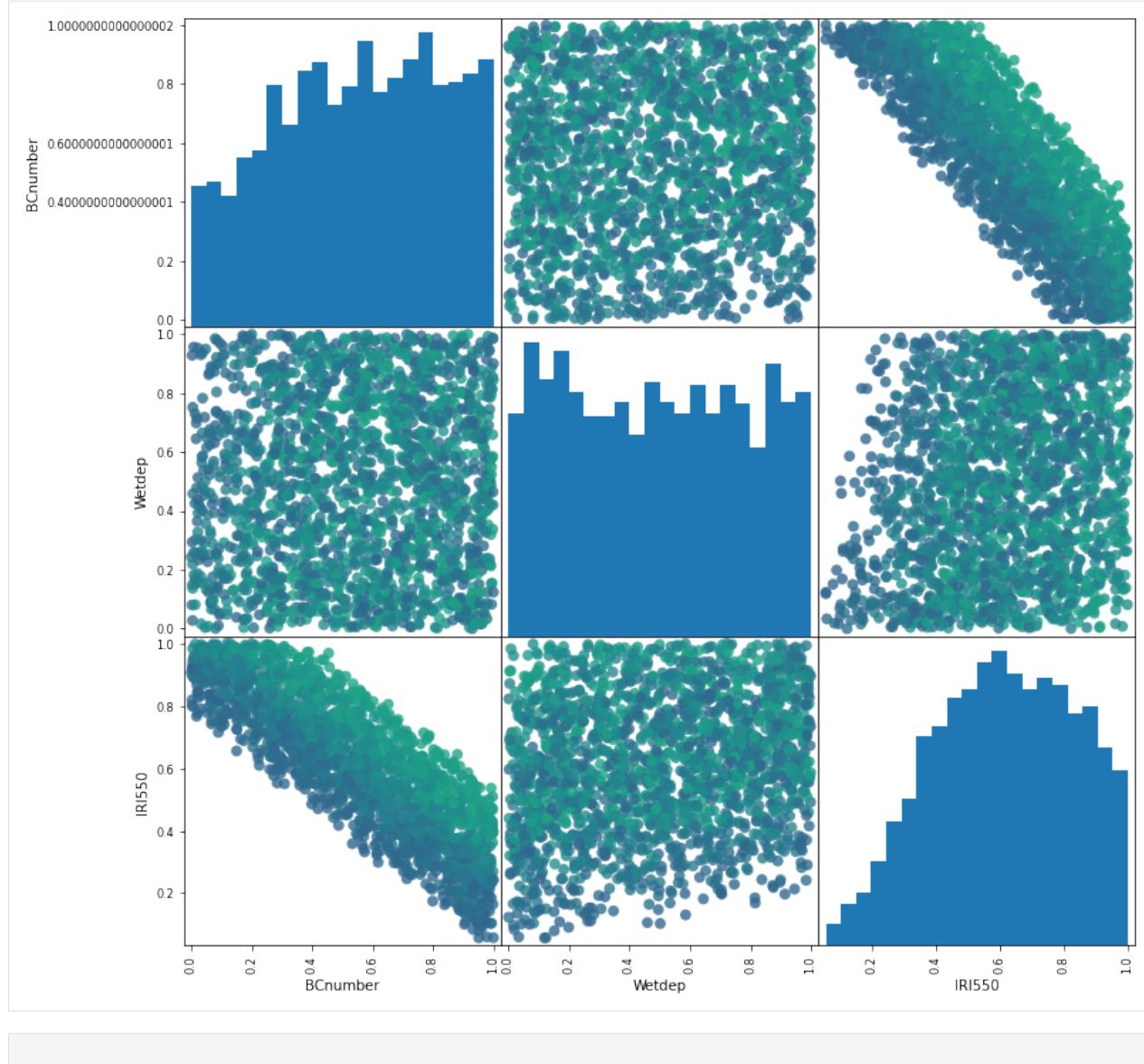
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Sample std dev: 0.0003013221458724632

```
[23]: plot_parameter_space(valid_points, target_df=X_test.iloc[0])
```



```
[24]: grr = pd.plotting.scatter_matrix(valid_points, c=Zs, figsize=(12, 12), marker='o',
                                     hist_kwds={'bins': 20}, s=60, alpha=.8, vmin=1e-3, v
                                     max=5e-3)
```



[ ]:

## 5.6 CMIP6 Emulation

```
[1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from esem import gp_model
from esem.utils import validation_plot
%matplotlib inline
```

```
[2]: df = pd.read_csv('CMIP6_scenarios.csv', index_col=0).dropna()
```

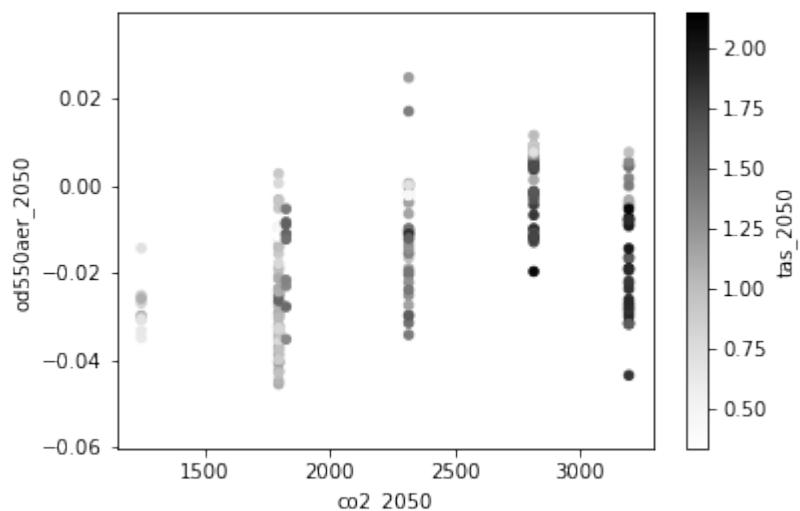
```
[3]: # These are the models included
df.model.unique()
```

```
[3]: array(['CanESM5', 'ACCESS-ESM1-5', 'ACCESS-CM2', 'MPI-ESM1-2-HR',
       'MIROC-ES2L', 'HadGEM3-GC31-LL', 'UKESM1-0-LL', 'MPI-ESM1-2-LR',
       'CESM2', 'CESM2-WACCM', 'NorESM2-LM'], dtype=object)
```

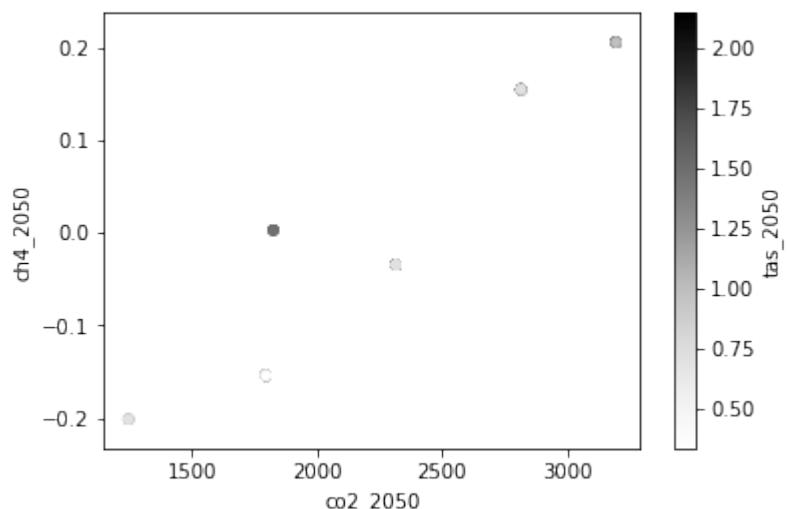
```
[4]: # And these scenarios
df.scenario.unique()
```

```
[4]: array(['ssp126', 'ssp119', 'ssp245', 'ssp370', 'ssp585', 'ssp434'],
       dtype=object)
```

```
[5]: ax = df.plot.scatter(x='co2_2050', y='od550aer_2050', c='tas_2050')
```



```
[6]: ax = df.plot.scatter(x='co2_2050', y='ch4_2050', c='tas_2050')
```



[7]: # Collapse ensemble members

```
df = df.groupby(['model', 'scenario']).mean()
df
```

[7]:

model	scenario	tas_2050	od550aer_2050	tas_2100	od550aer_2100	\
ACCESS-CM2	ssp126	1.000582	-0.025191	1.407198	-0.038552	
	ssp245	1.187015	-0.012222	2.489487	-0.027589	
	ssp370	1.103402	0.006077	3.791109	0.001188	
	ssp585	1.478602	-0.008842	5.016310	-0.016805	
ACCESS-ESM1-5	ssp126	0.819904	-0.013947	0.967981	-0.015895	
	ssp245	1.078853	-0.004727	2.052867	-0.009796	
	ssp370	1.047218	0.003862	3.422980	0.000495	
	ssp585	1.412020	-0.002210	4.096812	-0.001356	
CESM2	ssp126	0.974787	-0.003323	1.134402	-0.010735	
	ssp245	1.119462	0.000371	2.303586	0.000121	
	ssp370	1.154434	0.007554	3.501974	0.016237	
	ssp585	1.453782	0.002178	4.980513	0.016340	
CESM2-WACCM	ssp126	0.770853	0.000561	1.053462	-0.006278	
	ssp245	1.118329	0.000520	2.252389	0.006330	
	ssp370	1.022903	0.011563	3.526784	0.035894	
	ssp585	1.311378	0.005453	5.010599	0.038768	
CanESM5	ssp119	0.641977	-0.030857	0.333723	-0.036624	
	ssp126	0.989167	-0.029769	0.953098	-0.041451	
	ssp245	1.318495	-0.016138	2.454336	-0.033573	
	ssp370	1.724648	-0.004288	4.744037	-0.021374	
	ssp434	1.304015	-0.023741	1.863815	-0.040477	
	ssp585	1.877571	-0.025367	5.984793	-0.040852	
HadGEM3-GC31-LL	ssp126	0.768553	-0.026508	1.314000	-0.037692	
	ssp245	1.262740	-0.014397	2.651430	-0.024341	
	ssp585	1.606117	-0.007902	5.547793	-0.015396	
MIROC-ES2L	ssp119	0.703733	-0.014334	0.557255	-0.020002	
	ssp126	0.785037	-0.017767	0.572429	-0.028761	
	ssp245	0.789146	-0.010066	1.589075	-0.019236	
	ssp370	1.131194	0.001314	2.623115	0.006749	
	ssp585	1.117971	-0.003265	3.748485	-0.003236	
MPI-ESM1-2-HR	ssp126	0.370221	-0.009666	0.351424	-0.014088	
	ssp245	0.735819	-0.001871	1.352010	-0.008150	
	ssp370	0.819827	0.008950	2.616101	0.003430	
	ssp585	0.946618	0.004603	3.153594	-0.004407	
MPI-ESM1-2-LR	ssp126	0.518146	-0.009654	0.358184	-0.014067	
	ssp370	0.877299	0.008940	2.610222	0.003437	
	ssp585	1.029724	0.004613	3.278956	-0.004377	
NorESM2-LM	ssp126	0.331449	-0.011588	0.376486	-0.017219	
	ssp245	0.533042	-0.001380	1.295670	-0.005419	
	ssp370	0.704050	0.007691	2.552758	0.022834	
	ssp585	0.983765	0.007667	2.957163	0.009652	
UKESM1-0-LL	ssp119	0.918189	-0.027682	0.970074	-0.035182	
	ssp126	1.284645	-0.024166	1.563342	-0.036149	
	ssp245	1.604553	-0.010934	3.040434	-0.022142	
	ssp370	1.749074	0.004618	5.046036	0.007180	
	ssp434	1.511831	-0.009184	2.371976	-0.027848	
	ssp585	2.028151	-0.007564	6.038747	-0.006429	

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model	scenario	co2_2050	co2_2100	so2_2050	so2_2100	\
ACCESS-CM2	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
ACCESS-ESM1-5	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
CESM2	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
CESM2-WACCM	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
CanESM5	ssp119	1247.788346	905.867767	-0.069425	-0.080790	
	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp434	1825.355018	1612.733274	-0.055372	-0.077382	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
HadGEM3-GC31-LL	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	3192.373467	10283.292188	-0.028185	-0.059536	
	ssp585	1795.710867	1848.864201	-0.064020	-0.082066	
MIROC-ES2L	ssp119	1247.788346	905.867767	-0.069425	-0.080790	
	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
MPI-ESM1-2-HR	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
MPI-ESM1-2-LR	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
NorESM2-LM	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
UKESM1-0-LL	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	
	ssp119	1247.788346	905.867767	-0.069425	-0.080790	
	ssp126	1795.710867	1848.864201	-0.064020	-0.082066	
ch4_2050	ssp245	2314.385253	3932.717046	-0.035997	-0.059775	
	ssp370	2813.146604	6912.965613	0.004142	-0.020179	
	ssp434	1825.355018	1612.733274	-0.055372	-0.077382	
	ssp585	3192.373467	10283.292188	-0.028185	-0.059536	

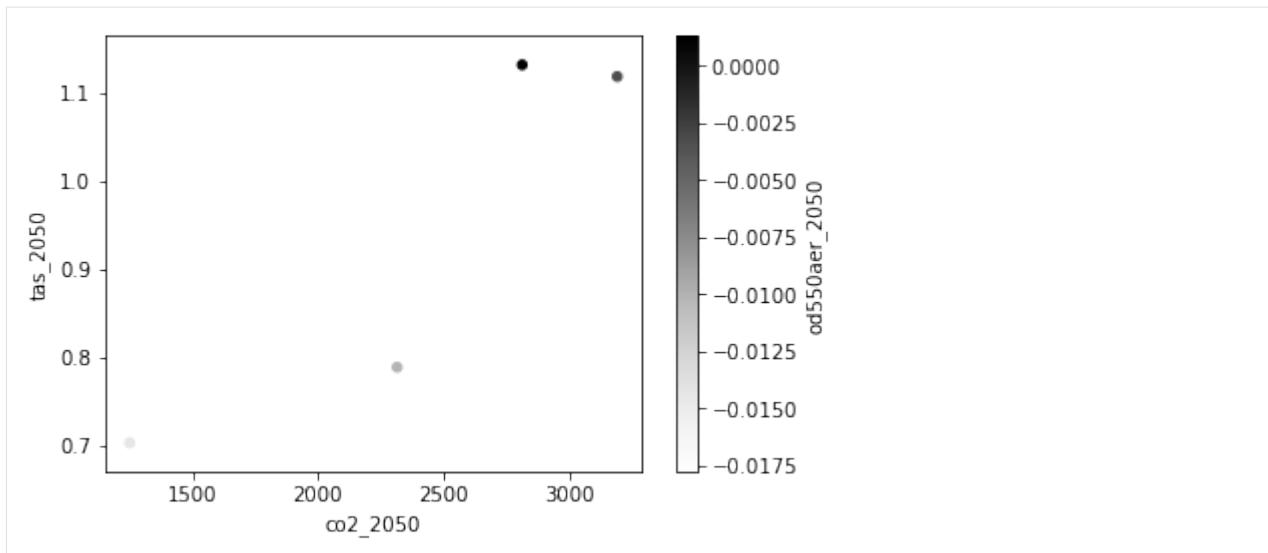
ch4\_2100

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model	scenario			
ACCESS-CM2	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp585	0.205365	0.104140	
ACCESS-ESM1-5	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp585	0.205365	0.104140	
CESM2	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp585	0.205365	0.104140	
CESM2-WACCM	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp585	0.205365	0.104140	
CanESM5	ssp119	-0.201275	-0.257405	
	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp434	0.002907	-0.032744	
	ssp585	0.205365	0.104140	
HadGEM3-GC31-LL	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp585	0.205365	0.104140	
MIROC-ES2L	ssp119	-0.201275	-0.257405	
	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp585	0.205365	0.104140	
MPI-ESM1-2-HR	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp585	0.205365	0.104140	
MPI-ESM1-2-LR	ssp126	-0.153999	-0.241390	
	ssp370	0.154481	0.368859	
	ssp585	0.205365	0.104140	
NorESM2-LM	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp585	0.205365	0.104140	
UKESM1-0-LL	ssp119	-0.201275	-0.257405	
	ssp126	-0.153999	-0.241390	
	ssp245	-0.034673	-0.092845	
	ssp370	0.154481	0.368859	
	ssp434	0.002907	-0.032744	
	ssp585	0.205365	0.104140	

```
[8]: ax = df.query("model == 'MIROC-ES2L'").plot.scatter(x='co2_2050', y='tas_2050', c=
    ↴'od550aer_2050')
```



```
[9]: from utils import normalize
# Merge the year columns in to a long df
df=pd.wide_to_long(df.reset_index(), ["tas", "od550aer", "co2", "ch4", "so2"], i=['model',
    'scenario'], j="year", suffix='_(\d+)')
# Choose only the 2050 data since the aerosol signal is pretty non-existent by 2100
df = df[df.index.isin(["_2050"], level=2)]
```

```
[10]: df.describe()
```

	tas	od550aer	co2	ch4	so2
count	47.000000	47.000000	47.000000	47.000000	47.000000
mean	1.085538	-0.006851	2415.708964	0.024789	-0.035145
std	0.381735	0.011775	613.880074	0.152382	0.025320
min	0.331449	-0.030857	1247.788346	-0.201275	-0.069425
25%	0.804487	-0.014140	1795.710867	-0.153999	-0.064020
50%	1.047218	-0.004727	2314.385253	-0.034673	-0.035997
75%	1.307697	0.003020	2813.146604	0.154481	-0.028185
max	2.028151	0.011563	3192.373467	0.205365	0.004142

```
[11]: # Do a 20/80 split of the data for test and training
msk = np.random.rand(len(df)) < 0.8
train, test = df[msk], df[~msk]
```

### 5.6.1 Try a few different models

```
[12]: from esem.utils import leave_one_out, prediction_within_ci
from scipy import stats

# Try just modelling the temperature based on cumulative CO2
res = leave_one_out(df[['co2']], df[['tas']].values, model='GaussianProcess', kernel=[

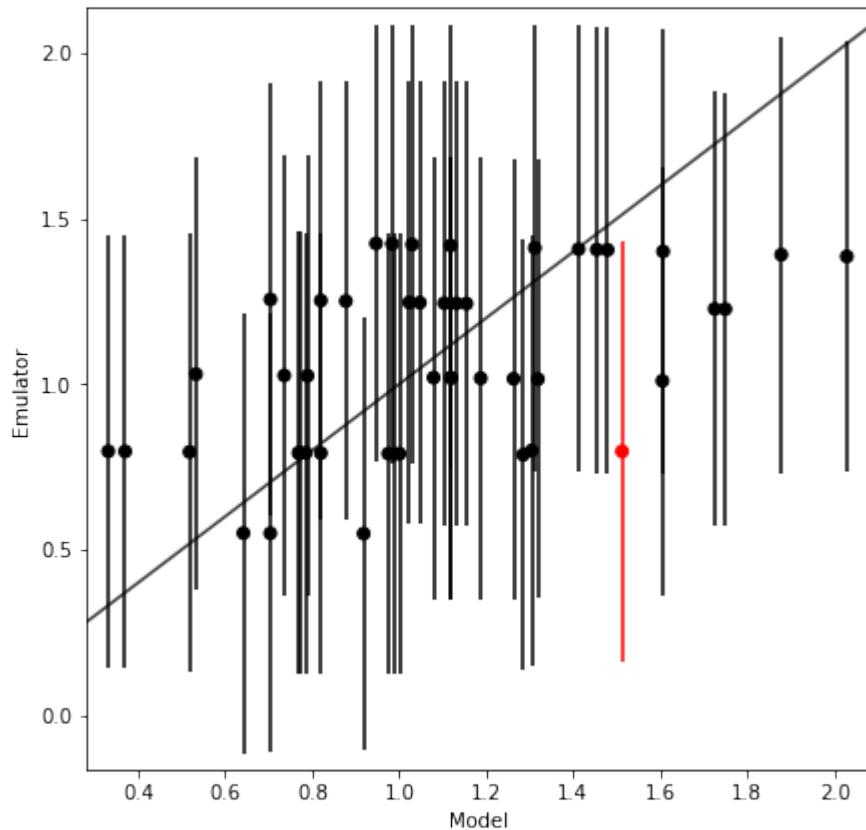
r2_values = stats.linregress(*np.squeeze(np.asarray(res, dtype=float)).T[0:2])[2]**2
```

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```
print("R^2: {:.2f}".format(r2_values))
validation_plot(*np.squeeze(np.asarray(res, dtype=float)).T)

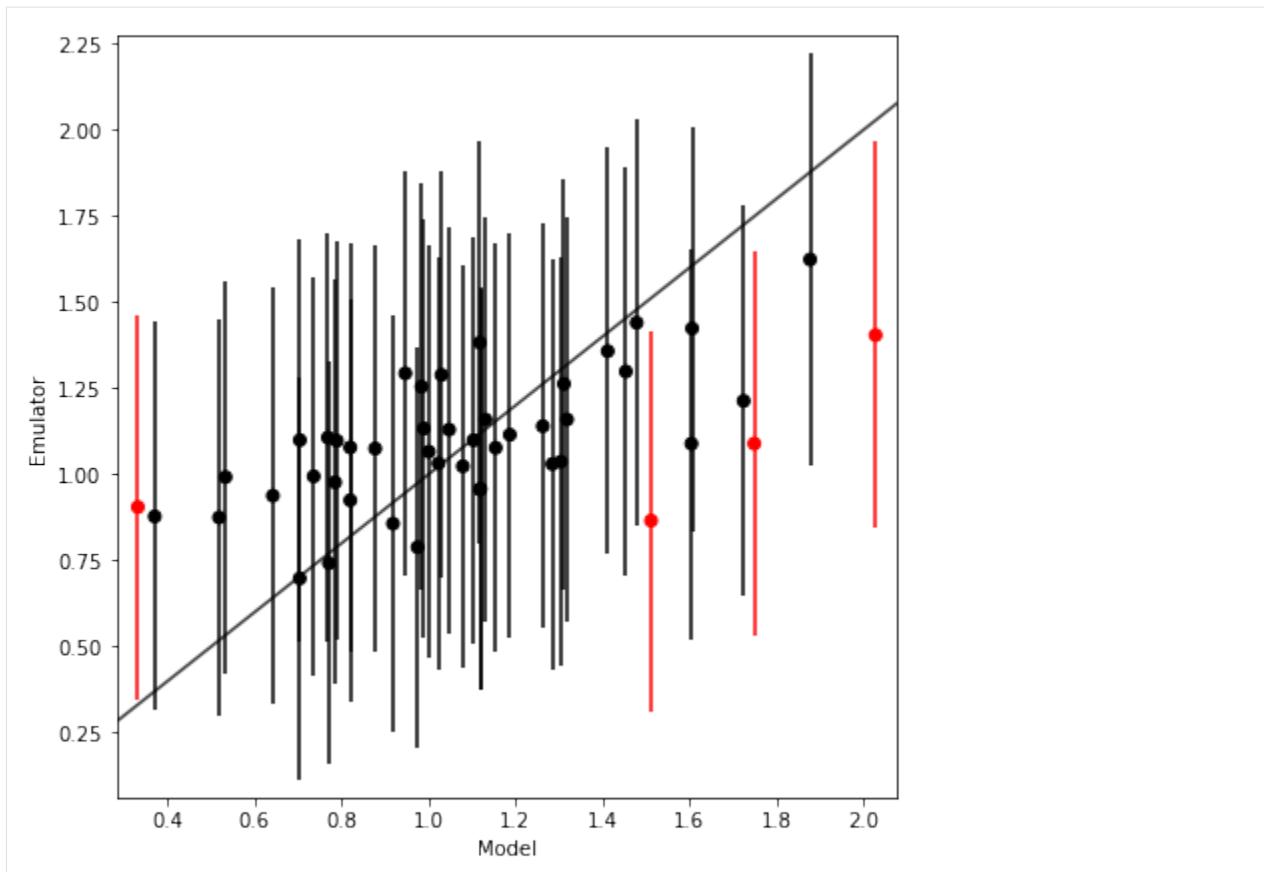
R^2: 0.25
Proportion of 'Bad' estimates : 2.13%
```



```
[13]: # This model still doesn't do brilliantly, but it's better than just CO2
res = leave_one_out(df[['co2', 'od550aer']], df[['tas']].values, model='GaussianProcess',
                     kernel=['Linear'])

r2_values = stats.linregress(*np.squeeze(np.asarray(res, dtype=float)).T[0:2])[2]**2
print("R^2: {:.2f}".format(r2_values))
validation_plot(*np.squeeze(np.asarray(res, dtype=float)).T)

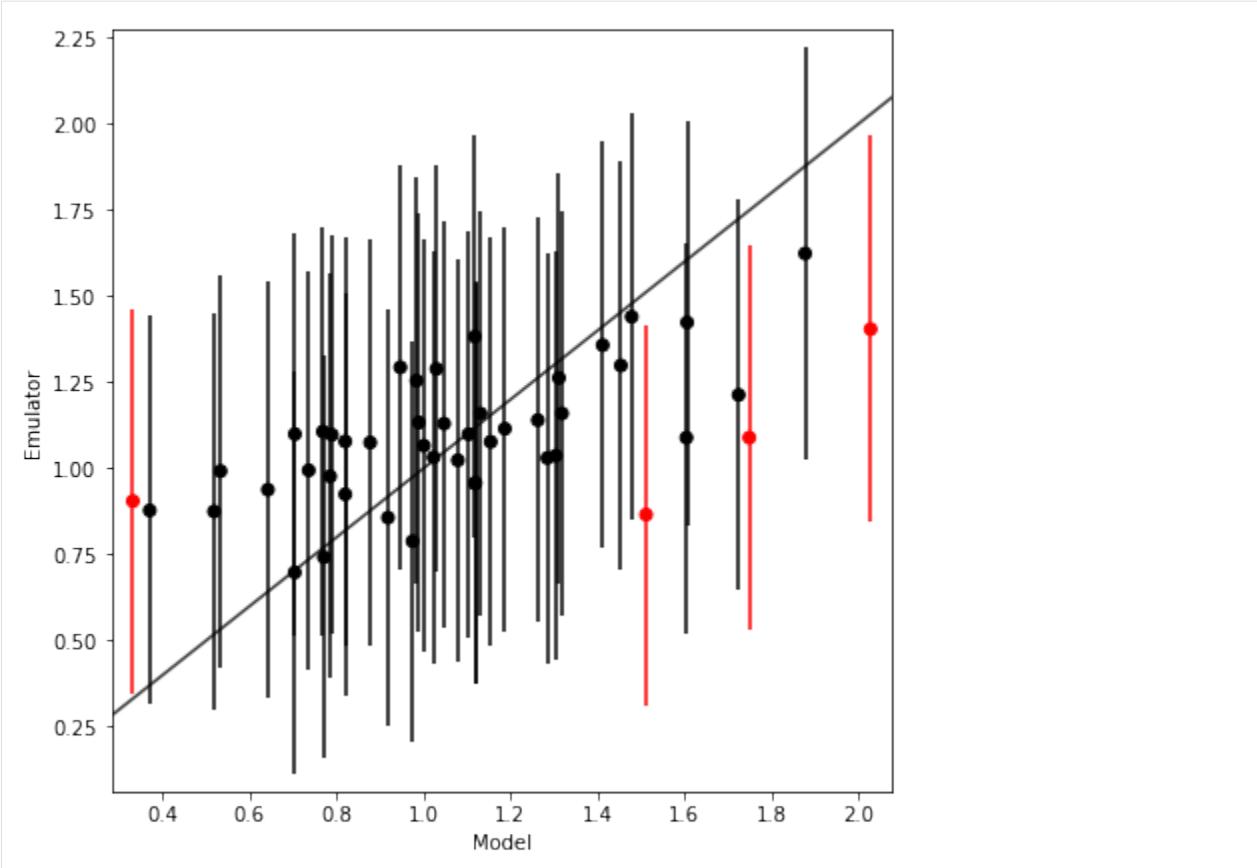
R^2: 0.40
Proportion of 'Bad' estimates : 8.51%
```



```
[14]: # Adding Methane doesn't seem to improve the picture
res = leave_one_out(df[['co2', 'od550aer', 'ch4']], df[['tas']].values, model=
    ↪'GaussianProcess', kernel=['Linear', 'Bias'])

r2_values = stats.linregress(*np.squeeze(np.asarray(res, dtype=float)).T[0:2])[2]**2
print("R^2: {:.2f}".format(r2_values))
validation_plot(*np.squeeze(np.asarray(res, dtype=float)).T)

R^2: 0.40
Proportion of 'Bad' estimates : 8.51%
```



## 5.6.2 Plot the best

```
[15]: m = gp_model(df[['co2', 'od550aer']], df[['tas']].values, kernel=['Linear'])
m.train()
```

```
[16]: # Sample a large AOD/CO2 space using the emulator
xx, yy = np.meshgrid(np.linspace(0, 4000, 25), np.linspace(-.05, 0.05, 20))
X_new = np.stack([xx.flat, yy.flat], axis=1)
Y_new, Y_new_sigma = m.predict(X_new)
```

```
[17]: # Calculate the scenario mean values for comparison
scn_mean = train.groupby(['scenario']).mean()
```

```
[18]: import matplotlib

scale = 1.5
matplotlib.rcParams['font.size'] = 12 * scale
matplotlib.rcParams['lines.linewidth'] = 1.5 * scale
matplotlib.rcParams['lines.markersize'] = 6 * scale

plt.figure(figsize=(12, 6))
```

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```

norm = matplotlib.colors.Normalize(vmin=-2.5,vmax=2.5)
p = plt.contourf(xx, yy, Y_new.reshape(xx.shape), norm=norm, levels=30, cmap='RdBu_r')

plt.scatter(train.co2, train.od550aer, c=train.tas, norm=norm, edgecolors='k', cmap=
    'RdBu_r', marker='x')
plt.scatter(scn_mean.co2, scn_mean.od550aer, c=scn_mean.tas, norm=norm, edgecolors='k',_
    cmap='RdBu_r', marker='s')

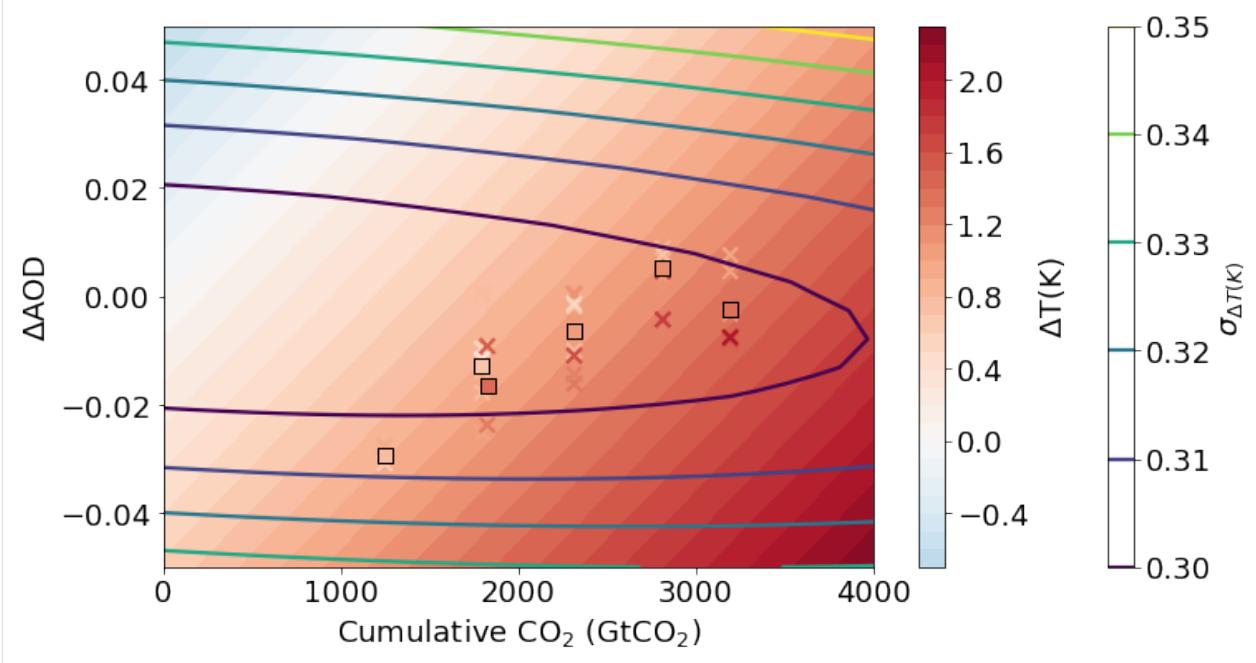
c = plt.contour(xx, yy, np.sqrt(Y_new_sigma.reshape(xx.shape)), cmap='viridis', levels=6)

plt.setp(plt.gca(), xlabel='Cumulative CO$_2$ (GtCO$_2$)', ylabel='$\Delta AOD$')

plt.colorbar(c, label='$\sigma_{\Delta T(K)}$')
plt.colorbar(p, label='$\Delta T(K)$')

# Cumulative CO2, delta T and delta AOD all relative to a 2015-2020 average. Each point_
# represents a single model integration for different scenarios in the CMIP6 archive.
plt.savefig('CMIP6_emulator_paper_v1.1.png', transparent=True)

```



### 5.6.3 Sample emissions for a particular temperature target

```

[19]: from esem.sampler import MCMCSampler

# The MCMC algorithm works much better with a normalised parameter range, so recreate_
# the model
m = gp_model(pd.concat([df[['co2']] / 4000, (df[['od550aer']] + 0.05) / 0.1], axis=1), df[['tas']]
    .values, kernel=['Linear'])
m.train()

```

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```
# Target 1.2 degrees above present day (roughly 2 degrees above pre-industrial)
sampler = MCMCSampler(m, np.asarray([1.2], dtype=np.float64))
samples = sampler.sample(n_samples=8000, mcmc_kwarg=dict(num_burnin_steps=1000) )

Acceptance rate: 0.9614173964786951
```

[20]: # Get the emulated temperatures for these samples  
new\_samples = pd.DataFrame(data=samples, columns=['co2', 'od550aer'])  
Z, \_ = m.predict(new\_samples.values)

[21]: fig = plt.figure(figsize=(9, 6))

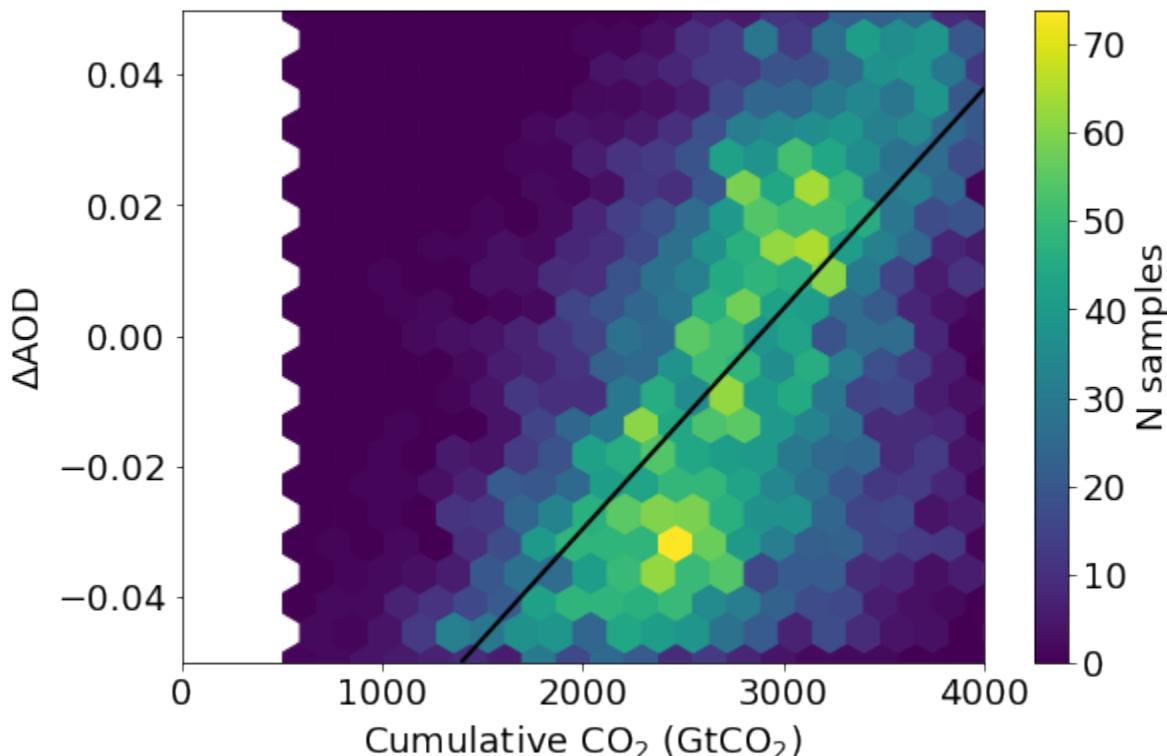
cl = plt.contour(xx, yy, Y\_new.reshape(xx.shape), levels = [1.2],
 colors='k', linestyles='--', linewidths=2)

cl=plt.hexbin(new\_samples.co2\*4000, new\_samples.od550aer\*0.1-0.05, gridsize=20)

plt.setp(plt.gca(), xlabel='Cumulative CO\$\_2\$ (GtCO\$\_2\$)', ylabel='\$\Delta AOD\$')

plt.colorbar(cl, label='N samples')
plt.setp(plt.gca(), ylim=[-0.05, 0.05], xlim=[0, 4000])

plt.savefig('CMIP6\_emulator\_sampled.png', transparent=True)



[ ]:

## 5.7 Create paper emulation figure

```
[1]: import os
## Ignore my broken HDF5 install...
os.putenv("HDF5_DISABLE_VERSION_CHECK", '1')
```

```
[2]: import iris

from utils import get_bc_ppe_data

from esem import cnn_model, gp_model
from esem.utils import get_random_params

import iris.quickplot as qplt
import iris.analysis.maths as imath
import matplotlib.pyplot as plt
%matplotlib inline
```

### 5.7.1 Read in the parameters and data

```
[3]: ppe_params, ppe_aaod = get_bc_ppe_data()

/Users/watson-parris/miniconda3/envs/gcem/lib/python3.8/site-packages/iris/__init__.py:
  ← 249: IrisDeprecation: setting the 'Future' property 'netcdf_promote' is deprecated and
  ← will be removed in a future release. Please remove code that sets this property.
    warn_deprecated(msg.format(name))
/Users/watson-parris/miniconda3/envs/gcem/lib/python3.8/site-packages/iris/__init__.py:
  ← 249: IrisDeprecation: setting the 'Future' property 'netcdf_promote' is deprecated and
  ← will be removed in a future release. Please remove code that sets this property.
    warn_deprecated(msg.format(name))
```

```
[4]: ## Ensure the wetdepnumbertime dimension is last - this is treated as the color 'channel'
## ppe_aaod.transpose((0,2,3,1))
ppe_aaod = ppe_aaod.collapsed('time')[0]

WARNING:root:Creating guessed bounds as none exist in file
WARNING:root:Creating guessed bounds as none exist in file
WARNING:root:Creating guessed bounds as none exist in file
```

```
[5]: n_test = 5

X_test, X_train = ppe_params[:n_test], ppe_params[n_test:]
Y_test, Y_train = ppe_aaod[:n_test], ppe_aaod[n_test:]
```

```
[6]: Y_train
[6]: <iris 'Cube' of Absorption optical thickness - total 550nm / (1) (job: 34; latitude: 96;
   ← longitude: 192)>
```

## 5.7.2 Setup and run the models

[7]: nn\_model = cnn\_model(X\_train, Y\_train)

[8]: nn\_model.model.summary()

Model: "decoder"

Layer (type)	Output Shape	Param #
input_1 (InputLayer)	[None, 3]	0
dense (Dense)	(None, 221184)	884736
reshape (Reshape)	(None, 96, 192, 12)	0
conv2d_transpose (Conv2DTran	(None, 96, 192, 12)	2172
conv2d_transpose_1 (Conv2DTr	(None, 96, 192, 1)	181
Total params:	887,089	
Trainable params:	887,089	
Non-trainable params:	0	

[9]: nn\_model.train()

```
Epoch 1/100
4/4 [=====] - 1s 284ms/step - loss: 0.8766 - val_loss: 0.4800
Epoch 2/100
4/4 [=====] - 0s 115ms/step - loss: 1.0869 - val_loss: 0.4778
Epoch 3/100
4/4 [=====] - 0s 100ms/step - loss: 1.1900 - val_loss: 0.4759
Epoch 4/100
4/4 [=====] - 0s 106ms/step - loss: 1.0656 - val_loss: 0.4741
Epoch 5/100
4/4 [=====] - 0s 99ms/step - loss: 1.2058 - val_loss: 0.4699
Epoch 6/100
4/4 [=====] - 0s 112ms/step - loss: 1.2211 - val_loss: 0.4672
Epoch 7/100
4/4 [=====] - 0s 105ms/step - loss: 1.0138 - val_loss: 0.4624
Epoch 8/100
4/4 [=====] - 0s 101ms/step - loss: 0.9261 - val_loss: 0.4645
Epoch 9/100
4/4 [=====] - 0s 112ms/step - loss: 0.9354 - val_loss: 0.4569
Epoch 10/100
4/4 [=====] - 0s 100ms/step - loss: 1.1767 - val_loss: 0.4494
Epoch 11/100
4/4 [=====] - 0s 101ms/step - loss: 1.1077 - val_loss: 0.4310
Epoch 12/100
4/4 [=====] - 0s 101ms/step - loss: 0.8575 - val_loss: 0.4343
Epoch 13/100
4/4 [=====] - 0s 101ms/step - loss: 0.8162 - val_loss: 0.4476
```

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```

Epoch 14/100
4/4 [=====] - 0s 102ms/step - loss: 0.8983 - val_loss: 0.4181
Epoch 15/100
4/4 [=====] - 1s 104ms/step - loss: 0.8968 - val_loss: 0.4003
Epoch 16/100
4/4 [=====] - 0s 100ms/step - loss: 0.8450 - val_loss: 0.3989
Epoch 17/100
4/4 [=====] - 0s 103ms/step - loss: 0.7290 - val_loss: 0.4107
Epoch 18/100
4/4 [=====] - 0s 100ms/step - loss: 0.8785 - val_loss: 0.3785
Epoch 19/100
4/4 [=====] - 0s 106ms/step - loss: 0.7236 - val_loss: 0.3918
Epoch 20/100
4/4 [=====] - 0s 103ms/step - loss: 0.8572 - val_loss: 0.3857
Epoch 21/100
4/4 [=====] - 0s 102ms/step - loss: 0.8973 - val_loss: 0.3543
Epoch 22/100
4/4 [=====] - 0s 101ms/step - loss: 0.9261 - val_loss: 0.3282
Epoch 23/100
4/4 [=====] - 0s 102ms/step - loss: 0.5686 - val_loss: 0.3123
Epoch 24/100
4/4 [=====] - 0s 100ms/step - loss: 0.9107 - val_loss: 0.3041
Epoch 25/100
4/4 [=====] - 0s 101ms/step - loss: 0.7340 - val_loss: 0.2931
Epoch 26/100
4/4 [=====] - 0s 100ms/step - loss: 0.7675 - val_loss: 0.2825
Epoch 27/100
4/4 [=====] - 0s 105ms/step - loss: 0.5598 - val_loss: 0.2795
Epoch 28/100
4/4 [=====] - 0s 101ms/step - loss: 0.6441 - val_loss: 0.2640
Epoch 29/100
4/4 [=====] - 0s 100ms/step - loss: 0.8158 - val_loss: 0.2579
Epoch 30/100
4/4 [=====] - 0s 102ms/step - loss: 0.7065 - val_loss: 0.2447
Epoch 31/100
4/4 [=====] - 0s 105ms/step - loss: 0.6948 - val_loss: 0.2351
Epoch 32/100
4/4 [=====] - 0s 125ms/step - loss: 0.7187 - val_loss: 0.2226
Epoch 33/100
4/4 [=====] - 0s 122ms/step - loss: 0.5605 - val_loss: 0.2154
Epoch 34/100
4/4 [=====] - 0s 104ms/step - loss: 0.5716 - val_loss: 0.2117
Epoch 35/100
4/4 [=====] - 0s 107ms/step - loss: 0.6035 - val_loss: 0.1991
Epoch 36/100
4/4 [=====] - 0s 107ms/step - loss: 0.6084 - val_loss: 0.1892
Epoch 37/100
4/4 [=====] - 0s 103ms/step - loss: 0.6416 - val_loss: 0.1786
Epoch 38/100
4/4 [=====] - 0s 100ms/step - loss: 0.4608 - val_loss: 0.1749
Epoch 39/100
4/4 [=====] - 0s 100ms/step - loss: 0.3582 - val_loss: 0.1674

```

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```

Epoch 40/100
4/4 [=====] - 0s 113ms/step - loss: 0.4616 - val_loss: 0.1607
Epoch 41/100
4/4 [=====] - 0s 117ms/step - loss: 0.5972 - val_loss: 0.1558
Epoch 42/100
4/4 [=====] - 0s 121ms/step - loss: 0.3961 - val_loss: 0.1471
Epoch 43/100
4/4 [=====] - 0s 111ms/step - loss: 0.5399 - val_loss: 0.1404
Epoch 44/100
4/4 [=====] - 0s 100ms/step - loss: 0.4118 - val_loss: 0.1385
Epoch 45/100
4/4 [=====] - 0s 126ms/step - loss: 0.3920 - val_loss: 0.1316
Epoch 46/100
4/4 [=====] - 0s 121ms/step - loss: 0.4126 - val_loss: 0.1237
Epoch 47/100
4/4 [=====] - 1s 247ms/step - loss: 0.3554 - val_loss: 0.1202
Epoch 48/100
4/4 [=====] - 0s 129ms/step - loss: 0.3429 - val_loss: 0.1145
Epoch 49/100
4/4 [=====] - 1s 128ms/step - loss: 0.3157 - val_loss: 0.1075
Epoch 50/100
4/4 [=====] - 1s 117ms/step - loss: 0.4217 - val_loss: 0.1051
Epoch 51/100
4/4 [=====] - 0s 112ms/step - loss: 0.4561 - val_loss: 0.1078
Epoch 52/100
4/4 [=====] - 0s 121ms/step - loss: 0.4475 - val_loss: 0.0932
Epoch 53/100
4/4 [=====] - 0s 123ms/step - loss: 0.2861 - val_loss: 0.0896
Epoch 54/100
4/4 [=====] - 0s 115ms/step - loss: 0.3469 - val_loss: 0.0886
Epoch 55/100
4/4 [=====] - 0s 126ms/step - loss: 0.3332 - val_loss: 0.0920
Epoch 56/100
4/4 [=====] - 0s 117ms/step - loss: 0.3290 - val_loss: 0.0793
Epoch 57/100
4/4 [=====] - 1s 136ms/step - loss: 0.3292 - val_loss: 0.0792
Epoch 58/100
4/4 [=====] - 0s 104ms/step - loss: 0.2757 - val_loss: 0.0760
Epoch 59/100
4/4 [=====] - 0s 110ms/step - loss: 0.2637 - val_loss: 0.0720
Epoch 60/100
4/4 [=====] - 0s 123ms/step - loss: 0.4001 - val_loss: 0.0729
Epoch 61/100
4/4 [=====] - 0s 110ms/step - loss: 0.4507 - val_loss: 0.0666
Epoch 62/100
4/4 [=====] - 1s 131ms/step - loss: 0.3879 - val_loss: 0.0695
Epoch 63/100
4/4 [=====] - 0s 106ms/step - loss: 0.2597 - val_loss: 0.0610
Epoch 64/100
4/4 [=====] - 0s 111ms/step - loss: 0.3288 - val_loss: 0.0725
Epoch 65/100
4/4 [=====] - 0s 102ms/step - loss: 0.2841 - val_loss: 0.0575

```

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```

Epoch 66/100
4/4 [=====] - 0s 101ms/step - loss: 0.2730 - val_loss: 0.0544
Epoch 67/100
4/4 [=====] - 0s 101ms/step - loss: 0.2694 - val_loss: 0.0517
Epoch 68/100
4/4 [=====] - 0s 110ms/step - loss: 0.3846 - val_loss: 0.0573
Epoch 69/100
4/4 [=====] - 0s 100ms/step - loss: 0.3429 - val_loss: 0.0528
Epoch 70/100
4/4 [=====] - 0s 106ms/step - loss: 0.2360 - val_loss: 0.0478
Epoch 71/100
4/4 [=====] - 0s 100ms/step - loss: 0.3651 - val_loss: 0.0486
Epoch 72/100
4/4 [=====] - 0s 107ms/step - loss: 0.2351 - val_loss: 0.0469
Epoch 73/100
4/4 [=====] - 0s 101ms/step - loss: 0.2993 - val_loss: 0.0476
Epoch 74/100
4/4 [=====] - 0s 105ms/step - loss: 0.2739 - val_loss: 0.0523
Epoch 75/100
4/4 [=====] - 0s 103ms/step - loss: 0.3909 - val_loss: 0.0438
Epoch 76/100
4/4 [=====] - 0s 109ms/step - loss: 0.2526 - val_loss: 0.0423
Epoch 77/100
4/4 [=====] - 0s 103ms/step - loss: 0.2566 - val_loss: 0.0485
Epoch 78/100
4/4 [=====] - 0s 101ms/step - loss: 0.2969 - val_loss: 0.0411
Epoch 79/100
4/4 [=====] - 0s 99ms/step - loss: 0.2602 - val_loss: 0.0413
Epoch 80/100
4/4 [=====] - 0s 126ms/step - loss: 0.1863 - val_loss: 0.0382
Epoch 81/100
4/4 [=====] - 0s 101ms/step - loss: 0.3251 - val_loss: 0.0394
Epoch 82/100
4/4 [=====] - 0s 109ms/step - loss: 0.3038 - val_loss: 0.0376
Epoch 83/100
4/4 [=====] - 0s 105ms/step - loss: 0.3729 - val_loss: 0.0405
Epoch 84/100
4/4 [=====] - 0s 107ms/step - loss: 0.2542 - val_loss: 0.0389
Epoch 85/100
4/4 [=====] - 0s 109ms/step - loss: 0.2379 - val_loss: 0.0403
Epoch 86/100
4/4 [=====] - 0s 117ms/step - loss: 0.2620 - val_loss: 0.0350
Epoch 87/100
4/4 [=====] - 0s 108ms/step - loss: 0.2459 - val_loss: 0.0356
Epoch 88/100
4/4 [=====] - 0s 107ms/step - loss: 0.1902 - val_loss: 0.0368
Epoch 89/100
4/4 [=====] - 1s 134ms/step - loss: 0.3153 - val_loss: 0.0362
Epoch 90/100
4/4 [=====] - 0s 116ms/step - loss: 0.3270 - val_loss: 0.0371
Epoch 91/100
4/4 [=====] - 1s 123ms/step - loss: 0.3058 - val_loss: 0.0393

```

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```

Epoch 92/100
4/4 [=====] - 0s 115ms/step - loss: 0.3535 - val_loss: 0.0359
Epoch 93/100
4/4 [=====] - 0s 102ms/step - loss: 0.3056 - val_loss: 0.0379
Epoch 94/100
4/4 [=====] - 0s 106ms/step - loss: 0.3331 - val_loss: 0.0354
Epoch 95/100
4/4 [=====] - 0s 99ms/step - loss: 0.2074 - val_loss: 0.0364
Epoch 96/100
4/4 [=====] - 0s 99ms/step - loss: 0.1920 - val_loss: 0.0364
Epoch 97/100
4/4 [=====] - 0s 101ms/step - loss: 0.3333 - val_loss: 0.0329
Epoch 98/100
4/4 [=====] - 0s 99ms/step - loss: 0.2105 - val_loss: 0.0340
Epoch 99/100
4/4 [=====] - 0s 105ms/step - loss: 0.3683 - val_loss: 0.0355
Epoch 100/100
4/4 [=====] - 0s 100ms/step - loss: 0.3073 - val_loss: 0.0361

```

[10]: `## Linear model: 0.3566 - val_loss: 0.0867`

[11]: `nn_prediction, _ = nn_model.predict(X_test.values)`

[12]: `gp_model_ = gp_model(X_train, Y_train, kernel=['Bias', 'Linear'])  
gp_model_.train()`

[13]: `gp_prediction, _ = gp_model_.predict(X_test.values)`

[14]: `import matplotlib  
import cartopy.crs as ccrs  
import iris.plot as iplt`

`plt.figure(figsize=(30, 10))  
matplotlib.rcParams['font.size'] = 24`

`plt.subplot(2,3,1, projection=ccrs.Mollweide())  
plt.annotate("(a)", (0.,1.), xycoords='axes fraction')  
iplt.pcolormesh(imath.log10(Y_test[0]), vmin=-4, vmax=-1)  
plt.gca().set_title('Truth')  
plt.gca().coastlines()`

`plt.subplot(2,3,2, projection=ccrs.Mollweide())  
plt.annotate("(b)", (0.,1.), xycoords='axes fraction')  
iplt.pcolormesh(imath.log10(gp_prediction[0]), vmin=-4, vmax=-1)  
plt.gca().set_title('GP')  
plt.gca().coastlines()`

`plt.subplot(2,3,3, projection=ccrs.Mollweide())  
plt.annotate("(c)", (0.,1.), xycoords='axes fraction')  
im=iplt.pcolormesh(imath.log10(nn_prediction[0]), vmin=-4, vmax=-1)`

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```

plt.gca().set_title('CNN')
plt.colorbar(im, fraction=0.046, pad=0.04, label='log(AAOD)')
plt.gca().coastlines()

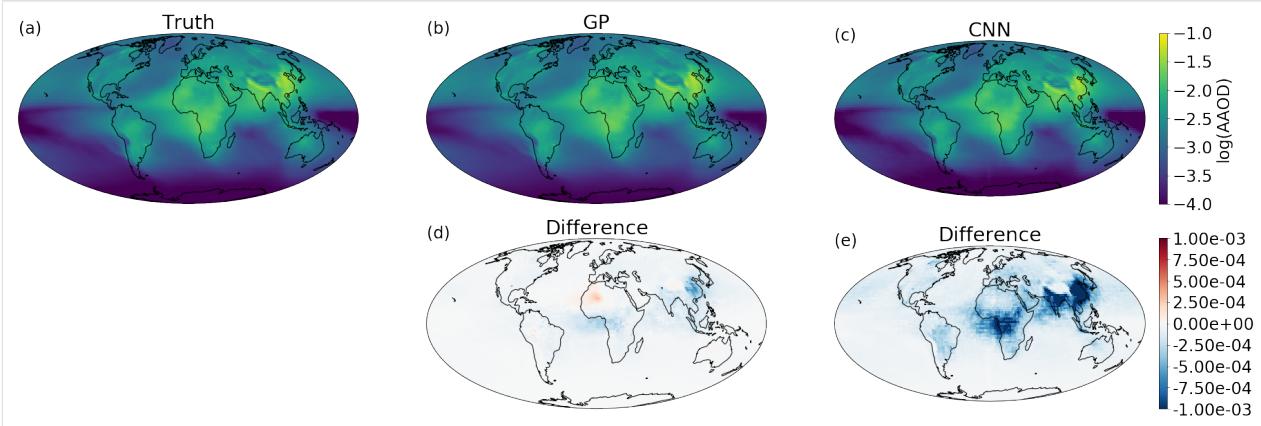
plt.subplot(2,3,5, projection=ccrs.Mollweide())
plt.annotate("(d)", (0.,1.), xycoords='axes fraction')
iplt.pcolormesh(gp_prediction.collapsed(['sample'], iris.analysis.MEAN)-Y_test.
    ↵ collapsed(['job'], iris.analysis.MEAN)), cmap='RdBu_r', vmin=-0.001, vmax=0.001)
plt.gca().coastlines()
plt.gca().set_title('Difference')

plt.subplot(2,3,6, projection=ccrs.Mollweide())
plt.annotate("(e)", (0.,1.), xycoords='axes fraction')
im=iplt.pcolormesh(nn_prediction.collapsed(['sample'], iris.analysis.MEAN)-Y_test.
    ↵ collapsed(['job'], iris.analysis.MEAN)), cmap='RdBu_r', vmin=-1e-3, vmax=1e-3)
cb = plt.colorbar(im, fraction=0.046, pad=0.04)
cb.ax.set_yticklabels(["{:2e}".format(i) for i in cb.get_ticks()]) ## set ticks of your
    ↵ format
plt.gca().coastlines()
plt.gca().set_title('Difference')

plt.savefig('BCPPE_emulator_paper.png', transparent=True)

/Users/watson-parris/miniconda3/envs/gcem/lib/python3.8/site-packages/iris/coords.py:
    ↵ 1410: UserWarning: Collapsing a non-contiguous coordinate. Metadata may not be fully
    ↵ descriptive for 'sample'.
    warnings.warn(msg.format(self.name()))
/Users/watson-parris/miniconda3/envs/gcem/lib/python3.8/site-packages/iris/coords.py:
    ↵ 1410: UserWarning: Collapsing a non-contiguous coordinate. Metadata may not be fully
    ↵ descriptive for 'sample'.
    warnings.warn(msg.format(self.name()))

```



[15]:

```

COLOR = 'white'
matplotlib.rcParams['text.color'] = COLOR
matplotlib.rcParams['axes.labelcolor'] = COLOR
matplotlib.rcParams['xtick.color'] = COLOR
matplotlib.rcParams['ytick.color'] = COLOR
matplotlib.rcParams['font.size'] = 20

```

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```

plt.figure(figsize=(30, 10))

plt.subplot(2,3,1, projection=ccrs.Mollweide())
plt.annotate("(a)", (0.,1.), xycoords='axes fraction')
iplt.pcolormesh(imath.log10(Y_test[0]), vmin=-4, vmax=-1)
plt.gca().set_title('Truth')
plt.gca().coastlines()

plt.subplot(2,3,2, projection=ccrs.Mollweide())
plt.annotate("(b)", (0.,1.), xycoords='axes fraction')
iplt.pcolormesh(imath.log10(gp_prediction[0]), vmin=-4, vmax=-1)
plt.gca().set_title('GP')
plt.gca().coastlines()

plt.subplot(2,3,3, projection=ccrs.Mollweide())
plt.annotate("(c)", (0.,1.), xycoords='axes fraction')
im=iplt.pcolormesh(imath.log10(nn_prediction[0]), vmin=-4, vmax=-1)
plt.gca().set_title('CNN')
plt.colorbar(im, fraction=0.046, pad=0.04, label='log(AAOD)')
plt.gca().coastlines()

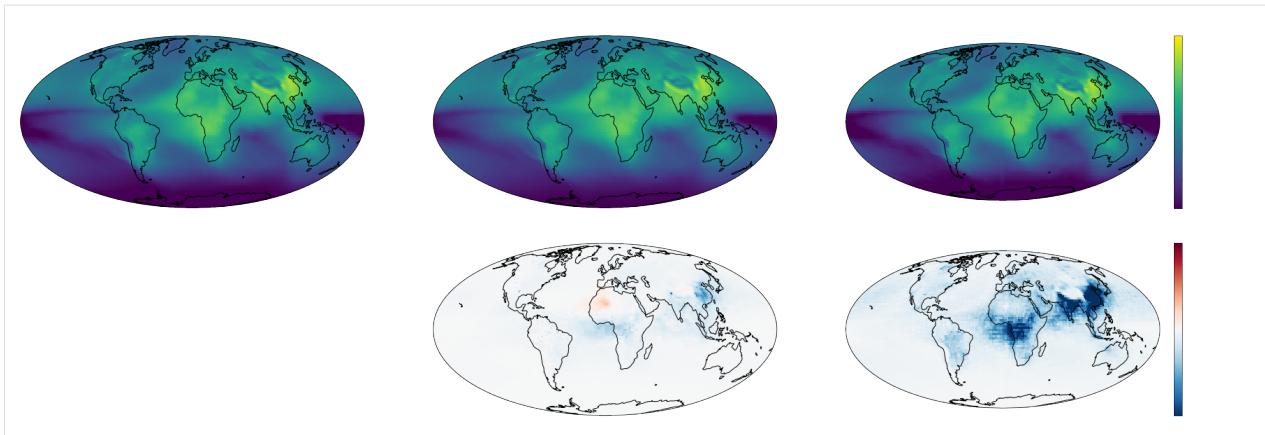
plt.subplot(2,3,5, projection=ccrs.Mollweide())
plt.annotate("(d)", (0.,1.), xycoords='axes fraction')
iplt.pcolormesh((gp_prediction.collapsed(['sample']), iris.analysis.MEAN)-Y_test.
    ↴collapsed(['job'], iris.analysis.MEAN)), cmap='RdBu_r', vmin=-0.001, vmax=0.001)
plt.gca().coastlines()
plt.gca().set_title('Difference')

plt.subplot(2,3,6, projection=ccrs.Mollweide())
plt.annotate("(e)", (0.,1.), xycoords='axes fraction')
im=iplt.pcolormesh((nn_prediction.collapsed(['sample']), iris.analysis.MEAN)-Y_test.
    ↴collapsed(['job'], iris.analysis.MEAN)), cmap='RdBu_r', vmin=-1e-3, vmax=1e-3)
cb=plt.colorbar(im, fraction=0.046, pad=0.04)
cb.ax.set_yticklabels(["{:e}".format(i) for i in cb.get_ticks()]) ## set ticks of your
    ↴format
plt.gca().coastlines()
plt.gca().set_title('Difference')

plt.savefig('BCPPE_emulator_talk.png', transparent=True)

/Users/watson-parris/miniconda3/envs/gcem/lib/python3.8/site-packages/iris/coords.py:
    ↴1410: UserWarning: Collapsing a non-contiguous coordinate. Metadata may not be fully
    ↴descriptive for 'sample'.
    warnings.warn(msg.format(self.name()))
/Users/watson-parris/miniconda3/envs/gcem/lib/python3.8/site-packages/iris/coords.py:
    ↴1410: UserWarning: Collapsing a non-contiguous coordinate. Metadata may not be fully
    ↴descriptive for 'sample'.
    warnings.warn(msg.format(self.name()))

```



[ ]:



## API REFERENCE

This page provides an auto-generated summary of xarray's API. For more details and examples, refer to the relevant chapters in the main part of the documentation.

### 6.1 Top-level functions

This provides the main interface for ESEm and should be the starting point for most users.

<a href="#"><code>gp_model</code></a>	Create a Gaussian process (GP) based emulator with provided <i>training_params</i> (X) and <i>training_data</i> (Y) which assumes independent inputs (and outputs).
<a href="#"><code>cnn_model</code></a>	Create a simple two layer Convolutional Neural Network Emulator using Keras.
<a href="#"><code>rf_model</code></a>	Create a simple Random Forest Emulator using sklearn.

#### 6.1.1 `esem.gp_model`

`esem.gp_model(training_params, training_data, data_processors=None, kernel=None, kernel_op='add', active_dims=None, noise_variance=1.0, name='', gpu=0, **kwargs)`

Create a Gaussian process (GP) based emulator with provided *training\_params* (X) and *training\_data* (Y) which assumes independent inputs (and outputs).

The *kernel* is a key parameter in GP emulation and care should be taken in choosing it.

##### Parameters

- **training\_params** (`DataFrame`) – The training parameters
- **training\_data** (`xarray.DataArray` or `iris.Cube` or `array_like`) – The training data - the leading dimension should represent training samples
- **data\_processors** (`list` of `esem.data_processors.DataProcessor`) – A list of ‘DataProcessor` to apply to the data transparently before training. Model output will be untransformed before being returned from the Emulator.
- **kernel** (`gpflow.kernels.Kernel` or `list` of `str` or `None`) – The GP kernel to use. A GPFLOW kernel can be specified directly, or a list of kernel names can be provided which will be initialised using the default values (of the correct shape) and combined using *kernel\_op*. Alternatively no kernel can be specified and a default will be used.
- **kernel\_op** (`{'add', 'mul'}`) – The operation to perform in order to combine the specified *kernel*'s. *Only used if kernel* is a list of strings.

- **noise\_variance** (float) – The noise variance to initialise the GP regression model
- **active\_dims** (list of int or slice or None) – The dimensions to train the GP over (by default all of the dimensions are used)
- **name** (str) – An optional name for the emulator
- **gpu** (int) – The GPU to use (only applicable for multi-GPU) machines
- **kwargs** (dict) – Dict of optional keyword arguments for *gpflow.models.GPR*, e.g., *mean\_function*

**Returns** Emulator – An esem emulator object which can be trained and sampled from

## 6.1.2 esem.cnn\_model

```
esem.cnn_model(training_params, training_data, data_processors=None, filters=12, learning_rate=0.001,
                decay=0.01, kernel_size=(3, 5), loss='mean_squared_error', activation='tanh',
                optimizer='RMSprop', name='', gpu=0)
```

Create a simple two layer Convolutional Neural Network Emulator using Keras.

Note that X should include both the train and validation data

### Parameters

- **training\_params** (pd.DataFrame) – The training parameters
- **training\_data** (xarray.DataArray or *iris*.cube.Cube or *array\_like*) – The training data - the leading dimension should represent training samples
- **data\_processors** (list of *esem.data\_processors.DataProcessor*) – A list of *DataProcessor* to apply to the data transparently before training. Model output will be untransformed before being returned from the Emulator.
- **filters** (int) – The dimensionality of the first convolutional layer output space
- **learning\_rate** (float) – The learning rate to use with the chosen optimizer
- **decay** (float) – Any decay to apply to the learning rate
- **kernel\_size** (tuple of int) – The convolutional kernel size
- **loss** (str) – The loss function to train against (see <https://keras.io/api/losses/>)
- **activation** (str) – The activation function to use in the final CNN layer (see <https://keras.io/api/layers/activations/>)
- **optimizer** ({'RMSprop', 'Adam'}) – The optimizer to train the model with
- **name** (str) – An optional name for the emulator
- **gpu** (int) – The GPU to use (only applicable for multi-GPU) machines

**Returns** Emulator – An esem emulator object which can be trained and sampled from

## Notes

The Keras model is compiled but not trained until `train` is called on the returned `Emulator` object.

### 6.1.3 esem.rf\_model

`esem.rf_model(training_params, training_data, data_processors=None, name='', gpu=0, *args, **kwargs)`  
Create a simple Random Forest Emulator using sklearn.

Note that because a Random Forest is just a recursive binary partition over the training data, there is no need to normalize/standardize the inputs.

i.e. At least in theory, Random Forests are invariant to monotonic transformations of the independent variables

#### Parameters

- `training_params` (`pd.DataFrame`) – The training parameters
- `training_data` (`xarray.DataArray` or `iris.cube.Cube` or `array_like`) – The training data  
- the leading dimension should represent training samples
- `data_processors` (`list` of `esem.data_processors.DataProcessor`) – A list of `DataProcessor` to apply to the data transparently before training. Model output will be untransformed before being returned from the Emulator.
- `name` (`str`) – An optional name for the emulator
- `gpu` (`int`) – The GPU to use (only applicable for multi-GPU) machines
- `args` (`list`) – List of optional arguments for `sklearn.ensemble.RandomForestRegressor`
- `kwargs` (`dict`) – Dict of optional keyword arguments for `sklearn.ensemble.RandomForestRegressor`

**Returns** `Emulator` – An esem emulator object which can be trained and sampled from

## 6.2 Emulator

<code>Emulator</code>	A class wrapping a statistical emulator
<code>Emulator.train</code>	Train on the training data
<code>Emulator.predict</code>	Make a prediction using a trained emulator
<code>Emulator._predict</code>	The (internal) predict interface used by e.g., a sampler.
<code>Emulator.batch_stats</code>	Return mean and standard deviation in model predictions over samples, without storing the intermediate predictions in memory to enable evaluating large models over more samples than could fit in memory

## 6.2.1 esem.emulator.Emulator

```
class esem.emulator.Emulator(model, training_params, training_data, name='', gpu=0)
```

A class wrapping a statistical emulator

### training\_data

A wrapped representation of the training data

Type `esem.wrappers.DataWrapper`

### model

The underlying model which performs the emulation

Type `ModelAdaptor`

### name

A human-readable name for the model

Type `str`

```
__init__(model, training_params, training_data, name='', gpu=0)
```

### Parameters

- **model** (`ModelAdaptor`) – The (compiled but not trained) model to be wrapped
- **training\_params** (`pd.DataFrame` or *array-like*) – The training parameters (X)
- **training\_data** (`esem.wrappers.DataWrapper` or `xarray.DataArray` or `iris.Cube` or *array-like*) – The training data - the leading dimension should represent training samples (Y)
- **name** (`str`) – Human readable name for the model
- **gpu** (`int`) – The machine GPU to assign this model to

### Methods

---

```
__init__(model, training_params, training_data)
```

#### Parameters

- **model** (`ModelAdaptor`) – The (compiled but not trained) model to be wrapped

---

```
batch_stats(sample_points[, batch_size])
```

Return mean and standard deviation in model predictions over samples, without storing the intermediate predictions in memory to enable evaluating large models over more samples than could fit in memory

---

```
predict(x, *args, **kwargs)
```

Make a prediction using a trained emulator

---

```
train([verbose])
```

Train on the training data

## 6.2.2 esem.emulator.Emulator.train

`Emulator.train(verbose=False, **kwargs)`

Train on the training data

**Parameters** `verbose` (bool) – Print verbose training output to screen

## 6.2.3 esem.emulator.Emulator.predict

`Emulator.predict(x, *args, **kwargs)`

Make a prediction using a trained emulator

**Parameters**

- `x` (pd.DataFrame or *array-like*) – The points at which to make predictions from the model
- `args` – The specific arguments needed for prediction with this model
- `kwargs` – Any keyword arguments that might need to be passed through to the model

**Returns** Emulated prediction and variance with the same type as `self.training\_data`

## 6.2.4 esem.emulator.Emulator.\_predict

`Emulator._predict(x, *args, **kwargs)`

The (internal) predict interface used by e.g., a sampler. It is still in tf but has been post-processed to allow comparison with obs.

**Parameters**

- `x` (*array-like*) – The points at which to make predictions from the model
- `args` – The specific arguments needed for prediction with this model
- `kwargs` – Any keyword arguments that might need to be passed through to the model

**Returns** Emulated prediction and variance as either np.ndarray or tf.Tensor

## 6.2.5 esem.emulator.Emulator.batch\_stats

`Emulator.batch_stats(sample_points, batch_size=1)`

Return mean and standard deviation in model predictions over samples, without storing the intermediate predictions in memory to enable evaluating large models over more samples than could fit in memory

**Parameters**

- `sample_points` (pd.DataFrame or *array-like*) – The parameter values at which to sample the emulator
- `batch_size` (int) – The number of samples to calculate in each batch. This can be optimised to fill the available (GPU) memory

**Returns** The batch mean and standard deviation with the same type as `self.training\_data`

## 6.3 Sampler

This class defines the sampling interface currently used by the ABC and MCMC sampling implementations.

<code>Sampler</code>	A class that efficiently samples a Model object for posterior inference
<code>Sampler.sample</code>	This is the call that does the actual inference.

### 6.3.1 esem.sampler.Sampler

```
class esem.sampler.Sampler(model, obs, obs_uncertainty=0.0, interann_uncertainty=0.0,  
                           repres_uncertainty=0.0, struct_uncertainty=0.0, abs_obs_uncertainty=0.0,  
                           abs_interann_uncertainty=0.0, abs_repres_uncertainty=0.0,  
                           abs_struct_uncertainty=0.0)
```

A class that efficiently samples a Model object for posterior inference

```
__init__(model, obs, obs_uncertainty=0.0, interann_uncertainty=0.0, repres_uncertainty=0.0,  
        struct_uncertainty=0.0, abs_obs_uncertainty=0.0, abs_interann_uncertainty=0.0,  
        abs_repres_uncertainty=0.0, abs_struct_uncertainty=0.0)
```

#### Parameters

- **model** ([esem.emulator.Emulator](#))
- **obs** (`iris.cube.Cube` or `array-like`) – The objective
- **obs\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty in observations
- **repres\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty due to the spatial and temporal representitiveness of the observations
- **interann\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty introduced when using a model run for a year other than that the observations were measured in.
- **struct\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty in the model itself.
- **abs\_obs\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty in observations
- **abs\_repres\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty due to the spatial and temporal representitiveness of the observations
- **abs\_interann\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty introduced when using a model run for a year other than that the observations were measured in.
- **abs\_struct\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty in the model itself.

## Methods

---

`__init__(model, obs[, obs_uncertainty, ...])`

**Parameters**

- `model` (`esem.emulator.Emulator`)

---

`sampLe([prior_x, n_samples])`

This is the call that does the actual inference.

---

### 6.3.2 esem.sampler.Sampler.sample

`Sampler.sample(prior_x=None, n_samples=1)`

This is the call that does the actual inference.

It should call `model.sample` over the prior, compare with the objective, and then output samples from the posterior distribution

**Parameters**

- `prior_x` (`tensorflow_probability.distribution`) – The distribution to sample parameters from. By default it will uniformly sample the unit N-D hypercube
- `n_samples` (`int`) – The number of samples to draw

**Returns** `np.array` – Array of samples

### 6.3.3 MCMCSampler

---

`MCMCSampler`

Sample from the posterior using the TensorFlow Markov-Chain Monte-Carlo (MCMC) sampling tools.

---

`MCMCSampler.sample`

This is the call that does the actual inference.

---

## esem.sampler.MCMCSampler

`class esem.sampler.MCMCSampler(model, obs, **kwargs)`

Sample from the posterior using the TensorFlow Markov-Chain Monte-Carlo (MCMC) sampling tools. It uses a HamiltonianMonteCarlo kernel.

### Notes

Note that NaN observations will create ill-defined likelihoods.

`__init__(model, obs, **kwargs)`

**Parameters**

- `model` (`esem.emulator.Emulator`)
- `obs` (`iris.cube.Cube` or `array-like`) – The objective
- `obs_uncertainty` (`float`) – Fractional, relative (1 sigma) uncertainty in observations

- **repres\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty due to the spatial and temporal representitiveness of the observations
- **interann\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty introduced when using a model run for a year other than that the observations were measured in.
- **struct\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty in the model itself.
- **abs\_obs\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty in observations
- **abs\_repres\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty due to the spatial and temporal representitiveness of the observations
- **abs\_interann\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty introduced when using a model run for a year other than that the observations were measured in.
- **abs\_struct\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty in the model itself.

## Methods

---

`__init__(model, obs, **kwargs)`

**Parameters**

- **model** ([esem.emulator.Emulator](#))

---

`sample([prior_x, n_samples, kernel_kwargs, ...])` This is the call that does the actual inference.

---

### **esem.sampler.MCMCSampler.sample**

`MCMCSampler.sample(prior_x=None, n_samples=1, kernel_kwargs=None, mcmc_kwargs=None)`

This is the call that does the actual inference.

It should call `model.sample` over the prior, compare with the objective, and then output a posterior distribution

**Parameters**

- **prior\_x** (`tensorflow_probability.distribution`) – The distribution to sample parameters from. By default it will uniformly sample the unit N-D hypercube
- **n\_samples** (int) – The number of samples to draw
- **kernel\_kwargs** (dict) – kwargs for the MCMC kernel
- **mcmc\_kwargs** (dict) – kwargs for the MCMC sampler

**Returns** `np.array` – Array of samples

### 6.3.4 ABCSampler

<code>ABCSampler</code>	Sample from the posterior using Approximate Bayesian Computation (ABC).
<code>ABCSampler.sample</code>	Sample the emulator over <code>prior_x</code> and compare with the observations, returning <code>n_samples</code> of the posterior distribution (those points for which the model is compatible with the observations).
<code>ABCSampler.get_implausibility</code>	Calculate the implausibility of the provided sample points, optionally in batches.
<code>ABCSampler.batch_constrain</code>	Constrain the supplied sample points based on the tolerance threshold, optionally in batches.

#### esem.abc\_sampler.ABCSampler

```
class esem.abc_sampler.ABCSampler(model, obs, obs_uncertainty=0.0, interann_uncertainty=0.0,
                                   repres_uncertainty=0.0, struct_uncertainty=0.0,
                                   abs_obs_uncertainty=0.0, abs_interann_uncertainty=0.0,
                                   abs_repres_uncertainty=0.0, abs_struct_uncertainty=0.0)
```

Sample from the posterior using Approximate Bayesian Computation (ABC). This is a style of rejection sampling.

#### Notes

Note that emulator samples compared to NaN observations are always treated as ‘plausible’.

```
__init__(model, obs, obs_uncertainty=0.0, interann_uncertainty=0.0, repres_uncertainty=0.0,
        struct_uncertainty=0.0, abs_obs_uncertainty=0.0, abs_interann_uncertainty=0.0,
        abs_repres_uncertainty=0.0, abs_struct_uncertainty=0.0)
```

#### Parameters

- **model** (`esem.emulator.Emulator`)
- **obs** (`iris.cube.Cube` or `array-like`) – The objective
- **obs\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty in observations
- **repres\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty due to the spatial and temporal representitiveness of the observations
- **interann\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty introduced when using a model run for a year other than that the observations were measured in.
- **struct\_uncertainty** (float) – Fractional, relative (1 sigma) uncertainty in the model itself.
- **abs\_obs\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty in observations
- **abs\_repres\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty due to the spatial and temporal representitiveness of the observations
- **abs\_interann\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty introduced when using a model run for a year other than that the observations were measured in.
- **abs\_struct\_uncertainty** (float) – Fractional, absolute (1 sigma) uncertainty in the model itself.

## Methods

---

<code>__init__(model, obs[, obs_uncertainty, ...])</code>	<b>Parameters</b> <ul style="list-style-type: none"><li>• <b>model</b> (<a href="#">esem.emulator.Emulator</a>)</li></ul>
<code>batch_constraint(sample_points[, tolerance, ...])</code>	Constrain the supplied sample points based on the tolerance threshold, optionally in batches.
<code>get_implausibility(sample_points[, batch_size])</code>	Calculate the implausibility of the provided sample points, optionally in batches.
<code>sample([prior_x, n_samples, tolerance, ...])</code>	Sample the emulator over <i>prior_x</i> and compare with the observations, returning <i>n_samples</i> of the posterior distribution (those points for which the model is compatible with the observations).

---

### [esem.abc\\_sampler.ABCSampler.sample](#)

`ABCampler.sample(prior_x=None, n_samples=1, tolerance=0.0, threshold=3.0)`

Sample the emulator over *prior\_x* and compare with the observations, returning *n\_samples* of the posterior distribution (those points for which the model is compatible with the observations).

#### Parameters

- **prior\_x** (`tensorflow_probability.distribution` or `None`) – The distribution to sample parameters from. By default it will uniformly sample the unit N-D hypercube
- **n\_samples** (`int`) – The number of samples to draw
- **tolerance** (`float`) – The fraction of samples which are allowed to be over the threshold
- **threshold** (`float`) – The number of standard deviations a sample is allowed to be away from the obs

**Returns** `ndarray[n_samples]` – Array of samples conforming to the specified tolerance and threshold

### [esem.abc\\_sampler.ABCSampler.get\\_implausibility](#)

`ABCampler.get_implausibility(sample_points, batch_size=1)`

Calculate the implausibility of the provided sample points, optionally in batches.

*Note* this calculates an array of shape (*n\_sample\_points*, *n\_obs*) and so can easily exceed available memory if not used carefully.

#### Parameters

- **sample\_points** (`ndarray` or `DataFrame`) – The sample points to calculate the implausibility for
- **batch\_size** (`int`) – The size of the batches in which to calculate the implausibility (useful for large samples)

**Returns** `Cube` – A cube of the implausibility of each sample against each observation

## `esem.abc_sampler.ABCSampler.batch_constraint`

`ABCampler.batch_constraint(sample_points, tolerance=0.0, threshold=3.0, batch_size=1)`

Constrain the supplied sample points based on the tolerance threshold, optionally in batches.

Return a boolean array indicating if each sample meets the implausibility criteria:

$$I < T$$

Return True (for a sample) if the number of implausibility measures greater than the threshold is less than or equal to the tolerance

### Parameters

- **sample\_points** (`ndarray`) – An array of sample points which are to be emulated and compared with the observations
- **tolerance** (`float`) – The fraction of samples which are allowed to be over the threshold
- **threshold** (`float`) – The number of standard deviations a sample is allowed to be away from the obs
- **batch\_size** (`int`) – The size of the batches in which to perform the constraining (useful for large samples)

**Returns** `ndarray` – A boolean array which is true where the (emulated) samples are compatible with the observations and false otherwise

## 6.4 Wrappers

<code>ProcessWrapper</code>	This class handles applying any data pre- and post-processing by any provided DataProcessor
<code>DataWrapper</code>	Provide a unified interface for numpy arrays, Iris Cube's and xarray DataArrays.
<code>DataWrapper.name</code>	
<code>DataWrapper.data</code>	
<code>DataWrapper.dtype</code>	
<code>DataWrapper.wrap</code>	Wrap back in a cube if one was provided
<code>CubeWrapper</code>	
<code>DataArrayWrapper</code>	

### 6.4.1 esem.wrappers.ProcessWrapper

```
class esem.wrappers.ProcessWrapper(data, data_processors=None)
```

This class handles applying any data pre- and post-processing by any provided DataProcessor

```
__init__(data, data_processors=None)
```

Initialize self. See help(type(self)) for accurate signature.

#### Methods

<code>__init__(data[, data_processors])</code>	Initialize self.
<code>post_process(mean, variance)</code>	Any necessary reshaping or un-weightings are performed here
<code>pre_process(data)</code>	Any necessary rescaling or weightings are performed here

#### Attributes

---

```
data
```

---

### 6.4.2 esem.wrappers.DataWrapper

```
class esem.wrappers.DataWrapper(data, data_processors=None)
```

Provide a unified interface for numpy arrays, Iris Cube's and xarray DataArrays. Emulation outputs will be provided based on the provided input type, preserving appropriate metadata.

```
__init__(data, data_processors=None)
```

Initialize self. See help(type(self)) for accurate signature.

#### Methods

<code>__init__(data[, data_processors])</code>	Initialize self.
<code>name()</code>	
<code>wrap(data[, name_prefix])</code>	Wrap back in a cube if one was provided

#### Attributes

---

```
data
```

---

```
dtype
```

---

### 6.4.3 esem.wrappers.DataWrapper.name

`DataWrapper.name()`

### 6.4.4 esem.wrappers.DataWrapper.data

`property DataWrapper.data`

### 6.4.5 esem.wrappers.DataWrapper.dtype

`property DataWrapper.dtype`

### 6.4.6 esem.wrappers.DataWrapper.wrap

`DataWrapper.wrap(data, name_prefix='Emulated')`

Wrap back in a cube if one was provided

#### Parameters

- `data (np.array)` – Model output to wrap
- `name_prefix (str)` –

#### Returns

### 6.4.7 esem.wrappers.CubeWrapper

`class esem.wrappers.CubeWrapper(cube, data_processors=None)`

`__init__(cube, data_processors=None)`

Initialize self. See help(type(self)) for accurate signature.

#### Methods

<code>__init__(cube[, data_processors])</code>	Initialize self.
<code>name()</code>	
<code>wrap(data[, name_prefix])</code>	Wrap back in a cube if one was provided

### Attributes

---

data

---

dtype

---

## 6.4.8 esem.wrappers.DataArrayWrapper

```
class esem.wrappers.DataArrayWrapper(dataarray, data_processors=None)
```

```
__init__(dataarray, data_processors=None)
```

Initialize self. See help(type(self)) for accurate signature.

### Methods

<code>__init__(dataarray[, data_processors])</code>	Initialize self.
<code>name()</code>	
<code>wrap(data[, name_prefix])</code>	Wrap back in a xr.DataArray if one was provided

### Attributes

---

data

---

dtype

---

## 6.5 ModelAdaptor

<code>ModelAdaptor</code>	Provides a unified interface for all emulation engines within ESEm.
<code>SKLearnModel</code>	A wrapper around scikit-learn models.
<code>KerasModel</code>	A wrapper around Keras models
<code>GPFlowModel</code>	A wrapper around GPFlow regression models

### 6.5.1 esem.model\_adaptor.ModelAdaptor

```
class esem.model_adaptor.ModelAdaptor(model)
```

Provides a unified interface for all emulation engines within ESEm. Concrete classes must implement both `train()` and `predict()` methods.

See the [API documentation](#) for a list of concrete classes implementing this interface.

```
__init__(model)
```

Initialize self. See `help(type(self))` for accurate signature.

#### Methods

<code><u>__init__</u>(model)</code>	Initialize self.
<code>predict(*args, **kwargs)</code>	This is either the tf model which can be called directly, or a generator over the <code>model.predict</code> (in tf, so it's quick).
<code>train(training_params, training_data[, verbose])</code>	Train on the training data :return:

### 6.5.2 esem.model\_adaptor.SKLearnModel

```
class esem.model_adaptor.SKLearnModel(model)
```

A wrapper around [scikit-learn](#) models.

```
__init__(model)
```

Initialize self. See `help(type(self))` for accurate signature.

#### Methods

<code><u>__init__</u>(model)</code>	Initialize self.
<code>predict(*args, **kwargs)</code>	This is either the tf model which can be called directly, or a generator over the <code>model.predict</code> (in tf, so it's quick).
<code>train(training_params, training_data[, verbose])</code>	Train the RF model.

### 6.5.3 esem.model\_adaptor.KerasModel

```
class esem.model_adaptor.KerasModel(model)
```

A wrapper around [Keras](#) models

```
__init__(model)
```

Initialize self. See `help(type(self))` for accurate signature.

**Methods**

<code>__init__(model)</code>	Initialize self.
<code>predict(*args, **kwargs)</code>	This is either the tf model which can be called directly, or a generator over the model.predict (in tf, so it's quick).
<code>train(training_params, training_data[, ...])</code>	Train the Keras model.

## 6.5.4 esem.model\_adaptor.GPFlowModel

```
class esem.model_adaptor.GPFlowModel(model)
    A wrapper around GPFlow regression models
    __init__(model)
        Initialize self. See help(type(self)) for accurate signature.
```

**Methods**

<code>__init__(model)</code>	Initialize self.
<code>predict(*args, **kwargs)</code>	This is either the tf model which can be called directly, or a generator over the model.predict (in tf, so it's quick).
<code>train(training_params, training_data[, ...])</code>	Train on the training data :return:

## 6.6 DataProcessor

<code>DataProcessor</code>	A utility class for transparently processing (transforming) numpy arrays and un-processing TensorFlow Tensors to aid in emulation.
<code>Log</code>	Return $\log(x + c)$ where $c$ can be specified.
<code>Whiten</code>	Scale the data to have zero mean and unit variance
<code>Normalise</code>	Linearly scale the data to lie between [0, 1]
<code>Flatten</code>	Flatten all dimensions except the leading one
<code>Reshape</code>	Ensure the training data is the right shape for the ConvNet
<code>Recast</code>	Cast the data to a given type

## 6.6.1 esem.data\_processors.DataProcessor

```
class esem.data_processors.DataProcessor
    A utility class for transparently processing (transforming) numpy arrays and un-processing TensorFlow Tensors to aid in emulation.

    See the API documentation for a list of concrete classes implementing this interface.

    __init__()
        Initialize self. See help(type(self)) for accurate signature.
```

**Methods**


---

<code>__init__()</code>	Initialize self.
<code>process(data)</code>	
<code>unprocess(mean, variance)</code>	

---

**6.6.2 esem.data\_processors.Log**

```
class esem.data_processors.Log(constant=0.0)
    Return log(x + c) where c can be specified.

__init__(constant=0.0)
    Initialize self. See help(type(self)) for accurate signature.
```

**Methods**


---

<code>__init__([constant])</code>	Initialize self.
<code>process(data)</code>	
<code>unprocess(mean, variance)</code>	

---

**6.6.3 esem.data\_processors.Whiten**

```
class esem.data_processors.Whiten
    Scale the data to have zero mean and unit variance

__init__()
    Initialize self. See help(type(self)) for accurate signature.
```

**Methods**


---

<code>__init__()</code>	Initialize self.
<code>process(data)</code>	
<code>unprocess(mean, variance)</code>	

---

## 6.6.4 esem.data\_processors.Normalise

```
class esem.data_processors.Normalise
    Linearly scale the data to lie between [0, 1]

    __init__()
        Initialize self. See help(type(self)) for accurate signature.
```

### Methods

<code>__init__()</code>	Initialize self.
<code>process(data)</code>	
<code>unprocess(mean, variance)</code>	

---

## 6.6.5 esem.data\_processors.Flatten

```
class esem.data_processors.Flatten
    Flatten all dimensions except the leading one

    __init__()
        Initialize self. See help(type(self)) for accurate signature.
```

### Methods

<code>__init__()</code>	Initialize self.
<code>process(data)</code>	
<code>unprocess(mean, variance)</code>	

---

## 6.6.6 esem.data\_processors.Reshape

```
class esem.data_processors.Reshape
    Ensure the training data is the right shape for the ConvNet

    __init__()
        Initialize self. See help(type(self)) for accurate signature.
```

## Methods

<code>__init__()</code>	Initialize self.
<code>process(data)</code>	
<code>unprocess(mean, variance)</code>	

## 6.6.7 esem.data\_processors.Recast

```
class esem.data_processors.Recast(new_type)
    Cast the data to a given type
    __init__(new_type)
        Initialize self. See help(type(self)) for accurate signature.
```

## Methods

<code>__init__(new_type)</code>	Initialize self.
<code>process(data)</code>	
<code>unprocess(mean, variance)</code>	

## 6.7 Utilities

A collection of associated utilities which might be of use when performing typical ESEm workflows.

<code>plot_results</code>	Validation plot for LeaveOneOut
<code>validation_plot</code>	
<code>plot_parameter_space</code>	
<code>get_uniform_params</code>	Slightly convoluted method for getting a flat set of points evenly
<code>get_random_params</code>	Get points randomly sampling a (unit) N-dimensional space
<code>ensemble_collocate</code>	Efficiently collocate (interpolate) many ensemble members on to a set of (un-gridded) observations
<code>leave_one_out</code>	Function to perform LeaveOneOut cross-validation with different models.
<code>get_param_mask</code>	Determine the most relevant parameters in the input space using a regularised linear model and either the Aikake or Bayesian Information Criterion.

### 6.7.1 esem.utils.plot\_results

```
esem.utils.plot_results(ax, truth, pred, title)
    Validation plot for LeaveOneOut
```

### 6.7.2 esem.utils.validation\_plot

```
esem.utils.validation_plot(test_mean, pred_mean, pred_var, figsize=(7, 7), minx=None, miny=None,
                           maxx=None, maxy=None, ax=None)
```

### 6.7.3 esem.utils.plot\_parameter\_space

```
esem.utils.plot_parameter_space(df, nbins=100, target_df=None, smooth=True, xmins=None, xmaxs=None,
                                 fig_size=(8, 6))
```

### 6.7.4 esem.utils.get\_uniform\_params

```
esem.utils.get_uniform_params(n_params, n_samples=5)
```

Slightly convoluted method for getting a flat set of points evenly sampling a (unit) N-dimensional space

#### Parameters

- **n\_params** (int) – The number of parameters (dimensions) to sample from
- **n\_samples** (int) – The number of uniformly spaced samples (in each dimension)

**Returns** ndarray –  $n\_samples^{**}n\_params$  parameters uniformly sampled

### 6.7.5 esem.utils.get\_random\_params

```
esem.utils.get_random_params(n_params, n_samples=5)
```

Get points randomly sampling a (unit) N-dimensional space

#### Parameters

- **n\_params** (int) – The number of parameters (dimensions) to sample from
- **n\_samples** (int) – The number of parameters to (randomly) sample

### 6.7.6 esem.utils.ensemble\_collocate

```
esem.utils.ensemble_collocate(ensemble, observations, member_dimension='job')
```

Efficiently collocate (interpolate) many ensemble members on to a set of (un-gridded) observations

---

**Note:** This function requires both Iris and CIS to be installed

---

### Parameters

- **ensemble** (GriddedData) – The ensemble of (model) samples to interpolate on to the observations
- **observations** (UngriddedData) – The observations on to which the observations will be sampled
- **member\_dimension** (str) – The name of the dimension which represents the ensemble members in *ensemble*

**Returns** `col_ensemble` (iris.cube.Cube) – The ensemble values interpolated on to the observation locations, with the ensemble members along the leading dimension.

## 6.7.7 esem.utils.leave\_one\_out

`esem.utils.leave_one_out(Xdata, Ydata, model='RandomForest', **model_kwargs)`

Function to perform LeaveOneOut cross-validation with different models.

### Parameters

- **Xdata** (*array-like* of shape (n\_samples, n\_features)) – Parameter values
- **Ydata** (*array-like* of shape (n\_samples,)) – Target values.
- **model** ({'RandomForest', 'GaussianProcess', 'NeuralNet'}, *default* 'RandomForest')
- **model\_kwargs** (dict) – More arguments to pass to the model.

**Returns** `output` (list of n\_samples (truth, prediction, variance) tuples) – which can then be passed to `validation_plot()`

## 6.7.8 esem.utils.get\_param\_mask

`esem.utils.get_param_mask(X, y, criterion='bic', **kwargs)`

Determine the most relevant parameters in the input space using a regularised linear model and either the Aikake or Bayesian Information Criterion.

### Parameters

- **X** (*array-like* of shape (n\_samples, n\_features)) – Parameter values
- **y** (*array-like* of shape (n\_samples,)) – target values.
- **criterion** ({'bic', 'aic'}, *default* 'bic') – The information criteria to apply for parameter selection. Either Aikake or Bayesian Information Criterion.
- **kwargs** (dict) – Further arguments for sklearn.feature\_selection.SelectFromModel

**Returns** `mask` (ndarray) – A boolean array of shape [# input features], in which an element is True iff its corresponding feature is selected for retention.



## ESEM DESIGN

Here we provide a brief description of the main architectural decisions behind the design for ESEm in order to hopefully make it easier for contributors and users alike to understand the various components and how they fit together.

### 7.1 Emulation

We try to provide a seamless interface for users whether they provide iris Cube's, xarray DataArray's or numpy ndarrays. This is done using the `esem.wrappers.DataWrapper` and associated subclasses, which keep a copy of the provided object but only exposes the underlying numpy array to the emulation engines. When the data is requested from this wrapper using the `esem.wrappers.DataWrapper.wrap()` method then it will return a copy of the input object (Cube or DataArray) with the data replaced by the emulated data.

This layer will also ensure the underlying (numpy) data is wrapped in a `esem.wrappers.ProcessWrapper`. This class transparently applies any requested `esem.data_processors.DataProcessor` in sequence.

The user can then create an `esem.emulator.Emulator` object by providing a concrete `esem.model_adaptor.ModelAdaptor` such as a `esem.model_adaptor.KerasModel`. There are two layers of abstraction here: The first to deal with different interfaces to different emulation libraries; and the second to apply the pre- and post-processing and allow a single `esem.emulator.Emulator.batch_stats()` method. The `esem.emulator.Emulator._predict()` provides an important internal interface to the underlying model which reverts any data-processing but leaves the emulator output as a TensorFlow Tensor to allow optimal sampling.

The top-level functions `esem.gp_model()`, `esem.cnn_model()` and `esem.rf_model()` provide a simple interface for constructing these emulators and should be sufficient for most users.

### 7.2 Calibration

We try and keep this interface very simple; a `esem.sampler.Sampler` should be initialised with an `esem.emulator.Emulator` object to sample from, some observations and associated uncertainties. The only method it has to provide is `esem.sampler.Sampler.sample()` which should provide sample  $\theta$  from the posterior.

Wherever possible these samplers should take advantage of the fact that the `esem.emulator.Emulator._predict()` method returns TensorFlow tensors and always prefer to use them directly rather than using `esem.emulator.Emulator.predict()` or calling `.numpy()` on them. This allows the sampling to happen on GPUs where available and can substantially speed-up sampling.

The `esem.abc_sampler.ABCSampler` extends this interface to include both `esem.abc_sampler.ABCSampler.get_implausibility()` and `esem.abc_sampler.ABCSampler.batch_constraint()` methods. The first allows inspection of the effect of different observations on the constraint and the second allows a streamlined approach for rejecting samples in batch, taking advantage of the large amounts of memory available on modern GPUs.



---

**CHAPTER  
EIGHT**

---

**GLOSSARY**

**array\_like** Any object that can be treated as a numpy array, i.e. can be indexed to retrieve numerical values. Typically not a tensorflow Tensor.



---

**CHAPTER  
NINE**

---

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