

## Recap of where we are (were from previous lecture)

- Given a pipeline / model we "can" tune each of the hyperparameters by constructing a validation curve.
- Problems . . . (manual approach is not practical anything but small problems)
  - Can have a huge (100s 1,000s) number of parameters.
  - A sequence of 1-dimensional searches is not the same as one *d*-dimensional search interplay between hyper-parameters.
- Approaches / Techniques . . .
  - Grid Search Systematic, regular, predetermined deterministic sample of parameter space.
  - Random Search Non-adaptive, random sample of parameter space.
  - Bayesian Search Adaptive, random sample of parameter space.

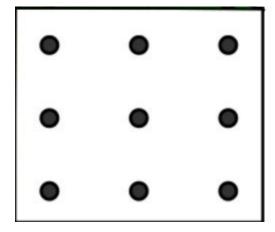
## Grid Search

Validation curves are important but examining at each parameter individually is time consuming and does not take into account interactions between parameters — in effect it is a series of 1D semi-manual searches.

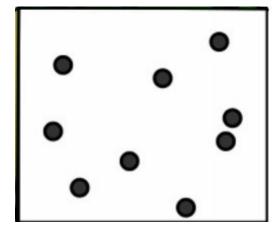
#### Grid Search

Try all available parameters combinations, 1 by 1, and choose the one with the best cross validation results.

Grid Search



Random Search



## Grid Search (sklearn.model\_selection.GridSearchCV)

#### Grid Search

Try all available parameters combinations, 1 by 1, and choose the one with the best cross validation results.

### >Drawbacks >

- Still very slow trying ALL combinations of ALL parameters with no modification of search based on results found to date.
- ALL combinations  $\implies$  every additional hyperparameter to vary multiplies the number of iterations you need to complete.
- It can work only with discrete values.

  If the global optimum is on n\_estimators=550, but you are doing GridSearchCV from 100 to 1000 with step 100, you will never reach the optimal point.

### Stratagies

- You need know / guess the approximate region of the optimum to start.
- To mitigate against drawbacks, do multiple lower dimensional grid searches, or repeat search with narrower grids and smaller step sizes.

## Grid Search Example 1

```
from sklearn.svm import SVC
from sklearn.model_selection import GridSearchCV
pipeline = Pipeline([
    ('scl', StandardScaler()),
     ('clf', SVC(random_state=SEED))])

param_range = np.logspace(-4,4,10)
param_grid = [{
        'clf__gamma': param_range,
        'clf__C': param_range,
        'clf__kernel': ['rbf']
}]
gs = GridSearchCV(estimator=pipeline,
        return_train_score=True,
        param_grid=param_grid, scoring='accuracy', cv=10, n_jobs=-1)
```

- Switched from LogisticRegression to SVM more hyperparameters to tweak.
- How many parameter combinations were generated/used?

## Grid Search Example 1

To perform grid search, we call the fit method as usual ...

```
gs = gs.fit(X_train, y_train)
print(gs.best_score_)
print(gs.best_params_)

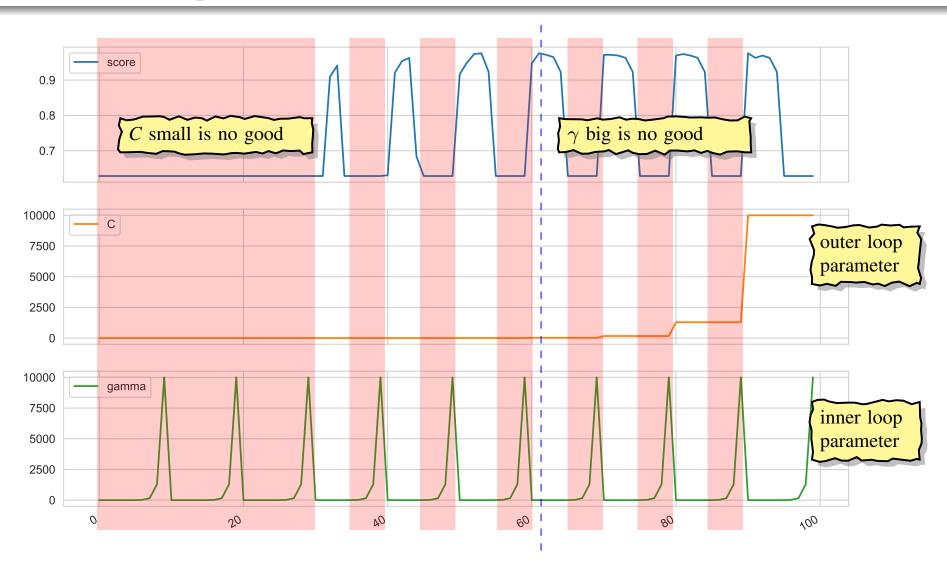
0.9757971014492753
{'clf__C': 2.782559402207126, 'clf__gamma': 0.046415888336127774, 'clf__kernel': 'rbf'}
```

- CV score improved from 0.971 with default parameter values to 0.978.
- GridSearchCV with option return\_train\_score=True, returns results of each combination determine effect of hyperparameters to refine grid search.

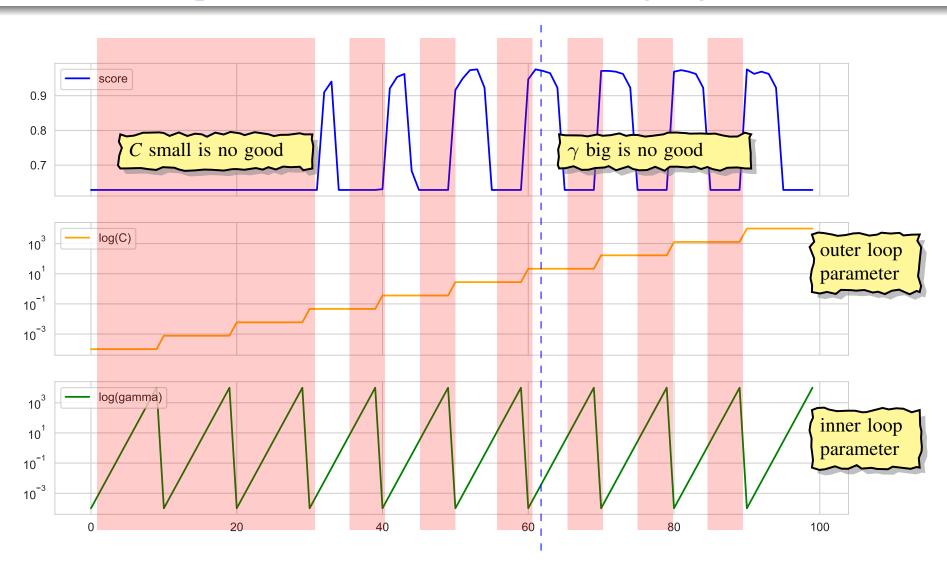
```
df_gs = pd.DataFrame(np.transpose([
    gs.cv_results_["mean_test_score"],
    gs.cv_results_["param_clf__C"].data,
    gs.cv_results_["param_clf__gamma"].data]),
    columns=['score', 'C', 'gamma'])

df_gs.plot(subplots=True,figsize=(12, 8))
plt.savefig("gs__svm__C_gamma.pdf", bbox_inches="tight")
plt.show()
```

# Grid Search Example 1 (C = 10, $\gamma = 0.01$ )



# Grid Search Example 1 (C = 10, $\gamma = 0.01$ ) (Using log scales)



## Grid Search Example 2

GridSearch can accept a list of dictionaries so that incompatible hyperparameter combinations can still be searched. For example, different SVM kernels have different parameters:

```
pipeline = Pipeline([
    ('scl', StandardScaler()),
    ('clf', SVC(random_state=SEED))
1)
from sklearn.model_selection import GridSearchCV
param\_range = [0.0001, 0.001, 0.01, 0.1, 1.0, 10.0, 100.0, 1000.0]
param_grid = [
        'clf__C': param_range,
        'clf kernel': ['linear']
    },
        'clf__gamma': param_range,
        'clf__C': param_range,
        'clf__kernel': ['rbf']
    }
                                    0.9780676328502415
                                    {'clf__C': 545.5594781168514, 'clf__gamma': 0.0001}
```

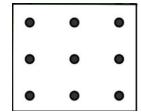
• How many hyperparameter combinations are now generated?

### Random Search

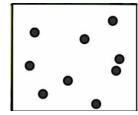
#### Random Search

Try random hyperparameter combinations and choose the one with the best cross validation results.

#### Grid Search



#### Random Search



### Advantages

- On every step random search varies all parameters  $\Rightarrow$  doesn't spend time on meaningless parameters.
- On average finds near optimal parameters much faster than Grid search.
- It is not limited by grid when optimising continuous parameters.

### Disadvantages )

- It may not find the global optimal parameter on a grid.
- All steps are independent. Does not use information about the results gathered to inform search.

### Random Search

```
pipeline = Pipeline([
   ('scl', StandardScaler()),
   ('clf', SVC(kernel='rbf', random_state=SEED))
])
from sklearn.model_selection import RandomizedSearchCV
from scipy.stats import randint
param_range = np.logspace(-4,4,20)
param_grid = {'clf__C': param_range,
             'clf__gamma': param_range}
rs = RandomizedSearchCV(estimator=pipeline,
   param_distributions=param_grid,
   n_iter = 100, random_state=SEED,
   return_train_score=True,
   scoring='accuracy', cv=10, n_jobs=-1)
```

• This is the same setup as used in GridSearch but doubled the number of parameter values in both C and  $\gamma$ , so search space is 400 points (not 100). Sampling 100 points.

## Random Search Example

To perform grid search, we call the fit method as usual ...

```
rs.fit(X_train, y_train)
print(rs.best_score_)
print(rs.best_params_)

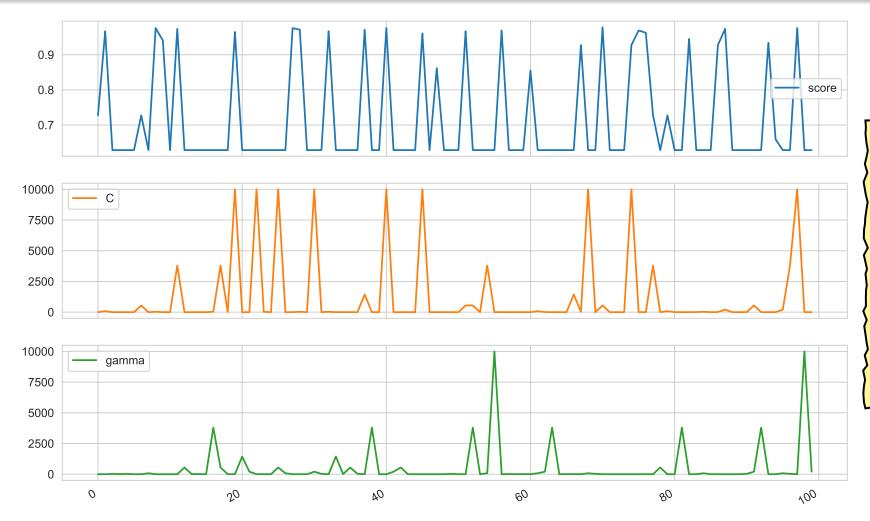
0.9780676328502415
{'clf__gamma': 0.0001, 'clf__C': 545.5594781168514}
```

• Sampling half the number of points (50) in a search space 4 times larger, RandomSearch outperforms GridSearch.

```
df_rs = pd.DataFrame(np.transpose([
    rs.cv_results_["mean_test_score"],
    rs.cv_results_["param_clf__C"].data,
    rs.cv_results_["param_clf__gamma"].data]),
    columns=['score', 'C', 'gamma'])

df_rs.plot(subplots=True, figsize=(12, 8))
plt.savefig("rs__example_1.pdf", bbox_inches="tight")
plt.show()
```

# Random Search Example ( $C = 10000, \gamma = 0.0002636$ )



Every step is completely random. ⇒ does not spent time on useless parameters, but does not use the information gathered on the first steps to improve outcomes of the latter ones.

### The Problem

Both GridSearch and RandomSearch ignore information gained during the search, by using such information can searches improve?

- What information would be useful?
  - Function value
  - Function derivative (gradient) gives direction in which the function is decreasing (locally).
- Why should we do this?
  - It seems reasonable that lower function values would be clustered so focus search near best function values found to date.
  - Making multiple decisions based on local information could locate global minimum.
- What can go wrong?
  - Multiple local minimum can get stuck in a side valley.
    - There will be situations in which we should make decisions against current function value information see Simulated Annealing.
  - Gradient could be flat (so no direction to go in), or worse almost flat (some algorithms become unstable), or non-existent.

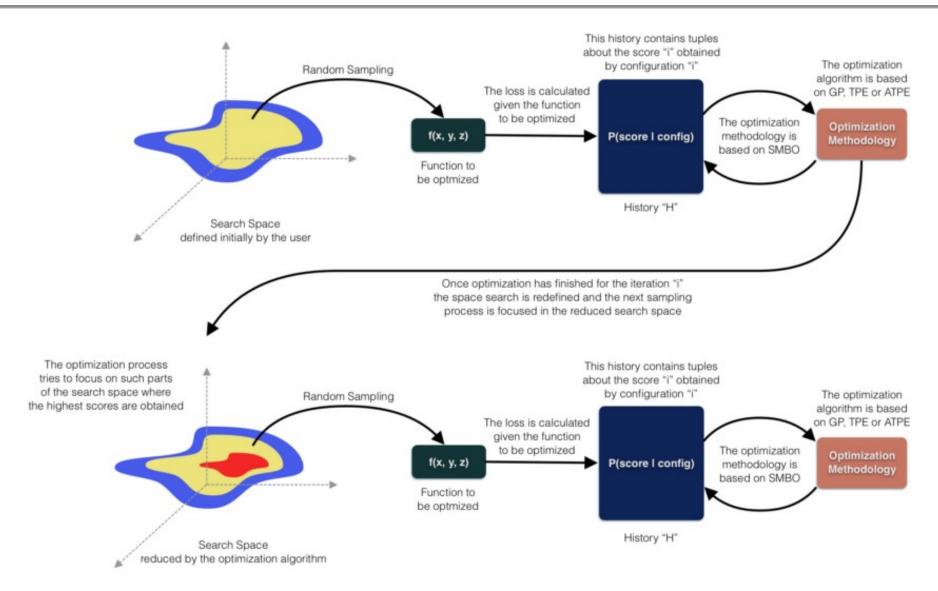
## **Bayesian Optimisation**

### **Sequential model-based optimisation (SMBO)**

(also known as Bayesian optimisation) is a general technique for function optimisation that uses information from earlier function evaluations to refine the optimisation search — with the aim of minimising number of function calls.

- For sufficiently complex functions, SMBO are some of the most call-efficient (in terms of function evaluations) optimisation methods currently available.
- Compared with standard optimisation strategies such as conjugate gradient descent methods, model-based optimisation algorithms invest more time between function evaluations in order to reduce the number of function evaluations overall.
- Takes advantage of smoothness without analytic gradient.
- Supports real-valued, discrete, and conditional variables.
- Performs relatively will in high dimensions —- hundreds of variables, even with budget of just a few hundred function evaluations, i.e., number of function calls is  $\mathcal{O}(\text{dimensions})$  not  $\mathcal{O}(e^{\text{dimensions}})$ .

# Sequential model-based optimisation (in HyperOpt)



## Hyperopt

### Hyperopt: Distributed Asynchronous Hyper-parameter Optimisation

• Python library for optimising over awkward search spaces with real-valued, discrete, and conditional dimensions.

github.com/hyperopt/hyperopt

- Install (Anaconda) or use pip conda install -cy conda-forge hyperopt
- Supports parallel evaluation
  - Uses MongoDB to share results of functions evaluations..

### hyperopt-sklearn

• Hyperopt-based model selection algorithms in scikit-learn

github.com/hyperopt/hyperopt-sklearn

Install using pip.

Purpose of hyperopt is not to always to completely optimise the search space but simply to do better than people.

— J Bergstra, SciPy 2013

### Resources

• Tune Hyperparameters for Classification Machine Learning Algorithms

```
machinelearningmastery.com/
hyperparameters-for-classification-machine-learning-algorithms/
List of important hyper-parameters for skilearn classifiers
```

- Hyperopt: A Python Library for Optimizing the Hyperparameters of Machine Learning Algorithms conference.scipy.org/proceedings/scipy2013/pdfs/bergstra\_hyperopt.pdf SCIPY Conference 2013 Paper
- A Conceptual Explanation of Bayesian Hyperparameter Optimization for Machine Learning

```
towardsdatascience.com/
a-conceptual-explanation-of-bayesian-model-based-hyperparameter-optimization-for-machine Relatively high level comparison of hyper-parameter tuning approaches.
```