

2957

# Chapter 1

2958

## Time series analysis

2959

### 1.1 catch22 features

2960

Table 1.1 lists *catch22* features.

2961

### 1.2 UMAP hyperparameters

2962

UMAP has several hyperparameters, of which four have major effects on the embedding:

2963

2964

- The *number of neighbours* ( $n$ ) to consider when approximating the local metric controls how the method balances local and global structure in the data. With low values of this parameter, the algorithm concentrates on very local structure, potentially to the detriment of the big picture. As the value increases, the algorithm ‘glues’ more nodes together to form clusters.

2965

2966

2967

2968

2969

- The *largest embedding dimension* ( $d$ ) controls the number of dimensions the data is reduced to. In other words, it controls whether the resulting map is one-dimensional, two-dimensional, three-dimensional, or of higher dimensions.

2970

2971

2972


brief intro to  
explain what UMAP  
does ("dimensionality reduction")  
+ references

Feature name	Description
DN_HistogramMode_5	Mode of z-scored distribution (5-bin histogram)
DN_HistogramMode_10	Mode of z-scored distribution (10-bin histogram)
SB_BinaryStats_mean_longstretch1	Longest period of consecutive values above the mean
DN_OutlierInclude_p_001_mdrmd	Time intervals between successive extreme events above the mean
DN_OutlierInclude_n_001_mdrmd	Time intervals between successive extreme events below the mean
first_1e_ac	First 1/e crossing of autocorrelation function
firstMin_acf	First minimum of autocorrelation function
SP_Summaries_welch_rect_area_5_1	Total power in lowest fifth of frequencies in the Fourier power spectrum
SP_Summaries_welch_rect_centroid	Centroid of the Fourier power spectrum
FC_LocalSimple_mean3_stderr	Mean error from a rolling 3-sample mean forecasting
CO_trev_1_num	Time-reversibility statistic, $\langle (x_{t+1} - x_t)^3 \rangle_t$
CO_HistogramAMI_even_2_5	Automutual information, $m = 2, \tau = 5$
IN_AutoMutualInfoStats_40_gaussian_fmfi	First minimum of the automutual information function
MD_hrv_classic_pnn40	Proportion of successive differences exceeding $0.04\sigma$
SB_BinaryStats_diff_longstretch0	Longest period of successive incremental decreases
SB_MotifThree_quantile_hh	Shannon entropy of two successive letters in equiprobable 3-letter symbolization
FC_LocalSimple_mean1_ttauresrat	Change in correlation length after iterative differencing
CO_Embed2_Dist_tau_d_expfit_meandiff	Exponential fit to successive distances in 2-d embedding space
SC_FluctAnal_2_dfa_50_1_2_logi_prop_r1	Proportion of slower timescale fluctuations that scale with DFA (50% sampling)
SC_FluctAnal_2_rsrangefit_50_1_logi_prop_r1	Proportion of slower timescale fluctuations that scale with linearly rescaled range fits
SB_TransitionMatrix_3ac_sumdiagcov	Trace of covariance of transition matrix between symbols in 3-letter alphabet
PD_PeriodicityWang_th0_01	Periodicity measure of Wang et al. (2007)

**Table 1.1:** *catch22* features, adapted from Lubba et al. (2019).

- 2973 • The *minimal distance* (`min_dist`) controls the desired separation between  
2974 close points in the embedding space. Specifically, this parameter controls  
2975 how tightly the algorithm is allowed to pack points together. With low  
2976 values, the visualisation forms ‘clumps’.
- 2977 • The previous hyperparameters are numerical, but the *metric* hyperparam-  
2978 eter instead specifies the distance metric that is used to compute distances  
2979 in the ambient space of the input data. For example, this metric can be the  
2980 Euclidean distance, the cosine distance, or other metrics used to compute  
2981 the distances between two vectors of numerical data.

## 2982 1.3 Classification pipeline

2983 In machine learning, classification is defined as the process of identifying a cate-  
2984 gory that a piece of input data belongs to. In this section, the classification task  
2985 is identifying whether a time series (input data) is oscillatory (belongs to one  
2986 category of two) or non-oscillatory (belongs to the other category of two). 

2987 A typical classification pipeline can be described by the following steps:

- 2988 1. *Pre-processing of data*: Input data is cleaned or normalised. For example,  
2989 to classify oscillatory time series, the input time series may be normalised  
2990 to give similar dynamic ranges.
- 2991 2. *Labelling*: Each piece of input data has a label assigned to it to denote  
2992 which category it belongs to. For example, to classify oscillatory time series,  
2993 a human can subjective assign the label ‘0’ for non-oscillatory time series  
2994 and ‘1’ for oscillatory time series, for a total of two categories.

3. *Featurisation*: Input data converted to feature vectors in the process of featurisation. This process uses domain knowledge related to the type or origin of the data to define characteristics of the data that may be useful for classification.
4. *Train-test split*: The input data set is then randomly divided into a training data set and a test data set.
5. *Training of model*: The machine learning model is then fit on the (featurised) training data set and its labels to fit parameters in the model.
6. *Evaluation of model on test dataset*: The model, trained on the training dataset, is used to predict the labels of data in the (featurised) test data set. The performance of the model is then evaluated on the test data set. This evaluation is based on computing quantities that express how well the model assigns labels to data, compared to the labels defined earlier.

## 1.4 Gillespie noise

To define the Gillespie algorithm, consider such a system with  $M$  reactions  $R_1, \dots, R_j, \dots, R_M$  involving  $N$  species  $S_1, \dots, S_i, \dots, S_N$  in a fixed volume  $V$  at thermal equilibrium. Let  $X_i(t)$  represent the number of molecules of  $S_i$  at time  $t$ , and the state vector

$$\mathbf{X}(t) := [X_1(t), \dots, X_N(t)] \quad (1.1)$$

thus gives the state of the system at any given time  $t$ .

Each reaction  $R_j$  is described by two quantities:

1. A state-change vector  $\mathbf{v}_j := [v_{1,j}, \dots, v_{N,j}]$  which defines how the stoichiometry of the system changes if the reaction occurs.  $v_{i,j}$  represents the change in the stoichiometry of  $S_i$  when  $R_j$  occurs.
2. A propensity function  $a_j$ , which gives the probability, given a the state  $\mathbf{X}(t) = \mathbf{x}$ , that one  $R_j$  reaction occurs in the volume  $V$  within the following short time interval  $[t, t + dt)$ . This function is defined by

$$a_j(\mathbf{x})dt = k_j \prod_{n=1}^N \mathbf{v}_n S_n \quad (1.2)$$

*Handwritten notes: A red arrow points from the text "short time interval" to the  $dt$  in the equation. A red circle is drawn around the  $dt$  with a question mark above it. Another red circle is drawn around the  $\mathbf{v}_n$  term. To the right, red text says "not sure this is correct".*

where  $k_j$  is the rate constant of reaction  $R_j$ .

The Gillespie algorithm aims to estimate the state vector given the initial state  $\mathbf{X}(t_0) = \mathbf{x}_0$ . It does so by iteratively choosing the next reaction that occurs, based on its propensity probability, and then choosing its firing time based on a probability distribution. Combining these simulations gives a trajectory of state vectors across the time course of interest. In detail, the direct Gillespie algorithm can be defined as stated in algorithm 1 (Gillespie, 2007):

*Handwritten notes: A red arrow points from the text "propensity" to the word "propensity" in the text. A red arrow points from the text "predict the temporal evolution of" to the word "propensity".*

---

**Algorithm 1:** Direct method of the Gillespie algorithm

---

**Input:** Stochastic model (with species  $S_1, \dots, S_i, \dots, S_N$  and reactions  $R_1, \dots, R_j, \dots, R_M$ , along with a state-change vector  $\mathbf{v}_j$  and a rate constant  $k_i$  for each reaction  $R_j$ ); initial time  $t_0$ ; and initial model state  $\mathbf{X}(t_0) = \mathbf{x}_0$

**Output:** Trajectory of state vectors  $\mathbf{X}(t)$ , with  $t$  taking discrete values in  $[t_0, t_{\max}]$

**while**  $t < t_{\max}$  **do**

    Calculate the propensities  $a_j(\mathbf{x})$  based on the current state  $\mathbf{x}$ ;

    Calculate the combined propensity  $a_0(\mathbf{x}) = \sum_j a_j(\mathbf{x})$ ;

    Generate two random numbers  $r_1$  and  $r_2$ , both from the uniform distribution  $U(0, 1)$ ;

    Choose the next reaction  $R_j$ , with  $j$  given by the smallest integer that satisfies  $\sum_{j'}^j a_{j'}(\mathbf{x}) > r_1 a_0(\mathbf{x})$ ;

    Calculate the time to the next reaction  $\tau = \frac{1}{a_0(\mathbf{x})} \ln(\frac{1}{r_2})$ ;

    Simulate the next reaction by updating the state vector  $\mathbf{x} \leftarrow \mathbf{x} + \mathbf{v}_j$  and store the new vector in  $\mathbf{X}(t)$ ;

    Update the time by  $t \leftarrow t + \tau$  and store the new time;

**end**

**return** Trajectory of state vectors  $\mathbf{X}(t)$  for a vector of times  $t$ ;

---