

## CH5115 PROJECT

Q1a) Goal of the Paper:

The main goal of the paper is to detect 'change-points' <sup>online</sup> in the generative parameters of a data sequence adopting a Bayesian approach.

They make the assumption that the parameters before and after a change point are independent of each other. They approach this problem by estimating the length of the current "run".

Contribution of the paper and Why is it useful

~~The authors state that~~

The authors state that most Bayesian approaches prior (pur not intended) to their approach use retrospective and offline approach to change point detection. If that is so, then <sup>work</sup> ~~this work contribution~~ is of immense value.

Many applications essentially require online detection, let's say a plant where certain vital states of <sup>system</sup> are being estimated or consider a medical application where ~~the~~ <sup>some</sup> signal indicating the health of the vital system of a man is processed. Immediate detection of abnormalities might prove to be necessary.

The authors also cite an example of brightness change detection in vision systems which ~~entirely~~ <sup>mandate</sup> the requirement of online detection.

Although frequentist approaches have been developed, Bayesian gives the additional advantages it always carries with it: i) use of prior information and a

ii) probabilistic distribution of the required parameters. The latter advantage might prove to be useful in many scenarios.

The approach is versatile in the sense that there is a clear cut separation between the implementation of the change-point algorithm and the implementation of the model. This means we can use this algorithms across <sup>processes with</sup> different models. This project <sup>comes</sup> <sup>^</sup>

itself is a nice example for this:

→ In q1) we apply the algorithm to predict parameters (mean) of a distribution.

→ In q2)

→ In q1) we use it to find change points in which the parameter of the distribution is changing

→ In q2) we use it to find change points in an AR(1) model where the variance of the endogenous white-noise driving force changes (variance)

So the algorithm works for 2 very different scenarios!

### How the Algorithm Works.

To detect changepoints the authors try to compute the run length distribution (conditioned on the data)

Run length is defined as the number of datapoints which has been following a specific generative process. Alternatively, it is the time since the last change point occurred

By its very definition  $r_{t+1} = \begin{cases} r_t + 1 & \text{probability } p \\ 0 & \text{probability } 1-p \\ \text{any other value} & 0 \end{cases}$   
where  $p$  is a valid probability





[Example of  
sum length  
values]

So by predicting  $r_t$  we will know whether  
change point has occurred at the datapoint  $x_t$ ,  
because at change point  $r_t$  will be 0

negd. value :  $P(r_t | x_{1:t}) \neq r_t$

we need  $\frac{P(r_t | x_{1:t})}{P(x_{1:t})}$

(Note :  $r_t \leq n$  where  
 $n$  is the index of the  
current data pt)

to get note that  $P(x_{1:t}) = \sum_{r_t \neq r_t} P(r_t | x_{1:t})$   
marginal.

So we need the joint distribution.

For that, after some neat simplifications (and  
assumption of Markov Property - for predicting  $r_t$   
we need only  $r_{t-1}$ ) the authors derive a

recursive eqn of the form:

$$P(r_t | x_{1:t}) = \sum P(r_t | r_{t-1}) P(x_t | r_{t-1}, x_{t-1}^{(r)}) P(r_{t-1}, x_{1:t-1})$$

$P(r_t | r_{t-1})$  can be modelled using a hazard function  
(like geometric distribution)

Now we need the term:  $P(x_t | r_{t-1}, x_t^{(r)})$

We can write it as some

$$P(x_{t+1} | x_{1:t}) = \int_{\Theta} P(x_{t+1} | \theta) P(\theta | x_{1:t}) d\theta$$

where  $P(x_{t+1} | \theta)$  is the so called predictive and  $\theta$  is the parameter(s) involved in the DGP of  $x$ . And  $P(\theta | x_{1:t})$  is the posterior

Here the author's usual exponential family likelihoods and conjugate exponential prior  $P(\theta | X, V)$  which ensures an exponential posterior.

Because this allows modularity wherein we can update the ~~param~~ hyperparameters  $X$  and  $V$  independently of run length predictions (thus ensuring the adaptation of change point algorithm and model).

Thus, For initial conditions we have the paper considers 2 cases!

- ①  $P(r_0 = 0) = 1$  (change point occurred a priori before the first datum.) of data
- ② Some recent observations of a subset is used to get a prior over the initial run length as the normalized survival function.

$$P(r_t = T) = \frac{1}{Z} S(T) ; S(T) = \sum_{t=T-11}^{\infty} \frac{1}{2} g_{\text{net}}(g=t)$$

$Z \rightarrow$  normalising agent

Thus, putting together all these pieces, the algorithm is complete! (we get a value for  $P(r_t | x_{1:t})$ )

### PERFORMANCE & SCOPE FOR IMPROVEMENT

- Three datasets are considered in the paper and the algorithm performs reasonably well in all 3 cases. It predicts the changepoints within a little error
- It detects changes in data mean (NMR well log data) and variance (DOW TONES RETURNS data)
- The algorithm is computationally very friendly (as long as  $P(x_{k+1} | r_t)$  is evaluated to be a simple function / exact solution for the integral exists)  
So it can easily keep up pace with the arrival of new data.
- However despite its very good performance in the project data as well as the original datasets the authors have used, there are some concerns.



### (i) Underflow error:

( $< 1$ )

As we keep on multiplying probabilities, the values of the joint probabilities get closer and closer to zero & eventually become so insignificant the computer can't evaluate resulting in the failure of the algorithm.

One alternative is to take logarithms of probabilities and work with those instead. But other ~~as well~~ more robust solutions are needed.

A closely related issue is that the  $\mathbf{r}(r_c, \mathbf{r}_p)$  vector (along  $\mathbf{r}_c$ ) is likely to be sparse, so ~~some~~ some approximations or modification in the implementation needs to be made to ensure optimal ~~at~~ space usage (Extend as well since this is  $O(N)$  complexity).

### (ii) Initial conditions:

This is an issue which we face in Bayesian estimation - choice of prior & hyperparameters.

One needs to test for the robustness of this algorithm in case there are errors in the prior or the model or the zero length assumption ( $P(L_0 = 0) = 1$ )

→ Further improvements could also be made for approximations of the predictive in case the exact solution is unavailable

As a closing remark, I would like to state that  
the algorithm is immensely useful <sup>is</sup> as long as  
the applications are chosen carefully and <sup>is</sup> definitely  
a big contribution for the field of online  
Bayesian changepoint detection