

User Guide of Tracer 1.6.0

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25/02/11

The new function of tracer allows analyzing values in 3 different trace types:

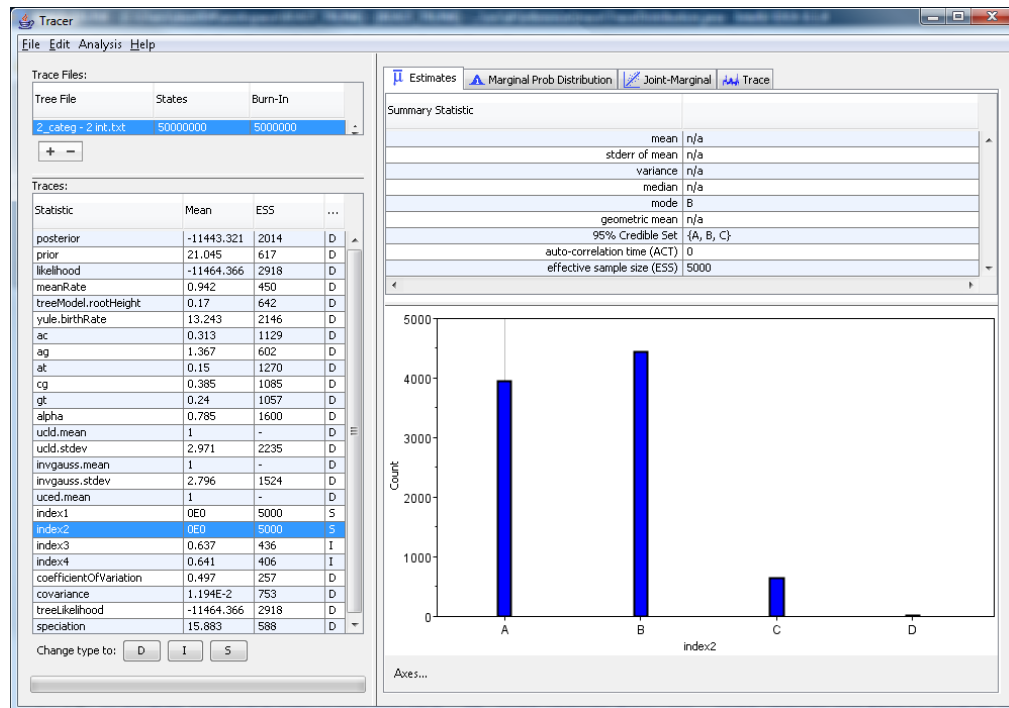
- Double (D),
- Integer (I),
- String (S).

1. Define Trace Type in Log File

The real (D) type is default. For other two types, the user needs to define in the top comment section of given log files. The format starts at keyword, *integer* or *string*, and follows by the set of trace names to be defined as this type, which are delimited by tab. For example:

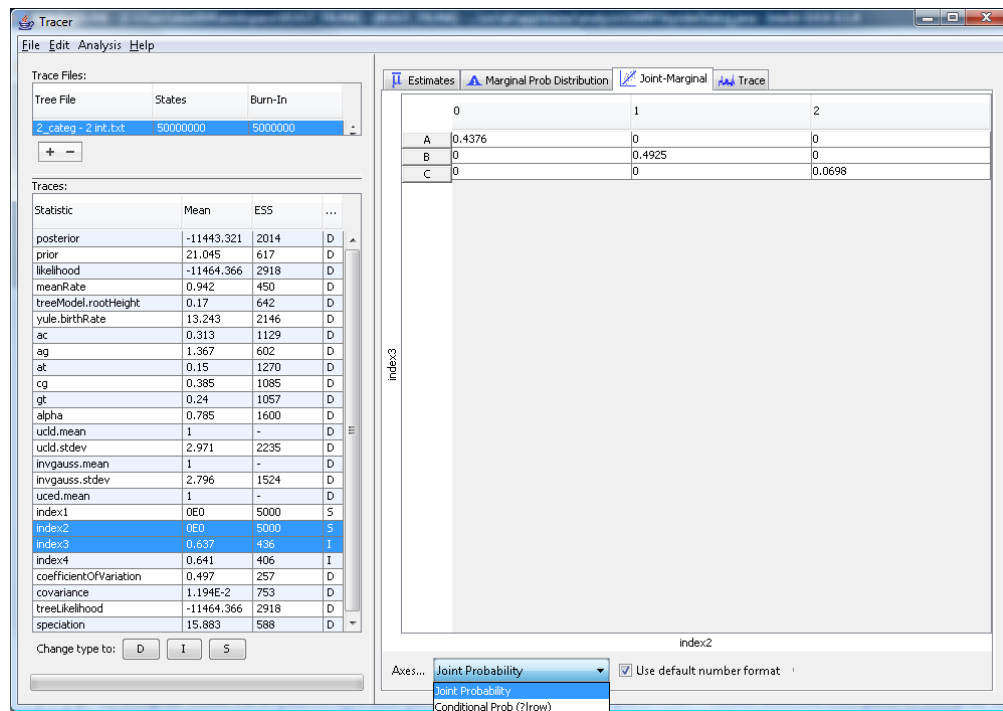
```
# ... ..  
# string      index1      index2  
# integer     index3      index4  
state posterior      ...      index1      index2      index3      index4  
0      -18107.89569      ...      A      B      0      0  
5000   -11757.36321      ...      C      C      2      2  
10000  -11682.44755      ...      C      A      2      2  
15000  -11652.15645      ...      C      A      2      1  
... ..
```

Alternatively, the trace type can be changed by clicking buttons: D, I, S on the bottom of **Statistic** table on the left side.



2. New Plot and Table on Joint-Marginal Panel

Go to **Joint-Marginal** panel and choose two traces in **Statistic** table on the left side, if their trace types are both discrete (*integer* or *string*), then a probability table will replace the plot. It also provides calculations for the marginal, joint, conditional probability and frequency count.

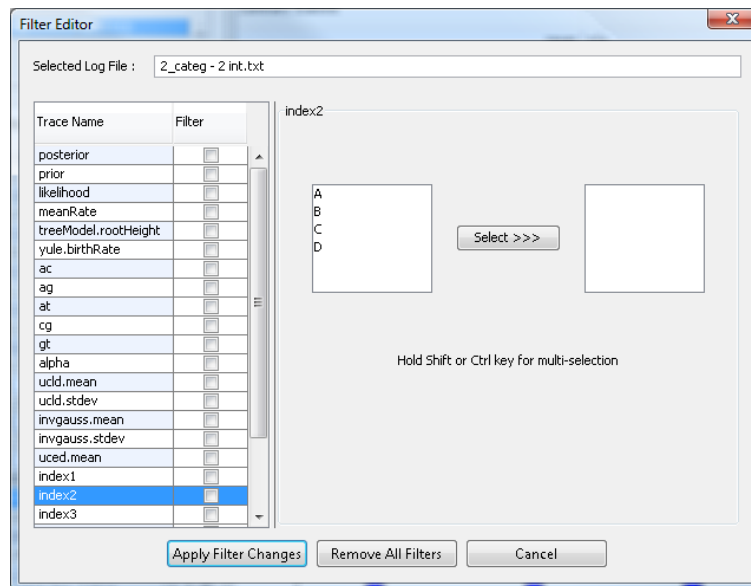


3. Filter To Find Conditional Posterior Distribution

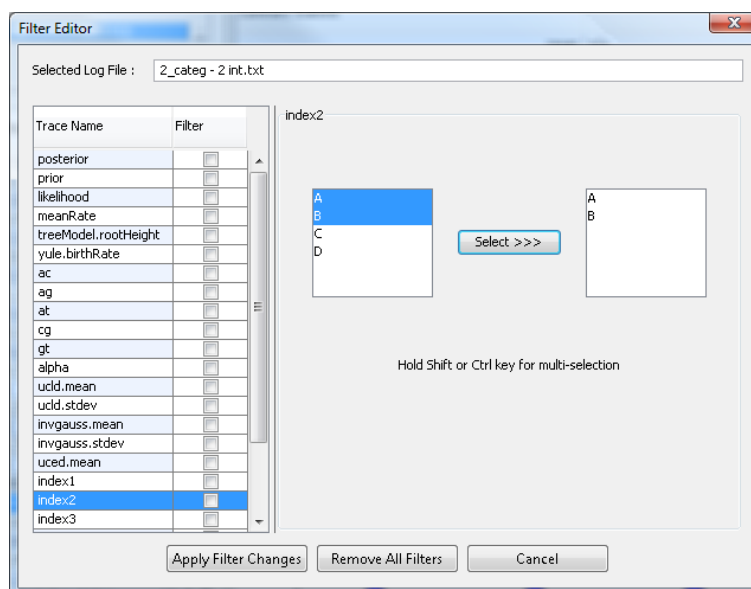
The purpose of filtering in Tracer is to provide:

- 1) Fairly general solution to looking at conditional posterior distributions;
- 2) Specifically, support for BSSVS forms of model averaging, in which some parameters are only in the likelihood when their submodel is "indicated" by some indicator function that is usually a discrete, integer or boolean variable. In that case the posterior of the parameter should not include the states when it was sampled only in the prior because the submodel it belonged to was "turned off". This is relevant for EBSF, Random Local Clocks model, microsatellite model averaging and relaxed clock model averaging methods all from my group in last couple of years. Also relevant for BSSVS in phylogeography as well depending on how the state is logged.

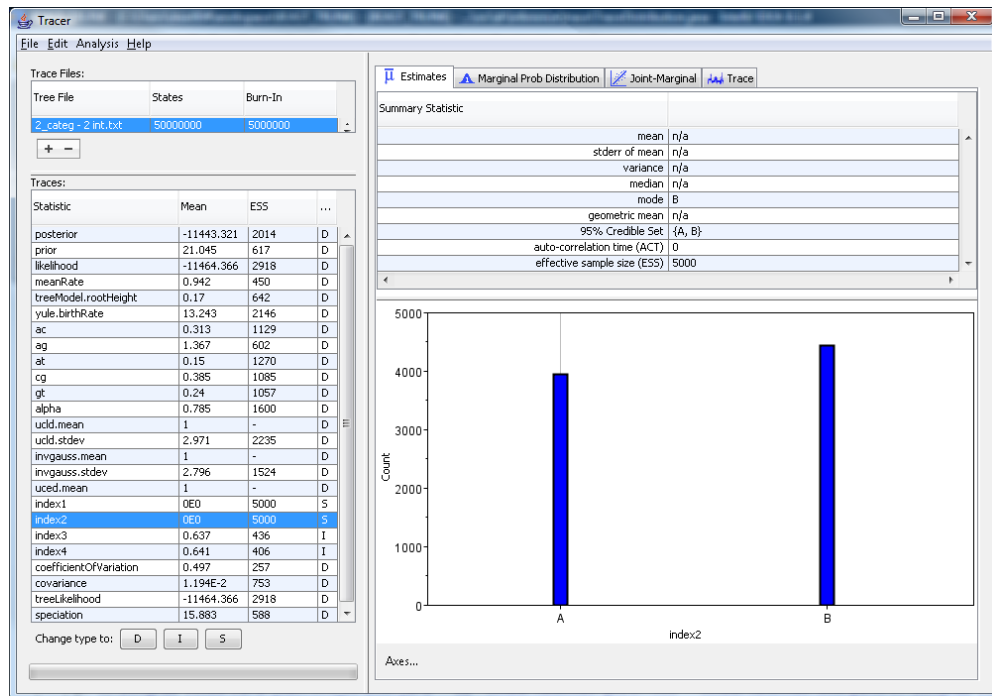
To create or edit filter(s), user needs to select one trace file each time and choose the option **Find Conditional Posterior Distribution ...** in **Analysis** menu. And then **Filter Editor** dialog will be popped out.



Hold *Shift* or *Ctrl* key to select both **A** and **B**, and click **Apply Filter Changes**.



As it can be seen, trace **index2** only shows its value **A** and **B**. In fact, only the state whose trace **index2** contains value **A** or **B** is allowed to participate in the following analysis, and the analysis result is recalculated based on these states whose trace **index2** contains value **C** and **D** are filtered out.



Developer Note of Tracer 1.6.0

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Class Diagram

