

Quantitative
Biology >

Download:

- [PDF](#)
- [Other formats](#)

Current browse context:

q-bio.BM

[< prev](#) | [next >](#)

[new](#) | [recent](#) | [1010](#)

Change to browse by:

[q-bio](#)

References & Citations

- [NASA ADS](#)

Bookmark([what is this?](#))



Biomolecules

**Exploring
the
Energy
Landscapes
of Protein
Folding
Simulations
with
Bayesian**

Computation

Nikolas S.

Burkoff, David L.

Wild

*(Submitted on 22
Oct 2010)*

Nested
sampling
is a
technique
developed
to
explore
probability
distributions
localised
in an
exponentially
small
area of
the
parameter
space.

The
algorithm
provides
both
posterior
samples
and an
estimate
of the
evidence

