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**integrative
analysis
of cancer
gene
expression
studies
using
Bayesian
latent
factor**

modeling

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*(Submitted on 6
Oct 2010)*

We
present
an
applied
study in
cancer
genomics
for
integrating
data and
inferences
from
laboratory
experiments
on
cancer
cell lines
with
observational
data
obtained
from
human
breast
cancer
studies.

The
biological