# Personalized Health and Africa 

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## Outline

Diversity of Data

Massively Missing Data

## Not the Scale it's the Diversity



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Big Data Proving to Be A Real Challenge for Data Scientists
July 2.2014 Writien by. Furhaad Shah Leave a reply
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In a recent survey conducted by Paradigm4, a computational database company, it was revealed that big data was proving to be a chalienge for data scientists - but not because of the amount, or volume, of data being produced, but rather the varlety. and diverse types of data these protessionals have to handie
surver -The increasing variety of data sources is forcing data scientists into shortcuts that leave data and money on the table,

Top Stories


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\section*{Outline}

\section*{Diversity of Data}

Massively Missing Data

\section*{Massive Missing Data}
- If missing at random it can be marginalized.
- As data sets become very large (39 million in EMIS) data becomes extremely sparse.
- Imputation becomes impractical.

\section*{Imputation}
- Expectation Maximization (EM) is gold standard imputation algorithm.
- Exact EM optimizes the log likelihood.
- Approximate EM optimizes a lower bound on log likelihood.
- e.g. variational approximations (VIBES, Infer.net).
- Convergence is guaranteed to a local maxima in log likelihood.

\section*{Expectation Maximization}

Require: An initial guess for missing data

\section*{Expectation Maximization}

Require: An initial guess for missing data repeat

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Update model parameters
(M-step)

\section*{Expectation Maximization}

Require: An initial guess for missing data repeat

Update model parameters
Update guess of missing data
(M-step)
(E-step)

\section*{Expectation Maximization}

Require: An initial guess for missing data repeat

Update model parameters
Update guess of missing data until convergence
(M-step)
(E-step)

\section*{Imputation is Impractical}
- In very sparse data imputation is impractical.
- EMIS: 39 million patients, thousands of tests.
- For most people, most tests are missing.
- M-step becomes confused by poor imputation.

\section*{Direct Marginalization is the Answer}
- Perhaps we need joint distribution of two test outcomes,
\[
p\left(y_{1}, y_{2}\right)
\]
- Obtained through marginalizing over all missing data,
\[
p\left(y_{1}, y_{2}\right)=\int p\left(y_{1}, y_{2}, y_{3}, \ldots, y_{p}\right) \mathrm{d} y_{3}, \ldots \mathrm{~d} y_{p}
\]
- Where \(y_{3}, \ldots, y_{p}\) contains:
1. all tests not applied to this patient
2. all tests not yet invented!!

\section*{Magical Marginalization in Gaussians}

\section*{Multi-variate Gaussians}
- Given 10 dimensional multivariate Gaussian, \(\mathbf{y} \sim \mathcal{N}(\mathbf{0}, \mathrm{C})\).
- Generate a single correlated sample \(\mathbf{y}=\left[y_{1}, y_{2} \ldots y_{10}\right]\).
- How do we find the marginal distribution of \(y_{1}, y_{2}\) ?

\section*{Gaussian Marginalization Property}


Figure : A sample from a 10 dimensional correlated Gaussian distribution.

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(a) A 10 dimensional sample

(b) colormap showing covariance between dimensions.

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(a) A 10 dimensional sample

(b) covariance between \(y_{1}\) and \(y_{2}\).

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\section*{Gaussian Marginalization Property}

(a) A 10 dimensional sample

(b) correlation between \(y_{1}\) and \(y_{2}\).

Figure : A sample from a 10 dimensional correlated Gaussian distribution.

\section*{Avoid Imputation: Marginalize Directly}

- Our approach: Avoid Imputation, Marginalize Directly.
- Explored in context of Collaborative Filtering.
- Similar challenges:
- many users (patients),
- many items (tests),
- sparse data
- Implicitly marginalizes over all future tests too.

Work with Raquel Urtasun (Lawrence and Urtasun, 2009) and ongoing work with Max Zwießele and Nicoló Fusi.

\section*{Marginalization in Bipartite Undirected Graph}


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\section*{Marginalization in Bipartite Undirected Graph}


For massive missing data, how many additional latent variables?

\section*{Methods that Interrelate Covariates}
- Need Class of models that interrelates data, but allows for variable \(p\).
- Common assumption: high dimensional data lies on low dimensional manifold.
- Want to retain the marginalization property of Gaussians but deal with non-Gaussian data!

\section*{Example: Prediction of Malaria Incidence in Uganda}
- Work with John Quinn and Martin Mubaganzi (Makerere University, Uganda)
- See http://air.ug/research.html.

\section*{Malaria Prediction in Uganda}


\section*{Malaria Prediction in Uganda}

Nagongera / Tororo (Multiple output model)


\section*{Malaria Prediction in Uganda}

Mubende



\section*{GP School at Makerere}


\section*{Early Warning Systems}



\section*{Early Warning Systems}

2010: week 46


\section*{Deep Models}


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\section*{Deep Models}


\section*{Deep Gaussian Processes}


Damianou and Lawrence (2013)
- Deep architectures allow abstraction of features (Bengio, 2009; Hinton and Osindero, 2006; Salakhutdinov and Murray, 2008).
- We use variational approach to stack GP models.

\section*{Deep Health}


\section*{Summary}
- Intention is to deploy probabilistic machine learning for assimilating a wide range of data types in personalized health:
- Social networking, text (clinical notes), survival times, medical imaging, phenotype, genotype, mobile phone records, music tastes, Tesco club card
- Requires population scale models with millions of features.
- May be necessary for early detection of dementia or other diseases with high noise to signal.
- Major issues in privacy and interfacing with the patient.
- But: the revolution is coming. We need to steer it.

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