

A Selective Review of Sufficient Dimension Reduction

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Sufficient dimension reduction

Basic regression (supervised learning) setup:

- study the conditional distribution of $Y \in \mathbb{R}^r$ given $X \in \mathbb{R}^p$
- find a $p \times d$ matrix $\gamma = (\gamma_1, \dots, \gamma_d)$, $d \leq p$, such that

$$Y \perp\!\!\!\perp X | \gamma^T X \Leftrightarrow Y|X = Y|\gamma^T X \Leftrightarrow X | (\gamma^T X, Y) = X | \gamma^T X$$

- replace X with $\gamma^T X = (\gamma_1^T X, \dots, \gamma_d^T X)$ without losing any regression information of $Y|X$
- $(\gamma_1^T X, \dots, \gamma_d^T X)$ are called the *sufficient predictors*
- γ is not unique!



Key concepts

Central subspace:

$$Y|X = Y|\gamma^T X \Rightarrow \mathcal{S}_{DRS} = \text{Span}(\gamma) \Rightarrow \mathcal{S}_{Y|X} = \cap \mathcal{S}_{DRS}$$

Examples:

$$Y = f(\gamma_1^T X) + \sigma \varepsilon$$

$$Y = f_1(\gamma_1^T X) + f_2(\gamma_2^T X) \times \varepsilon$$

$$\text{logit} = \gamma_1^T X, \text{ where } \text{logit} = \log \left\{ \frac{P(Y = 1|X)}{1 - P(Y = 1|X)} \right\}$$



Key concepts

Central mean subspace:

$$E(Y|X) = E(Y|\gamma^T X) \Rightarrow \mathcal{S}_{E(Y|X)}$$

For many models, $\mathcal{S}_{Y|X} = \mathcal{S}_{E(Y|X)}$

Examples:

$$Y = f_1(\gamma_1^T X) + \dots + f_d(\gamma_d^T X) + \varepsilon$$

$$Y = f_1(\gamma_1^T X) + f_2(\gamma_2^T X) \times \varepsilon$$



Estimation approaches

Inverse moment based:

- sliced inverse regression (Li, 1991) and many variants: $E(X|Y)$
- sliced average variance estimation (Cook and Weisberg, 1991)
 $\text{Cov}(X|Y)$
- directional regression (Li and Wang, 2007)

Kernel smoothing based:

- minimum average variance estimation (Xia et al. 2002): estimation of the derivative of $E(Y|X)$
- variants: Xia (2007), Wang and Xia (2008)

Others: (not complete)

- ordinary least squares (Li and Duan, 1991)
- reproducing kernel Hilbert space (Fukumizu, Bach and Jordan, 2004, 2009)
- contour based (Li, Zha and Chiramonte, 2005, Li, Artemious and Li, 2010)



Estimation approaches

Comparison:

- Inverse moment based:
 - very easy and fast to compute
 - requires a relatively large sample size
 - requires conditions on the distribution of X (*linearity condition*)
- Kernel smoothing based:
 - works well for small sample size
 - requires no condition on X
 - requires kernel smoothing
 - relatively slow



Sliced inverse regression

Foundation: under the linearity condition,

$$\Sigma_x^{-1} E\{X - E(X)|Y\} \in \mathcal{S}_{Y|X}$$

Spectral decomposition formulation:

$$\Sigma_{x|y} \gamma_j = \lambda_j \Sigma_x \gamma_j, \quad j = 1, \dots, p,$$

where $\Sigma_{x|y} = \text{Cov}[E\{X - E(X)|Y\}]$ and $\Sigma_x = \text{Cov}(X)$.

- obtain the first d eigenvectors $(\gamma_1, \dots, \gamma_d)$ corresponding to the largest d positive eigenvalues $\lambda_1 \geq \dots \geq \lambda_d > 0$, then $\text{Span}(\gamma_1, \dots, \gamma_d) \subseteq \mathcal{S}_{Y|X}$
- assumes Y is categorical or slice Y to estimate $E(X|Y)$
- asymptotic test / permutation test / BIC to determine d



Sliced inverse regression

The linearity condition:

- $E(X|\gamma^T X)$ is a linear function of $\gamma^T X$ for a $\mathcal{S}_{Y|X}$ basis γ
- X is elliptically symmetric; X is normally distributed
- approximately true as $p \rightarrow \infty$ with a fixed d
- involves no Y or $Y|X$, so *nonparametric* or *model-free*

Some important variants:

- canonical correlation analysis: $\max \text{Corr}^2\{h(Y), b^T X\}$ over $h(\cdot)$ and b
- letting $\beta \equiv E[h(Y)\Sigma_X^{-1}E\{X - E(X)|Y\}] = \Sigma_X^{-1}\text{Cov}\{h(Y), X\}$, then $\beta \in \mathcal{S}_{Y|X}$

Beyond SIR:

- sliced average variance estimation: 2nd inverse moment; exhaustive
- directional regression: 1st and 2nd inverse moments; exhaustive



Other dimension reduction approaches

Principal components analysis:

- spectral decomposition of Σ_X
- unsupervised; linear combinations of X

Partial least squares:

- at the population level, PLS = OLS; under the linearity condition, PLS estimates $S_{E(Y|X)}$ (Li, Cook and Tsai, 2007)
- supervised; linear combinations of X

Multidimensional scaling and nonlinear dimension reduction:

- unsupervised; nonlinear combinations of X

Independent components analysis:

- unsupervised; linear combinations of X



Classification

Discriminant analysis:

- directly applicable to categorical Y
- at the population level, SIR \Leftrightarrow LDA \Leftrightarrow Fisher's discriminant analysis; SAVE \Leftrightarrow QDA
- SIR/SAVE produce sufficient predictors instead of classification rule; LDA/QDA produce probability estimate of $Y = g|X$ and a classification rule
- SIR/SAVE require the linearity condition (normality) on X ; LDA/QDA require the normality assumption on $X|Y$

Why useful:

- of course ...



Variable selection

Basic ideas:

- rewrite the SDR estimation in least squares, then apply L_1 type penalty (adaptive group Lasso, SCAD)
- foundation: $Y \perp\!\!\!\perp X_A | X_I \Leftrightarrow$ corresponding rows of $\gamma = 0$
- differ from most model-based variable selection approaches in that no parametric model on $Y|X$ is imposed

Consistency in selection:

- fixed p , $n \rightarrow \infty$ (Ni, Cook and Tsai, 2005, Bondell and Li, 2009)
- diverging $p \rightarrow \infty$, $n \rightarrow \infty$, $p < n$ (Wu and Li, 2010)
- $p = o(a^n)$ for any fixed $a > 1$ (Zhu, Li, Li and Zhu, 2010)

Why useful:

- help interpretation, e.g., identifying regions of brain that are relevant to phenotype



Multivariate and complex responses

Basic ideas:

- dimension reduction is still on X instead of Y
- key observations:

$$\begin{aligned}\mathcal{S}_{E(Y|X)} &= \mathcal{S}_{E(Y_1|X)} \oplus \dots \oplus \mathcal{S}_{E(Y_r|X)} \\ \mathcal{S}_{Y|X} &\supseteq \mathcal{S}_{Y_1|X} \oplus \dots \oplus \mathcal{S}_{Y_r|X}\end{aligned}$$

- multivariate reduced rank model (Cook and Setodji, 2003): one response at a time
- projective sampling (Li, Wen and Zhu, 2008): sample a on a unit ball $O(n)$ times, and regress $a^T Y$ on X

Why useful:

- e.g., voxel-wise imaging genetics (ignore the spatial information)
- what if Y has structures, such as spatial information in MRI, or positive definiteness in DTI? — open question



Predictors with structures

Basic ideas:

- predictors have group structure, and dimension reduction (linear combinations) should be within groups (Li, 2009, Li, Li and Zhu, 2010)
- direct sum structure: $\gamma_1 \oplus \dots \oplus \gamma_g$
- partial dimension reduction, e.g., genetic / imaging information plus clinical / demographical information

Why useful:

- fusion of different data modalities
- what if X has, e.g., network structures? — open question



Matrix or array valued predictors

Basic ideas:

- predictor is a matrix or an array instead of a vector, and dimension reduction wishes to preserve interpretation
- dimension folding (Li, Kim and Altman, 2010):

$$\gamma^T X \eta = (\eta \otimes \gamma)^T \text{vec}(X)$$

- tensor PCA / tensor ICA

Why useful:

- MRI or fMRI



Functional predictors

Basic ideas:

- predictor is a functional curve (dense / sparse)
- sliced inverse regression in functional space (Ferré and Yao, 2003, 2005, Hsing and Ren, 2009, Li and Hsing, 2010):
- functional PCA

Why useful:

- common nowadays in genetics and imaging data



Nonlinear sufficient dimension reduction

Basic ideas:

- map X to $\phi(X)$, then do linear SDR in the $\phi(X)$ space (Wu, Liang and Mukherjee, 2008, Zhu and Li, 2010)
- the optimal separating hyperplane (Li, Artemious and Li, 2010)

Why useful:

- categorical predictors
- $n < p$
- predictors with complex structures
- how to do variable selection in this setup? — open question



Conclusion and discussion

- application of existing SDR solutions to imaging data
- motivate new methodology development for dimension reduction



Thank You!

