

Mathematical Statistics Team

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Statistics and Machine Learning: Methodology and Applications

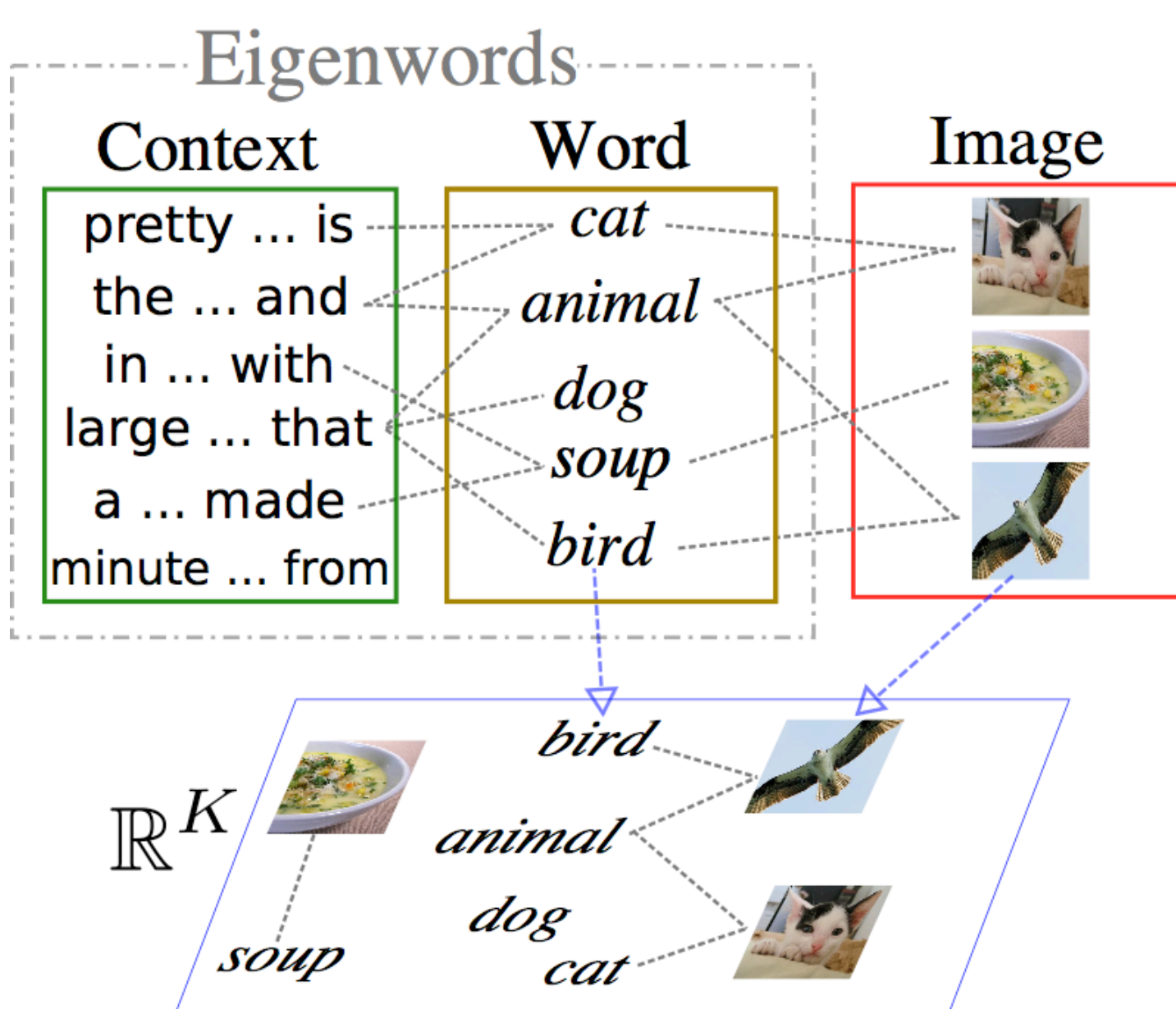
- Inductive inference, Resampling methods, Information geometry
- Generalization error under Missing, Covariate-shift, etc.
- Multi-modal data representation, Graph Embedding, and Multivariate Analysis
- Network growth mechanism (Preferential attachment and fitness)
- Phylogenetics, Gene expression (hierarchical clustering)
- Image, word embedding (search and reasoning)

Multimodal Eigenwords (Fukui, Oshikiri and Shimodaira, Textgraphs 2017)

- A multimodal word embedding that jointly embeds words and corresponding visual information
- We employed **Cross-Domain Matching Correlation Analysis** (CDMCA; Shimodaira 2016) for extending a CCA-based word embedding (Dhillon et al. 2015) to deal with complex associations

Our proposed method:

- Feature learning via graph embedding
- Feature vectors reflect both semantic and visual similarities
- The method enables **multimodal vector arithmetic** between images and words



Multimodal vector arithmetic:



Probabilistic Multi-view Graph Embedding (PMvGE)

(Okuno, Hada and Shimodaira, arXiv:1802.04630)

Multi-view feature learning with many-to-many associations via neural networks for predicting new associations

$$w_{ij}^{(de)} \sim \text{Po}(\alpha^{(de)} \exp(\langle f_{qb}^{(d)}(x_i^{(d)}), f_{qb}^{(e)}(x_j^{(e)}) \rangle))$$

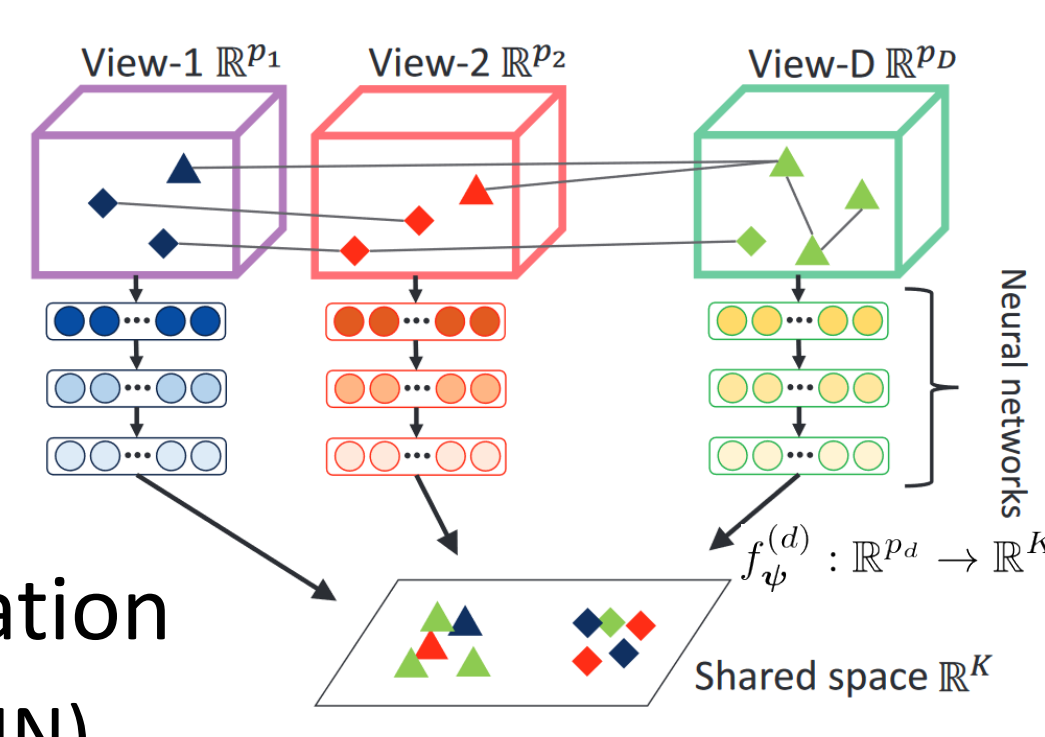
Theorem (universal approximation):

Inner product + Neural networks

≈ Arbitrary similarity + Arbitrary transformation
(Mercer's theorem + universal approximation of NN)

Advantages:

- (1) PMvGE with neural networks is proved to be **highly expressive**.
- (2) PMvGE **approximately non-linearly generalizes CDMCA** (Shimodaira 2016, Neural Networks) which already generalizes various existing methods such as CCA, LPP, and Spectral graph embedding.
- (3) Likelihood-based estimation of neural networks is **efficiently computed** by mini-batch Stochastic Gradient Descent.

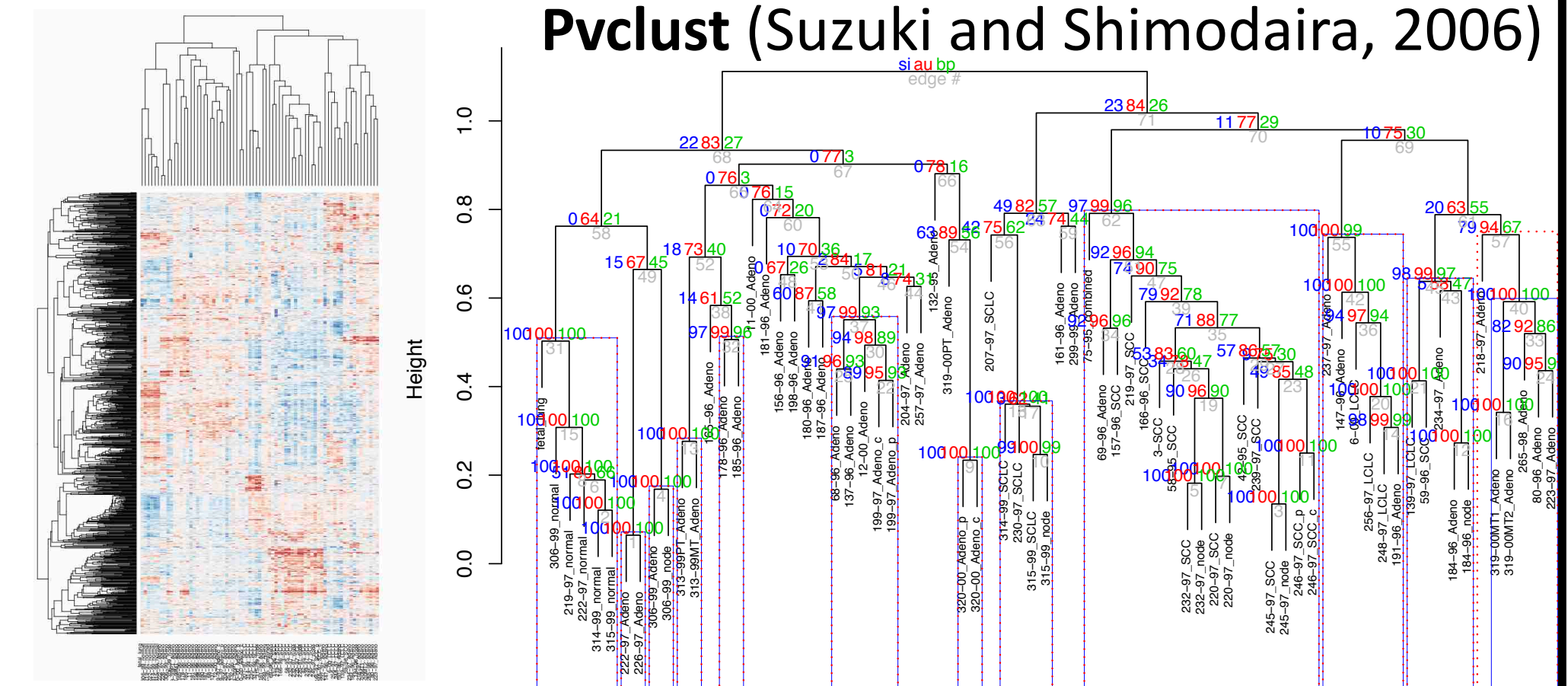


Selective inference (Terada and Shimodaira, arXiv:1711.00949)

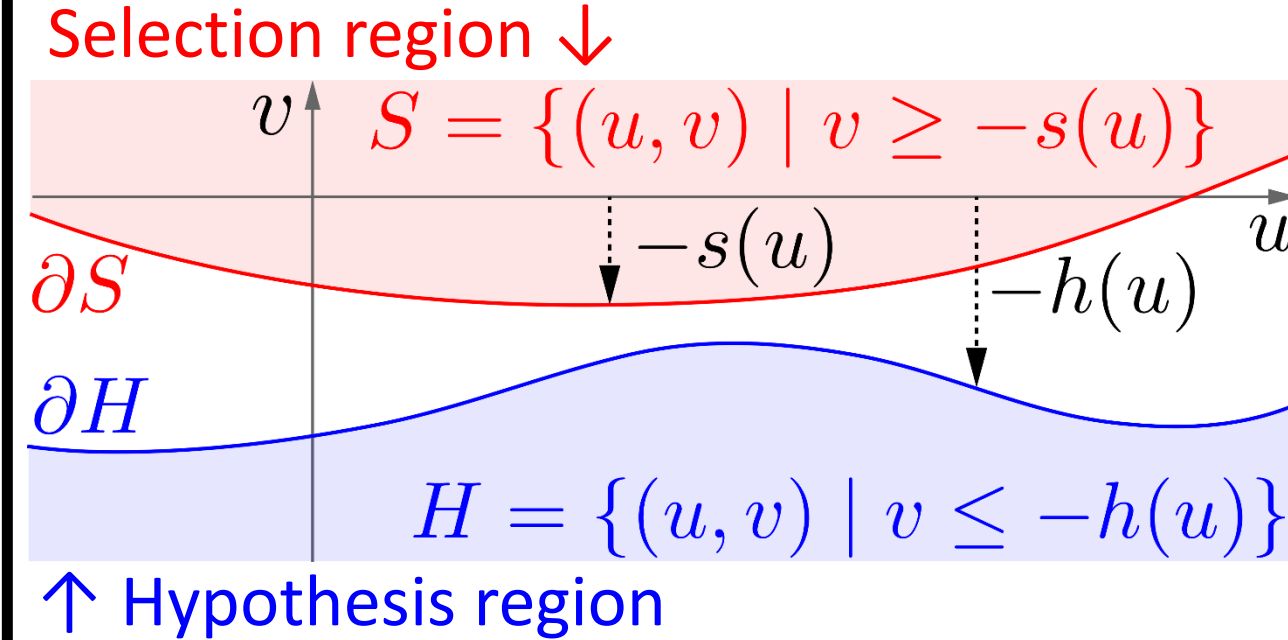
- **Motivation** : Assessing the confidence of each **obtained cluster**
- Consider approx. unbiased p -values as frequentist confidence measures
Null hypothesis = obtained cluster is NOT true
- Example : lung data (73 tumors, 916 genes; Garber et al., 2001)

The hypothesis is tested **only for** the clusters appeared in the obtained tree

Need selective inference for pvclust



- Two asymptotic theories



Theorem (large sample theory):

Boundary surfaces of H and S are smooth and "nearly parallel",
⇒ The proposed p -value is **second order accurate with error $O(n^{-1})$**

Theorem (nearly flat surfaces):

Boundary surfaces are "nearly flat", approaching flat but allowing non-smooth such as cones and polyhedra
⇒ the proposed p -value is justified **as unbiased with error $O(\lambda^2)$**

Key point: **Multiscale Bootstrap** (Shimodaira, 2002; 2004)

- ✓ Low computational cost : $O(B)$
- ✓ Double bootstrap method has same accuracy but high comp. cost $O(B^2)$

- (1) Large sample theory with "nearly parallel surfaces"

$$h(u) = h_0 + h_{1j}u_j + h_{2jk}u_ju_k + h_{3jkl}u_ju_ku_l + \dots, \quad n \rightarrow \infty$$

$h_0 = O(1), h_{1j} = O(n^{-1/2}), h_{2jk} = O(n^{-1/2}), h_{3jkl} = O(n^{-1}), \dots$ Smooth surface
(u, v) are $O(\sqrt{n})$, but $h_0 = O(n^{1/2}), h_i = O(1)$ are multiplied by $O(n^{-1/2})$

- (2) Asymptotic theory of "nearly flat surfaces"

$$\sup_{u \in \mathbb{R}^m} |h(u)| = O(\lambda), \quad \int_{\mathbb{R}^m} |h(u)| du < \infty, \quad \int_{\mathbb{R}^m} |\mathcal{F}h(u)| d\omega < \infty$$

(Shimodaira 2008) non-smooth surface $\lambda \rightarrow 0$

Algorithm : Computing approx. unbiased selective p -values

- 1: Specify several $n' \in \mathbb{N}$ values, and set $\sigma^2 = n/n'$. Set the number of bootstrap replicates B , say, 1000.
- 2: For each n' , perform bootstrap resampling to generate Y^* for B times and compute $\alpha_{\sigma^2}(H|y) = C_H/B$ and $\alpha_{\sigma^2}(S|y) = C_S/B$ by counting the frequencies $C_H = \#\{Y^* \in H\}$ and $C_S = \#\{Y^* \in S\}$. (We actually work on X^* , instead of Y^* .) Compute $\psi_{\sigma^2}(H|y) = \sigma\Phi^{-1}(\alpha_{\sigma^2}(H|y))$ and $\psi_{\sigma^2}(S|y) = \sigma\Phi^{-1}(\alpha_{\sigma^2}(S|y))$.
- 3: Estimate parameters $\beta_H(y)$ and $\beta_S(y)$ by fitting models **Model fitting to psi**
 $\psi_{\sigma^2}(H|y) = \varphi_H(\sigma^2|\beta_H)$ and $\psi_{\sigma^2}(S|y) = \varphi_S(\sigma^2|\beta_S)$,
respectively. The parameter estimates are denoted as $\hat{\beta}_H(y)$ and $\hat{\beta}_S(y)$. If we have several candidate models, apply above to each and choose the best model based on AIC value.
- 4: Approximately unbiased p -values of selective inference (p_{SI}) and non-selective inference (p_{AU}) are computed by one of (A) and (B) below.
(A) Extrapolate $\psi_{\sigma^2}(H|y)$ and $\psi_{\sigma^2}(S|y)$ to $\sigma^2 = -1$ and 0 , respectively, by **Extrapolation** $z_H = \varphi_H(-1|\hat{\beta}_H(y))$ and $z_S = \varphi_S(0|\hat{\beta}_S(y))$, and then compute p -values by
Selective p-value $p_{SI}(H|S, y) = \frac{\Phi(z_H)}{\Phi(z_H + z_S)}$ and $p_{AU}(H|y) = \Phi(z_H)$ **Non-selective p-value**
(B) Specify $k \in \mathbb{N}$, $\sigma_0^2, \sigma_1^2 > 0$ (e.g., $k = 3$ and $\sigma_1^2 = \sigma_0^2 = 1$). Extrapolate $\psi_{\sigma^2}(H|y)$ and $\psi_{\sigma^2}(S|y)$ to $\sigma^2 = -1$ and 0 , respectively, by
 $z_{H,k} = \varphi_{H,k}(-1|\hat{\beta}_H(y), \sigma_0^2)$ and $z_{S,k} = \varphi_{S,k}(0|\hat{\beta}_S(y), \sigma_0^2)$,
where the Taylor polynomial approximation of φ_H at $\tau^2 > 0$ with k terms is:
$$\varphi_{H,k}(\sigma^2|\hat{\beta}_H(y), \tau^2) = \sum_{j=0}^{k-1} \frac{(\sigma^2 - \tau^2)^j}{j!} \frac{\partial^j \varphi_H(\sigma^2|\hat{\beta}_H(y))}{\partial(\sigma^2)^j} \Big|_{\sigma^2=\tau^2}$$

and that of φ_S is defined similarly. Then compute p -values by
 $p_{SI,k}(H|S, y) = \frac{\Phi(z_{H,k})}{\Phi(z_{H,k} + z_{S,k})}$ and $p_{AU,k}(H|y) = \Phi(z_{H,k})$.

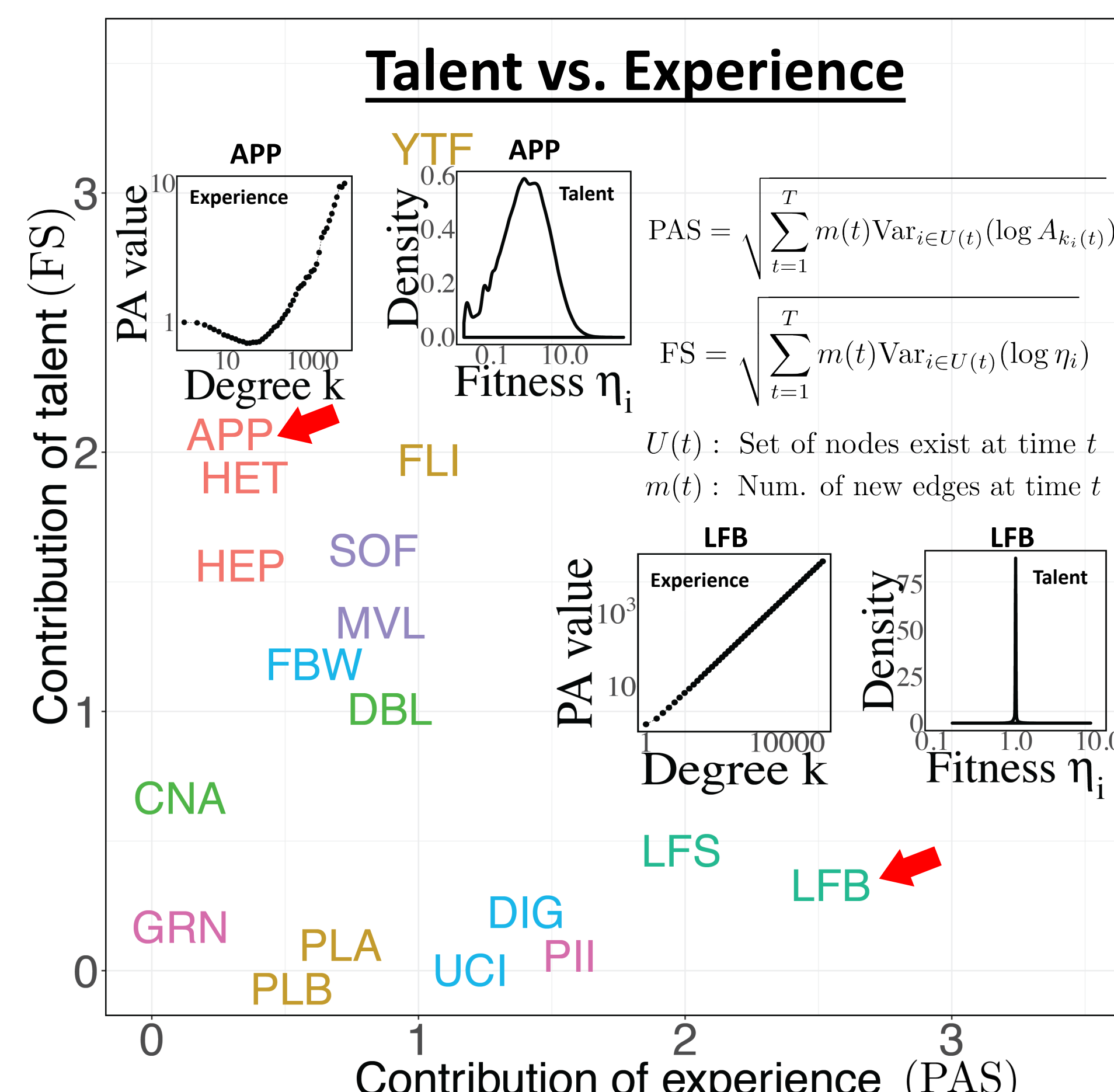
PAFit: an R Package for Estimating Preferential Attachment and Node Fitness in Temporal Complex Networks

(Pham, Sheridan and Shimodaira, arXiv:1704.06017)

- What drives the growth of real-world networks across diverse fields?
- Using two **interpretable and universal** mechanisms: (1) **Talent**: the intrinsic ability of a node to attract connections (fitness η_i) and (2) **Experience**: the preferential attachment (PA) function A_k governing the extent to which having more connections makes a node more/less attractive for forming new connections in the future.

Probability that node v_i gets a new edge at time $t = A_{k_i}(t)\eta_i$

Key finding: Although both talent and experience contribute to the growth process, the ratios of their contributions vary greatly.



1. Citation networks
HEP: arXiv hep-ph papers
HET: arXiv hep-th papers
APP: APS journal papers
2. Social networks
YTF: YouTube followership
PLB: Prosper loan before Lehman
PLA: Prosper loan after Lehman
FLI: Flickr friendship
3. Co-author networks
DBL: dblp authors
CNA: complex network authors
4. Interaction networks
LFS: last.fm song
LFB: last.fm band
5. Communication networks
UCI: UC Irvine forum message
FBW: Facebook wall-post
DIG: Digg message
6. Rating networks
SOF: Stackoverflow favourite
MVL: MovieLens rating
7. Biological networks
GRN: Cancer gene
PII: Human PPI