Necessary and Sufficient Conditions for High-Dimensional Salient Feature Subset Recovery

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ISIT (Jun 14, 2010)

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Salient Feature Subset Recovery

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- Number of variables $d \approx 10^6$.
- But only $k \approx 30$ are salient for assessing susceptibility to asthma.
- Identification of these salient features is important.

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- What is the meaning of saliency?
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- What are the fundamental limits for recovery of the salient set?
- Are there any efficient algorithms for special classes of features?

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 We define the salient set by appealing to the error exponents in hypothesis testing.

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- Sufficiency: We show that if for all n sufficiently large,

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• Necessity: Under certain conditions, for $\lambda \in (0, 1)$, if

$$n < \lambda \cdot C_3 \cdot \log\left(\frac{d}{k}\right)$$

then the error probability $\geq 1 - \lambda$.



System Model

- Let the alphabet for each variable be \mathcal{X} , a finite set.
- High-dimensional setting where d and k grow with n.
- Two sequences of unknown *d*-dimensional distributions

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- IID samples $(\mathbf{x}^n, \mathbf{y}^n) := (\{\mathbf{x}^{(l)}\}_{l=1}^n, \{\mathbf{y}^{(l)}\}_{l=1}^n)$ drawn from $P^{(d)} \times Q^{(d)}$.
- Each pair of samples $\mathbf{x}^{(l)}, \mathbf{y}^{(l)} \in \mathcal{X}^d$.



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Chernoff-Stein Lemma: If $Pr(\hat{H}_1|H_0) < \alpha$, then

$$\Pr(\hat{H}_0|H_1) \doteq \exp(-nD(P^{(d)}||Q^{(d)}))$$

Chernoff Information:

$$\Pr(\text{err}) = \pi_0 \Pr(\hat{H}_1 | H_0) + \pi_1 \Pr(\hat{H}_0 | H_1) \doteq \exp(-nC(P^{(d)}, Q^{(d)}))$$

where

$$C(P,Q) := -\min_{t \in [0,1]} \log \sum_{\mathbf{z}} P(\mathbf{z})^t Q(\mathbf{z})^{1-t}$$



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• Definition: A set $S_d \subset \{1, \dots, d\}$ is KL-divergence-salient if

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• Definition: A set $S_d \subset \{1, \dots, d\}$ is Chernoff Information-salient if

$$C(P^{(d)}, Q^{(d)}) = C(P^{(d)}_{S_d}, Q^{(d)}_{S_d}).$$

Characterization of Saliency

Lemma

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$$P^{(d)} = P_{S_d}^{(d)} \cdot \textcolor{red}{\mathbf{W}_{S_d^c \mid S_d}} \qquad Q^{(d)} = Q_{S_d}^{(d)} \cdot \textcolor{red}{\mathbf{W}_{S_d^c \mid S_d}}.$$

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What are the scaling laws on (n, d, k) so that the error probability can be made arbitrarily small?

Definition of Achievability

 A decoder is a set-valued function that maps samples to subsets of size k, i.e.,

$$\psi_n: (\mathcal{X}^d)^n \times (\mathcal{X}^d)^n \to {1 \choose k}.$$

 The decoder is given the true value of k, the number of salient features.

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Definition: The tuple of model parameters (n, d, k) is achievable for $\{P^{(d)}, O^{(d)}\}_{d\in\mathbb{N}}$ if there exists a sequence of decoders $\{\psi_n\}$ such that

$$q_n(\psi_n) := \Pr(\psi_n(\mathbf{x}^n, \mathbf{y}^n) \neq S_d) < \epsilon, \quad \forall n > N_{\epsilon}.$$



Three Assumptions on Distributions $P^{(d)}$, $Q^{(d)}$

• Saliency: For every $P^{(d)}$, $Q^{(d)}$ there exists a salient set S_d of known size k, i.e.,

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• η -Distinguishability: There exists a constant $\eta > 0$ such that

$$D(P_{S_d}^{(d)} || Q_{S_d}^{(d)}) - D(P_{T_d}^{(d)} || Q_{T_d}^{(d)}) \ge \eta$$

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• L-Boundedness: There exists a constant $L \in (0, \infty)$ such that

$$\log \left[\frac{P_{S_d}^{(d)}(\mathbf{z}_{S_d})}{Q_{S_d}^{(d)}(\mathbf{z}_{S_d})} \right] \in [-L, L]$$

for all states $\mathbf{z}_{S_d} \in \mathcal{X}^k$.



Achievability Result

Theorem

If there exists an $\delta > 0$ such that for some B > 0

$$n > \max \left\{ \frac{k}{B} \log \left(\frac{d-k}{k} \right), \exp \left(\frac{2k \log |\mathcal{X}|}{1-\delta} \right) \right\},$$

then there exists a sequence of decoders ψ_n^* that satisfies

$$q_n(\psi_n^*) = O(\exp(-nE)),$$

for some exponent E > 0.

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Let $k = k_0$ be a constant and $R \in (0, B/k_0)$. Then if

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$$n > \frac{\log d}{R} \qquad \Rightarrow \qquad q_n(\psi_n^*) = O\left(\exp(-nE)\right).$$

Converse Result

We assume that the salient set S_d is chosen uniformly at random over all subsets of size k.

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Theorem

If for some $\lambda \in (0,1)$,

$$n < \frac{\lambda \cdot k \cdot \log\left(\frac{d}{k}\right)}{H(P^{(d)}) + H(Q^{(d)})}$$

then

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for all decoders ψ_n .



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$$n < \underbrace{\frac{\lambda}{H(P^{(d)}) + H(Q^{(d)})}}_{O(d)} \cdot k \cdot \log\left(\frac{d}{k}\right)$$

 However, converse is interesting if distributions have additional structure on their entropies.

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- However, converse is interesting if distributions have additional structure on their entropies.
- Assume most of the features are processed or redundant.
- Example: there could be two features, "BMI" and "isObese". One is a processed version of the other.

Converse Result

Corollary

Assume that there exists a $M < \infty$ such that the conditional entropies satisfy

$$\max\left\{H\big(P_{S_d^c|S_d}^{(d)}\big),H\big(Q_{S_d^c|S_d}^{(d)}\big)\right\}\leq M\cdot k.$$

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If the number of samples satisfies

$$n < \frac{\lambda}{2(M + \log |\mathcal{X}|)} \cdot \log \left(\frac{d}{k}\right)$$

then

$$q_n(\psi_n) \geq 1 - \lambda$$
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- Conversely, there is another rate $R_{\rm conv}$ so that if $R > R_{\rm conv}$, then recovery is not possible.



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- Provided necessary and sufficient conditions for salient set recovery.
- Number of samples n can be much smaller than d, total number of variables.
- In the paper, we provide a computationally efficient and consistent algorithm to search for S_d when $P^{(d)}$ and $Q^{(d)}$ are Markov on trees.