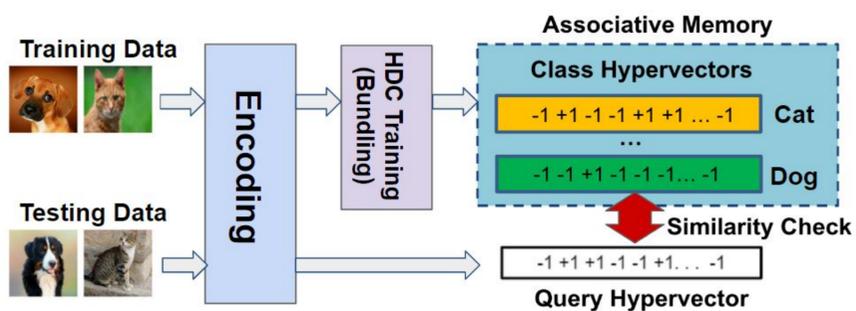


Unleashing Hyperdimensional Computing with Nyström Method based Encoding

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Background and Motivations



Success of HDC based machine learning approaches is heavily dependent on the encoding function that maps raw data to high-dimensional space

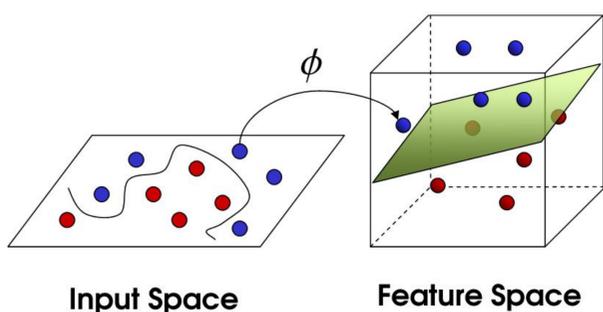
Hyperdimensional Computing (HDC)

- Lightweight and efficient computing paradigm capable of ML tasks such as classification, regression.
- Amenable to highly-parallel circuitry, and require low-precision processing.

Limitations of Existing HDC encoding methods

- Tend to only capture basic notion of similarities, which may be suboptimal when dealing with data with complex structure (e.g. strings, graphs)
- Existing kernel method literature has wide variety of kernel functions (similarity functions)

Kernel methods and HDC



- Inner-products in HD space should be reflective of some salient notion of similarity on ambient space.
- Idea: Construct HD encoding functions using suitable kernel functions.

Random Fourier Features (RFF)

- Commonly referred as “non-linear” encoding in HDC literature.
- Capable of modeling shift-invariant kernels in HDC (e.g. the Gaussian kernel, polynomial kernel)

Why Nyström method ?

$$\hat{G}_{ij} \approx \phi_{nys}(x_i)^T \phi_{nys}(x_j) = \left(\Lambda^{-\frac{1}{2}} Q^T C^{(i)} \right)^T \left(\Lambda^{-\frac{1}{2}} Q^T C^{(j)} \right)$$

- RFF only works with shift-invariant kernels on a Euclidean space, which many useful kernels do not satisfy (i.e. kernels on graphs and strings)

- Kernel method solves “nonlinear” task using linear model with the help of kernel functions.
- Just like in HDC, kernel methods work by embedding data into a high-dimensional space wherein similarities are measured using inner-products

Nyström Method based HDC Encoding & Main Results

- “Top down” approach where embedding directly approximates some data-appropriate notion of similarity.

Theorem 1 Define $\Theta_i = \Lambda^{-\frac{1}{2}} Q^T C^{(i)}$ and $\Theta_i \in \mathbb{R}^n$. Based on above HDC encoding algorithm, the encoding function $\phi : \mathcal{X} \rightarrow \mathcal{H}$ can be write as following:

$$\phi(x_i) = \sqrt{\frac{\pi}{2d}} \text{sign}(P_{rp} \Theta_i) \quad (3)$$

We pose no restriction on Kernel K , the following holds up to a first order approximation:

$$\mathbb{E}[\langle \phi(x_i), \phi(x_j) \rangle] \approx \frac{\hat{G}_{ij}}{\sqrt{\hat{G}_{ii} \hat{G}_{jj}}} \quad \text{i.e. normalized kernel} \quad (4)$$

Where \hat{G}_{ij} is estimated kernel value between x_i and x_j produced by Nyström method and the expectation is taken with respect to randomness and orthogonality in P_{rp} .

- our proposed encoding method preserves the kernel in HD space of some user defined kernel..

Experimental setup:

Task	dataset	# of training samples	# of testing samples	# of classes
String	Protein sequences [39]	721	181	6
	SMS Spam collection [1]	4459	1115	2
Graph	ENZYME ¹ [3]	480	120	6
	NCII ² [45]	3288	822	2
Image	MNIST [23]	60000	10000	10
	FashionMNIST [47]	60000	10000	10

Key Results:

Task	Dataset	Ours	N-gram [13]	GraphHD [31]	LinearRP [34, 12]	NonlinearRP [44]
String	Protein sequence	99%	81%	-	-	-
	SMS Spam collection	96%	93%	-	-	-
Graph	ENZYME	63%	-	26%	-	-
	NCII	72%	-	62%	-	-
Image	MNIST	96%	-	-	96%	97%
	FashionMNIST	86%	-	-	85%	86%

Summary & Acknowledgement

- In summary, we propose a new way to generate embeddings for HDC which can turn any user-defined positive-semidefinite similarity function into an equivalent embedding.
- This work allows future HDC works to exploit the power of kernel methods while still conforming to the general formalism and benefits of HDC.

Discussion & Future Work

- We recognize that the improvements in our proposed HDC encoding methods also come with additional computation costs in the form of kernel evaluation.
- How to achieve the best efficiency-accuracy trade-offs for HDC applications are non-trivial problems that need further investigations.

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