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# Laelaps: An Energy-Efficient Seizure Detection Algorithm from Long-term Human iEEG Recordings without False Alarms

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**Abstract**—We propose Laelaps, an energy-efficient and fast learning algorithm with no false alarms for epileptic seizure detection from long-term intracranial electroencephalography (iEEG) signals. Laelaps uses end-to-end binary operations by exploiting symbolic dynamics and brain-inspired hyperdimensional computing. Laelaps’s results surpass those yielded by state-of-the-art (SoA) methods [1], [2], [3], including deep learning, on a new very large dataset containing 116 seizures of 18 drug-resistant epilepsy patients in 2656 hours of recordings—each patient implanted with 24 to 128 iEEG electrodes. Laelaps trains 18 patient-specific models by using only 24 seizures: 12 models are trained with one seizure per patient, the others with two seizures. The trained models detect 79 out of 92 unseen seizures *without any false alarms across all the patients* as a big step forward in practical seizure detection. Importantly, a simple implementation of Laelaps on the Nvidia Tegra X2 embedded device achieves  $1.7\times\text{--}3.9\times$  faster execution and  $1.4\times\text{--}2.9\times$  lower energy consumption compared to the best result from the SoA methods. Our source code and anonymized iEEG dataset are freely available at <http://ieeg-swez.ethz.ch>.

**Index Terms**—Hyperdimensional computing, symbolic analysis.

## I. INTRODUCTION

Epilepsy is a highly prevalent chronic neurological disorder affecting 1–2% of the world’s population [4]. One third of patients with epilepsy continue to suffer from seizures despite pharmacological therapy. For improved monitoring of these patients with drug-resistant epilepsy—e.g., during presurgical diagnostics—machine learning methods are developed for automatic seizure detection. These SoA methods are based on extracting useful features followed by traditional supervised machine learning methods such as K-nearest neighbor [5], random forest [6], [4], and support vector machines (SVMs) [1], [7]; more recent methods exploit automatic feature extraction algorithms based on deep learning such as convolutional neural networks (CNNs) [2] and long short-term memories (LSTMs) [3].

Most seizure detection methods assume that the patients with epilepsy experience two distinct states of brain activity—i.e., interictal (between seizures) and ictal (during seizures)—that can be detected by EEG. Recent methods propose to use unobtrusive non-EEG signals such as heart rate and accelerometry that despite the distance from the seizure generating brain regions reflect extracerebral changes associated with specific types of seizures [8]. However, these methods suffer from high false alarm rates as high as  $1.2\text{ h}^{-1}$  [9], compared to lower (e.g.,  $0.47\text{ h}^{-1}$  [5] to  $0.27\text{ h}^{-1}$  [7]) yielded by EEG-based methods; the false alarms may increase anxiety in epilepsy patients and are thus likely to further decrease their already impaired quality of life [10]. Enabling even more proximate monitoring of the seizure generating regions, intracranial EEG (iEEG, or ECoG) currently

provides the best spatial resolution and highest signal-to-noise ratio to record electrical brain activity [4]. Using such invasive recordings can reduce the false alarm rate to  $0.13\text{ h}^{-1}$  [11].

However, one major challenge with iEEG signals is to realize seizure detection algorithms with low complexity suitable for execution on implantable devices with limited battery size for long-term or even lifelong operation. Another challenge—common to all methods—is to reliably detect seizures from a small number of examples due to the uneven distribution with much longer interictal than ictal time periods, and the patient-specific seizure dynamics [7], [12]. To address these challenges, one promising option is to develop computationally efficient and fast learning algorithms by combining methods from symbolic dynamics [5], [13], [14], [15], [1] and brain-inspired hyperdimensional computing [16], [17]. In this paper, we describe the following contributions:

(1) We propose Laelaps<sup>1</sup>, an energy-efficient algorithm for both learning and inference from long-term iEEG recording of patients having large heterogeneity implanted with 24 to 128 electrodes. Laelaps operates with end-to-end binary operations to avoid otherwise expensive fixed- or floating-point arithmetic. Laelaps exploits local binary patterns (Sec. II-A) to map a sequence of iEEG samples into 6-bit codes as abstract symbols. These symbols are further combined over time and across all the electrodes to generate a compact representation for encoding the brain state of interest. Such a *holographic* representation is effectively constructed by using brain-inspired hyperdimensional computing (Sec. II-B) that enables fast learning with binary operations on  $d$ -bit vectors where  $d \in [1000, 10000]$  (see Sec. III).

(2) For each of the 18 patients, Laelaps trains a patient-specific model with one or maximally two seizure examples from the assigned patient. Remarkably, these models detect 79 out of 92 unseen seizures *with no false alarms over 1357 hours of recording*. Using the same setup, the SoA methods detect fewer seizures with higher false alarm rates: 69 detected seizures with  $0.31\text{ h}^{-1}$  using the SVM, 51 detected seizures with  $0.36\text{ h}^{-1}$  using the CNN, and 69 detected seizures with  $0.54\text{ h}^{-1}$  using the LSTM. The SoA methods also reach lower sensitivity averaged across patients, except for the LSTM (88.4% vs. 85.5%). See Sec. IV.

(3) GPU implementation of Laelaps on the Nvidia Tegra X2 (TX2) achieves  $1.7\times\text{--}3.9\times$  faster execution and  $1.4\times\text{--}2.9\times$  lower energy consumption, depending on the number of electrodes,

<sup>1</sup>Laelaps was a dog in Greek mythology who never failed to catch what she was hunting.

compared to the SVM as the most energy-efficient implementation among the SoA methods (Section V).

## II. BACKGROUND

We first provide a background in the symbolization that transforms the iEEG signals into abstract symbols. Then we introduce brain-inspired hyperdimensional computing that combines these abstract symbols to compute useful representations for the brain state of interest.

### A. Symbolization using Local Binary Patterns (LBP)

A class of data analysis methods is referred to as symbolization, which describes the process of transforming experimental measurements into a series of discrete symbols. Symbolization is particularly interesting for iEEG analysis [14], [15], because it faithfully preserves dominant dynamical signal characteristics while significantly increasing the efficiency of detecting and quantifying information contained in real-world time series [13]. Symbolization may be efficiently achieved by mapping a sequence of iEEG samples into a bit string, i.e. a one-dimensional local binary pattern (LBP) [5], [1]. Here a LBP code reflects the relational aspects between consecutive values of the iEEG signals, i.e., whether their amplitudes increase or decrease. We find that LBP codes are more efficient than other symbolization methods, e.g., directed horizontal graphs [15] that assign an integer input and output degree to each time point.

A LBP code can be computed in two steps: (1) The iEEG samples are converted into a bit string depending on the sign of the temporal difference of two adjacent samples. If the difference is positive, we assign a 1 to the sampling point, otherwise a 0. (2) A LBP code of length  $\ell$  is associated with every sampling point by concatenating its bit with the successive  $\ell - 1$  bits. We observe that the distribution of LBP codes is significantly different between the ictal and the interictal states [12]. More specifically, during the interictal state the histogram of LBP codes is flattened out (i.e., the counts are almost evenly distributed over all the possible codes). In contrast, the ictal state has a predominant portion of a single LBP code and many LBP codes never occur due to the relatively slower and more asymmetric iEEG oscillations typically emerging during seizures.

### B. Hyperdimensional (HD) Computing

Inspired by the size of the brain's circuits, we can model neural activity patterns with points of a hyperdimensional space. Hyperdimensional (HD) computing [16] explores this idea by computing with *hypervectors*—very high dimensional ( $d$ ) random vectors with independent and identically distributed (i.i.d.) components—in the following simply referred to as vectors. HD computing starts by assigning an *atomic* vector to every symbol defined in the system in a so-called item memory (IM) (see Fig. 1). Atomic vectors are  $d$ -bit vectors that are drawn from the binomial distribution with  $p = 0.5$ ,  $n = d$ , and  $d/2$  as the mean (i.e., a generated vector contains i.i.d. components of equally probable 1s and 0s). They serve as *seeds* for constructing representations of more complex objects.

When the dimensionality is in the thousands, e.g.  $d \geq 1000$ , there exist a very large number of nearly orthogonal atomic vectors [16]. This allows HD computing to combine two atomic vectors into a new *complex* fixed-width vector using well-defined vector space operations, while keeping the information of the two atomic vectors with high probability. Computing with such fixed-width ( $d$ ) vectors offers fast [18] and energy-efficient [19]

learning by providing a novel perspective on data representations and associated operations [17].

In the following, we describe how the  $d$ -bit vectors can be combined using HD arithmetic operations. We focus on two main operations of HD computing: bundling and binding. Bundling, or addition, is defined as the componentwise majority function: the input vectors are summed together followed by a normalization that retains them inside the binary space. It is essentially a bitwise thresholded sum of  $k$  vectors that results in 0 when  $k/2$  or more arguments are 0, and 1 otherwise. Bundling of three vectors  $A, B, C$  is denoted as  $[A + B + C]$  where the brackets  $[\dots]$  stand for normalization. Binding, or multiplication, is defined as the componentwise Exclusive OR (XOR). Likewise, it is denoted as  $A \oplus B \oplus C$ . Both operations work on an arbitrary number of input vectors and produce a fixed-width  $d$ -bit vector with an important distinction: bundling produces a vector that is *similar* to the input vectors, whereas binding produces a *dissimilar* vector. The similarity between vectors can be measured by Hamming distance ( $\eta$ ) defined as the number of components at which they differ. See [17] for a more comprehensive overview.

HD computing for learning and classification tasks is composed of four main steps: 1) mapping symbols to atomic vectors; 2) combining atomic vectors with the arithmetic operations inside an *encoder* to produce a complex vector representing an event of interest; 3) combining the complex vectors from the same category of events to produce a *prototype* vector representing the entire class or category (i.e., learning); 4) finally comparing the prototype vectors with a query vector for categorization (i.e., inference/classification).

## III. LAELAPS: LBP CODES AND HD COMPUTING

In this section, we present the main contribution of the paper. We show how the LBP feature extractor and HD computing can be combined to efficiently learn from iEEG recordings, and then detect novel seizures. Our proposed algorithm, named Laelaps, exploits LBP codes to directly symbolize the iEEG signal of each electrode. Laelaps then applies HD computing to construct a holographic  $d$ -bit representation that captures the statistics of the LBP codes across all the electrodes and over time. This representation is then used for learning and classification in the HD space, followed by straightforward postprocessing. Laelaps is computationally efficient and extracts symbols for analyzing the occurrence of patterns, a process even somewhat similar to classical EEG reading by a human expert who tries to integrate visually detectable local and global characteristics of the iEEG signals into a coherent interpretation; then Laelaps quickly learns from one or few examples of these patterns per patient. The entire processing chain (Fig. 1) is described in the following.

### A. Preprocessing and LBP Feature Extraction

After filtering and downsampling the raw iEEG signals, an  $\ell$ -bit LBP code is computed for every sampling point. The LBP codes with different lengths ( $\ell \in [4, 8]$ ) produce almost similar performance [12], hence we set  $\ell = 6$  because using larger code sizes impairs the applicability to non-stationary iEEG signals. In addition, larger codes increase the delay of classification since the code size determines the minimum size of the statistical analysis window, i.e., the size of such a window should be large enough that all symbols can at least theoretically occur once [14]. With  $\ell = 6$ , the analysis window can be as short as 1s (containing 512 samples) that meets  $512 > 2^\ell$ . Our LBP code moves by one

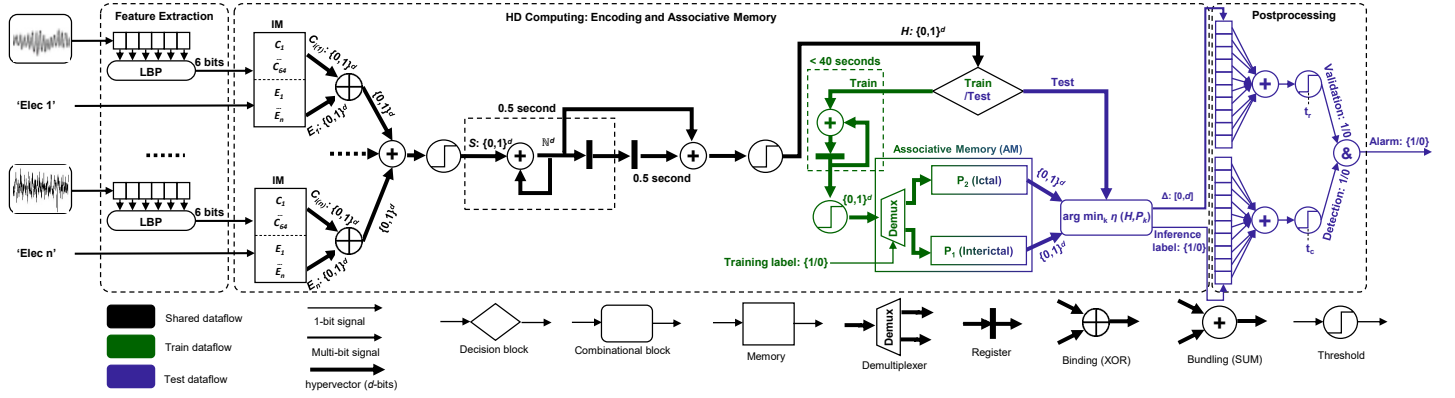


Fig. 1: Processing chain of Laelaps: (1) Feature extraction generates a 6-bit LBP code for each electrode; (2) HD computing maps these codes to  $d$ -dimensional atomic vectors and constructs a complex vector  $H$  to represent the histogram of 1 s recording. During training the associative memory (AM) learns from this vector, and during inference provides a label; (3) As postprocessing, a patient-dependent ( $t_r$ ) voting decides based on the last 10 labels and their distance scores ( $\Delta$ ).

sample, and generates  $2^6$  different symbols that are fed into the next stage for learning and classification (see Fig. 1).

### B. Learning and Classification in HD Space

HD computing first applies random embeddings of the LBP codes in the HD space via the IM, which assigns a nearly orthogonal i.i.d.  $d$ -bit vector to every LBP code, i.e.,  $C_1 \perp C_2 \dots \perp C_{64}$ . To combine these vectors across all electrodes, HD computing generates a spatial record ( $S$ ), in which an electrode *name* is treated as a field, and its LBP code as the value of this field. The IM also maps the names of the electrodes to random embeddings nearly orthogonal binary vectors:  $E_1 \perp E_2 \dots \perp E_n$  for a patient with  $n$  electrodes. This allows to bind ( $\oplus$ ) the name of each electrode ( $E_j \mid j \in [1, n]$ ) to its corresponding code ( $C_{i(j)} \mid i \in [1, 64]$ ). This binding ( $E_j \oplus C_{i(j)}$ ) generates a new set of nearly orthogonal vectors to represent LBP codes per electrode that effectively reduces the size of IM from  $64 \times n$  vectors to  $64 + n$  vectors. The spatial record ( $S$ ) is then constructed by bundling the bound vectors of all electrodes:  $S = [E_1 \oplus C_{i(1)} + E_2 \oplus C_{i(2)} + \dots + E_n \oplus C_{i(n)}]$ . The bundling operation is well suited for representing sets and multisets.

The  $d$ -bit vector  $S$  is computed for every new sample, and holographically represents the spatial information about the LBP codes of all the electrodes. The next step is to compute the histogram of LBP codes for a moving window of 1 s with 0.5 s overlap. To estimate the histogram of LBP codes inside this window, a multiset of temporally generated  $S$  vectors is computed as  $H = [S^1 + S^2 + \dots + S^{512}]$ . The bundling operation is applied in the temporal domain through accumulation (i.e., componentwise addition) of  $S^t$  vectors  $t \in \{1, \dots, 512\}$ , that are produced within the window, and then thresholding at half (i.e., normalization). As mentioned in Sec. III-A, the interictal and ictal states are characterized by different distributions of LBP codes that are reflected by  $H$ .

The output of the encoding is  $H$ , a  $d$ -bit vector that is updated every 0.5 s. Note that representation of  $H$ , as a composite structure, is constructed directly from representations of the atomic vectors by applying the operations in the encoder *without requiring any learning*. To train our classifier, we use vector  $H$  to build an associative memory (AM) containing two *prototype* vectors representing ictal and interictal labels. To train

the interictal prototype, all  $H$  vectors computed over an interictal state of 30 s are accumulated (summed), and then thresholded (normalized) to be stored in the AM as a  $d$ -bit prototype vector ( $P_1$ ). Similarly, an ictal prototype vector ( $P_2$ ) is generated from an ictal state that can last 10 s to 30 s depending on seizures' duration. For classification, every 0.5 s, the label of an unseen window is determined by comparing its  $H$  to  $P_1$  and  $P_2$  in the AM. The prototype that results in the minimum Hamming distance ( $\arg \min_k \eta(H, P_k)$ ) is the label.

### C. Postprocessing

The last part of Laelaps postprocesses the labels and distances produced by the HD classifier for the final decision. For each labeled window, we associate a  $\Delta$  score as the absolute difference between two prototype distances:  $\Delta = |\eta(H, P_1) - \eta(H, P_2)|$ . We consider a postprocessing window that takes into account the last 10 labels and their corresponding  $\Delta$  scores (shifting them every 0.5 s). The final decision is made based on two thresholds ( $t_c$ ,  $t_r$ ), the former considers the classification labels and the latter checks their  $\Delta$  scores. Laelaps flags an alarm (as the seizure onset) only if the following two conditions are met: the number of ictal labels inside the postprocessing window  $\geq t_c$ , and the mean of  $\Delta$  of those labels exceeds  $t_r$ .

We set  $t_c = 10$  for all the patients, thus flagging a seizure alarm after at least 10 consecutive output labels indicating an ictal state. This increases the detection delay but filters out many false alarms. In contrast,  $t_r$  is tuned for each patient  $P_i$ . If there are no false alarms after filtering based on the hard classification results using  $t_c$ , we set  $t_r^{(P_i)} = \min \Delta_{\text{ictal}}^{(P_i)}$  to maximize robustness to false alarms without affecting the sensitivity. Otherwise, we select  $t_r$  as the highest integer multiple of  $\max \Delta_{\text{interictal}}^{(P_i)}$ , such that it remains lower than  $\max \Delta_{\text{ictal}}^{(P_i)} - \alpha$ , where  $\alpha$  is the difference between the mean  $\Delta_{\text{ictal}}$  across the samples used to train the HD classifier and the remaining samples in the training set, averaged across all patients. This is used to compensate for the higher confidence of the classifier for the samples on which it was trained.



TABLE I: Used abbreviations/symbols: Elect.: number of electrodes, Seiz.: total number of seizures, Rec.: total duration of the recording, TrS: number of seizures used in training,  $\ell$ : delay of seizure onset detection, FDR: false detection rate, Sen.: sensitivity, and n.a. not applicable. Note that FDR\* is computed only on 20 h that is randomly selected from interictal test set.

Patient Information					Laelaps				LBP + Linear SVM [1]			LSTM [3]			STFT + CNN [2]		
ID	Elect. [#]	Seiz. [#]	Rec. [h]	TrS [#]	$\ell$ [s]	FDR [h <sup>-1</sup> ]	Sen. [%]	$d$ [kbit]	$\ell$ [s]	FDR* [h <sup>-1</sup> ]	Sen. [%]	$\ell$ [s]	FDR* [h <sup>-1</sup> ]	Sen. [%]	$\ell$ [s]	FDR* [h <sup>-1</sup> ]	Sen. [%]
P1	88	2	293	1	28.5	0.00	100.0	3	10.0	0.00	100.0	8.0	0.10	100.0	8.0	0.00	100.0
P2	66	2	235	1	16.5	0.00	100.0	10	8.0	0.75	100.0	17.0	0.40	100.0	3.0	0.75	100.0
P3	64	4	158	1	17.0	0.00	100.0	7	7.0	0.05	100.0	5.8	0.20	100.0	2.0	0.00	100.0
P4	32	14	41	2	19.8	0.00	66.7	6	30.0	0.65	50.0	22.1	1.20	91.7	n.a.	0.00	0.0
P5	128	4	110	1	5.3	0.00	100.0	1	2.7	0.25	100.0	5.8	0.30	100.0	2.0	0.15	66.7
P6	32	8	146	1	17.9	0.00	85.7	10	10.0	0.20	85.7	12.4	0.20	100.0	0.8	1.90	42.9
P7	75	4	69	2	17.2	0.00	50.0	1	26.5	1.15	50.0	9.2	1.45	100.0	26.0	0.00	100.0
P8	61	4	144	2	11.0	0.00	100.0	10	2.0	1.30	100.0	8.5	1.05	100.0	16.3	1.20	100.0
P9	48	23	41	2	8.6	0.00	81.0	6	16.3	0.10	38.1	n.a.	0.05	0.0	n.a.	0.00	0.0
P10	32	17	42	1	17.4	0.00	100.0	3	3.6	0.10	100.0	25.9	1.60	100.0	37.0	1.00	93.8
P11	32	2	212	1	19.5	0.00	100.0	3	12.0	0.40	100.0	7.0	0.05	100.0	5.0	0.20	100.0
P12	56	9	191	2	36.3	0.00	100.0	1	27.6	0.00	100.0	28.4	1.15	100.0	7.0	0.00	100.0
P13	64	7	104	2	21.1	0.00	80.0	2	11.3	0.00	100.0	6.2	0.90	100.0	1.3	0.40	100.0
P14	24	2	161	1	n.a.	0.00	0.0	1	n.a.	0.00	0.0	n.a.	0.00	0.0	n.a.	0.00	0.0
P15	98	2	196	1	20.0	0.00	100.0	1	3.0	0.15	100.0	2.5	0.05	100.0	5.0	0.00	100.0
P16	34	5	177	1	20.4	0.00	100.0	10	9.0	0.55	100.0	8.8	0.80	100.0	7.0	0.20	100.0
P17	60	2	130	1	19.0	0.00	100.0	1	13.0	0.00	100.0	3.5	0.10	100.0	16.0	0.45	100.0
P18	42	5	205	1	25.7	0.00	75.0	1	26.3	0.00	75.0	19.0	0.15	100.0	11.0	0.20	75.0
mean					18.9	0.00	85.5	4.3	12.8	0.31	83.3	11.9	0.54	88.4	9.8	0.36	76.6

#### IV. DATASET AND EXPERIMENTAL RESULTS

##### A. Long-term Human iEEG Dataset

Our anonymized dataset comprises 18 patients (P1–P18) from the epilepsy surgery program of the Inselspital Bern. For each patient, the number of seizures varies from 2 to 23, and the total durations of interictal recordings from 41 h to 293 h. Two patients out of 18 (P8 and P14) show a high number of fast and short seizures (70 and 60, respectively), hence we only consider 4 and 2 seizures as the lead seizures [20] for them. The dataset includes a total of 2656 h of recording, with 116 seizures marked by an experienced and board-certified epileptologist (K.S.). The first part of Tbl. I shows the number of electrodes, the number of seizures, and the total hours of recordings for every patient.

##### B. Results: Sensitivity, False Alarms, and Detection Delay

To evaluate the performance of Laelaps, we assess the following metrics given a few trained seizure examples: (1) sensitivity, defined as number of detected seizures out of the number of test seizures; (2) false detection rate (FDR), defined as the number of false alarms that occurred during an hour; (3) delay of detection, defined as the time distance between the algorithm detection and the expert marked seizure onset. Note that this is measured as the *working* delay of algorithms, and not the implementation delay. Implementation results including speed of execution and energy consumption are discussed in Sec. V-C.

We evaluate the performance for a practical clinical setting with the goal of reducing *time-to-functioning* of an algorithm to get it reliably operational as soon as the first few seizures are observed. Hence, we partition the dataset in chronological order to a training set, covering continuous recording until the end of the first or second seizure depending on the patient (see TrS in Tbl. I), that is followed by a longer testing set until the end of recording. The training set includes 37.7% of total recording on average among the patients. We train patient-specific models by using one or two ictal state(s), ranging from 10 s to 30 s, and one 30 s interictal state that is chosen to be 10 min before the first seizure onset. This training generates the content of AM in Laelaps and weights for other SoA. We also use the rest of

interictal and ictal samples to tune  $t_r$  available in the training set at least until the first seizure.

As shown in Tbl. I, Laelaps achieves 85.5 % mean sensitivity (100 % median) among the patients, and an ideal 0.0 h<sup>-1</sup> FDR on 1357 h of test set. In addition, for the majority of patients (11 out of 18), Laelaps achieves perfect sensitivity (100 %). Note that we have already done cross-validation on a short-time iEEG dataset in [12], and we consistently observed superior sensitivity, specificity, and fast learning benefits; however, cross-validation with this long-term dataset is highly impractical especially for the SoA methods, e.g., the classification time of the LSTM is 2.1× slower than the real-time on an Intel Xeon CPU at 2.4 GHz.

We compare Laelaps with the SoA methods in seizure detection including LBP+SVM [1], and LSTM networks [3]. Besides, we consider CNNs coupled with short time Fourier transform (STFT) that is proposed as a universal method for seizure prediction [2]. We apply these methods to our seizure detection task using the same setup but  $t_r = 0$ . These methods do not attain ideal FDR: SVM comes closest and performs ideally on 4 out of 18 patients (0.0 h<sup>-1</sup> FDR and 100 % sensitivity), but achieves 0.31 h<sup>-1</sup> FDR and 83.3 % mean sensitivity among all the patients. The other SoA methods suffer from significantly higher FDR, 0.36 h<sup>-1</sup> with the CNN, and 0.54 h<sup>-1</sup> with the LSTM, which limits their application in long-term recording. Interestingly, Laelaps still maintains a lower FDR of 0.15 h<sup>-1</sup> even with  $t_r = 0$  (i.e., without any tuning). Note that we were able to compute the FDR for the SoA methods only on 20 h of randomly selected interictal recordings due to their slow inference time. Overall, Laelaps detects 79 out of 92 unseen seizures, while the other methods perform worse: 69, 51, and 69 detected seizures using the SVM, the CNN, and the LSTM, respectively. Noteworthy, only for two patients (P7 and P14), Laelaps shows a very low sensitivity (50% and 0%, respectively). For P7, it could be due to the LBP features as LBP+SVM also reaches the same sensitivity, and for P14, all the SoA methods fail as well.

The average delay of seizure onset detection of Laelaps is slightly larger than the one yielded by the SoA methods (18.89 s vs. 12.84 s of the SVM). Other iEEG-based methods reach a

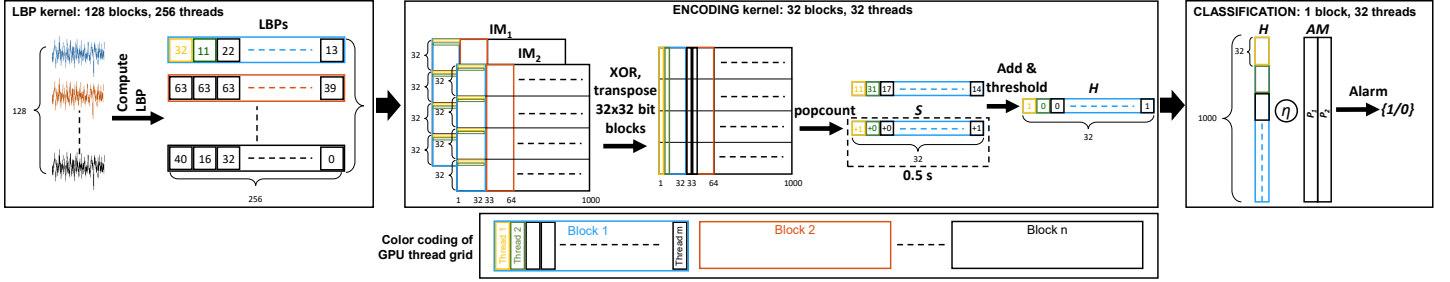


Fig. 2: GPU implementation of Laelaps using three kernels. Different colors indicate different threads and thread blocks.

similar average delay of 13.8 s on 289 h of recording with an FDR of  $0.13 \text{ h}^{-1}$  [11]. Such a detection delay below 20 s is still suited for several important applications considering that iEEG seizure onset often precedes clinical onset by more than 20 s [21].

Finally, we tune the dimension ( $d$ ) of the vectors separately for each patient. First, we perform the experiments with  $d = 10 \text{ kbit}$  to build a golden model, and then we reduce  $d$  as long as the same performance is maintained. As shown in Tbl. 1,  $d$  is reduced for 14 patients, on average to  $d = 4.3 \text{ kbit}$ , without any performance loss while saving memory, energy, and time.

## V. IMPLEMENTATION ON THE TEGRA X2 PLATFORM

In this section, we introduce the Nvidia Tegra X2 platform and describe the GPU implementation of Laelaps. We have chosen Tegra X2 to be able to compare the energy consumption and execution time of Laelaps with the SoA methods in the same hardware fabric. However, HD computing naturally fits in ultra-low power nodes ( $\approx 2 \text{ mW}$  at  $28 \text{ nm}$  [19]) as well as emerging 3D nanoscalable fabrics [17].

### A. Platform Overview

Nvidia's Tegra X2 platform is an embedded computing SoC targeted at AI workloads. It consists of a dual-core Denver2 ARMv8 CPU, a 4-core ARM Cortex-A57, and a 256-core GPU based on the Pascal architecture, which provides a performance of 750 GFLOPS (single-precision) at a power consumption of around 15 W with a memory bandwidth of 58.4 GB/s. The TX2 has several power modes, among them Max-Q, which allows for the maximum energy efficiency by running the ARM cluster at 1.2 GHz and the GPU at 0.85 GHz. We use this mode for all our experiments on the Jetson TX2 development board, and the on-board sensors for power measurements.

### B. GPU Implementation of Laelaps

We provide an overview of the GPU implementation in Fig. 2 and discuss the individual computing kernels in the following paragraphs. Readers inexperienced in GPU programming can refer to [22].

a) *LBP Kernel*. The first kernel of Laelaps maps an iEEG window of 256 samples (0.5 s) to its LBP codes. Each LBP value is computed by an independent thread: one thread block per electrode (e.g. 128), and one thread per block for each LBP (i.e. 256). This allows us to exploit data locality within each thread block by copying the corresponding electrode's data samples to shared memory before computing the LBPs.

b) *HD Encoding Kernel*. For implementation of the IM we consider two physical IM<sub>1</sub> and IM<sub>2</sub>. The computed LBPs are used to index the IM<sub>1</sub> for mapping the LBPs to the vectors. Similarly

the electrode index is used to find the corresponding vector in the IM<sub>2</sub>. The vectors ( $d = 1 \text{ kbit}$ ) stored in the IMs are packed into 32 integer variables with 32-bit each (padded, if necessary). With 64 possible LBPs, the IM<sub>1</sub> occupies 64 kbit and the IM<sub>2</sub> is 128 kbit. They thus fit entirely into the 64 kB of shared memory available per multiprocessor on the TX2 for fast access, even for the largest model configurations considered herein.

The processing is done by 32 thread blocks, one for each 32-bit chunk of the vector independently, with 32 threads each. After loading the IMs to the shared memory, the threads within each block: 1) load the vector related to the LBP code, 2) load the vector for the corresponding electrode, and 3) perform the XOR operation on them. Each of these steps takes one cycle on the TX2's GPU. The threads are then synchronized.

At this point and for each thread block, we have 32 threads, each storing a single 32-bit word. This  $32 \times 32$ -bit matrix is now transposed (using bit-masking and `__ballot_sync` instructions). A single popcount instruction is then used to sum the contribution of 32 electrodes in each thread. This is repeated until all electrodes' contributions are accumulated (e.g., 4 times for 128 electrodes).

Once this execution is completed, the results are binarized producing  $S$  that is further accumulated over 256 time steps (0.5 s), and stored as a partial sum. This result is added to the one of the previous iteration (i.e. creating the window of 1 s) and then again binarized, forming the  $d$ -bit vector  $H$  that can be used for learning and classification.

c) *Classification Kernel*. In the last kernel, the vector  $H$  is compared to the two prototype vectors in the AM, determining the Hamming distance to each of them, and the aforementioned (Sec. III-C) postprocessing is applied by the master thread. This is a computationally negligible step and is only parallelized across a single thread block of 32 threads.

### C. TX2 Measurements: Inference Speed and Energy

To measure the speed and energy of different methods discussed in Section IV, we consider an identical scenario: each model is trained offline and is loaded into the TX2 memory together with a 30 min interictal state followed by one seizure event. The model is then run on this data sequence, and the execution time and energy consumption for a single classification event are measured.

The SoA methods are implemented using optimized libraries and frameworks, specifically Keras for deep learning approaches (relying itself on Nvidia's highly optimized cuDNN library) and Scikit-learn for the SVM. We use various combinations of resources (GPU and/or CPU) to find the best performance for

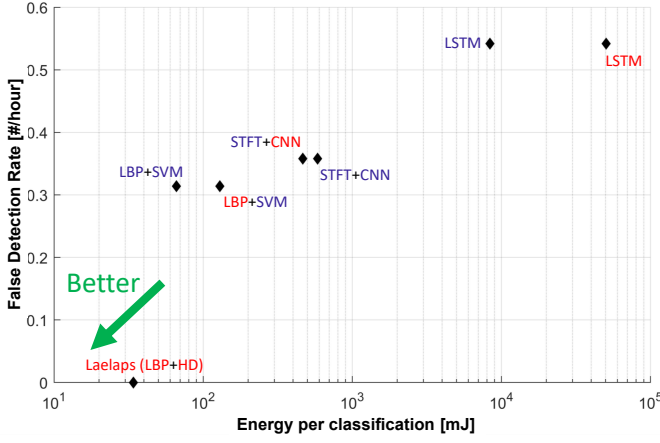


Fig. 3: Comparing the methods in terms of false detection rates and energy consumption with 64 electrodes on the TX2 in the Max-Q power mode. Legend: CPU, GPU.

TABLE II: Energy and inference time of Laelaps, SVM, CNN, and LSTM for one classification event. Legend: CPU, GPU.

Feature Extr. Classifier	LBP HD	LBP SVM	STFT CNN	n.a. LSTM
<b>#Electrodes: 128</b>				
time [ms]	13 (1×)	51 (3.9×)	213 (16×)	6333 (487×)
energy [mJ]	35 (1×)	103 (2.9×)	556 (16×)	16224 (464×)
<b>#Electrodes: 24</b>				
time [ms]	13 (1×)	21 (1.7×)	53 (4.2×)	1416 (113×)
energy [mJ]	32 (1×)	45 (1.4×)	131 (4.1×)	3980 (124×)

each of the SoA methods. Fig. 3 compares the mean FDR of all the patients, and energy consumption per every 0.5 s classification event using 64 electrodes as the median number electrodes among the patients. It highlights that the SVM requires up to 2 orders of magnitude less energy with lower FDR compared to the deep learning methods, confirming its usage in the wearable EEG-based seizure detection devices [7]. Laelaps remarkably surpasses the SVM by  $1.9\times$  lower energy and zero false alarms.

In Tbl. III, we further analyze these aspects for two electrode configurations: 24 electrodes as the minimum number of electrodes among patients (P14), and 128 as the maximum (P5). We show the results of a single implementation of each method, either on the CPU, GPU, or a combination of them, that results in minimal energy consumption (note that LSTM is memory bound while CNN is compute bound).

With the 24-electrode model, Laelaps takes 12.5 ms and 32 mJ achieving  $1.7\times$  faster classification and  $1.4\times$  lower energy consumption compared to the SVM. Laelaps achieves a speed-up of  $4.2\times$ – $113\times$  and an energy saving of  $4.1\times$ – $124\times$  with respect to deep learning algorithms.

We also evaluate the scalability by increasing the number of electrodes to 128. Laelaps’s savings (in both time and energy) are magnified:  $3.9\times$  ( $2.9\times$ ) faster execution (lower energy) with respect to the SVM, and  $16\times$ – $487\times$  ( $16\times$ – $464\times$ ) compared to the deep learning methods. These results indicate almost constant execution time and energy consumption of Laelaps with respect to the number of electrodes (12.5 ms with 24 electrodes vs. 13.0 ms with 128 electrodes, and 32.0 mJ vs. 35.0 mJ). In contrast, the SoA methods exhibit a linear growth at least in both these metrics

(e.g., 20.8 ms vs. 51.0 ms, and 44.8 mJ vs. 103.0 mJ in SVM), thus implying better scalability of Laelaps for more electrodes.

## VI. CONCLUSION

Laelaps—on our very large dataset with 116 seizures in 2656 h of continuous recording from 18 epilepsy patients implanted with 24 to 128 intracranial EEG electrodes—simultaneously reduces the following metrics compared to LBP+SVM, STFT+CNN, and LSTM: the number of undetected seizures, false alarms (to zero), execution time and energy consumption for classification on a TX2 embedded device. Our future work will focus on designing specialized hardware for Laelaps and on reducing the delay.

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