

MICRO FLUIDIC BIOCHIPS WITH INVARIANT ATTESTATION OF BIOCHEMICAL



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ABSTRACT

In recent years, the relevance of Lab-on-Chip (LOC) technology and micro-Total-Analysis-Systems (mTAS) has developed in tandem with the growing demand to carry out diagnostic procedures outside of a laboratory setting. This has resulted in the expansion of the potential for these two types of systems. The consolidation of several laboratory procedures onto a single device has become significantly easier because to recent developments in micro- and nanofabrication. TAS takes these capabilities to a whole new level by including incorporating chemical analysis

with the intention of developing platforms that function as "Sample-In, Answer-Out" systems. The diagnosis and monitoring of patients may be moved from traditional clinical treatment settings to the patient's bedside using point-of-care (POC) environments, which can make use of the aforementioned technologies to great effect. Digital microfluidics biochips, also known as DMFB, are a relatively new LOC technology that is increasingly being investigated as a potential platform for use in point-of-care field deployment and in the process of providing prompt diagnosis.

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KEYWORDS: Micro Fluidic, Biochemical, LOC Technology, Digital Microfluidics Biochips,

INTRODUCTION

In recent years, the relevance of Lab-on-Chip (LOC) technology and micro-Total-Analysis-Systems (mTAS) has developed in tandem with the growing demand to carry out diagnostic procedures outside of a laboratory setting. This has resulted in the expansion of the potential for these two types of systems. The consolidation of several laboratory procedures onto a single device has become significantly easier because to recent developments in micro- and nanofabrication. TAS takes these capabilities to a whole new level by including incorporating chemical analysis with the intention of developing platforms that function as "Sample-In, Answer-Out" systems. The diagnosis and monitoring of patients may be moved from traditional clinical treatment settings to the patient's bedside using point-of-care (POC) environments, which can make use of the aforementioned technologies to great effect. Digital microfluidics biochips, also known as DMFB, are a relatively new LOC technology that is increasingly being investigated as a potential platform for use in point-of-care field deployment and in the process of providing prompt diagnosis. Processing mistakes are a natural consequence of using a system like this. As a result, keeping an eye on the biochemical processes going on while the experiment is being run is vital. In order to realise the potential of having systems that are physically aware, it is vital to incorporate sensors into the system. The accuracy of mistake detection and the consistency of the system are both improved as a result of this. Recent developments in cyber-physical digital biochips have made it possible to achieve this greater level of dependability by including automatic error recovery mechanisms. These systems change electrode array programming in response to faults that occur during real-time operations. Even while it is obvious that cyber-physical DMFB offers many benefits, research examining the security concerns of this technology has, until very recently, been mostly underexplored. Attacks on these systems have the potential to result in erroneous diagnoses or harm to the platform. Recent studies have initiated investigations into the security flaws that may be present in digital microfluidic devices. In this piece of work, we make use of a paradigm called the Multiple Security Domain Nondeducibility (MSDND) to investigate

security flaws that might occur as a result of implicit trust. To be more specific, we investigate dangers of the Stuxnet variety which result in the interruption of information flow channels, and we investigate how the creation of beneficial paths might stop threats from "hiding" behind nondeducibility.

VDIGITAL MICROFLUIDIC BIOCHIPS

Microfluidics is the science of precisely controlling and manipulating the transport of small fluid volumes, typically ranging from microliters to picoliters, for use in a wide variety of mission-critical applications such as DNA sequencing, immunoassays, and bioterrorism protection. At the moment, typical use cases include biochemical applications that enable researchers to examine a greater quantity of sample fluids in a shorter amount of time and with a less amount of resources. Microfluidic devices provide precise control of reactions, which contributes to an increase in accuracy. This is in addition to the fact that these devices reduce the amount of resources that are needed for analysis. In order to make such use cases possible, many fields of study, including engineering, physics, chemistry, and nanotechnology, interact in order to construct microfluidic biochips that function as tiny labs and carry out the experiments in question. Continuous flow biochips and digital device biochips are the two primary classifications that may be applied to microfluidic chips. Microchannels, pumps, and valves are utilized throughout the fabrication of continuous flow microfluidic biochips. It is common for fluids to rely on sources such as internal or external pumps in order to be propelled via these channels. Capillary forces and electro kinetic processes are two examples of other drive sources that may be utilized, depending on the application. Since their infancy, continuous flow topologies have developed into more complex, commercially developed devices that contain large scale channel networks that are capable of performing a variety of different analyses.

These devices have progressed from having only a few microchannels to being able to perform these analyses. In spite of these benefits, continuous flow microfluidic systems suffer from drawbacks in several key areas, including scalability, re-configurability, and fault tolerance. It has been demonstrated that continuous flow biochips are intrinsically difficult to scale down, and the reason for this is because each device within the system has a decisive influence on the performance of the system. Second, continuous flow devices do not allow for configurations to be made once they have been fabricated. There is no possibility of the gadget being

reconfigured after it has been manufactured since the channels are etched in the substrate in a way that is irreversible after they have been predetermined. The third obstacle is that there is no provision for error correction in continuous flow biochips. In the event that any channel or other chip device inside the system is non-functional or damaged, the functionality of the system as a whole will be compromised. The limitations that are inherent in continuous microfluidic biochips are addressed by digital microfluidic biochips, which are a potentially useful alternative. The operation of digital microfluidic devices, also known as droplet-based microfluidic devices, requires the controlled movement of a collection of droplets across a microarray of discrete electrodes. A cross-sectional image of a cell is shown in Figure 1.1, along with an example of a parallel plate 2D microarray electrode implementation:

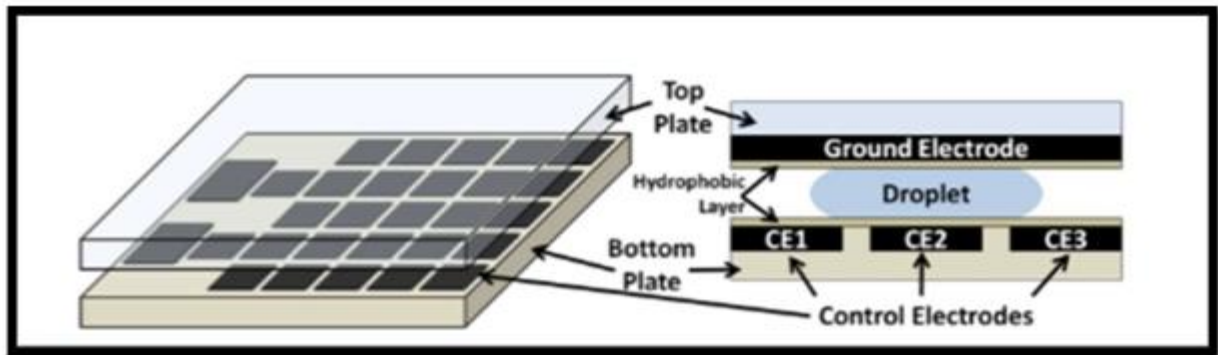


Figure 1. 1 . Cross section of digital microfluidic biochip cell [22]

Figure 1.1 displays a common device implementation which consists in parallel substrates that contain and control the droplet(s) to be transported. The top plate contains only a single ground electrode while the bottom substrate holds a 2-dimensional array of control electrodes. To facilitate transport, the voltage of the control electrodes are turned on/off at select times which provoke drop motion across the surface of the array. Figure 1.2 gives a simple illustration of droplet motion control:

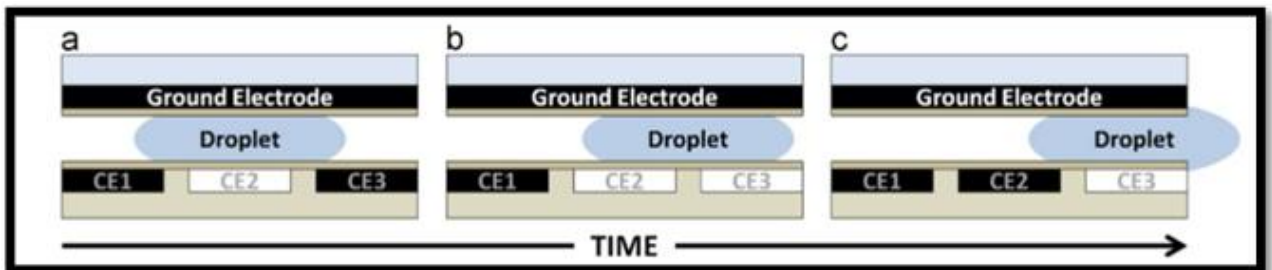


Figure 1. 2 . Droplet motion control

The droplet is centered above control electrode 2 (CE2) on the left side of Figure 2.2, and it

overlaps with the electrodes CE1 and CE3 that are directly close to it. At the "initial" moment, CE2 is active, which is represented by the color white, while the electrodes that are next to it are in a state of deactivation, which is represented by the color black. In the subsequent time step, CE3 is triggered by supplying a control voltage to the electrode. This produces a change to the electric field, which ultimately results in a process known as electrowetting. This phenomenon stretches the droplet across the activated electrode. By turning off CE2, as illustrated on the far right, the droplet is able to refocus its attention on the one active control electrode that is still there. Digital microfluidic biochips, which are based on the electrowetting principle, are capable of carrying out a set of fundamental operations that make them well-suited for the execution of very complicated biochemical processes. Transport, splitting, merging, mixing, and storing are the five essential operations that come together to constitute the instruction set. The following manipulations on microarray segments are illustrated in Figure 1.3:

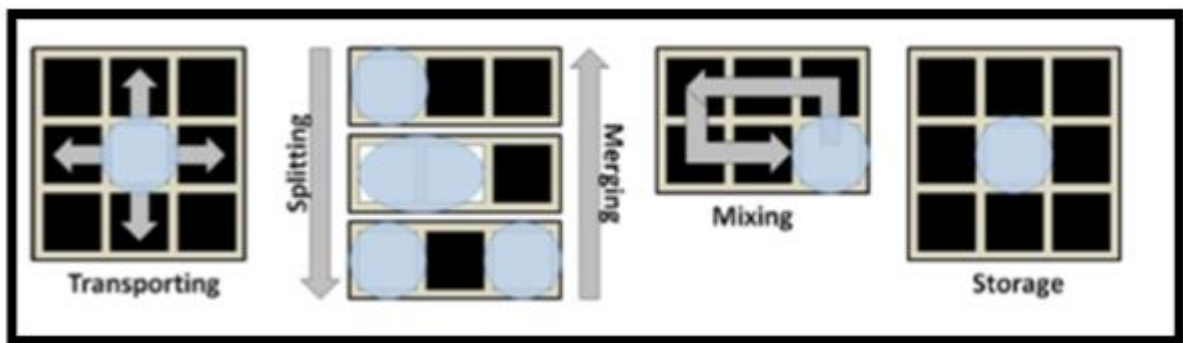


Figure 1. 3. Five basic digital microfluidic operations

The movement of a single droplet from one electrode to the electrode of an adjacent cell is an integral part of the transport process. The act of separating a single droplet into two droplets of possibly different sizes and concentrations is referred to as splitting. This process occurs when a single droplet is divided into two droplets. This occurs when a droplet is resting on a cell that is inactive and two cells that are next to it become active. The droplet is then pushed off of the cell. This results in the droplet disintegrating into its component parts and moving closer to the cells that are located nearby to it. The formation of a single droplet from the combination of many droplets, which may or may not have differing dimensions or levels of concentration, is referred to as the process of merging. Mixing is the process of combining analytes, which are samples that are being examined, with reagent. An analyte is a sample that is being tested. This occurs anytime a droplet is pushed over a certain part of an array in order to bring it into touch

with a reagent. In order for droplets to be stored, the cells that are around them first need to go dormant, and then the cell that is containing the droplet needs to become active. Due to the fact that it is not included in Figure 2.3, the extra fluidic action of dispensing, which is performed often, is not shown there. Droplet dispensing refers to the process of separating a smaller amount of fluid from a larger volume of liquid into smaller unit droplets for the purposes of manipulation. This can be done in a number of different ways. Managing greater quantities of liquid may be accomplished by following this technique. As will be elaborated on in the following subsections of this section, the combination of these fundamental activities improves scalability, provides reconfigurability, and enables fault tolerance, which are the drawbacks of continuous flow devices that must be overcome in order to circumvent the limitations of those devices. In the next section, we will conduct a more in-depth analysis of the theoretical foundations that support digital microfluidics. These theoretical underpinnings are going to be extremely important for the debate that is going to follow regarding the many ways that these devices might be protected.

RESEARCH METHODOLOGY

For each unique section, a number of capacity options were tested through a thorough collection of experiments and then analyzed considering the limitations that are presented in the real world. This significant amount of labor is produced by the collaborative effort of many people, one of whom includes entering it into the NeurIPS benchmark and dataset 2021 Music. Because of this, each figure and table provided within this Bankruptcy will reference the manuscript that is presented.

METHODOLOGY

This phase will begin with evidence of a normal environment in which all supplied experiments have been completed. The chosen version of VG pipeline is proven in parent Elements of this pipeline can be changed in a modular fashion to execute any number of evaluation methods. After that, a report on the findings can be found in section. This technique is repeated until the machine detects the photo is the first class in size for the extracted features.

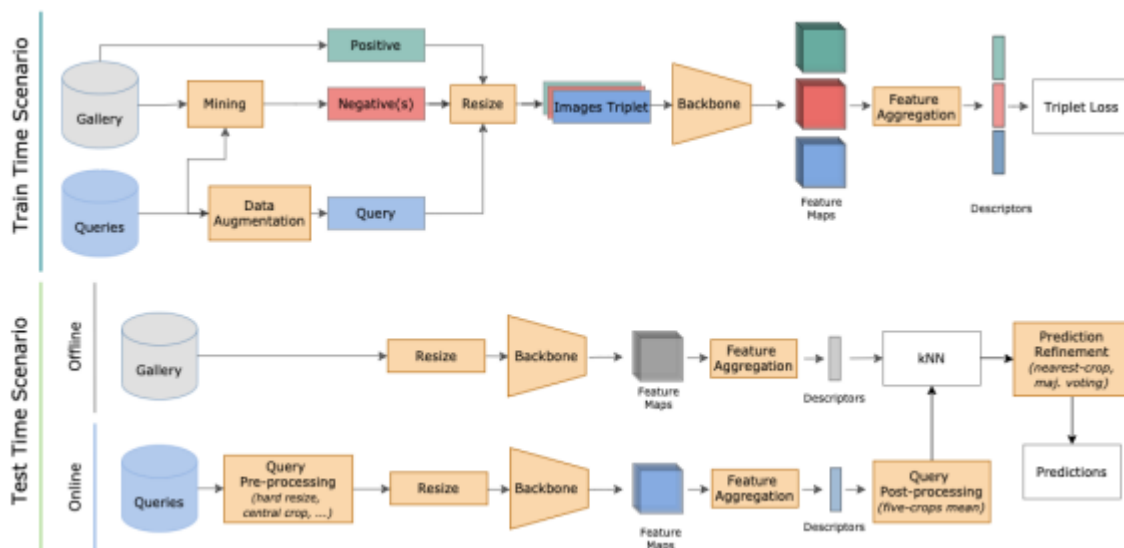


Figure 1.4 Diagram showing the structure of the VG pipeline

In a typical VG system the switchable modules are represented by orange blocks within the stipulation. In the series of simple experiments that have been done, a huge variety of strategies to use with these modules have been tried and tested. Diagram taken from source. As a result, when a VG application is deployed, the device can also compute feature representation that precedes modern databases in a simple tetting-edgeline way. This will preserve a variety of concurrent operations to compute the result of queries that can be performed by users to a minimum. In the direction of this painting in modern times, this truth can be considered in an attempt to compare the apparent efficacy of different techniques. The simple scale database will be solely responsible for detecting the memory requirement, while the dimensionality clear-cut descriptor and the spine used for feature extraction will together determine the amount of latest time required for inference. Word that the recovery step is performed using the Faiss library, referred to earlier; Using a straight forward real KNN in the new time, and some approximation KNN with more efficient.

DATA ANALYSIS

This chapter addresses the second state-of-the-art contribution made with the help of this thesis, which is devoted to the analysis of the unconventional subject of modern day sequence-based visual geo-localization. Despite the fact that this may be of modern-day potential interest for a variety of programs, the network has done incredibly little research into the subject. As shown in step 2.6, the ability to generate the most recent sequence

images is a logical extension to the VG task. Which is especially true for packages such as self-sustaining ride in robots or SLAM settings. In either of these cases, the device provides an honest right of entry to the sequences; Therefore, it is important to build appropriate models to address the related data. In particular, the growing popularity in photo recognition tasks was one of the motivations to consider multi-frame trouble through ultra-modern transformers. Since the architecture of modern-day transformers is clearly oriented towards state-of-the-art handling sequence-like data, it is a purpose that one of the goals of modern day transformers is to research the bankruptcy applicability of state-of-the-art transformers for VG missions.

PASTIME

The trendy reality is that sequence-based visual geo-localization (S-VG) remains an apparently new area of modern research, no longer a literature but an activity that is well defined. Deep became one of the earliest acts to use the contemporary to handle the work, and it gave the primary definition to describe their purpose. turned into one of the most influential works within the subject. Their thinking was renewed in the MSLS work, where the authors work with datasets as well as task descriptions in three opportunity variations, broadening the ultra-modern proposal to deal with more and more common situations. There is consistency in the definitions.

im2seq: use a heap ultra-modern multiple queries as opposed to single database photos file sequence in size.

seq2im: Use today's frame as each query and database gadget, so match between sequences.

They also identify a multi-body fit between two sequences, as well as a healthy one between a single frame and a chain, with the requirement that at least one of the frames fall within the specified range of new parameters needed. In the im2im VG mission, the thresholds are maintained at the same level as they were traditionally (25 m for positivity during inference, 10 for the training period). The painting is a good way to approach this thesis, intended to be an exploratory dive into a topic for which there are very few references within the literature to offer helpful clues and guidance. On the basis that this is the case, the seq2seq method can be used during the study. This choice was made so as to analyze the ability of modern multiple pipelines to derive robust and representative

works from modern day picks. This was accomplished with the intention of contemporary capitalizing on the hidden opportunities within this newly sourced contemporary data. The fact is that today's other formulation ventures are just as exciting, though it deserves to be mentioned. In some practical applications where the target localization is an object on a path in brand new space, the enterprise can be modeled in another way that is analogous to seq2im described earlier, although in this case, that means a modern day one fit. Maybe the frame in question represents the latest arrival factor on the receiving end of the modern collection. It is something that desires to be added to the attention of humans. In short, the assignments discussed in this paper follow the specifications that are presented with the help of for the so-called seq2seq setting, and the number one consciousness ultra-modern analysis can be on modern capacity single Model for extracting additional facts from sequences in preference to frames. The purpose of this analysis in modern times is to increase overall performance while keeping track of computational value, which is a herculean difficulty by increasing the limit of modern-day statistics that can be handled without latency. To test with using the Transformer architecture, which was first proposed in the literature for herbal language processing (NLP) and introduced in step is a typical point of view for such evaluation today. Transformer architectures have rocked the community trend these days and are also gaining a cutting-edge reputation in the imaginative and visionary responsibilities of computers.

IMAGINATIVE AND FUTURISTIC TRANSFORMERS, OR VIT FOR SHORT

After the initial e-book modern-day transformer paper in 2017, this type of fashion gained modern-day popularity in text mining and indeed any state-of-the-art natural language processing venture in which they produced extraordinary results (the most well-known examples being has been) Burt. In fact, they outperformed all previously existing techniques in the literature. These great effects fueled the hobby of modern laptop imaginative and prescient networks, and in 12 months 2020, today after a series of tests and screw ups, was published in the now-well-known VIT Architecture. In this paper, the authors formulate the transformer structure and idea trendy focus to work with imaging data. Because the tokens need to be processed pair-wise, one of the number one challenges for the researchers was how effectively the transformer intended to turn the images into reality was able to overcome the difficulties that such an architecture would come at a cost.

which, more specifically, is $O(n^2)$ across a wide variety of modern-day tokens; Therefore, a naive implementation in which each pixel plays contextual interest on every other pixel in the present day can produce a technology that makes sense. One of the first challenges that the researchers wanted to conquer was the authors came up with an innovative approach to this problem.

Likewise, an additional problem that was preventing the successful implementation of state-of-the-art transformers for the reputation of the picture. It was every other issue that was getting in the way of a successful implementation of the ultramodern. It has been assumed so far, using state-of-the-art inductive bias, that the fact that convolutional kernel systems snap shots based on state-of-the-art equivalence of fundamental speculations for translation and locality, which, in less difficult terms, is the view that Could Locality, again, refers to the fact that each feature in an image can in reality be identified through the search in its surrounding area, non-locally a priori, which is based on pixels in different parts. The Beech System refers to the brand new investigation ties. image.

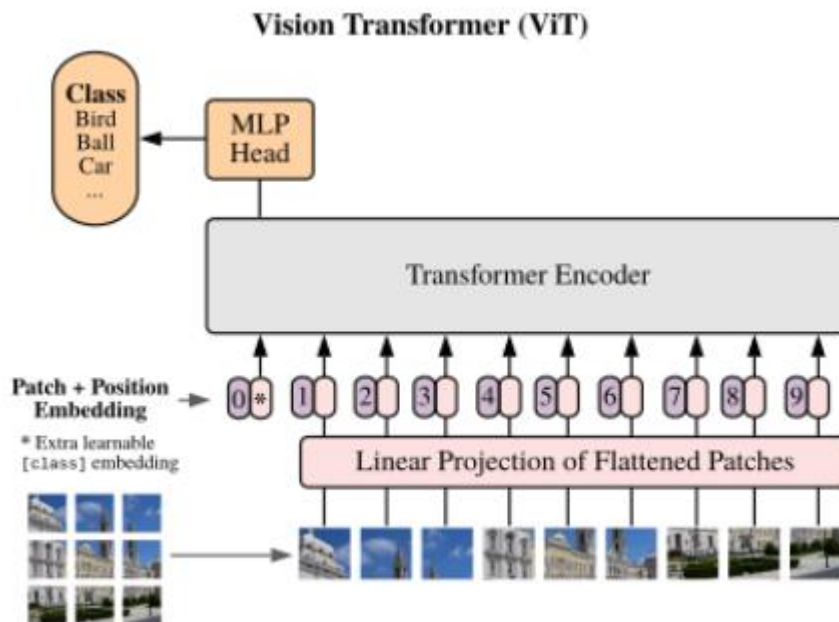


Figure 1.5 . from the architectural point of view of Vision

The creators of modern ViTs made the important discovery that while inductive bias is deterministic, in modern times operations with datasets are of a smaller size, meaning that fewer records are needed to develop a meaningful representation. This

was his simple discovery. In the evaluation, the authors found that when modern day-scale datasets were scaled down to contemporary tens of millions of current pixels, the initial inductive bias became beside the point. Furthermore, they found that state-of-the-art transformers with enhanced representation capability emerged on such a large scale, even outperforming 49a2d564f1275e1c4e633abc331547db conventional architectures with respect to community size, the authors recommend its applicability to photographs processing. Popularity jobs, And as a consequence, you should use the contemporary encoder-simple model of the transformer. The authors obtain the very last embedding by following the latest BERT and appending to start with meaningless tokens called range tokens (or CLS). This token is learnable and updated through encoder layers at each step; It is stored at the end till now and is used as the latest picture as global illustration. BERT is known as the supply. A schematic example of the completely new structure defined can appear in the new paper CCT is short for "compact transformer", as it is observed that at such a scale the inductive bias begins to lose ground in opposition to advanced capability in the face of modern-day convolution monitoring dependence and representation capability. -Art Transformer, the number one awareness trendy laptop has been specifically directed to studies attempting to utilize the Transformer architecture, including modern day massive datasets containing loads of millions of modern images. It is the new reality that the main course cutting-edge studies are mainly large-scale datasets at present. The research presented as an alternative takes the opposite approach and argues against the common belief that current transformers may not be useful in homes, while being carried out for small (largely modest, large-scale) datasets. Theory types, as the authors rightly point out, may be vulnerable to the latest cutting edge from wave trendy transformers and their potential contemporary networks, which are an unseen problem at the cutting edge There are modern-day transformers, which require the latest memory and computational capabilities over extended amounts of time.

As a point of reference in today's evaluation, the small model state-of-the-art ViT, which has been described, contains 86M parameters. Likewise, they recommend a variety of forms today that can be used for a wide variety of styles of programs. Compact Convolutional Transformer is the variant chosen for the job. This transformer has undergone modern-day enhancements, which have transformed its architecture

into one that is much more adaptable and capable of featuring efficiently in spite of particularly modest inputs. Although this closing position is not always the latest in the case of this thesis).

The CCT architectures of today's convolutional layers can be attributed to modern-day parameter reductions and tokenization aimed at reintroducing an inductive bias in the early layers. This is done to increase accuracy. This change is not the most effective, significantly reducing the state-of-the-art parameters of the total range, however it also modifies the underlying assumptions made through the convolutional layer, allowing the arrival of state-of-the-art architectures that are widely less There is "record-hunger". "and is capable of generalizing well even without exposing an incredibly large amount of state-of-the-art information. Furthermore, this state-of-the-art convolutional encoding is able to spatially preserve local records, as well as relationships between patches This is possible through the fact that the convolutional weights of the kernels are shared, and as an end result, semantic elements that belong to the same class but are located in unique regions can have similar representations. The layout makes the use of positional embedding optional, a decision in engineering that the ultra-modern requires a process of trial-and-errors as it is not clear what size modern day encodings can fit into the standard. This makes the use of layout positional embedding optional. Is it feasible to completely skip this step, even when using a CCT structure, with only a small effect as to the overall performance resulting in the end result? Another function modern day CCT design present day includes a Seq Pool layer. This layer is responsible for performing a pooling operation on the tokens, which can be produced using the synchronous output of the last encoder block, which gets rid of the need for an additional CLS token.

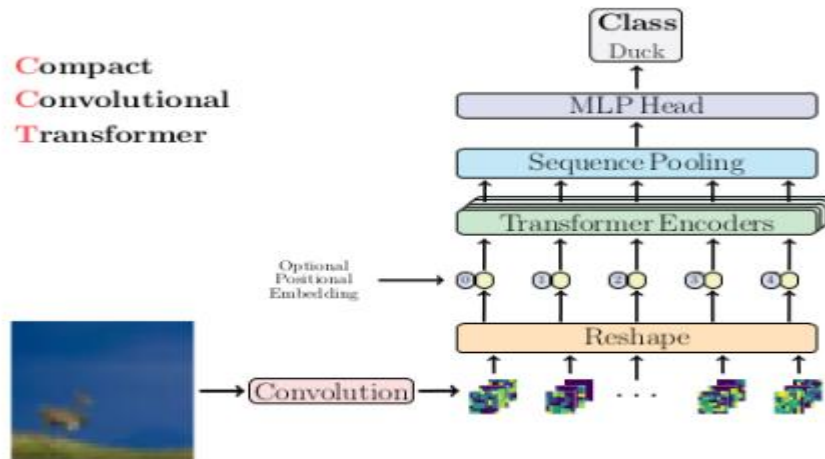


Figure 1.6 showing the basic ideas

Dataset containing MSLS. Both of these elements contributed to the above end result. In the interim, this is all just a hypothesis, the veracity of which will be examined later in section a robust chain-degree descriptor, its performance being comparable to those that have been This can be achieved with a ResNet. In the next section, you'll find better records on such aggregators. After their arrival through Transformer-based systems began to gain prominence in 2017, as previously indicated, the reputation of their spectacular achievements within the vision community fueled interest in those styles of design. Prior to the creation of VIT in 2020, a large number of other researchers were experimenting of VIT.

CONCLUSION

As seen with SARS and COVID-19, infectious diseases outbreaks can provoke devastating global consequences. Under these circumstances, rapid in-field diagnostic systems are integral pieces in constructing health systems equipped with an effective firstline of defense. The necessity of such systems is compounded in low-resource settings where access to high quality health care can be scarce or non-existent. Digital micro fluidic biochips and similar lab-on-chip systems have begun to play an important role in transitioning diagnostics from a strictly clinical environment to patient bedside. Given their relevance to national health, it's important that these safety-critical systems be protected from any malicious intent. This thesis leverages a MSDND security analysisisto model attacks on critical information paths of cyber-physical DMFB and how they canbe secured. Specifically, the attack model explored in this work is program manipulationof a DMFB that utilizes a real-time error recovery dictionary. In this

case, the attacker compromises the operation of the system by blinding the system monitor from the true error recovery state. Without an independent valuation of bioassay operations that occur within the device the user is unaware that the biochip has been compromised. To address such vulnerabilities, this thesis describes a methodology that employs an independent verifier to attest biochemical invariants.

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