

STORAGE DEVELOPER CONFERENCE



Fremont, CA  
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*BY Developers FOR Developers*

A  SNIA Event

# DNA Data Storage

DNA Data Storage Alliance - Rosetta Stone Initiative

Presented by

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A SNIA Technology Affiliate

# Helpful Links

- [Preserving our Digital Legacy – an Introduction to DNA Data Storage](#)

# Agenda

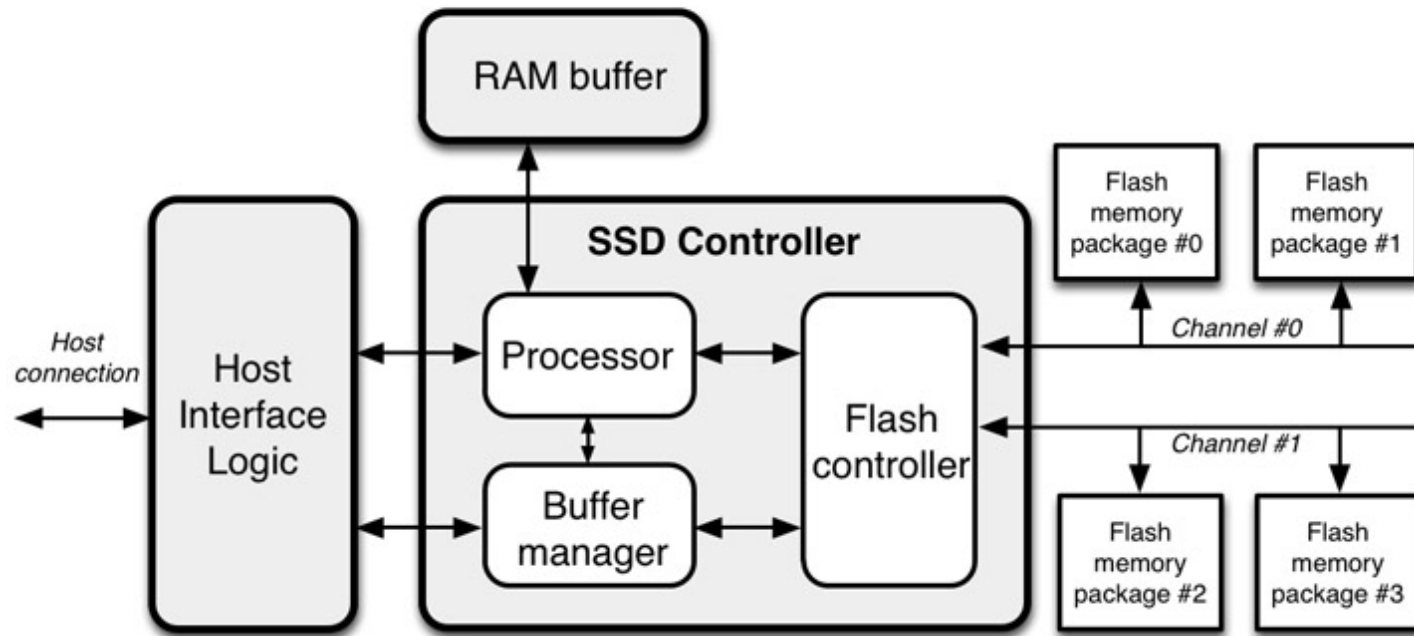
- Differences: DNA vs Traditional Media
- Overview of the Rosetta Stone
- How to Participate
- Summary

# Differences: DNA vs Traditional Media

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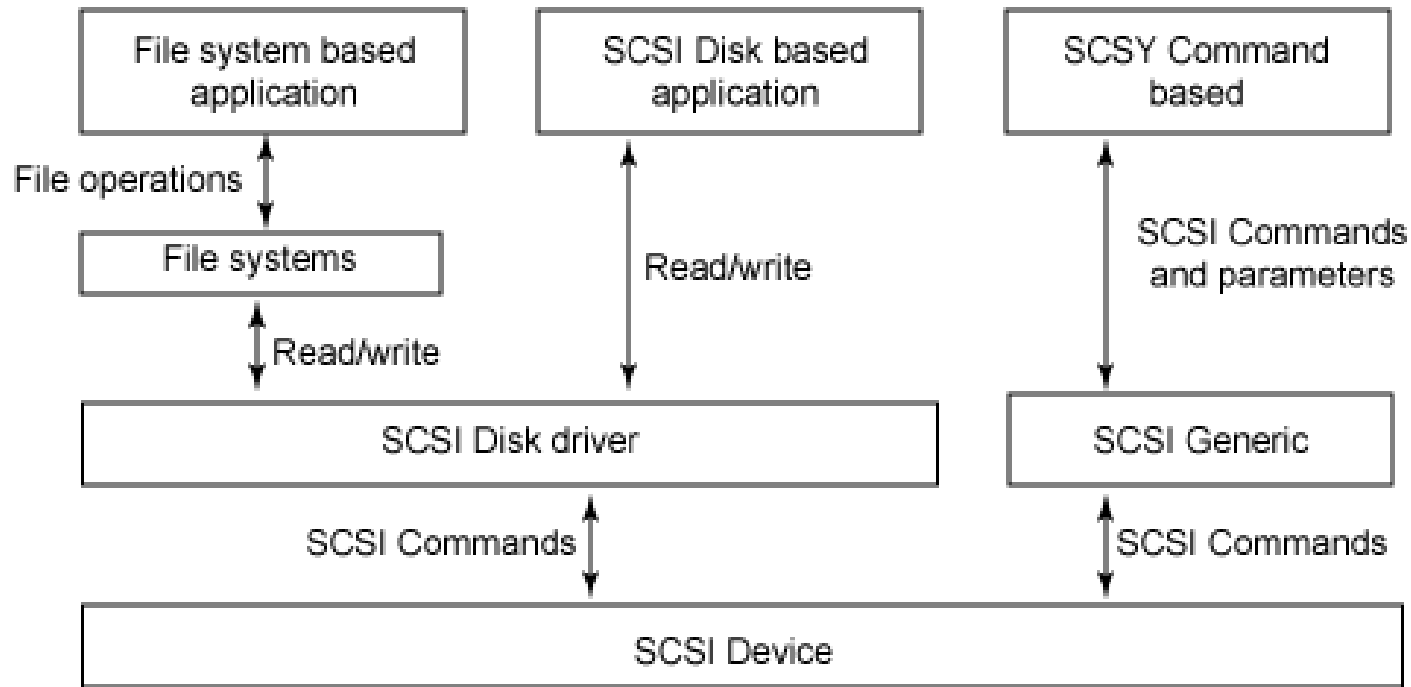
## 1. Exposing a device to the system

### Architecture of a solid-state drive



# Differences: DNA vs Traditional Media

## 2. Organizing abstractions to create filesystem storage



# Differences: DNA vs Traditional Media

## 3. Media without integrated controller



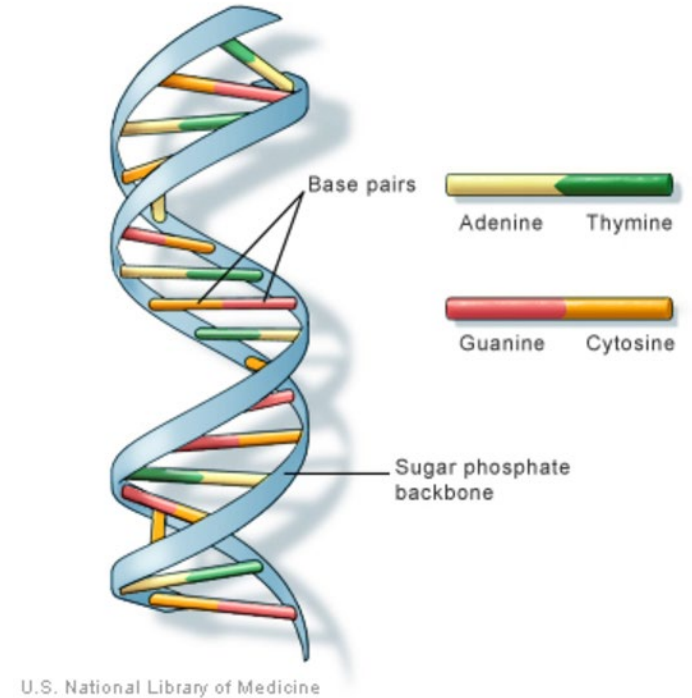
*Barcode with volume serial number, generation, and type of cartridge*



*Read LTFS from beginning of the tape*

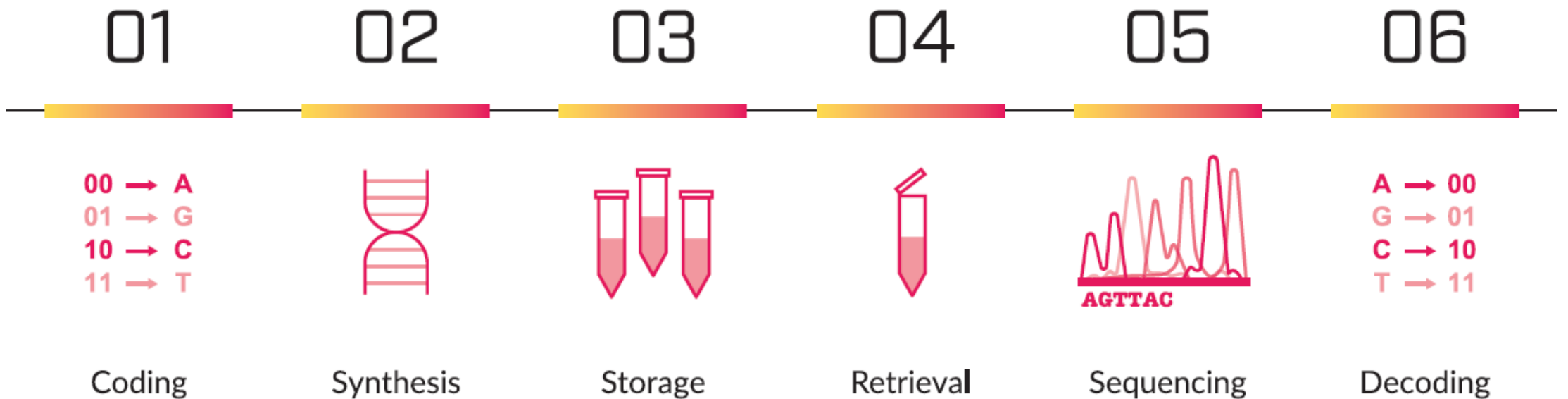
# A “Primer” on DNA Data Storage Media

- The fundamental unit of storage in DNA is an oligonucleotide (also called ‘oligo’)
  - Short, single strand of synthetic DNA or RNA
  - Often a sugar phosphate backbone
  - Base compounds Adenine, Cytosine, Thymine, Guanine
  - Base compounds attach to the strand ...
  - ... and to a mate on the opposing strand
  - Adenine bonds w/ Thymine, Guanine bonds w/ Cytosine
- A DNA molecule is a pair of strands (oligos), tightly wound around one another, held together by the bonds between the bases





# A “Primer” on DNA Data Storage Media

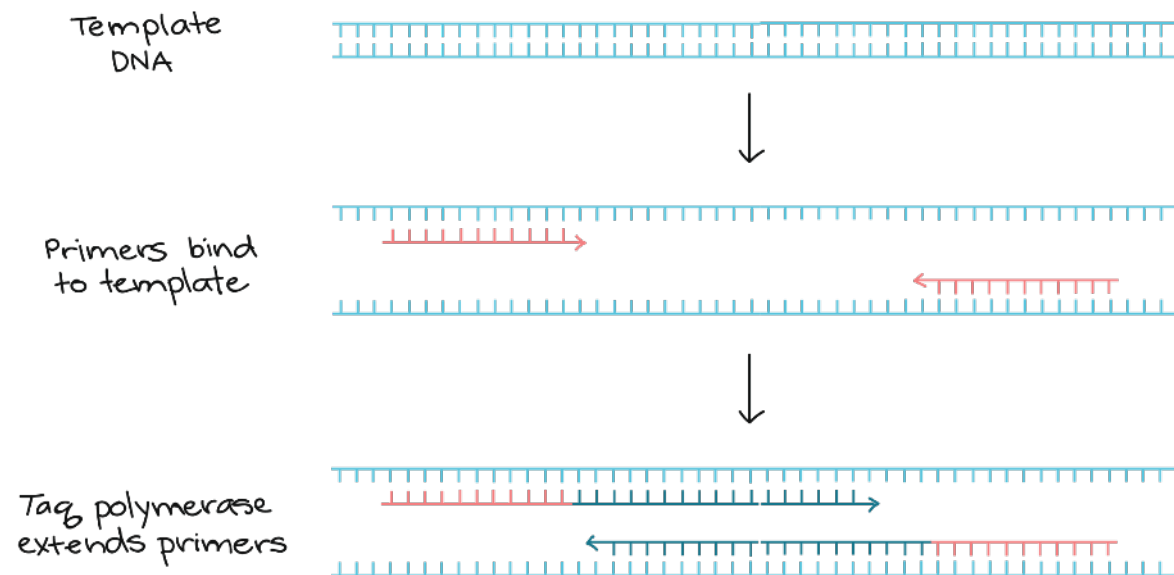


# A “Primer” on DNA Data Storage Media

- The process of storing binary data into DNA data storage media involves
  - Coding – conversion of binary numbers to ATGC base pairs (“bits to bases”)
  - Synthesis – creation of the strands and chemical bonds between the base compounds
  - Storage – placing constructed strands into sealed medium until contents are needed
- The process of retrieving binary data from DNA data storage media involves
  - Retrieval – accessing the sealed medium containing the required strands
  - Sequencing – discerning the bases found in a segment of DNA
  - Decoding – converting ATGC pairs into binary (“bases to bits”)
- DNA has neither addressable sectors (disk) or relative position (tape)
  - Locations and addresses must be encoded into the material itself

# A “Primer” on DNA Data Storage Media

- A **primer** is a short stretch of DNA targeting a unique sequence, generally to identify the sequence for **amplification**
- **Polymerase chain reaction (PCR)** is the process used to create one or many copies of the amplified DNA



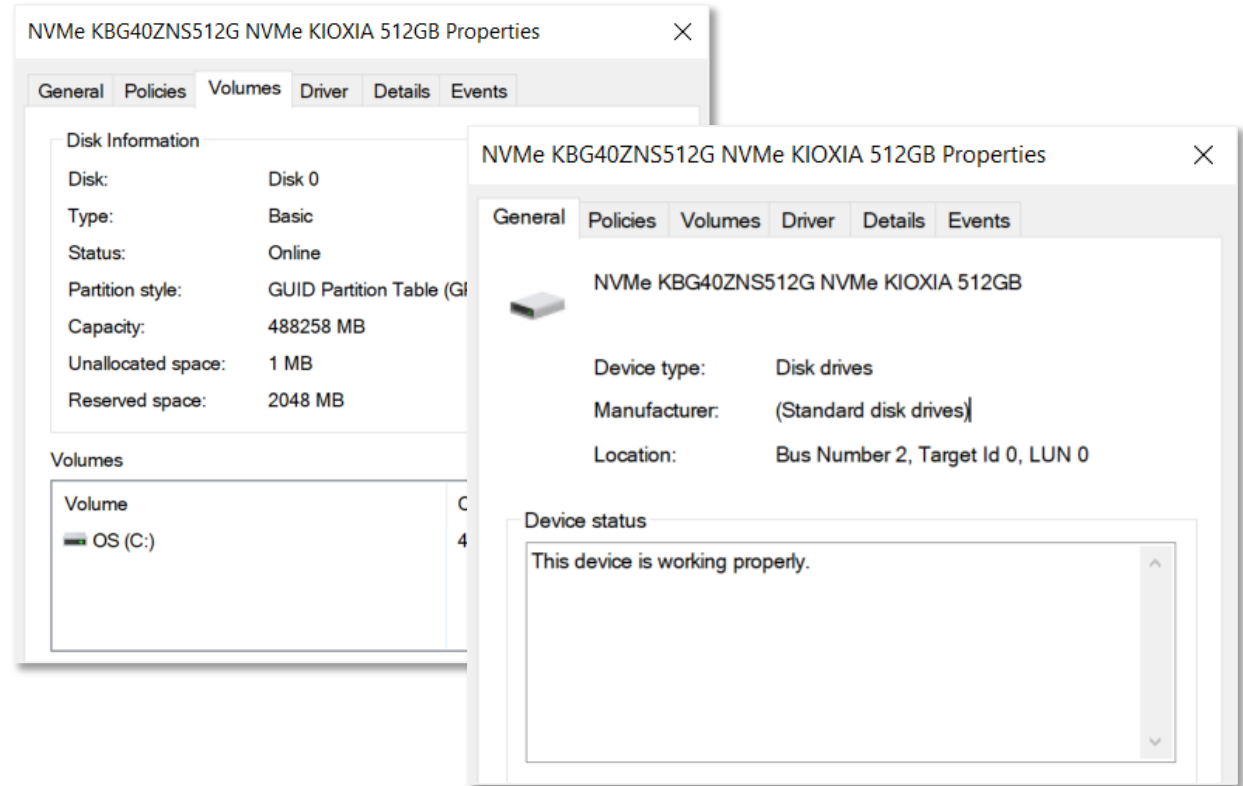
# Overview of the DNA Data Storage Rosetta Stone project (DARS)

# The problem

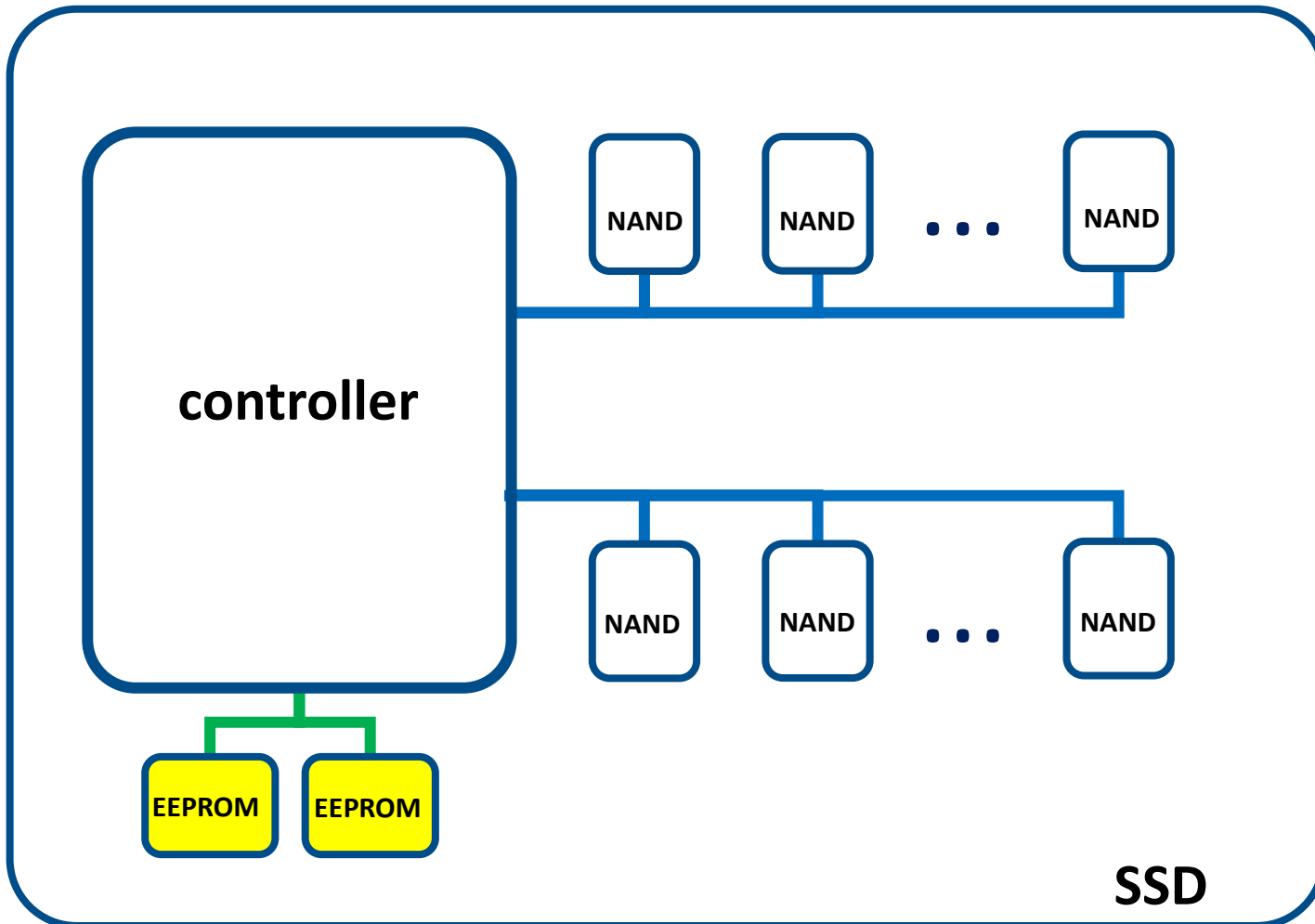
- DNA media does not share properties found in other storage media types
  - No built-in controller, or linear addressing of physical storage regions
  - No structured media topology
  - No built-in facilities for addressing specific parts of the media
  - Addresses (sectors) need to be encoded for later reading
- Multiple mechanisms (CODECs) exist for encoding data into DNA
  - CODEC must be discernable from within the media itself in a standard way
- With >100 year lifespan, we must anticipate technology evolution
  - Categories of innovation expected within DNA media and the value chain?
  - What is considered a safe assumption today that may not be one tomorrow?

# What is an archive “boot record”?

- With traditional media, controller knows where sector zero resides, packages device metadata for the consumer
  - Operating system connects to and initializes device for consumption
  - Manages translation of upper layer APIs (e.g. POSIX) into lower layer protocol primitives (e.g. SCSI)
  - Generally governed by an intermediary (e.g. filesystem)
- No controller within DNA media, no linear addressing within the media, and no file system

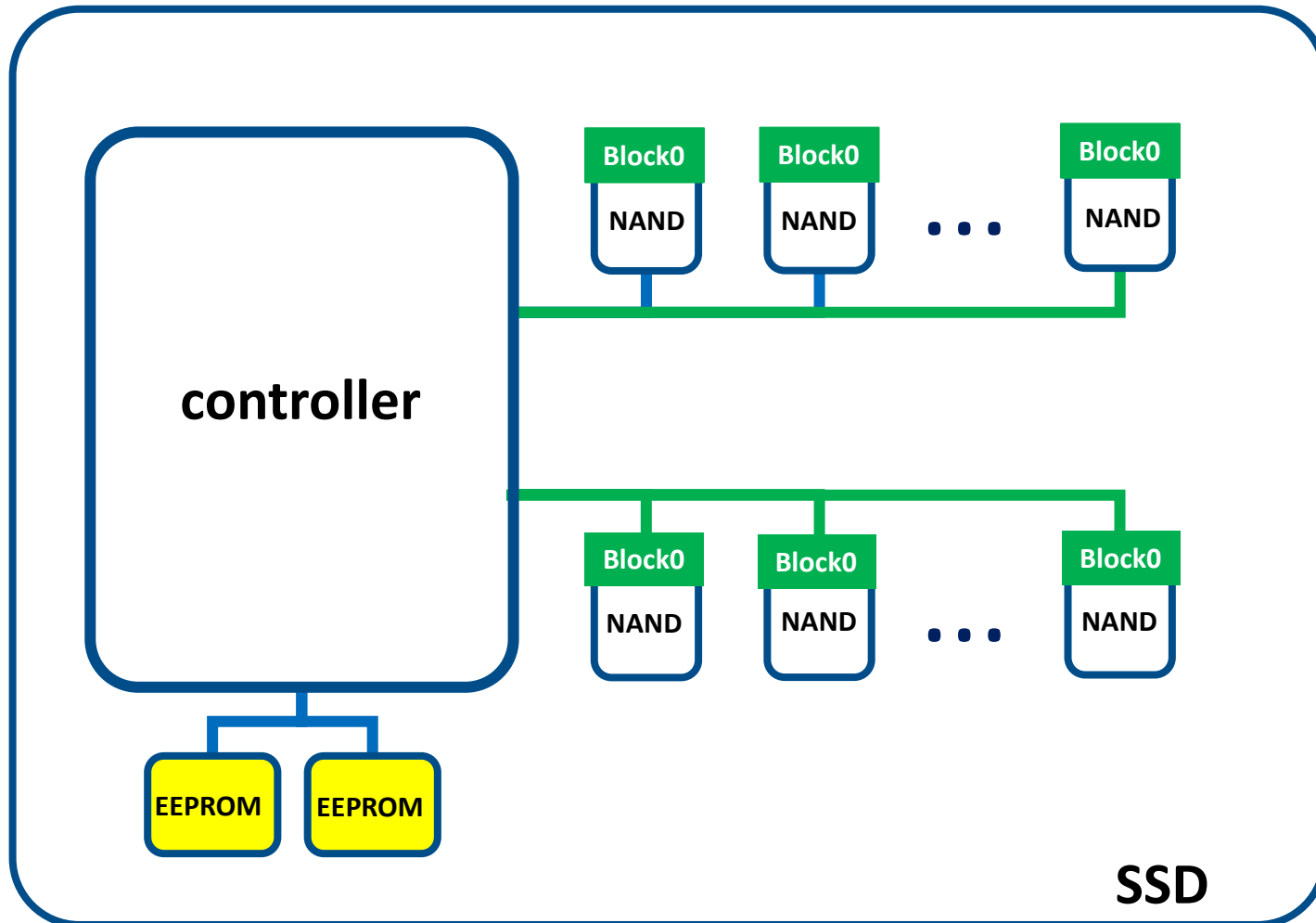


# Current State – Initializing an SSD



- Controller first reads information on E2PROM about HW configuration (type of NAND, timings, vendor ID, channel addressing, type of ECC used to load FW)
- Data read from E2PROM is protected by ECC to ensure reliability

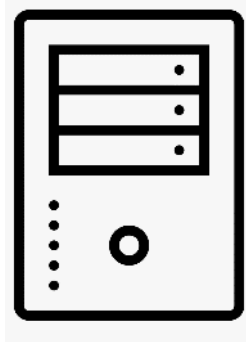
# Current State – Initializing an SSD



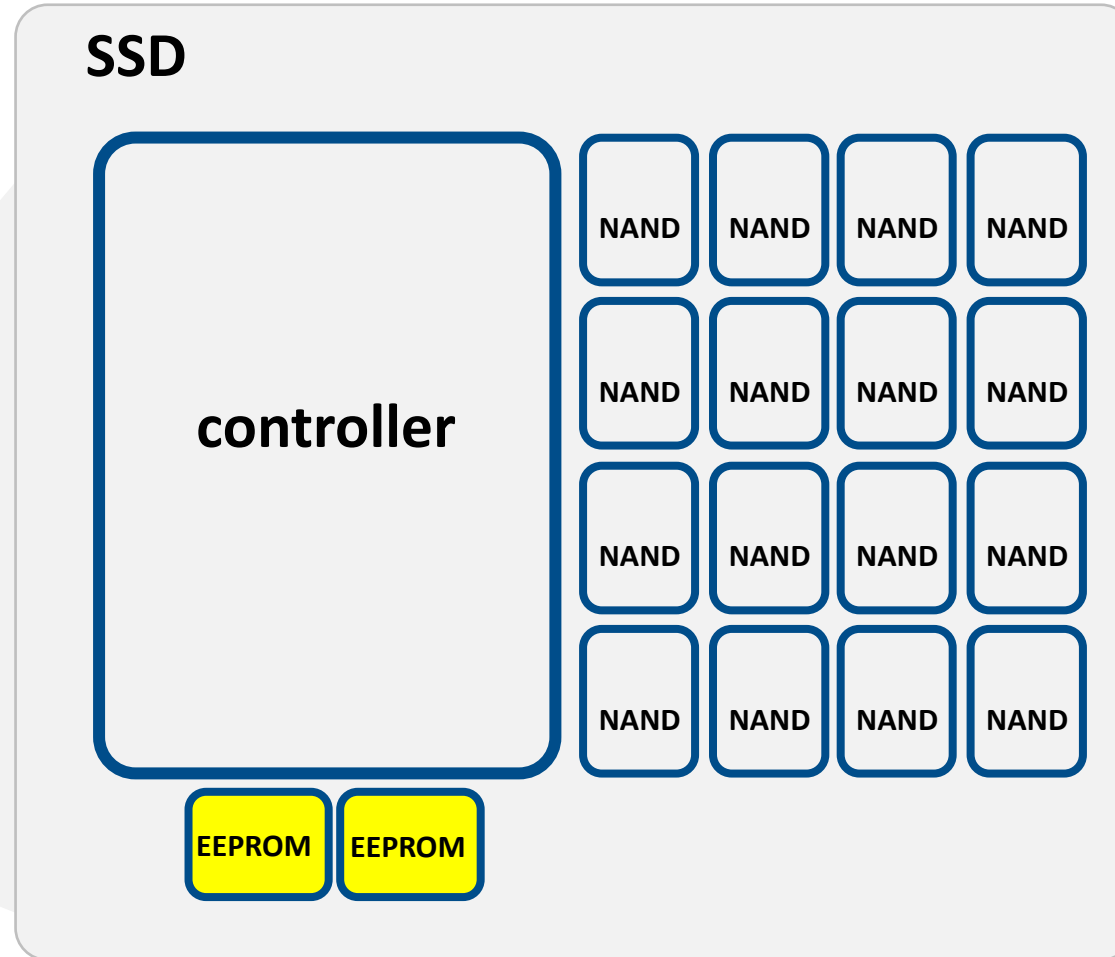
- Using previously read information, controller is able to read NANDs
- By reading block0 of NAND devices, controller loads the firmware
- Block0 is guaranteed good by NAND vendors for this purpose



# Booting a Machine from an SSD



File System
Sector 0 (MBR)
Sector 1
...
Sector n



# Booting a DNA Data Storage Archive



- Without a controller how can we read the archive?
- Where can we discover metadata such as vendor ID, CODEC used in the archive?
- This metadata is contained in the archive itself, but we need a way to discriminate it from other data

# Rosetta Stone Project (1/3)

- Part of DNA Data Storage Alliance
- Goals:
  - Agree on a common identifier format for universally bootstrapping any DNA Archive
  - Enable identification of the CODEC used to encode an archive, from within the archive
  - Enable innovation in DNA CODECs for the main archive by enabling a standard for discovering the CODEC that was used
  - Provide fast access to archive metadata

DNA Archive Rosetta Stone (DARS)



# Rosetta Stone Project (2/3)

## ■ Working Assumptions

- A generally-available specification document is accessible
- Archive boot record is built using natural DNA bases (ACTG)...
- ...but the archive may contain non-natural DNA bases
- Standard means of identifying the CODEC used within the archive is needed
- We assume a reader will have some form of Internet connectivity
- DNA will primarily be used as a write-once archival medium

# Rosetta Stone Project (3/3)

- Decisions to Make

- Agreement upon length of oligonucleotides
- ...error recovery metrics and mechanisms
- ...how many “sectors” are required

- Progress to Date

- Initial proposals drafted and discussed
- Covering sector zero implementation, identification
- Outlining payload contents and their meaning
- Discussions and tests around error modeling and recoverability

- Roadmap

- Reviewing future proposals
- Creation of and maintenance of a specification of a standard
- Build policy and procedure documentation
- External registry of CODECs

# Future State

- Rosetta Stone sets the stage for controllers, drivers, and ecosystem
  - Agreeing on decoding standard enables vendors to work on consumers of sector zero
  - Standard controller functions for management (e.g. SMART) may come from our error models
- Address space governance
  - CODEC ID / address issuance may work similar to IP addresses
  - DNA Data Storage Alliance could operate similar to ICANN
- Technology will always evolve
  - CODECs, address space will form part of the Alliance's industry roadmap
  - Working assumptions based on current technology
    - Advances may lead to review of assumptions, error model, number of codecs etc.
  - Synergy with other SNIA storage technologies i.e. computational
    - Exposing novel CODEC capabilities enabled by the DNA medium

# How to Participate

# How to Participate

- Standards only succeed when they consider and support the needs of a broad base of constituents with an eye toward the future
- Our working group is growing and diverse, and looking to
  - Increase representation from both public and private sector
  - Increase representation from a variety of markets and domains

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# Summary

# Summary

- DNA data storage media provides the promise of density, durability, and cost effectiveness to meet the challenges of data growth, retention, compliance, and climate change
- Writing data to DNA involves coding, synthesis, and storage, and conversely, reading data from DNA involves retrieval, sequencing, and decoding
- DNA as a storage media does not share properties found in other storage media types, e.g. no built-in controller, or linear addressing of physical storage regions
- Rosetta Stone aims to ensure that a DNA archive can be consumed in a consistent manner by making discoverable the structure and encoding of the archive

# Thank You

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