SNIA DEVELOPER CONFERENCE



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DNA Data Storage An End-To-End System Concept

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From Lab to Datacenter Use Case

Abstract Concept of a DNA Data Storage System

- Defining an Interoperability Standard
- Logical interfaces
- System Operations
- Type of DSS Racks

System Implementation

- Strategic Enablers
- Integration in Datacenter
- System Performance
- Current Status



DNA in the Datacenter

From here





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Datacenter Use Cases









Concept of a DNA Data Storage System

Defining an Interoperability Standard



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End-To-End DNA Data Storage System (DDSS)

DNA Data Storage System (DDSS)

Sub-Blocks: Encoder, Synthesizer, Storage & Retrieval, Sequencer, Decoder





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End-To-End System : Orchestrator



- Main brain controlling the DDSS Setup & Functional Phases
- Channel through which all writes & reads are issued and read-data returned
- Tracks & logs the status and error feedback from each DDSS Sub-block. Logging used for decision making - Retry / Alarm decisions
- Monitors Accumulated Error at the System Level



Defining an Interoperability Standard: Phases of Operation

- 1. System Assembly Phase
- 2. Setup / Initialization Phase



Logical Interfaces - Functional Interface for "Synthesizer"



Input Interface

Data	Description	Source - Destination
Encoded Header	Name, Identification of customer, Metadata encoded into ATCG format as per CODEC (Primers included)	Encoder -> Synth
Encoded Addressing	Address encoded into ATCG format as per CODEC (Primers included)	Encoder \rightarrow Synth
Encoded Customer Data	Customer Data encoded into ATCG (DNA) format as per CODEC (Primers included)	Encoder $ ightarrow$ Synth
Retry Signal	Orchestrator may issue a Retry signal in case of irreparable Error received from Synthesizer or Storage blocks	Orchestrator -> Synth



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Logical Interfaces - Functional Interface for "Synthesizer"



Output Interface

Output	Description	Source - Destination
DNA Sequences/Strands	Sequences containing Address, Metadata and Customer Data	Synth -> Storage
Status metrics	Cycle Done, Key metrics	Synth -> Orchestrator
Flag: Est. Error Rate > Spec (optional)	Error flag that fires when the actual error rate during synthesis is higher than the Specification Error Rate of the synthesizer	Synth -> Orchestrator
Flag: Strand Length (optional)	Strand Length check using Bioanalyzer or Qubit post synthesis - for sample	Synth -> Orchestrator
Flag - Generic	General Error flag for issue in flow/control, reagent levels that may be detected via monitoring.	Synth -> Orchestrator
Quality Check (optional)	Quality Check can be executed using Sequencer, Bioanalyzer, or Qubit	Synth -> Orchestrator
Reagent Levels	Report Current Reagent Levels in terms of Number of Runs possible	Synth -> orchestrator



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Defining an Interoperability Standard: System Operations

- Write / Read
- Modify
- Erase / Delete
 - Ability to delete part / all the previously written data. Implemented as:
 - 'Erase' at container granularity
 - 'Erase' partial data from a container
- Self-Check (optional)
 - During system IDLE, a known golden data file is written & read and checked for error rate.

Join us to work on the standards @SNIA: <u>www.snia.org/groups/snia-dna-technology-affiliate dnastoragealliance.org</u> @email: <u>info@dnastoragealliance.org</u>







DNA Data Storage Racks

Defining an Interoperability Standard



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Type of DSS Racks

Traditional Rack

All 5 sub-components of the DDSS, along with the power, plumbing and management accessories needed, fit into a traditional Rack by specification

'N' Racks act as Single Rack Unit

The 5 sub-components of the DDSS may occupy N non-identical racks of standard power, weight and spatial specs. This set of racks acts as a single unit and racks deployed only in multiples of N.

Disaggregated Rack Unit

The 5 sub-components of the DDSS occupy N non-identical racks. The system does not need to be deployed in multiples of N racks. A sub-portion of this Rack Unit can be scaled as required. Minimum N Racks footprint needed







System Implementation

Example of the Biomemory DNA Data Storage appliance



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Pioneering DNA in Data Centers









✓ Store data in DNA for 150+ years @
 room temp, with no energy consumption

✓ Suited for Data Centers

- Cost-effective
- ✤ Space-efficient
- Reliable
- Scalable
- ✤ Upgradable
- ✤ Resilient
- ✤ Interoperable
- Low-Power

Disclaimer:

This presentation is for informational purpose only. It is meant to provide a preview of Biomemory's products, architecture, features and targeted performance, as an example of implementation leveraging the recommendations from the DNA Data Storage Alliance.

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Strategic enablers



- Low-cost biosourced mass-production of nonhazardous biosafe DNA blocks used as consumables (low customer opex)
- < 1000 times lower cost than life-science oligonucleotide synthesis technologies
- Simple assembly of a small number of DNA blocks, compatible with high-speed and massive parallelization.
- Target > 1 billion reactions in parallel
- Low error-rate Write and Read processes, with efficient tolerance to errors (strong ECC)

90% error free process, with 100% error detection and recovery

Biotech adaptation to the system (not the contrary)



 Modular software-driven system architecture aligned to Data Center requirements, with dedicated OAM channel

Compatible with existing Data Centers operations

Industrialization and scalability, with focus on lowest risk, high-reliability and minimum custom hardware development

80% reuse and adaptation of industry-available components.

 Aligned to SNIA DNA Data Storage Alliance specifications and recommendations

Interoperability, portability, reliability



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Integration in Data Centers







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System Implementation



Hardware & Fluidics

 80+ % available or adapted from commercially-available components

Software / Firmware

 ~ 80 % to be developed or adapted



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Biotech

- ~ 80 % available in Biomemory lab, ready to be industrialized and scaled up.
- Continuous roadmap to adapt and optimize the biotech components for Enterprise-grade Data Storage



Clear path towards high-performance





September 2024 status



Miniaturization (Gen 1)

- DNA Assembly (Writing) with volume of 1nL.
- Validation = Writing / PCR / Reading cycle
- Enabling writing batches of ~1 Million sites (64 MB/batch)
- Moving towards 50 pL for ~10M sites/batch (640 MB/batch)

Reaction Acceleration (Gen1)

- Currently 1.3 day / batch
- Starting acceleration towards 3 batches/day (> 2GB/day by end of 2025)

System Automation (Gen 1)

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- Biowriter system design completed
- Bioreader solutions under evaluation







Going beyond cold storage





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Back-up, cloning and war chest raw data, pending AI readiness/analytics

- **Cloud Storage Services**
- Factory raw data for AI training
- Data Lakes for AI and Big Data
- IoT & rich media data
- Genomics data sets



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