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Title: Comments on Draft Text of ISO/IEC DIS 23092-1 Transport and Storage of

Genomic Information

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1 Introduction

At the 121st MPEG meeting, a Draft DIS of MPEG-G part 1 was produced [1]. This contribution provides a few comments on that document and an additional related comment.

2 Comments

- In Table 4: clarify that block is not a gen_info element (maybe with a different font).
- In Table 12: change "gen info" with "lbll".
- Issue with dscn: currently, it is not possible to know where the blocks end, as there might be a protection box (the sentence saying that blocks' size equal to dscn size dshd size has to be changed). The issue can be corrected by changing the position of all boxes to be prior to the blocks, or adding information on the total size of the blocks.
- Random access / representation of the file structure in memory would be simplified if Access unit header had always information about the sequence the information is mapped to (i.e. also when MIT_flag == 1). This would remove the need to test the AU_byte_offset to determine which AU is being indexed.
- In last sentence of 6.5.2.1 introduction: the special value is not only used for block_byte_offset, but also to indicate that the access unit is empty (see, for example, AU_START_POSITION). Maybe consider not having empty AUs.
- Issue with the use of "Descriptor Id" (for example in Dataset header or MIT): desc_ID as used in Dataset Header refers to the index in the list of available descriptors, not to the actual value of the descriptor id. Thid could be clarified by changing to desc_index, for example. Similar issue in MIT: desc_ID refers to the index in the list of available descriptors, but could be understood as the ID of the descriptor identifier as specified in Table 10 of ISO/IEC 23092-2.
- Stream container: remove metadata.
- Descriptor stream header: remove protection flag, which is outdated (it is a legacy from the GENIFF proposal).
- Table 9- Reference box syntax defines seq_count to be of type 32 bits, but in table 17 the seq_count is of type 16 bits: a dataset header could not use all the defined sequences. Consider which size is more appropriate and update all occurences.

3 Comments related to other parts

There are multiple names used for the concept of the access unit type (au_type, class id, class type). Consider which name is prefered and update all occurences.

- Part 1 defines the position in the genome to be 32 bits (Table 29 au_*_position, extended_AU_*_position, and same in auhd), but the mpegg_record defined in table 64 of part 2 defines the mapping position to be of type 64 bits. Consider which size is more appropriate and update all occurrences.

4 References

[1] ISO/IEC JTC 1/SC 29/WG 11, N17320 "Draft Text of ISO/IEC DIS 23092-1 Transport and Storage of Genomic Information" Gwangju, January 2018