# MPEG-G: The Emerging Standard for Genomic Data

Jan Voges • Jörn Ostermann {voges,office}@tnt.uni-hannover.de



## Performance

### Overall MPEG-G performance on human high-coverage WGS data

### Impact of MPEG-G quasi-lossless quality value compression on variant calling





#### Data:

- whole genome from individual NA12878
- re-sequenced with Illumina HiSeq @ 53x

- customized variant calling pipeline following the "GATK Best Practices" [2]
- comparison of the called variants to the NIST consensus set

#### Data:

- chromosomes 11 and 20 from individual NA12878
- re-sequenced with Illumina HiSeq 2000 @ 14x

# Standardization Workplan

October 2017	Committee Draft
July 2018	Draft International Standard (DIS)
January 2019	Final Draft International Standard (FDIS
International Standard (IS)	



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#### Leibniz Universität Hannover

**Institut für Informationsverarbeitung** Appelstr. 9A, 30167 Hannover, Germany

#### http://www.tnt.uni-hannover.de

