SiM



Standardizing Al Annotations The DICOM Way

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- Editor of the DICOM Standard (NEMA Contract)
- Owner of PixelMed Publishing, LLC
- Author of book on DICOM Structured Reporting
- Consulting for GE, Carestream, MDDX (Bioclinica), Curemetrix, HCTS, Hologic
- Supported by NIH U24CA180918 QIICR, NCI Leidos BOA 29XS219 Task Order #05



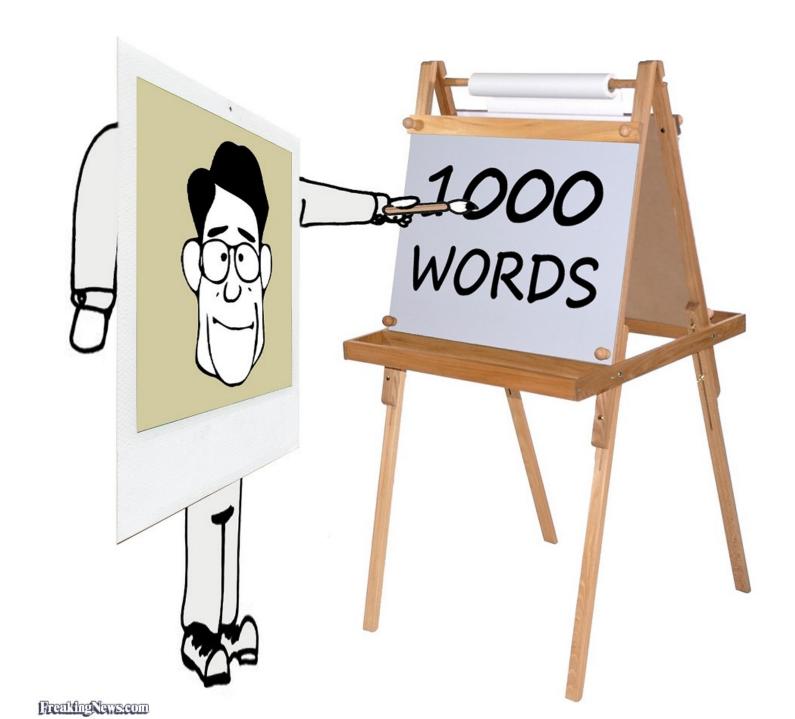
Outline

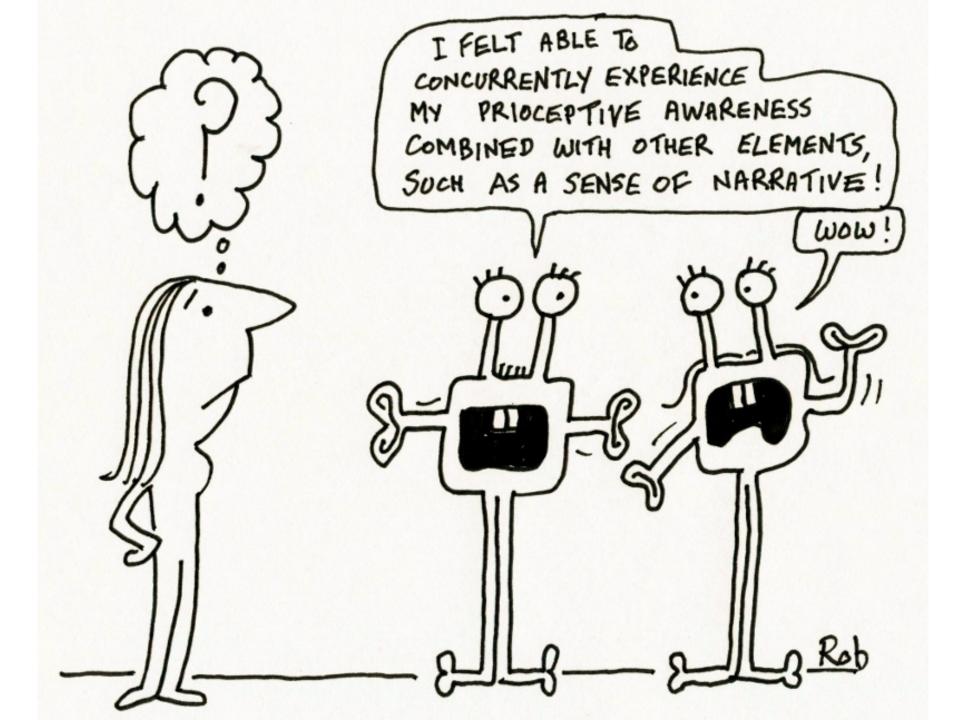


Background

- radiologists and annotations
- Annotation use cases
 - In general and for AI
 - definition and different types of annotation
- DICOM Encoding
- Workflow and DICOM
- Beyond radiology
- Gaps and future improvements









A Picture Is Worth A Thousand Words:

Needs Assessment for Multimedia Radiology Reports in a Large Tertiary Care Medical Center

Lina Nayak, MD, Christopher F. Beaulieu, MD, PhD, Daniel L. Rubin, MD, MS, Jafi A. Lipson, MD

Rationale and Objectives: Radiology reports are the major, and often only, means of communication between radiologists and their referring clinicians. The purposes of this study are to identify referring physicians' preferences about radiology reports and to quantify their perceived value of multimedia reports (with embedded images) compared with narrative text reports.

Materials and Methods: We contacted 1800 attending physicians from a range of specialties at large tertiary care medical center via e-mail and a hospital newsletter linking to a 24-question electronic survey between July and November 2012. One hundred sixty physicians responded, yielding a response rate of 8.9%. Survey results were analyzed using Statistical Analysis Software (SAS Institute Inc, Cary, NC).

Results: Of the 160 referring physicians respondents, 142 (89%) indicated a general interest in reports with embedded images and completed the remainder of the survey questions. Of 142 respondents, 103 (73%) agreed or strongly agreed that reports with embedded images could improve the quality of interactions with radiologists; 129 respondents (91%) agreed or strongly agreed that having access to significant images enhances understanding of a text-based report; 110 respondents (77%) agreed or strongly agreed that multimedia reports would significantly improve referring physician satisfaction; and 85 respondents (60%) felt strongly or very strongly that multimedia reports would significantly improve patient care and outcomes.

Conclusions: Creating accessible, readable, and automatic multimedia reports should be a high priority to enhance the practice and satisfaction of referring physicians, improve patient care, and emphasize the critical role radiology plays in current medical care.

Key Words: Multimedia reports; radiology reporting; digital images; communication; radiology practice.

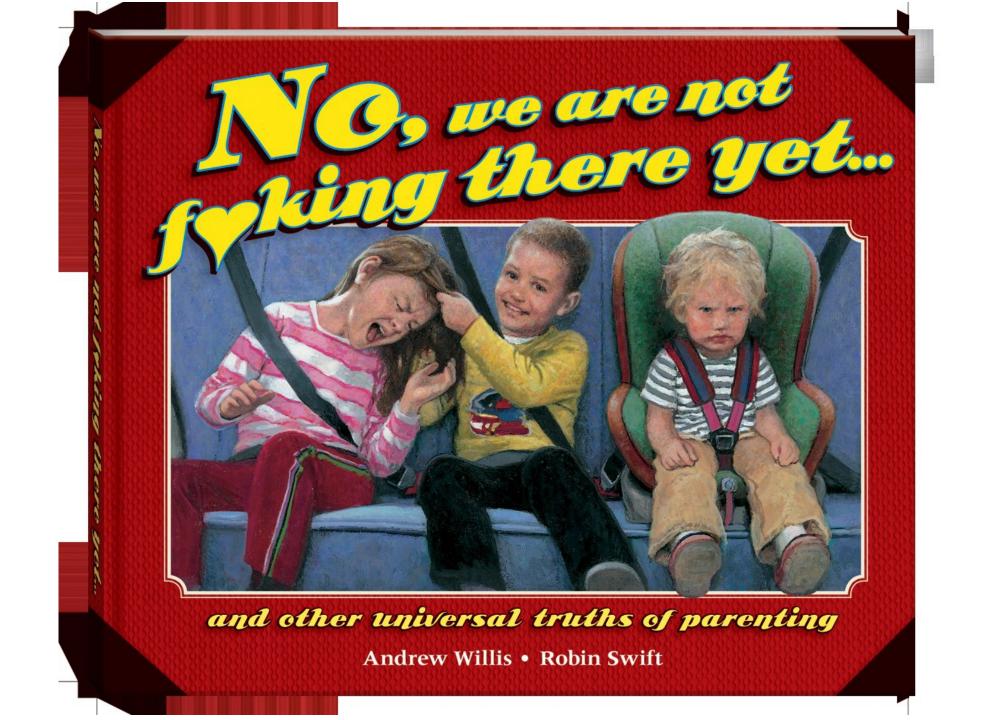
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Structured Radiology Reporting: Are We There Yet?¹

Curtis P. Langlotz, MD, PhD

G iven the prominent role that information technology will play in the future of health care delivery, the potential benefits of structured reporting systems now seem more relevant than ever. These systems may lead to rapid

cohort design. The same 25 brain magnetic resonance (MR) imaging cases were reviewed in two distinct phases by two separate groups of residents: a control group and an intervention group. The MR imaging cases contained a representative



HOW DO WE GET THERE FROM HERE?

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Use Cases – Non-Al or Al, Use versus Re-Use

Clinical Use Cases

- Non-Al
 - communicate rad-tech, rad-rad, radiologist-clinician, multidisciplinary team meetings
 - preserve state for priors (where, what, what size, ...)
 - legal record
- AI
 - automated result output prior to human read (traditional mammo CADe, triage, priority)
 - human defined selection for targeted AI

Re-use Use Cases

- Non-Al
 - retrospective and prospective research
 - education
- AI
 - Non-Al annotations used for Al training/testing
 - Al annotations used for a different Al application training/testing



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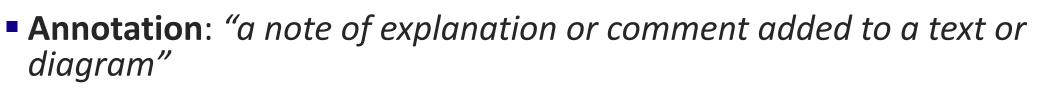
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Use Cases – What is an "Annotation" anyway?



- Label: "a classifying phrase or name applied to a person or thing [especially one that is inaccurate or restrictive ⁽²⁾]"
- <u>NOT</u> Markup: "a set of tags assigned to elements of a text to indicate their relation to the rest of the text or dictate how they should be displayed"

distinguish semantics from visual representation "meaning" for machines not just humans



- Patient/Case
- Imaging Study
- Series/Acquisition
- Image
- Frame (pixel data array at one place in space/time/...)
- Region ("of interest" ROI)
- Single point (label each/every voxel/pixel)



Use Cases – Qualitative, Quantitative

Qualitative

- categorical shape = round, ...
- Ordinal
 - roundness scale of 1 to 5

Quantitative

- morphology size (diameter, volume, ...)
- quantity signal intensity, attenuation coefficient, ...
- numerical features entropy of GLCM, fractal dimension, ...
- on transformed variants registered, resampled, filtered, wavelet, ...
- units absolute and relative (to what reference region, population)
- derivation mean, max, …
- method model, fitting, sampling, binning, ...



Use Cases – Concept Representation

Single concept

e.g., "round shape"; "42"

Name-value pair

e.g., "shape" = "round (generic)", "round shape"; "ultimate question" = "42"

Coded versus text

- SRT:M-02100 (SCT:42700002) v. languages: "qaab wareegsan", "圆形"
- synonyms ("round" v. "circular"), case, punctuation ("Shape, round"), ...

Same concepts in different classifications

- SCT:42700002, NCIt:C48348, RadLex:RID5799
- Metathesaurus (mapping): UMLS:C0332490



Use Cases – Localization Representation

- Patient/Case Patient/Specimen Identifier
- Study, Series, Image Unique Identifiers (UID, UUID)
- Frame UID + frame number/offset

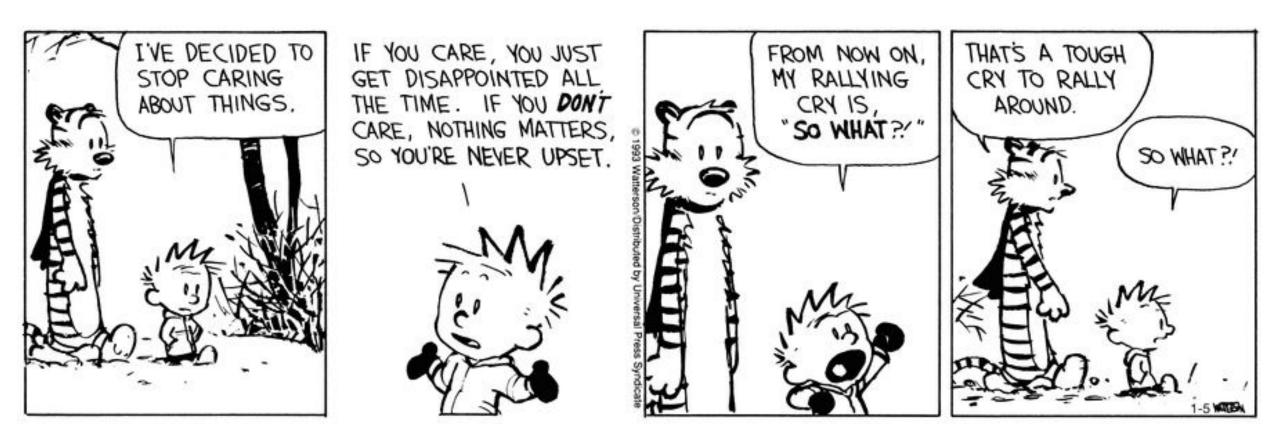
ROI

- contours image (2D) or space (patient, 3D) relative
- temporal coordinates (frame #, relative/absolute time)
- segmentations (bitmap, partial occupancy, probability)
- Single point (each/every voxel/pixel)
 - parametric maps (bitmap, scaled integer, floating point)
- Transformations
 - rigid (affine), non-rigid (deformation field, spline, ...)





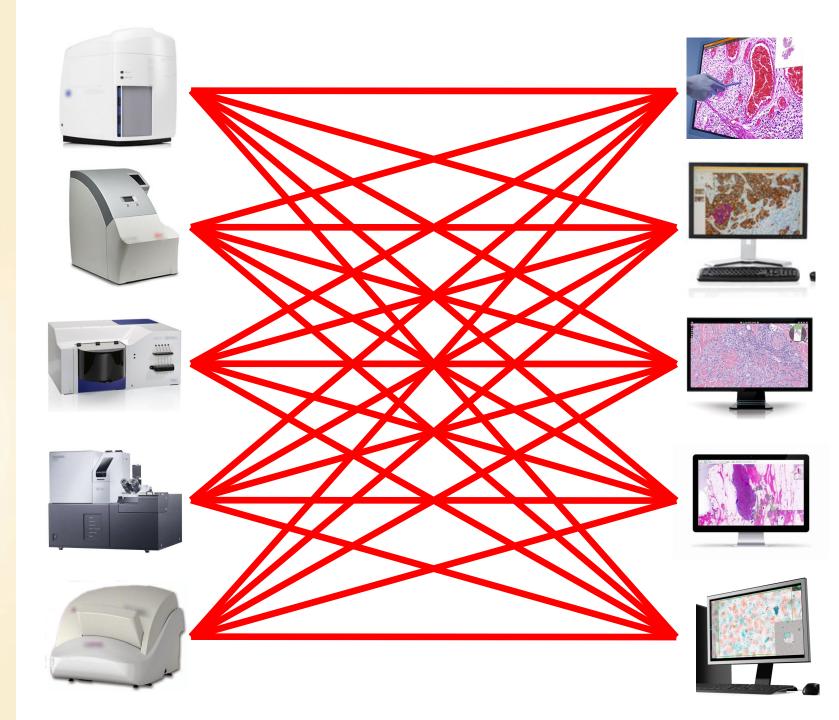
WHO CARES?



JOHN PALFREY AND URS GASSER

Interop The PROMISE and PERILS of HIGHLY INTERCONNECTED SYSTEMS

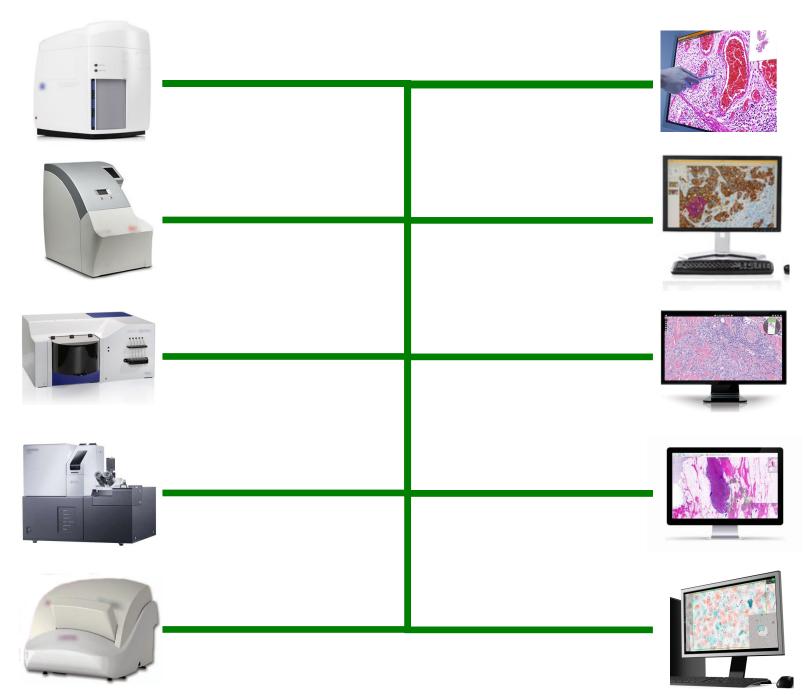






Digital Imaging and Communications in Medicine







DICOM "in"

- obviously all (radiology) images start out as DICOM
 - can be rearranged from multiple single files to single multi-frame DICOM file (happier algorithm developers)
- "metadata" in DICOM "headers" are "annotations" too
 - e.g., Series Description = "T1 axial post-Gd", B value = 1000
 - can be better structured/coded (retrospectively), e.g., Acquisition Contrast = T1
 - phase to cleanup/canonicalize multi-site data DICOM attributes
 - do not discard known-safe vendor private data elements during deidentification – may be useful for unanticipated re-use cases



Where DICOM fits in

DICOM "out"

- All types of annotation (including processed images) can be shared as DICOM
- Pros:
 - allows re-use of clinical imaging/annotation systems
 - can be stored/shared/indexed in off-the-shelf DICOM archives (e.g., TCIA)
 - can be created/viewed/analyzed by (some) OTS DICOM tools, viewers, ...
 - not just a bunch of poorly labeled/organized files on somebody's disk
 - self-describing/identifying contains (pseudonymous) identifiers
- Cons:
 - requires more attention to de-identification (if clinical origin)
 - requires use of DICOM toolkits/libraries to create/access
 - requires more attention to preservation/propagation of "composite context" (identities and UIDs) in processing pipeline (e.g., to restore identifiers from images to results)
 - more complex and arcane than making it up as you go along



Gonfessions of a Content Greator: Don't Care About Data



UNTHINKABLE.FM



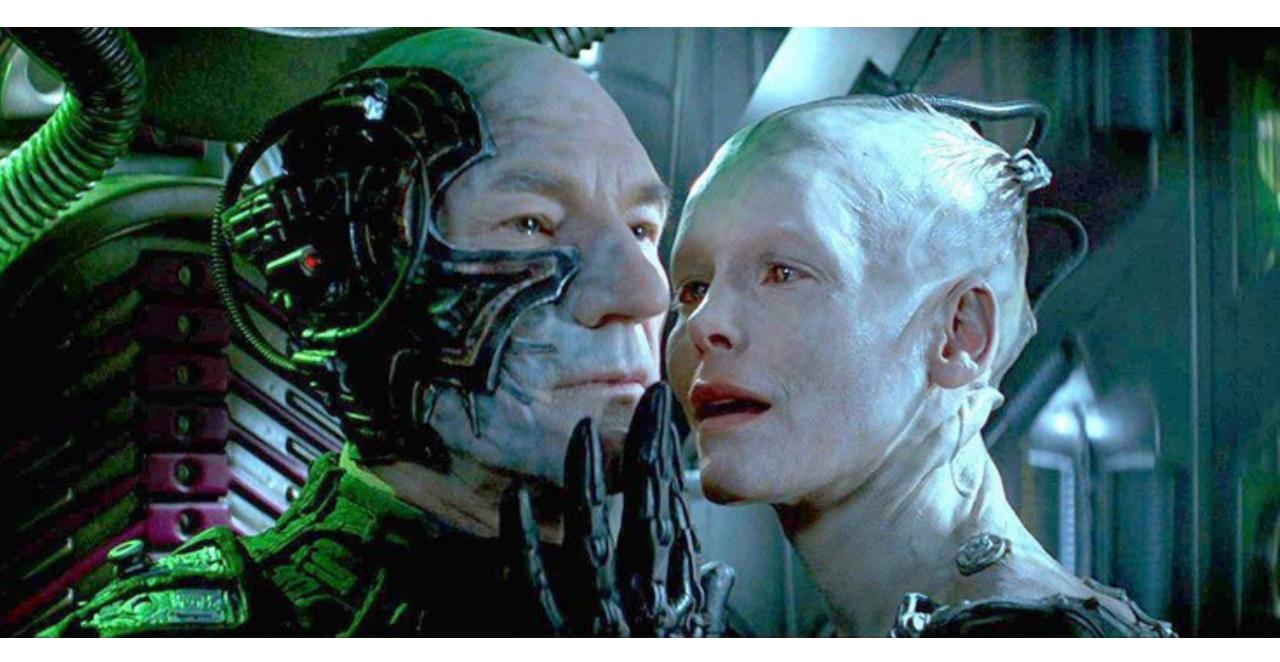
AI CHANGES THE GAME

Annotation (interoperability) matters now

Previously:

- little incentive to annotate
- few tools to create or view annotations
- annotation interoperability was a low priority for product managers
- presentation rather than semantics were the priority for annotation tools
- Now:
 - semantic annotations have (real monetary) value beyond primary use case
 - recognition of existence of unanticipated re-use cases
 - annotations are expensive to create/recreate retrospectively
 - more expensive to process if proprietary rather than OTS standard
 - Al-generated annotations need to be interoperable for display
 - "interactive" AI requires interoperable annotation exchange
 - AI vendors unlikely to be the same as scanner/PACS vendors mix and match





Bad ways in DICOM, historically:

- Burned in graphics and text
 - including screen shots
- Overlay graphics and text
 - in pixel data, header or separate object
- Presentation states
 - still only graphics and text (no semantics)
 - currently very popular in clinical PACS





Good ways in DICOM – standardized for ages, variable use

- Structured Reports (SR)
 - tree of codes, numbers, 2D, 3D & temporal coordinates, references, ...
 - basis for Ultrasound, Cardiovascular, Mammo CAD, radiation dose
 - Key Object Selection (KOS) flags key images with text/coded label
- Segmentation (SEG or DSO)
 - rasterized bitmap, probability, occupancy; coded property/anatomy; ROI, atlases (i.e., pixel level categorical annotation), ...
 - surface mesh (rarely used)
- Radiotherapy Structure Sets (RTSS)
 - 3D coordinates, some component semantics, few quantities (volume)
 - widely used in RT planning and re-used in workstations, e.g., for PET



Relatively new things in DICOM

- Real World Value Maps
 - coded way to describe voxel values (beyond Rescale Type)
 - retrofitted to all existing DICOM images
 - form of "annotation" that makes pixel values semantically meaningful
- Parametric Maps
 - RWVM combined with floating point or scaled integer pixels
- Second-generation Radiotherapy annotations
 - Conceptual Volumes "grammar" for combining contours, segmentations



DICOM and Annotations – Then to Now

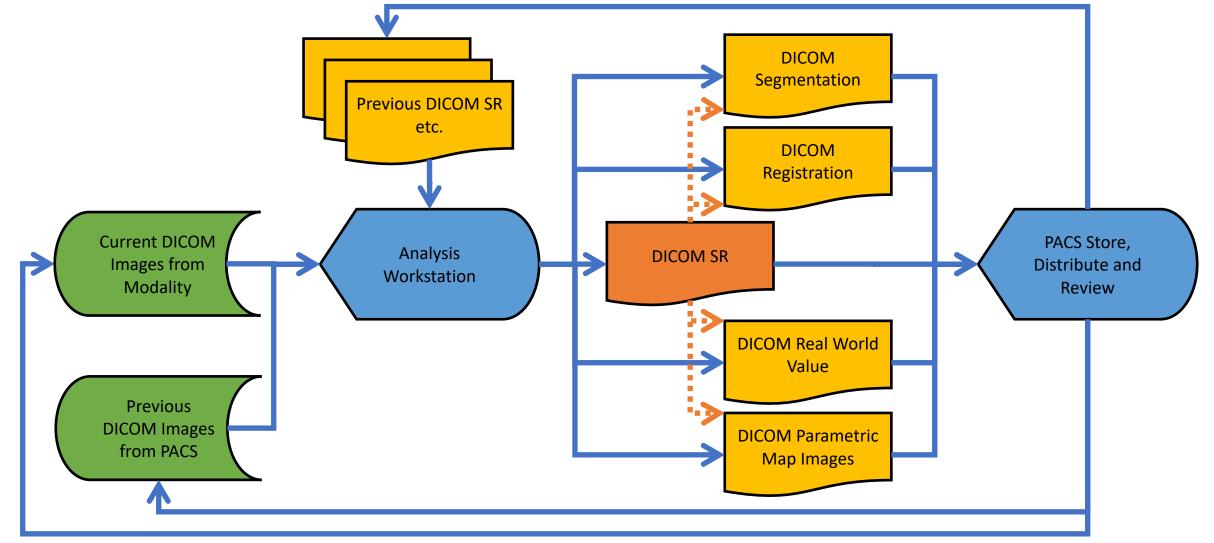
Related DICOM IODs

- Fiducials
 - markers with shape and location
- Registration
 - rigid
 - deformable
 - well-known frames of reference (e.g., atlases)



Putting it all together ...

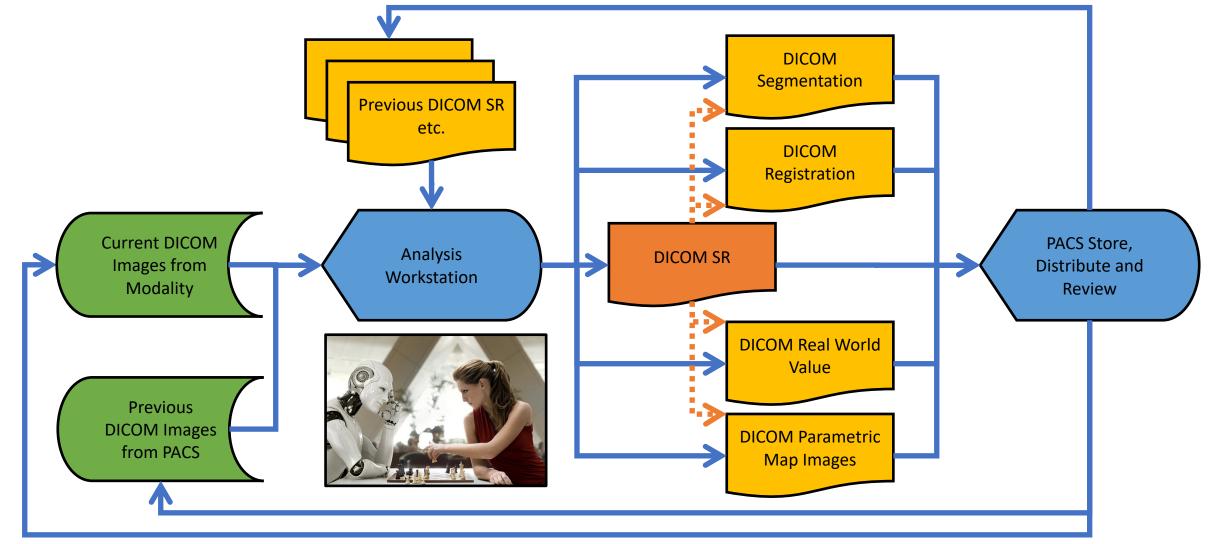






Putting it all together ...











This illustration is inspired by and in part derived from the work by Scott Simmerman, "The Square Wheels Guy" http://www.performancemanagementcompany.com/

Too busy? – No problem, we supply tools

- Store and regurgitate
 - OTS commercial and open source DICOM archives as well as PACS
 - most modern systems will handle any new or old DICOM object
 - unlike ancient PACS, which rejected things they could not view
 - this increasing flexibility driven by "vendor neutral archive", "universal viewer" and "deconstructed PACS" phenomena
- DICOM annotation objects are just like images
 - some actually are "images" (segmentations, parametric maps)
 - non-images all share common "composite context" encoding (e.g., patient/study/series/equipment identification/description)



Too busy? – No problem, we supply tools

Creation

- Iong history of ultrasound, mammo CAD, radiation dose, key object SR authoring
- increasing SEG and SR support in toolkits (esp. open source)
- extraction/propagation/merging of composite context
- merging descriptions of metadata created in XML or JSON into DICOM object content (included coded descriptions) – easier than hand-coding in programming language
- Transcoding
 - of other formats, proprietary, academic, and project-specific
 - other structured annotation formats into DICOM SR (e.g, AIM via PS3.21 mapping)
 - segmentations and label maps in other formats into DICOM SEG
 - parametric maps in other formats into DICOM Parametric Maps
 - single frame DICOM image sets into multi-frame single DICOM files



Too busy? – No problem, we supply tools



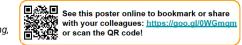
- annotation objects can be queried/retrieved using traditional DICOM network protocols widely supported by many toolkits
- DICOMweb (WADO-RS, QIDO-RS, ...) protocols increasingly supported by archives and are just like any other HTTP request (can use curl, postman, as well as DICOM toolkit utilities)
- DICOMweb payloads (XML and JSON metadata) and separate bulk data access increase accessibility and simplify parsing
- Viewing and Analysis
 - toolkit support for extracting metadata and structured content into generic XML, JSON, CSV
 - more viewers support display of DICOM SEG superimposed on underlying reference images
 - some viewers can display content of DICOM SR objects rendered into hierarchical plain text, HTML or PDF – few (if any) can "tabulate" content into user friendly form
 - voice dictation systems have long been able to ingest DICOM SRs, extract content and make available "merge fields" in dictation templates



Interoperable communication of quantitative image analysis results using DICOM standard



Andrey Fedorov¹, Daniel Rubin², Javashree Kalpathy-Cramer³, Justin Kirby⁴, David Clunie⁵, Michael Onken⁶, David Flade⁷, Pattanasak Mongkolwat⁸, Rajesh Venkateraman⁹, Jan Bertling¹⁰, Steve Pieper¹¹, Ron Kikinis¹ ¹Brigham and Women's Hospital. ²Stanford University, ³Massachusetts General Hospital. ⁴NCI Fredrick, ⁵PixelMed Publishina. ⁶OpenConnections GmbH, ⁷Brainlab, ⁸Mahidol University, ⁹Eigen Medical, ¹⁰Hermes Medical, ¹¹Isomics Inc



Introduction

As quantitative imaging (QI) is gaining momentum in research and commercial platforms, it becomes important to support its usage scenarios:

- Clinical workflows: storage of the analysis results on PACS alongside the imaging data; longitudinal followup of the patient with quantitative imaging across workstations.
- Research workflows: validation of imaging biomarker analysis tools; community repositories of the analysis results; secondary analysis of data.

Various types of derived data important in quantitative imaging research include image annotations (points, distance measurements, contours, labeling of image voxels), parametric maps and numeric results of the quantitative measurements.

Image segmentation is a key preprocessing task concerned with defining a region of interest for subsequent analysis and quantitation. It is therefore of critical importance to support interchange of the segmentation results.

Digital Imaging and Communication in Medicine (DICOM) is the standard used ubiguitously for communicating image data. Although DICOM provides the means to also describe *derived* image-related information, thus far it has found very limited acceptance in the quantitative imaging community.

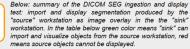
As a step towards improving QI analysis results interoperability, we investigate the use of DICOM Segmentation Storage SOP Class (DICOM SEG) for communicating image segmentation results.

Workstation Support of DICOM SEG

Workstations evaluated (commercial products are in italics):

- 3D Slicer X, http://slicer.org
- ePAD v1.7. http://epad.stanford.edu AIM on ClearCanvas v4.0.6.4, http://www ict.mahidol.ac.th/research/Imaging-
- Informatics
- Brainlab PDM 2.2
- Siemens syngo.via VA30A HF06







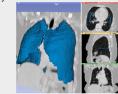
Disclaimer: the workstations evaluated do not necessarily represent the complete list of workstations that support DICOM SEG, although we attempted to contact a number of major vendors.

Image Segmentation and Quantitative Imaging

Image segmentation is concerned with labeling areas of the image into distinctive regions. These regions can correspond to pathology areas, organs, or identify regions of general interest, to support for example

- volumetric assessment of tumor burden
- quantification of the metabolic of functional activity within the ROI
- quantification of the image properties by means of the radiomics features





Segmentation tools vary in complexity, typically need to be customized to the specific problem, and can be manual, semi-automatic or automatic.



• Know of a workstation/toolkit supporting DICOM SEG but not listed? • Want to learn more or get help adopting DICOM SEG? • Have sample DICOM SEG datasets you would like to contribute? We would LOVE to hear from you! Please email andrey.fedorov@gmail.com

DCMTK - DICOM Toolkit (C++, free open source)

- high-level abstractions for initializing and interacting with SEG (functional groups, image frames)
- attribute-level validation
- helper APIs, propagation of the relevant attributes from the image dataset
- query and retrieval of SEGs with DICOM networking
- used by 3D Slicer

PixelMed Toolkit (Java, free open source)

http://www.dclunie.com/pixelmed/software/

- · classes providing abstractions to support interaction with SEG
- used by ePAD

dicom3tools (C++, free open source)

dciodvfy tool for validating DICOM SEG IOD compliance

DICOM for Image Segmentation Storage

DICOM SEG is the preferred way of communicating segmentations represented as labeled voxels. Some of the important features supported include:

- size efficiency with multi-frame storage and bit encoding
- structured terminology for encoding semantics
- binary and fractional segmentation (e.g., probability maps)
- encoding of the presentation (color)
- multiple voxel occupancy

Being part of the DICOM object "family", integrates with other types of data:

- patient and study composite contexts, frame of reference maintained
- references source image data
- can be referenced from the measurement documents (DICOM SR)



Sample Datasets and Community Adoption

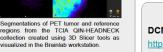
- Sample datasets used in this evaluation are publicly available
- DICOM SEG is the recommended format for communicating segmentation results generated by the teams participating in the NCI Quantitative Imaging Network (QIN).
- Several collections of the NCI Cancer Imaging Archive (TCIA, http: //thecancerimagingarchive.net) contain various segmentation results stored as DICOM SEG objects:
- QIN-HEADNECK: longitudinal PET/CT, head&neck cancer, tumor and lymph node segmentations
- LIDC-IDRI: CT, lung cancer, tumor segmentations obtained using various tools
- NSCLC Radiogenomics: CT, lung cancer, tumor segmentations
- · QIN Lung CT: CT, lung cancer, tumor segmentations obtained with different tools
- QIBA CT-1C: CT phantom segmentatins
- RIDER Lung CT (QIBA CT-1B Round 2)



Acknowledgments

This work was supported in part by the National Institutes of Health through the grants U24 CA180918 (QIICR), U01 CA142555, U01 CA190214 and U01 CA140206.





Support in Developer Toolkits

http://dcmtk.org



Practical implementation example



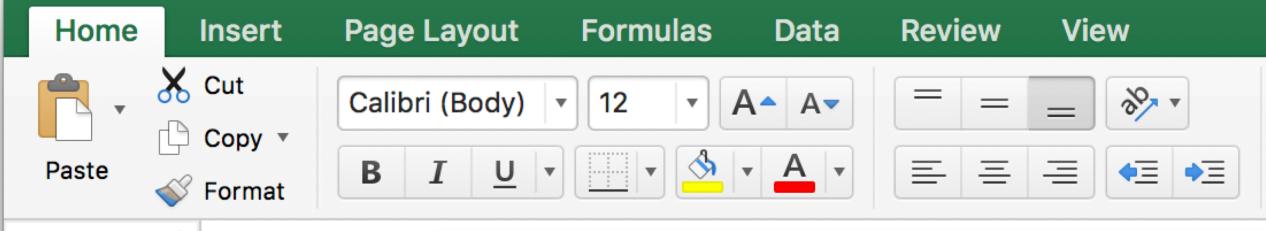
- Crowds Cure Cancer project at RSNA 2017 (http://doi.org/10.7937/K9/TCIA.2018.OW73VLO2)
- Booth and mobile app to find mass and draw longest diameter on liver, kidney, lung and ovarian tumors on TCIA DICOM images
- Sent a CSV file with all annotation coordinates and lengths, and metadata including patient, study, series, instance identifiers of images
- Created two XSLT stylesheets
 - extract cells from CSV table into XML files, one for each row (annotation) driven by column headers
 - convert extracted row cells into DICOM SR instance of DICOM TID 1500 Measurement Report encoded as PixelMed toolkit specific XML format
- Applied existing open source PixelMed toolkit XML to DICOM SR converter
- Validated DICOM SRs that had been created
 - ran automated DICOM SR validator in PixelMed toolkit (knows about TID 1500 since QIICR project)
 - visually inspected dumps of DICOM SR content with PixelMed and dicom3tools toolkit utilities and rendered SR coordinates and measurements on images using PixelMed tool
- One morning's work (leveraging familiar toolkits and techniques and stylesheets used from other projects as a starting point)
- Plan is to to put back into TCIA along with the already publicly accessible images



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	4 Renal	1.3.6.1.4.1.1 TO	GA-B8-55	1.3.6.1.4	4.1.1	41.721506	5 178.25374	6 319.352647	218.14985	294.801199	actual_gi	raff 5b6eb430)1d:	15 1511725334	11/26/:	17 1.3.6.1.4	.1.1 200	040317	84529.765	1.2.840.1000	08.5.1.4.1.1.	2
	5 Renal	1.3.6.1.4.1.1 TO	GA-B8-51	1.3.6.1.4	4.1.1 4	9.7632368	133.75424	6 376.127872	164.443556	329.582418	actual_gi	raff 5b6eb430)1d:	33 1511725358	11/26/	17 1.3.6.1.4	.1.1 200	031023	80726.031	1.2.840.1000	08.5.1.4.1.1.	2
	6 Renal	1.3.6.1.4.1.1 TO	GA-B8-A5	1.3.6.1.4	4.1.1 4	2.8640175	362.90109	9 368.455545	371.596404	317.306693	actual_gi	raff 5b6eb430)1d:	23 1511725400	11/26/3	17 1.3.6.1.4	.1.1 200	050803	95301.39	1.2.840.1000	08.5.1.4.1.1.	2
	7 Renal	1.3.6.1.4.1.14	GA-B8-51	1.3.6.1.4	4.1.1 1	8.0562563	92.835164	8 291.220779	113.806194	283.548452	actual_gi	raff 5b6eb430)1d:	20 1511725419	11/26/	17 1.3.6.1.4	.1.1 200	031228	83231.781	1.2.840.1000	08.5.1.4.1.1.	2
	8 Renal	1.3.6.1.4.1.14 TO	GA-B8-51	1.3.6.1.4	4.1.1 5	9.0727826	5 119.94405	6 311.68032	186.949051	343.392607	actual gi	raff 5b6eb430)1d:	37 1511725442	11/26/3	17 1.3.6.1.4	.1.1 200	031118	93309.25	1.2.840.1000	08.5.1.4.1.1.	2
	9 Renal	1.3.6.1.4.1.1 TO												29 1511725456		17 1.3.6.1.4				1.2.840.1000		
	10 Renal	1.3.6.1.4.1.14	GA-B8-46	1.3.6.1.4	4.1.1 4	6.0867891	311.75224	8 317.818182	369.55045	315.772228	actual gi	raff 5b6eb430)1d:	40 1511725469	11/26/3	17 1.3.6.1.4	.1.1 200	020509	83303.343	1.2.840.1000	08.5.1.4.1.1.	2
	11 Renal	1.3.6.1.4.1.14	GA-B8-41	1.3.6.1.4	4.1.1 6	3.6642731	218.66133	9 232.3996	189.506494	290.709291	actual gi	raff 5b6eb430)1d:	37 1511725494		17 1.3.6.1.4		030519	85121.125	1.2.840.1000	08.5.1.4.1.1.	2
	12 Renal	1.3.6.1.4.1.14	GA-B8-46	1.3.6.1.4	4.1.14	40.70546	309.70629	4 283.036963	346.021978	303.496504	actual gi	raff 5b6eb430)1d:	29 1511725510	11/26/	17 1.3.6.1.4	.1.1 200	031015	95443.281	1.2.840.1000	08.5.1.4.1.1.	2
	13 Renal	1.3.6.1.4.1.14												23 1511725550		17 1.3.6.1.4				1.2.840.1000		
	14 Renal	1.3.6.1.4.1.1 T												30 1511725562		17 1.3.6.1.4				1.2.840.1000		
	15 Renal	1.3.6.1.4.1.1 T												22 1511725588		17 1.3.6.1.4				1.2.840.1000		
	16 Renal	1.3.6.1.4.1.1 T												21 1511725597		17 1.3.6.1.4				1.2.840.1000		
	17 Renal	1.3.6.1.4.1.1 T												27 1511725617		17 1.3.6.1.4				1.2.840.1000		
	18 Renal	1.3.6.1.4.1.1 T												32 1511725629		17 1.3.6.1.4				1.2.840.1000		
	19 Renal	1.3.6.1.4.1.1					-							21 1511725640		17 1.3.6.1.4				1.2.840.1000		
	20 Renal	1.3.6.1.4.1.1												16 1511725654		17 1.3.6.1.4				1.2.840.1000		
	21 Renal	1.3.6.1.4.1.14												21 1511725663		17 1.3.6.1.4 17 1.3.6.1.4				1.2.840.1000		
	22 Renal	1.3.6.1.4.1.1												17 1511725675		17 1.3.6.1.4				1.2.840.1000		
	23 Renal	1.3.6.1.4.1.14												15 1511725688		17 1.3.6.1.4 17 1.3.6.1.4				1.2.840.1000		
	24 Renal	1.3.6.1.4.1.14												25 1511725699		17 1.3.6.1.4 17 1.3.6.1.4				1.2.840.1000		
	25 Renal	1.3.6.1.4.1.14												12 1511725710		17 1.3.6.1.4 17 1.3.6.1.4				1.2.840.1000		
	26 Renal	1.3.6.1.4.1.14												24 1511725721		17 1.3.6.1.4 17 1.3.6.1.4				1.2.840.1000		
	20 Renal						-							24 1511725721 28 1511725732		17 1.3.6.1.4 17 1.3.6.1.4				1.2.840.1000		
	27 Renal 28 Renal	1.3.6.1.4.1.1 ⁴ TO 1.3.6.1.4.1.1 ⁴ TO												28 1511725732 65 1511725748		17 1.3.6.1.4 17 1.3.6.1.4		00624				
	28 Renal	1.3.6.1.4.1.14 1.3.6.1.4.1.14												37 1511725760		17 1.3.6.1.4 17 1.3.6.1.4		900624		1.2.840.1000		
	30 Renal	1.3.6.1.4.1.14														17 1.3.6.1.4 17 1.3.6.1.4		900105				
	30 Renal	1.3.6.1.4.1.14												64 1511725783 36 1511725809		17 1.3.6.1.4 17 1.3.6.1.4		390927		1.2.840.1000		
	31 Renal																	390927 380816		1.2.840.1000		
	32 Renal	1.3.6.1.4.1.1 T												28 1511725826		17 1.3.6.1.4		911008		1.2.840.1000		
		1.3.6.1.4.1.1 T												31 1511725837		17 1.3.6.1.4				1.2.840.1000		
	34 Renal	1.3.6.1.4.1.1.1												18 1511725851		17 1.3.6.1.4		910226		1.2.840.1000		
	35 Renal	1.3.6.1.4.1.1.1												43 1511725870		17 1.3.6.1.4		01224		1.2.840.1000		
	36 Ovarian	1.3.6.1.4.1.14												55 1511905225		17 1.3.6.1.4				1.2.840.1000		
	37 Ovarian	1.3.6.1.4.1.1 T										-		99 1511905350		17 1.3.6.1.4		960913		1.2.840.1000		
	38 Ovarian	1.3.6.1.4.1.1 ⁴ TO										-		63 1511905392		17 1.3.6.1.4		960723		1.2.840.1000		
	39 Ovarian	1.3.6.1.4.1.1 ⁴ T(CGA-13-079	1.3.6.1.4	4.1.14 2	6.660773	304.08839	8 106.563536	333.507182	106.563536	adept du	gor 5b6eb430)1d:	35 1511905442	11/28/:	17 1.3.6.1.4	.1.14 199	951123 2	30125.687	1.2.840.1000)8.5.1.4.1.1.	2
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Courtesy of Jayashree Kalpathy-Cramer

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A1 \checkmark \times \checkmark f_x order

	А	В	С	D	E	F	G	Н
1	order	anatomy	seriesUID	patientID	instanceUID	length	start_x	start_y e
2	1	Renal	1.3.6.1.4.1.14	TCGA-BP-434	1.3.6.1.4.1.14	66.4385613	172.835359	270.064088
3	2	Renal	1.3.6.1.4.1.14	TCGA-B8-414	1.3.6.1.4.1.14	49.5155823	149.098901	287.128871
4	3	Renal	1.3.6.1.4.1.14	TCGA-B8-510	1.3.6.1.4.1.14	70.0270211	131.196803	338.789211
5	4	Renal	1.3.6.1.4.1.14	TCGA-B8-554	1.3.6.1.4.1.14	41.721506	178.253746	319.352647
6	5	Renal	1.3.6.1.4.1.14	TCGA-B8-510	1.3.6.1.4.1.14	49.7632368	133.754246	376.127872
7	6	Renal	1.3.6.1.4.1.14	TCGA-B8-A5	1.3.6.1.4.1.14	42.8640175	362.901099	368.455545
8	7	Renal	1.3.6.1.4.1.14	TCGA-B8-51	1.3.6.1.4.1.14	18.0562563	92.8351648	291.220779
•	0		4 2 6 4 4 4 4	TOCA DO EA	4 2 6 4 4 4 4	50 0707000	440 044050	244 60022

	DICOMDIR	🛛 🔴 🕘 TCGA-BP-4343[TCGA-BP-4343]:[19870620:Renal]:4578[SR:Crowds Cure Cancer Annotation as Measurement Report]:1:					
) 19870620 Renal } CT} 3 MIN DELAY {SR} Crowds Cure Cancer Annotation as Measurement Report	 CGA-BP-4343[ICGA-BP-4343]:[19870620:Renal]:4578[SR:Crowds Cure Cancer Annotation as Measurement Report]:1: CONTAINER: Imaging Measurement Report [SEPARATE] (DCMR.1500) HAS CONCEPT MOD: CODE: Language of Content Item and Descendants = English HAS CONCEPT MOD: CODE: Country of Language = United States HAS OBS CONTEXT: PNAME: Person Observer Name = accomplished_peafowl HAS CONCEPT MOD: CODE: Procedure reported = CT Abdomen CONTAINS: CONTAINER: Image Library Group [SEPARATE] CONTAINS: CONTEXT: CODE: Modality = Computed Tomography HAS ACQ CONTEXT: DATE: Study Date = 19870620 HAS ACQ CONTEXT: TIME: Study Time = 135823 CONTAINS: CONTAINER: Imaging Measurement Group [SEPARATE] CONTAINS: CONTAINER: Imaging Measurement Group [SEPARATE] CONTAINS: CONTAINER: Imaging Measurement Group [SEPARATE] AS ACQ CONTEXT: TIME: Study Time = 135823 CONTAINS: CONTAINER: Imaging Measurement Group [SEPARATE] HAS OBS CONTEXT: TEXT: Tracking Identifier = 5b6eb4301d3175942d2985a3d0fbb00 HAS OBS CONTEXT: UDREF: Tracking Unique Identifier = 1.3.6.1.4.1.5962.1.1.0.0.1535644357.22655.1 HAS CONCEPT MOD: CODE: Finding Site = Kidney CONTAINS: NUM: Length = 66.43856134 mm 					
CompletionFlag Content COMPLETE 2017112	26 224217 1 images\TCGA-BP-4343\	INFERRED FROM: SCOORD: = POLYLINE (172.835357666016,270.064086914062,133.798889160156,343.045318603516) SELECTED FROM: IMAGE: = 1.2.840.10008.5.1.4.1.1 2 : 1.3.6.1.4.1.14519.5.2.1.9203.4004.268018422288818573226516023762 ICGA-BP-4343]:[19870620:CT CH/AB/PEL KIDNEY PROTOCOL]:3[CT:]:57: Imaging in the imaging interment of the					

1: : CONTAINER: (126000, DCM, "Imaging Measurement Report") [SEPARATE] (DCMR, 1500) >1.1: HAS CONCEPT MOD: CODE: (121049,DCM,"Language of Content Item and Descendants") = (eng,RFC5646,"English") >>1.1.1: HAS CONCEPT MOD: CODE: (121046, DCM, "Country of Language") = (US, ISO3166_1, "United States") >1.2: HAS OBS CONTEXT: PNAME: (121008, DCM, "Person Observer Name") = "accomplished peafowl" >1.3: HAS CONCEPT MOD: CODE: (121058,DCM,"Procedure reported") = (41806-1,LN,"CT Abdomen") >1.4: CONTAINS: CONTAINER: (111028, DCM, "Image Library") [SEPARATE] >>1.4.1: CONTAINS: CONTAINER: (126200,DCM,"Image Library Group") [SEPARATE] >>>1.4.1.1: CONTAINS: IMAGE: = (1.2.840.10008.5.1.4.1.1.2,1.3.6.1.4.1.14519.5.2.1.9203.4004.268018422288818573226516023762) >>>>1.4.1.1.1: HAS ACQ CONTEXT: CODE: (121139,DCM, "Modality") = (CT,DCM, "Computed Tomography") >>>>1.4.1.1.2: HAS ACQ CONTEXT: DATE: (111060,DCM, "Study Date") = "19870620" >>>>1.4.1.1.3: HAS ACQ CONTEXT: TIME: (111061,DCM,"Study Time") = "135823" >1.5: CONTAINS: CONTAINER: (126010,DCM,"Imaging Measurements") [SEPARATE] >>1.5.1: CONTAINS: CONTAINER: (125007, DCM, "Measurement Group") [SEPARATE] >>>1.5.1.1: HAS OBS CONTEXT: TEXT: (112039,DCM,"Tracking Identifier") = "5b6eb4301d3175942d29985a3d0fbb00" >>>1.5.1.2: HAS OBS CONTEXT: UIDREF: (112040,DCM,"Tracking Unique Identifier") = "1.3.6.1.4.1.5962.1.1.0.0.0.1535644357.22655.1" >>>1.5.1.3: HAS CONCEPT MOD: CODE: (G-C0E3,SRT,"Finding Site") = (T-71000,SRT,"Kidney") >>>1.5.1.4: CONTAINS: NUM: (G-D7FE,SRT,"Length") = 66.43856134 (mm,UCUM,"mm") >>>>1.5.1.4.1: INFERRED FROM: SCOORD: = POLYLINE {172.835357666016,270.064086914062,133.798889160156,343.045318603516} >>>>1.5.1.4.1.1: SELECTED FROM: IMAGE: = (1.2.840.10008.5.1.4.1.1.2,1.3.6.1.4.1.14519.5.2.1.9203.4004.268018422288818573226516023762)

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L_1 (121000, DCH) I CI SON ODSCIVCI Name / - accomptished_pearowe
: (121058,DCM,"Procedure reported") = (41806-1,LN,"CT Abdomen")
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Terminal — -tcsh — 140×37

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Terminal — -tcsh — 140×37

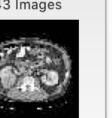
[graythin:NCI/CrowdsCureCancer/converttoDICOMSR] dclunie% dcdump dicomsrfiles/1.dcm (0x0002,0x0000) UL File Meta Information Group Length VR= VL=<0x0004> [0x000000bc] (0x0002.0x0001) OB File Meta Information Version VR = < OB > $VL = <0 \times 0002 > [0 \times 00, 0 \times 01]$ (0x0002,0x0002) UI Media Storage SOP Class UID VR=<UI> VL=<0x001e> <1.2.840.10008.5.1.4.1.1.88.22> VR=<UI> VL=<0x002e> <1.3.6.1.4.1.5962.1.1.0.0.0.1535642467.22502.1> (0x0002,0x0003) UI Media Storage SOP Instance UID (0x0002,0x0010) UI Transfer Syntax UID VR=<UI> VL=<0x0014> <1.2.840.10008.1.2.1> (0x0002,0x0012) UI Implementation Class UID VR=<UI> VL=<0x0016> <1.3.6.1.4.1.5962.99.2> (0x0002,0x0013) SH Implementation Version Name VR=<SH> VL=<0x0010> <PIXELMEDJAVA001 > (0x0008,0x0016) UI SOP Class UID VR=<UI> VL=<0x001e> <1.2.840.10008.5.1.4.1.1.88.22> (0x0008,0x0018) UI SOP Instance UID VR=<UI> VL=<0x002e> <1.3.6.1.4.1.5962.1.1.0.0.0.1535642467.22502.1> (0x0008,0x0020) DA Study Date VR=<DA> VL=<0x0008> <19870620> (0x0008,0x0021) DA Series Date VR=<DA> VL=<0x0000> <> (0x0008,0x0023) DA Content Date VR=<DA> VL=<0x0008> <20171126> VR=<TM> VL=<0x0006> <135823> (0x0008,0x0030) TM Study Time (0x0008,0x0033) TM Content Time VR=<TM> VL=<0x0006> <224217> (0x0008,0x0050) SH Accession Number VR=<SH> VL=<0x0000> <> VR=<CS> VL=<0x0002> <SR> (0x0008,0x0060) CS Modality (0x0008,0x0070) L0 Manufacturer VR = <LO >VL=<0x0008> <PixelMed> (0x0008,0x0080) LO Institution Name VR = <L0 >VL=<0x0000> <> (0x0008,0x0090) PN Referring Physician's Name VR=<PN> VL=<0x0000> <> (0x0008,0x1010) SH Station Name VR=<SH> VL=<0x0004> <NONE> (0x0008,0x1030) LO Study Description VR=<LO> VL=<0x0006> <Renal > VR = <LO> VL = <0x0034> <Crowds Cure Cancer Annotation as Measurement Report > (0x0008,0x103e) LO Series Description (0x0008,0x1090) LO Manufacturer's Model Name VR=<LO> VL=<0x0022> <XSLT from annotations_expanded.csv> (0x0008,0x1111) SQ Referenced Performed Procedure Step Sequence VR=<SQ> VL=<0xffffffff

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(0x0020,0x000d)	UI	Study Instance UID	VR= <ui></ui>	VL=<0x0040>	<1.3.6.1.4.1.14519.5.2.1.9203.4004.100545948082587796019777812429>
(0x0020,0x000e)	UI	Series Instance UID	VR= <ui></ui>	VL=<0x002c>	<1.3.6.1.4.1.5962.1.3.0.0.1535642467.22502.1>
(0x0020,0x0010)	SH	Study ID VR= <sh></sh>	VL=<0×	<0000> <>	
(0x0020,0x0011)	IS	Series Number	VR= <is></is>	VL=<0x0004>	<4578>
(0x0020,0x0013)	IS	Instance Number	VR= <is></is>	VL=<0x0002>	<1 >

СТ 101 Images



3 MIN DELAY 43 Images



Crowds Cure Cance... asurement Report 2 Pages

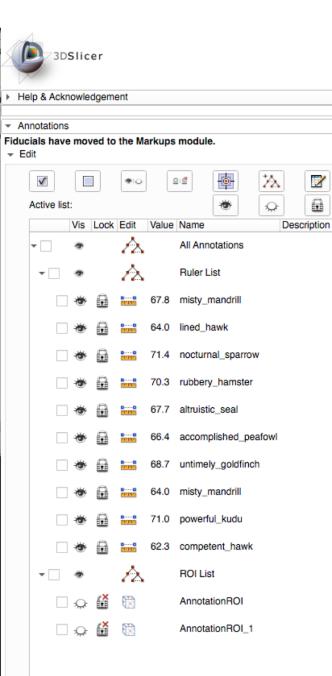


Image size: 1224 x 1584 View size: 1246 x 751 WL: 127 WW: 255 X: -623 px Y: 320 px Val	Patient: TCGA-BP-4343 (#TCGA-BP4343) Study: Renal Series: Crowds Cure Cancer Annotation as Measurement Report (#4578) Manufacturer: PixelMed (XSLT from annotations_expanded.csv, #9723613413261) Completion Flag: COMPLETE VeimOction for fore() IndVERIFIED Content Date/Time: 2017-11-26 22:42:17
	Imaging Measurement Report Concept Modifier: Language of Content Item and Descendants [Annex 1] Observation Context: Person Observer Name = accomplished_peafowl Concept Modifier: Procedure reported = CT Abdomen (41806-1, LN)
	Image Library
	Image Library Group
R	<u>CT image</u>
	Acquisition Context: Modality = Computed Tomography (CT, DCM) Acquisition Context: Study Date = 1987-06-20 Acquisition Context: Study Time = 13:58:23
	Imaging Measurements
	Measurement Group
	Observation Context: Tracking Identifier = "5b6eb4301d3175942d29985a3d0fbb00" Observation Context: Tracking Unique Identifier = 1.3.6.1.4.1.5962.1.1.0.0.0.1535644357.22655.1 Concept Modifier: Finding Site = Kidney (T-71000, SRT)
	Length: 66.43856134 mm
	Inferred from: Spatial Coordinates [Annex 2]
	Annex
Zoom: 50% Angle: 0	Annex 1
Im: 3/3 Uncompressed	Language of Content Item and Descendants:

TCGA-BP-4343 (-, -) Ct Ch-Ab-Pel Kidney Protocol 4578



11/26/17, 10:42:17 PM Made In Horos



Courtesy of Steve Pieper

accompl**htisty_mondrill:64.0mm** powerful_kuclu:71.0mm rubbery_hamster:**70.3mm**

B: 105: 3 MIN DELAY

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Annotation workflow and use cases

Traditional "box under the table/in back room" ("headless")

- acquire images -> process -> send both to PACS for reader
- "push workflow" ("unmanaged") familiar from Mammography CADe
- cloud variants and large data volumes add security/timing challenges
- data to algorithm or algorithm to data
- Interactive (virtually "embedded")
 - user views images -> selects task/frames/ROI -> requests CAD -> see result
 - user expects immediate response (or likely will not use)
 - responsiveness may depend on prearranged proximity of data & algorithm
 - standard payload not only for results but request as well
 - standard protocol for command and control of request/progress/complete

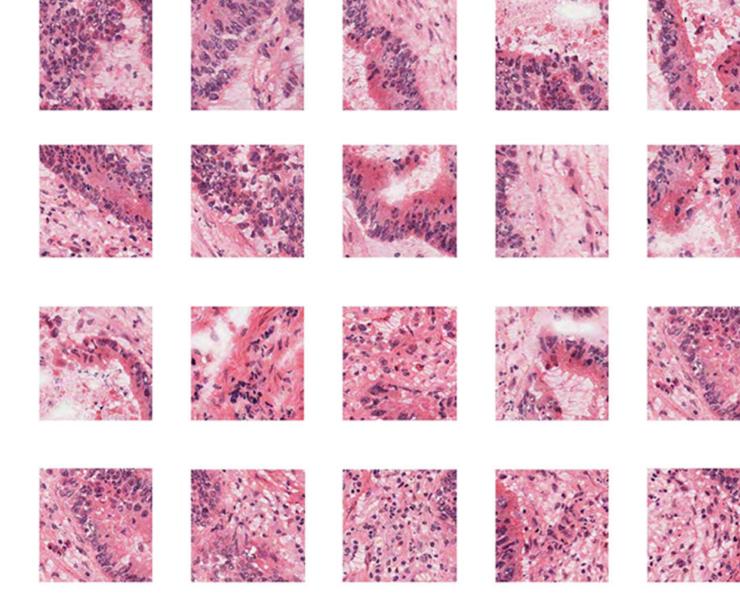


Annotation workflow standards

- Standard payload for images in/out & results (DICOM)
 - for visualization "pretty picture" screen shots and presentation states
 - semantically meaningful segmentations, parametric maps, SR
- Standard protocols to choose from:
 - DICOM Unified Procedure Step (UPS) and DICOMweb UPS-RS variant
 - IHE Invoke Image Display (IID) HTTP request to view specified study(ies)
 - HL7 synchronized applications CCOW and now FHIRcast
 - DICOM Application Hosting SOAP-based, never popular; revisit RESTfully?
- "Push" unmanaged request image timeout, UPS N-CREATE, IHE IID
- "Pull" from worklist UPS C-FIND or UPS-RS, do work and update status
- Fetch payload DICOM C-MOVE/C-GET or WADO-RS (metadata)
- Return result DICOM C-STORE or STOW-RS
- Security transport (TLS), authentication (Kerberos, SAML, JWT, OAUTH)

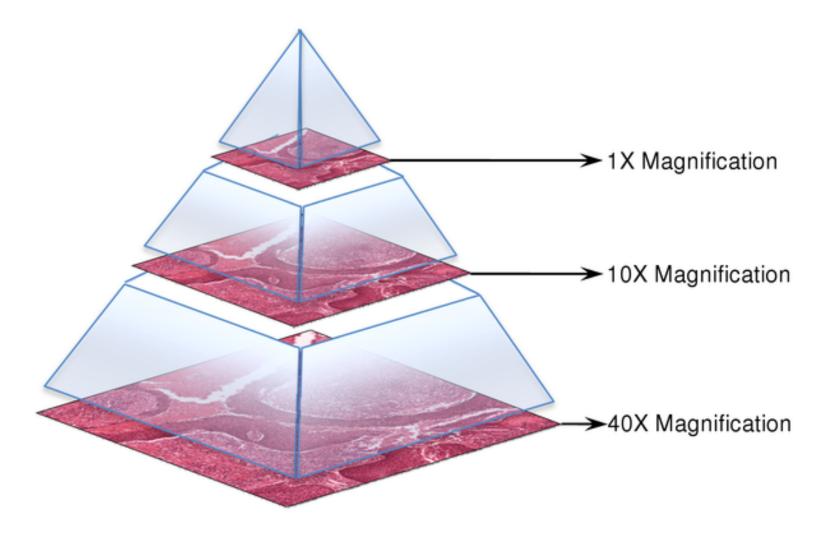




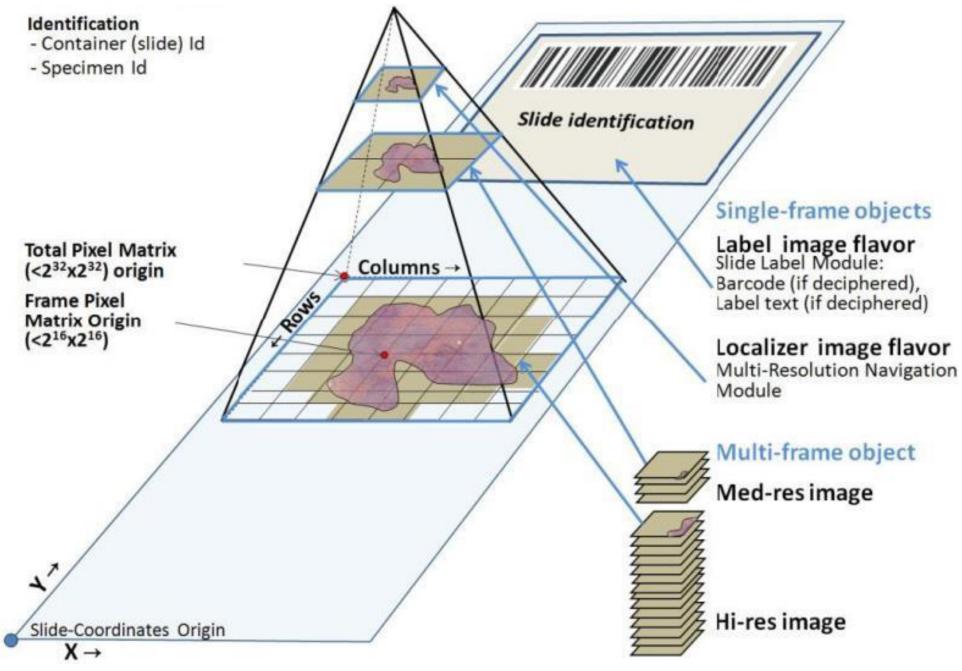


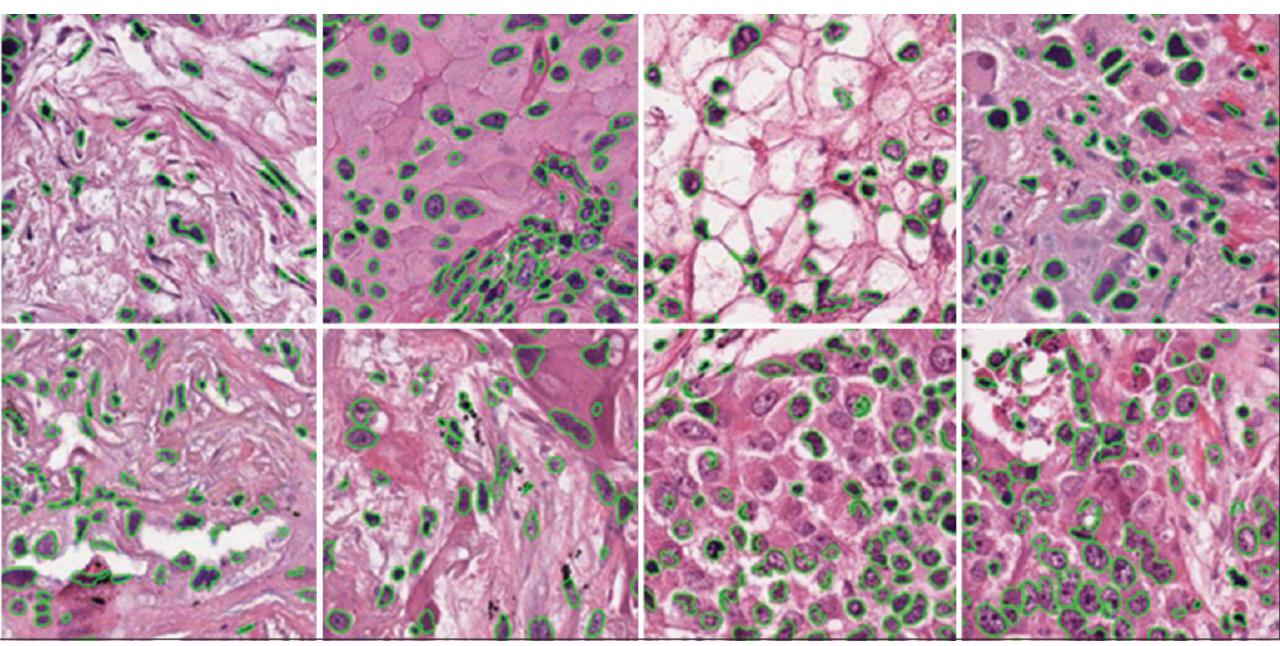
Yoon et al. Tumor Identification in Colorectal Histology Images Using a Convolutional Neural Network. J Digit Imaging. 2018 Jul 31;1–10.

An illustration of how digital slides are stored in a pyramid structure.



Wang Y, Williamson KE, Kelly PJ, James JA, Hamilton PW (2012) SurfaceSlide: A Multitouch Digital Pathology Platform. PLOS ONE 7(1): e30783. https://doi.org/10.1371/journal.pone.0030783 http://journals.plos.org/plosone/article?id=10.1371/journal.pone.0030783

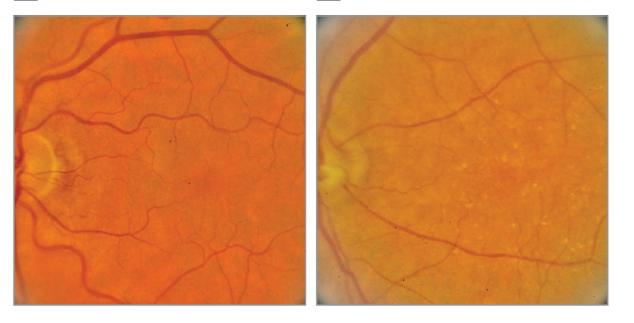




Wen et al. A methodology for texture feature-based quality assessment in nucleus segmentation of histopathology image. JPI. 2017.

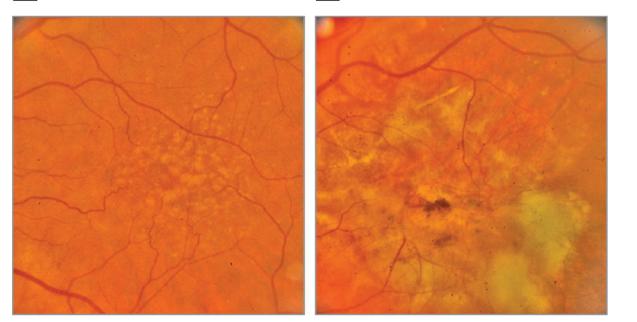
A Category 1

B Category 2



c Category 3

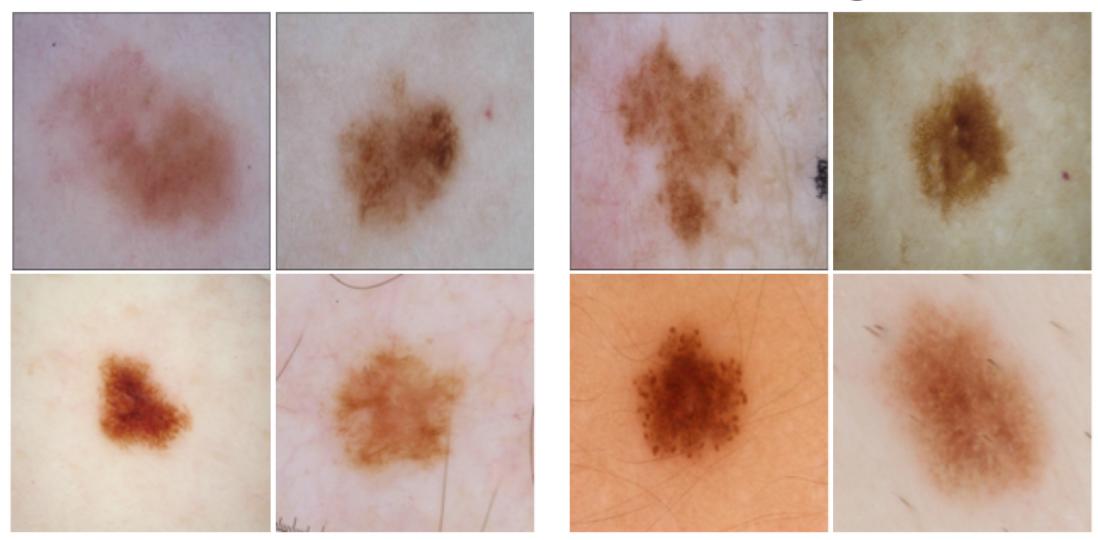
D Category 4



Burlina et al. Automated Grading of Age-Related Macular Degeneration From Color Fundus Images Using Deep Convolutional Neural Networks. JAMA Ophthalmol. 2017 Nov 1;135(11):1170–6.

Melanoma





Gutman D et al. Skin Lesion Analysis toward Melanoma Detection: A Challenge at the International Symposium on Biomedical Imaging (ISBI) 2016, hosted by the International Skin Imaging Collaboration (ISIC). arXiv:160501397.

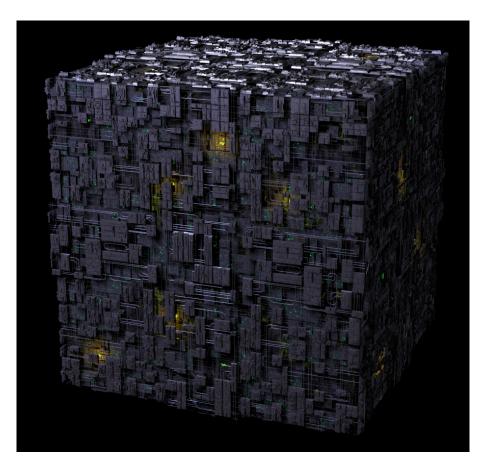
DICOM – Gaps and Future Improvements

- Extending Segmentation and Parametric Maps to tiled images for Whole Slide Imaging (CP 1830)
 - Plane Position (Slide) macro instead of Plane Position (Patient)
- DICOM query or DICOMweb RESTful API for annotation access?
 - "spatial" queries, e.g., all annotations that intersect defined frame/region
- More compact representation of very large numbers of contours?
 - e.g., all nuclei, all membranes in WSI (versus SEG bit-plane representation)
- Explicit Label Map rather than Segmentation bit planes?
 - in SEG object have one bit-plane per segment (label), each as a frame
 - in traditional label maps, one multi-valued voxel where each voxel is an index whose value represents a segment (label)
 - same semantics but "gratuitously" different representation convenience?
- Always need more coded concepts
 - e.g., more texture features all of IBSI (CP 1764)





How long until you are assimilated?





"we will add your biological and technological distinctiveness to our own"

"your culture will adapt to service us"

"resistance is futile"

