



Imaging Data Anonymisation Guideline

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Revision History

Version	Date	Summary
2.0	09 July 26	

1. Introduction

1.1 Purpose

The purpose of this guideline is to define the minimum anonymisation standards for DICOM imaging data shared externally via a secure, governed research environment (SGRE). This document aims to ensure all shared clinical trial imaging data complies with privacy regulations and ethical standards, thereby protecting the identities of clinical trial participants. It is intended for people involved in data processing, anonymisation, and external collaborations, as well as for external researchers who will access this anonymised data.

1.2 Scope

This guideline applies to all DICOM imaging data prepared for sharing with third-party researchers on an SGRE. While the primary focus is on DICOM data, the principles outlined may also inform anonymisation practices for other types of imaging data when applicable.

1.3 Secure, governed research environment (SGRE)

A secure, governed research environment (SGRE) is any controlled environment designed to support privacy-preserving access to research data.

This anonymisation guidance assumes that anonymised imaging data will be shared only within an SGRE that adheres to established principles for responsible data access. A commonly referenced model is the Five Safes framework, developed by the UK Office for National Statistics, which is widely used for enabling privacy-preserving research. The framework ensures:

- Safe People (only trained and authorised users may access the data)
- Safe Projects (research is ethically and scientifically justified)
- Safe Data (data is appropriately anonymised and prepared to minimise identification risk)
- Safe Settings (data can only be accessed within secure, technically controlled environments)
- Safe Outputs (all results leaving the environment undergo disclosure review).

Together, these safeguards ensure data can be used for meaningful research while maintaining privacy, confidentiality and public trust.

2. DICOM standard overview

DICOM (Digital Imaging and Communications in Medicine) is the internationally recognised standard format for storing, transmitting and managing medical imaging data developed and maintained by the National Electrical Manufacturers Association (NEMA). A DICOM file encapsulates both the medical image itself (pixel data) and a rich set of metadata that describes the image and its clinical context. This standardised structure supports interoperability across imaging devices, clinical systems and research platforms.

Due to its structural complexity and extensibility, the DICOM format has the potential to store personal data, including sensitive health data, across both metadata and pixel data, including nested sequences and vendor-specific fields.

The following overview focuses only on key, foundational aspects of the DICOM standard that are most relevant to anonymisation and external data sharing. It is not intended to be a comprehensive treatment of the full specification. For full technical details, implementers should consult the authoritative NEMA DICOM standard documentation ((NEMA), Digital Imaging and Communications in Medicine (DICOM) standard, 2025).

2.1 Metadata

DICOM metadata is a structured collection of **attributes** (also called "tags") that describe aspects of the imaging study ((NEMA), PS3.5 2025c - Data Structures and Encoding, 2025). Each attribute is uniquely identified by a tag in the form (group, element) – for example, (0010,0010) for Patient Name.

Each tag has a defined **value representation (VR)**, which specifies the data type and format of the values. VRs play a critical role in health data risk assessment by indicating the likelihood of a data element containing personal data. Please find the VR types, descriptions and health data risk assessment in Appendix A.

The importance and required presence of the attributes are defined in the Information Object Definitions (IODs) through a **Type system**:

- Type 1: Required and must have a valid, non-empty value
- Type 2: Required but may have an empty value
- Type 3: Optional attributes

2.1.1. Standard tags

Standard tags are defined in the DICOM standard and are consistent across vendors and systems ((NEMA), PS3.6 2025c - Data Dictionary, 2025). They include essential information required for interpretation, storage and clinical integration of imaging data. Many of these tags contain patient-related or study-specific data that may be considered personal data and require anonymisation.

2.1.2. Private tags

Private tags are non-standard attributes reserved for modality vendors, PACS systems, or institutions to store proprietary information not defined in the public DICOM dictionary. They occupy odd-numbered groups (e.g. 0019,xxxx or 0029,xxxx) and are organised into blocks allocated by a private creator element (e.g. (0019,0010)), which is a vendor-chosen label that declares ownership of the subsequent private elements and ensures different vendors do not collide when storing their custom metadata.

Because private tags are flexible by design, their **internal structure varies widely**: many use standard VRs such as LO, DS, IS or FD, but others contain complex or opaque content including binary arrays, mixed binary-text structures, proprietary encoded tables, XML or JSON fragments, or large textual blobs. Siemens CSA headers, GE ExamInfo structures and Philips private dictionaries are typical examples of where metadata may be embedded in formats that are not human-readable without vendor-specific parsers. **The type of information** stored in these private tags ranges from highly technical acquisition and reconstruction parameters to workflow-derived clinical or operational details. Scientifically useful content is common, especially in MRI and PET, because not all acquisition parameters are captured in standard DICOM attributes. Many advanced or experimental protocols also encode their internal sequence descriptors, reconstruction modes, or software versioning in private metadata. Some of these parameters might be essential for accurate preprocessing and reproducibility in quantitative pipelines.

However, private tags also carry **substantial re-identification risk** because vendors and workflow systems often copy clinical or operational details into these fields. They may include patient identifiers, accession numbers, operator or technologist names, site or department names, and precise acquisition timestamps. Free-text protocol notes, encoded binary structures and device-specific identifiers such as scanner serial numbers can also embed personal information or quasi-identifiers in ways that are difficult to detect without fully decoding the private metadata.

2.1.3. Sequences and nested elements

DICOM supports complex data structures through **sequences** (VR 'SQ'). A sequence tag contains a list of items, each of which is a nested set of DICOM attributes. Sequences are used to encode repeating or hierarchical information. Personal data can be present at any level of the sequence hierarchy. For example, a sequence may contain nested patient names referring to physician details, or institution identifiers.

2.1.4. Other structural considerations

In addition to tags and sequences, DICOM files may include:

- **Overlay and graphic data:** Elements such as (6000,3000), which may be used to annotate images with text or graphical content, potentially containing identifiers.
- **Encapsulated documents:** PDF or CDA documents embedded in tags such as (0042,0011), which may include clinical notes or identifiable content.
- **DICOM structured reports (SRs):** Specialised DICOM objects that encode clinical information – such as measurements, observations, interpretations and annotations – in a structured, machine-readable format. Unlike free-text narrative reports, SRs use a hierarchical data model built from coded entries and nested sequences, enabling both human readability and automated processing.

- **Waveform data:** For modalities such as ECG or EEG, PHI can be embedded in waveform annotation sequences or headers.
- **Radiation therapy objects (RTSTRUCT, RTPLAN, RTDOSE):** Radiation Therapy Structure Sets, Plans, and Dose objects may contain patient identifiers, anatomical labels, physician names, timestamps, and treatment-related annotations that require full review and anonymisation.
- **Presentation states (PRs):** Presentation state objects control visualisation parameters (e.g. windowing, overlays, annotations) and may contain burned-in text, labels, or graphical overlays that include PHI.
- **Segmentation objects (SEGs):** Segmentation objects encode derived mask data and may contain identifying labels, structure names, timestamps, and references to source images that require anonymisation.
- **CT dose reports:** For CT scans, dose report objects or secondary capture images may include displayed patient names, dates, institution identifiers, and operator information rendered directly into the image pixels.

2.2 Pixel data

Pixel data represents the raw image content, such as grayscale intensities in an MRI slice or coloured values in ultrasound. It is stored in tag (7FE0,0010) and is distinct from metadata. Pixel data is often large and can contain sensitive information embedded directly into the image.

Common forms of sensitive content within pixel data include:

- **Burned-in annotations:** These are alphanumeric characters or graphics permanently overlaid onto the image pixels, often by the imaging device or PACS system. They may include patient names, IDs, dates of birth, examination dates or institutional information. Burned-in annotations can appear in a variety of image types, particularly those that are non-volumetric, secondary or presentation-focused.
- **Identifiable anatomy:** Certain modalities, especially structural head imaging such as MRI or CT, capture detailed facial features or cranial geometry. These anatomical structures can be used to reconstruct a 3D facial surface or be matched with external photographs, making them a risk for biometric re-identification.

3. Anonymisation overview

3.1 Definitions

The terms de-identification, pseudonymisation and anonymisation are often used interchangeably in clinical research and data sharing, but they reflect different levels and approaches to reducing the risk of re-identifying individuals. Understanding these distinctions is essential, especially when working with sensitive imaging data.

- **De-identification** typically refers to the process of removing or masking direct identifiers (such as names, IDs and dates) to reduce the risk of re-identification. However, re-identification remains theoretically possible under certain conditions (e.g. via dataset linkage or background knowledge).
- **Pseudonymisation** involves replacing identifiable information (e.g. patient ID) with artificial identifiers (pseudonyms), while maintaining a reversible mapping (often stored separately and securely). This approach does not count as anonymisation under most privacy laws, as re-identification remains possible by those holding the key.
- **Anonymisation** is the process of transforming data so that individuals cannot be identified, either directly or indirectly, using any information that is reasonably available. It involves permanently removing or altering personal identifiers in a way that prevents reconstruction or linkage back to a specific person. Proper anonymisation preserves the usefulness of the dataset for analysis while ensuring no individual can be re-identified.

Below is a simplified comparison:

Aspect	De-identification	Pseudonymisation	Anonymisation
Goal	Minimise risk of re-identification	Mask direct identifiers, enable controlled linkage	Eliminate re-identification risk
Re-identification	Possible under certain conditions	Possible by trusted party with mapping key	Not reasonably possible
Identifiers	Removed or obfuscated	Replaced with pseudonyms	Removed or irreversibly transformed
Linkability	Partially retained	Retained via mapping	Lost
Scientific utility	Preserved	Preserved	May be partially reduced
Imaging challenges	May skip pixel or file-level identifiers	Same as de-id; mapping must be protected	Requires full metadata, pixel, and structure review

As such, the workstream's recommendation aligns more closely with the **anonymisation standard**, aiming to eliminate personal data to protect patient privacy while preserving the scientific utility of imaging data.

3.2 Regulatory standards: HIPAA, GDPR and cross-jurisdictional applicability

Two widely recognised frameworks govern anonymisation practices:

- **HIPAA (US): Safe Harbor and Expert Determination**
Under the Health Insurance Portability and Accountability Act (HIPAA), two primary methods are recognised for de-identifying protected health information (PHI):
 - o **Safe Harbor method:** This approach requires removing 18 specific identifiers from PHI. These include direct and indirect identifiers such as:
 - o Names
 - o All geographic subdivisions smaller than a state, including street address, city, county, precinct, ZIP code, and their equivalent geocodes, except for the initial three digits of the ZIP code if, according to the current publicly available data from the Bureau of the Census:
 - o The geographic unit formed by combining all ZIP codes with the same three initial digits contains more than 20,000 people
 - o The initial three digits of a ZIP code for all such geographic units containing 20,000 or fewer people is changed to 000.
 - o All elements of dates (except year) for dates that are directly related to an individual, including birth date, admission date, discharge date, death date, and all ages over 89 and all elements of dates (including year) indicative of such age, except that such ages and elements may be aggregated into a single category of age 90 or older
 - o Telephone numbers
 - o Vehicle identifiers and serial numbers, including licence plate numbers
 - o Fax numbers
 - o Device identifiers and serial numbers
 - o Email addresses
 - o Web universal resource locators (URLs)
 - o Social security numbers
 - o Internet protocol (IP) addresses
 - o Medical record numbers
 - o Biometric identifiers, including finger and voice prints
 - o Health plan beneficiary numbers
 - o Full-face photographs and any comparable images
 - o Account numbers

- o Any other unique identifying number, characteristic or code
- o Certificate/licence numbers.

If these elements are removed, and no actual knowledge of re-identification risk remains, the dataset is considered de-identified (Malin, 2010).

- o **Expert Determination method:** In this approach, a qualified statistical or scientific expert applies formal risk assessment techniques and determines that the risk of re-identification is very small, even if some elements not permitted under Safe Harbor are retained.

Neither method provides a foolproof, zero-risk de-identification solution. Rather, they represent pragmatic frameworks designed to enable the creation and sharing of healthcare information with a sufficiently low and manageable risk of re-identification.

- **GDPR (EU)**

The General Data Protection Regulation (GDPR) defines anonymisation more stringently. Anonymised data must be irreversibly stripped of personal identifiers, such that no individual can be re-identified by any means reasonably likely to be used. This does not require absolute impossibility of re-identification, but rather that the risk is no longer reasonably plausible. Pseudonymised data – where identifiers are replaced by codes, but re-identification is still possible through a key – is not considered anonymous under the GDPR (University College of London, 2025) (Sharp Cookie Advisors, 2020).

While the HIPAA and the GDPR provide important reference points for health data de-identification and anonymisation, they should not be treated as universal substitutes for jurisdiction-specific legal review. Other privacy and health data frameworks may apply depending on the country or region of data origin, the location of the data controller or processor, the recipient institution, and the intended data-sharing pathway. Examples include Canada's Personal Information Protection and Electronic Documents Act (PIPEDA) and provincial health privacy laws, Japan's Act on the Protection of Personal Information (APPI), the UK GDPR and Data Protection Act, Australia's Privacy Act, Singapore's Personal Data Protection Act (PDPA), Brazil's Lei Geral de Proteção de Dados (LGPD), China's Personal Information Protection Law (PIPL), South Korea's Personal Information Protection Act (PIPA), and India's Digital Personal Data Protection Act (DPDP Act).

Although these frameworks generally share a common concern with whether an individual remains identifiable, directly or indirectly, they differ in terminology, legal thresholds, treatment of pseudonymised data, research-use conditions, and cross-border transfer requirements. For example, under HIPAA, de-identification may be achieved through the Safe Harbor or Expert Determination methods, whereas GDPR distinguishes anonymous data from pseudonymised personal data, with pseudonymised data generally remaining within the scope of data protection law if re-identification is possible using additional information. Canadian and Japanese frameworks also use jurisdiction-specific concepts of de-identification, anonymisation and pseudonymisation. Therefore, these guidelines use HIPAA and GDPR as benchmark frameworks, while recognising that

international medical imaging data sharing may require additional jurisdiction-specific review.

For medical imaging data specifically, legal requirements should be applied together with imaging-specific technical standards such as DICOM/NEMA PS3.15. DICOM/NEMA provides the operational framework for addressing imaging-specific identifiers, including DICOM attributes, private tags, dates, UIDs, descriptors, structured content, burned-in annotations, and pixel data. However, DICOM/NEMA conformance alone does not determine whether a dataset is legally anonymised under every applicable jurisdiction. The final anonymisation assessment should consider both technical de-identification completeness and the applicable legal standard for identifiability.

3.3 Imaging data anonymisation standards

Imaging data anonymisation presents unique challenges due to its dual structure: metadata (tags) and pixel data (the image itself), both of which can contain personal data. For this reason, the medical imaging industry refers to established de-identification standards, notably the **NEMA DICOM de-identification standard** ((NEMA), PS3.15 E Attribute Confidentiality Profiles (Normative), 2025).

This standard outlines multiple 'de-identification profiles', each specifying which DICOM attributes should be removed, retained, replaced or blanked, depending on the intended use case. For example:

- **The Basic Profile** removes all directly identifying attributes and generalises or removes potentially identifying metadata.
- **The Retain Safe Private Option** allows some private attributes to remain, if they have been reviewed and deemed non-identifiable.
- **The Clean Descriptors Option** addresses descriptions that may inadvertently reveal identity, such as body part labels including patient names.

These profiles help ensure a consistent, auditable and standards-based approach to anonymising DICOM files, especially when sharing imaging data across institutions, countries, or with external researchers.

In addition to the NEMA standard, **The Cancer Imaging Archive (TCIA)** provides publicly available imaging datasets that follow strict de-identification procedures aligned with DICOM PS3.15, including detailed documentation of removed, retained and sanitised attributes; TCIA also maintains a comprehensive *Private Tag Dictionary* that helps identify risky vendor-specific metadata. Similarly, the **Radiological Society of North America (RSNA)** promotes best practices for imaging de-identification through initiatives such as the RSNA Clinical Trials Processor, educational resources, and guidelines that emphasise personal information removal, safe handling of DICOM metadata, and robust anonymisation workflows for multi-institutional research.

Despite the availability of established standards and reference practices, there remain significant challenges in defining explicit

anonymisation criteria and developing fully standardised, practical guidelines that address the complete spectrum of risks associated with DICOM anonymisation for clinical data sharing. As a result, organisations still face ambiguities when operationalising these standards – particularly when balancing regulatory compliance, scientific usability and interoperability across external collaborators. Continued harmonisation efforts and clearer consensus on operational best practices are therefore needed to support consistent, safe and effective anonymisation of imaging data in real-world clinical and research environments.

4. Anonymisation criteria

The workstream has conducted a comprehensive review of current anonymisation and de-identification standards, including the DICOM standard as maintained by NEMA, HIPAA Privacy Rule requirements, GDPR definitions of personal data and anonymisation, and practical frameworks adopted across leading research institutions such as RSNA guidance on imaging de-identification and the operational approaches implemented by TCIA, which provides large-scale, publicly accessible imaging datasets anonymised according to rigorous, transparent procedures.

Building on this foundation, the workstream has developed and continues to refine a unified set of imaging data anonymisation guidelines and workflow procedures that address the full range of challenges inherent to DICOM metadata and pixel data. This criteria provides clear, practical rules for removing, transforming and retaining information in a manner that maintains scientific and clinical utility while meeting the compliance expectations outlined earlier in this document.

To support consistency, traceability and scalability, our recommended approach incorporates automated detection, transformation and auditing of both metadata and pixel data, enabling reliable application of anonymisation rules across diverse modalities, vendors and study contexts. By grounding the anonymisation workflow in established regulatory, technical and community standards, the strategy provides a strong and defensible foundation for compliant and interoperable clinical data sharing with external collaborators.

4.1 Consistency across clinical and imaging data

Clinical and demographic data are essential complements to imaging datasets, and imaging data likewise provides critical context for clinical analyses. When these two data types are shared together, they must be anonymised using **harmonised transformation rules** to ensure both analytic consistency and protection against re-identification. Divergent anonymisation approaches can introduce inconsistencies, break linkages, or inadvertently expose personal data when the two datasets are joined.

To prevent these issues, the same anonymisation logic must be applied across all datasets intended for joint analysis, including:

- **Direct identifiers** (e.g. names, medical record numbers, full dates of birth, contact information, addresses) must

be **removed from both clinical and imaging datasets without exception**, as their presence in either domain would re-identify the joined dataset.

- **Quasi-identifiers** (e.g. demographic attributes, temporal information, geographic descriptors, institutional identifiers) may be retained only when supported by a documented risk assessment demonstrating that their inclusion does not meaningfully increase re-identification likelihood. When quasi-identifiers are scientifically relevant, they should follow an aligned transformation logic across datasets (consistent date shifting, age banding, or controlled vocabulary harmonisation, etc.).

For certain demographic elements that are not required for imaging-based analyses but may be relevant to clinical or epidemiological research (e.g. race, ethnicity, nationality, or county of residence), it may be appropriate to **retain these fields in the clinical dataset while omitting them from the imaging metadata**. This approach minimises exposure of sensitive attributes to the more broadly distributed imaging files while preserving information needed for clinical analyses conducted under stricter governance.

Consistent cross-dataset anonymisation ensures clinical and imaging data remain accurately joinable, analytically coherent, and privacy-preserving, enabling integrated research without introducing avoidable re-identification risks.

4.2 Metadata anonymisation

4.2.1. Standard tags

As part of the anonymisation workflow development, we have conducted a structured review of standard DICOM attributes to identify elements that may contain personal data or pose a re-identification risk. Based on this analysis, we have compiled a **detailed list of DICOM tags, along with the required anonymisation actions**, grounded in the NEMA DICOM standard and further informed by practical guidance from TCIA and RSNA. This tag-action mapping serves as the foundation of our pipeline and addresses both commonly used and less visible metadata elements (see Appendix C).

Below are several key attributes and the corresponding anonymisation strategies we suggest:

- **Patient ID:** Replace with an anonymised identifier generated using a salted cryptographic hash function. SHA-256 is recommended due to its strong collision resistance and wide adoption. A randomly generated salt is combined with the original identifier prior to hashing to prevent guessing or reconstruction. The resulting identifier should remain consistent for each participant across all imaging sessions to enable linkage within the dataset without retaining any real-world identifying information. The salt or any identifier mapping enabling re-identification must not be shared with data recipients, thereby preserving external anonymisation. However, such mappings may be retained internally under strict access controls, separate from the released dataset, to support an institutional crosswalk for operational purposes.

- **Person name (PN) attributes:** All VR 'PN' attributes should be cleared or removed, since names are direct identifiers and are unnecessary for image interpretation or research workflows.
 - Exception: **Patient name:** Replace with anonymised Patient ID value, providing internal dataset linkage while removing all name-related information.
- **Patient's birth date and age:** Patient Birth Date (0010,0030) is classified as a direct identifier and must be removed or replaced with a zero-length or placeholder value (e.g. "00000000"). DOB must not be date-shifted or replaced with a surrogate date, as doing so may still permit inference of the true age. Instead, age information may be retained in Patient's Age (0010,1010), provided it is generalised to prevent re-identification. Recommended generalisation rules are:
 - *Participants ≥90 years:* age aggregated into a single category (e.g. "90+" for clinical data, "090Y" for the Age tag) per Safe Harbor.
 - *All other participants (<90 years):* the degree of age rounding or dynamic banding (e.g. monthly bins for infants, broader bands for minors, or 5–10-year bands for adults) should be determined through a risk assessment that considers cohort size, study population characteristics, rarity of condition and other factors (Kniola, 2016).

These rules ensure DICOM objects retain clinically meaningful demographics while preventing reconstruction or inference of the original birth date.

- **Demographic fields:** Retain only demographic attributes needed for imaging analysis. Sex, weight and size may be kept when scientifically required. Remove race, ethnicity, nationality and country values, as they increase re-identification risk and are not needed for image interpretation. For any field considered for retention, ensure consistency with the clinical dataset and document a risk assessment justifying its inclusion.
- **Patient information (group 0010):** contains demographic and identity-related attributes, most of which are direct identifiers or high-risk quasi-identifiers. Aside from the fields explicitly addressed above, all other attributes in this group should be removed. These include patient address, telephone numbers, ethnic group, occupation, comments, and other descriptive fields that may reveal personal or contextual information about the subject.
- **Site ID:** When present (typically stored under the Clinical Trial Site ID (0012,0030) tag), replace with an anonymised site identifier generated using the same salted-hash logic applied to Patient ID. The resulting value should be irreversible and consistent for each site across the dataset.
- **Unique identifiers (UIDs)** (e.g. Study Instance UID, Series Instance UID, SOP Instance UID): Replace with newly generated, DICOM-compliant UIDs using a deterministic method to preserve linkages between related objects. New UIDs should be constructed under a valid, registered UID root, with a reproducible hashed or mapped suffix

derived from the original UID. This prevents linkage to source systems while maintaining consistent associations within the anonymised dataset. SOP Instance UID (0008,0018) must be unique across the dataset.

- **Dates:** Apply a subject-specific date offset, determined from the anonymised Patient ID, and use it consistently across all imaging exams and visits to preserve temporal relationships while masking true calendar dates. All retained DA and DT attributes should be shifted by the same offset within a subject, using a random value constrained to a ±30-day window to ensure both privacy protection and longitudinal consistency.
- **Time:** Time-of-day fields associated with patient workflow events (e.g. Study Time, Series Time, Acquisition Time) should be removed or zeroed, consistent with the DICOM Basic Application Confidentiality Profile, as they may enable linkage to operational logs. Technical timing parameters (e.g. TE, TR, TI, frame timing) that do not represent real-world clock times may be retained. For DT attributes, the date component must be shifted, and the time component removed or retained based on whether it contains identifying information.
- **Accession number and medical record numbers:** Remove or replace with non-identifiable placeholders, as they are direct patient or workflow identifiers and can be used to link imaging exam back to the originating clinical system.
- **Institution and device information fields:** Remove or replace with non-identifiable placeholders, as these attributes reveal the originating site, scanner location, or specific device hardware and can enable linkage back to the hospital or acquisition environment.
- **Free-text fields** (VRs LO, LT, SH, ST, UT): Free-text attributes pose a high risk of containing PHI, including manually entered names, initials, dates, site identifiers, or workflow comments. Because some free-text fields may also contain information useful for downstream scientific analysis, these attributes should undergo a tiered evaluation process:
 1. Apply automated detection using pattern-matching (e.g. regular expressions for dates, names, IDs), NLP-based PHI detection, and OCR when text is embedded in images or burned-in overlays.
 2. Apply targeted Manual Review only for fields marked as potentially retainable after automated analysis.
 3. Retain only when the field contains scientifically necessary information and is verified to be free of personal information.

Otherwise, remove the field or replace its content with a standardised, non-identifiable placeholder.

- **De-identification disclosure attributes:** Retain or populate these attributes to document that anonymisation was performed and to specify the methods applied. These fields contain no personal data and are recommended for transparency, reproducibility and regulatory compliance:
 - Patient Identity Removed (0012,0062)

- o De-identification Method (0012,0063)
- o De-identification Method Code Sequence (0012,0064)

DICOM Attribute Groups with higher risk of personal data disclosure:

- Group 0010 (Patient Information): Contains direct identifiers such as Patient Name, Patient ID, Birth Date, Sex, Weight/Size and patient comments.
- Group 0012 (Clinical Trial Attributes): Includes protocol IDs, site IDs, sponsor information, coordinating centre details and trial-specific subject identifiers.
- Group 0008 (General Study and Series Information): Often contains institution and department names, accession numbers, descriptions, operator and physician names, workflow timestamps, and protocol text.
- Group 0032 (Study Scheduling and Request Information): Includes requesting physician/service, procedure step IDs, request numbers and workflow-related timestamps.
- Group 0040 (Procedure and Observation Workflow): Contains performed procedure details, observer names, report timestamps, coded observations and free-text descriptions.
- Group 0020 (Study/Series/Image Identification): May include Study ID, frame-of-reference identifiers, referencing relationships, and vendor-specific IDs linking back to source systems.

4.2.2. Private tags

As mentioned in Section 2.1.2, DICOM private attributes allow modality vendors, PACS systems and software tools to store proprietary metadata that is not defined in the public DICOM dictionary. Because these attributes are flexible by design, vendors may encode information using standard text-based VRs or more complex formats such as binary arrays, mixed binary/text structures, nested sequences, or proprietary encoded tables. These structures are often undocumented and difficult to interpret without vendor-specific knowledge, and some can contain deeply embedded information not visible in simple header inspection.

Private tags frequently include workflow-related and operational details that may replicate personal information already present in standard tags. Examples include patient or operator names, accession numbers, internal identifiers, timestamps, device serial numbers, institutional descriptors, or free-text protocol comments. In addition, some private blocks mirror standard fields with personal information for internal workflow compatibility, meaning personal data can appear redundantly or in modified form. Because of this variability, lack of transparency and the possibility of encoded personal information, automated sanitisation is unreliable, and thorough manual decoding may not be feasible. For these reasons, **the default recommendation is to remove all private tags during anonymisation**, which is consistent with DICOM PS3.15 and common practice across major research consortia.

Despite these risks, private tags sometimes store *scientifically important acquisition and reconstruction parameters* that are not available in the public dictionary. These may include advanced sequence settings, gradient tables and diffusion directions, slice-

timing information, reconstruction kernels, motion or calibration parameters, PET SUV scaling and correction factors, proprietary sequence descriptors, or software versioning metadata. Such details can be essential for reproducible quantitative imaging workflows, especially in MRI and PET, where analysis pipelines rely on modality-specific metadata that vendors store privately.

Exception process:

In rare cases where specific private tags are requested by researchers (e.g. for algorithm development or analysis) or identified as technically necessary to preserve the scientific usability of the dataset, exceptions may be considered. Such requests will require a formal internal review, handled on a case-by-case basis. Any private tags approved for retention will be subject to:

- Reviewing specific manufacturer **conformance statements**, which provide details on private tag usage and content, helping determine whether a given tag can be kept, modified or removed during anonymisation.
- Consultation of **supporting resources**, including the NEMA “Retain Safe Private Option” ((NEMA), PS3.15 - Retain Safe Private Option, 2025), the TCIA Private Tag Dictionary, and vendor-specific dictionaries available through tools such as DCMTK, Pydicom and GDCM, to determine whether certain private attributes are documented, non-identifying, and potentially safe to retain. These resources help clarify private tag purpose, typical content, and associated risk, and can support evidence-based decisions on whether a private element can be kept, modified, or must be removed.
- When **explicit value representation (VR)** is available, it may also be used to evaluate potential risk (see Appendix A for VR and corresponding PHI risk assessment).
- **Manual** decoding and interpretation by the imaging data team.
- A **targeted** personal data review to assess and remove any identifying content before final inclusion.

4.2.3. Other elements

Sequence elements: Anonymisation must support full-depth traversal of items within each sequence and apply de-identification rules consistently across all levels. This approach is essential for properly anonymising complex objects such as structured reports, radiation therapy plans, and enhanced MR or CT objects, which heavily rely on nested sequences.

Encapsulated documents: Since personal data may be embedded within the document's content rather than in structured metadata, these files must be parsed and treated with the same level of scrutiny as burned-in annotations (see 4.3.1).

4.3 Pixel data anonymisation

Pixel data within DICOM files can contain embedded personal data, in the form of burned-in annotations, visible identifiers on objects within the field of view, or recognisable facial features.

4.3.1. Burned-in annotation

Burned-in annotation refers to text or graphics embedded directly in image pixels, such as patient names, medical record numbers, dates of birth, institutional identifiers, acquisition timestamps, and information displayed on jewellery, metal plates, wristbands, implants or external devices visible in the image. These require dedicated image processing techniques for detection and redaction.

- **Detection**

Burned-in text and visible identifiers may be detected using:

- o OCR-based methods
- o Vision-language models (VLMs), which may offer improved detection performance in complex imaging scenarios.

Both open-source and commercial solutions may be used, provided they are deployed in an HIPAA-compliant environment, including appropriate infrastructure controls, contractual safeguards, and access restrictions.

- **Redaction**

Detected personal data in the text should be redacted or obfuscated before data sharing, ensuring diagnostic value is preserved while eliminating identifiable information.

Analytically essential burned-in information (exception handling)

In limited cases, burned-in annotations may contain information critical for scientific interpretation (e.g. timestamps in ophthalmology time-series images generated from non-DICOM source formats such as JPG). In such cases, a **controlled exception process** must be applied, including:

- o Scientific justification for retention
- o Formal privacy risk review
- o Additional safeguarding measures
- o Explicit documentation of the decision.

Machine registration marks

Redaction workflows must preserve machine-generated graphical overlays that are essential for image interpretation, such as **laterality markers, scan plane indicators, scale bars and orientation labels**, unless these overlays themselves contain personal identifiers.

Residual risk after burned-in text redaction

As part of our evaluation, we reviewed the potential for recovering or reconstructing text information that has been redacted or obscured in DICOM pixel data.

- Our assessment indicates that when opaque masking or pixel replacement methods are applied correctly, the likelihood of reversing or revealing the original burned-in text is extremely low.
- Techniques such as blurring or pixelation alone may leave residual patterns that could, in theory, be partially recovered using advanced image enhancement or machine learning methods; therefore, these approaches are not recommended.

- Overall, proper opaque redaction – validated through visual and OCR-based checks – reduces the risk of re-identification to a negligible level. (Hill S., 2016) (Ho N.Z-Y., 2008) (Zhai H., 2025)

Recommendation for text redaction

Based on current evidence and industry best practices, opaque masking or region replacement should be used for removing burned-in text from DICOM pixel data, as these methods effectively eliminate recoverable signal and minimise re-identification risk, unlike blurring or pixelation techniques.

- **Exclusion**

If reliable redaction is not feasible or introduces complexity, any image found to contain burned-in text should be excluded from external transfer.

Exclusion logs

Any image excluded from external sharing due to burned-in annotations or failure of reliable redaction must be formally documented. Exclusions shall be captured in dataset-level exclusion logs and data transfer reports, including the reason for exclusion. This ensures transparency, traceability and auditability of all removed content.

4.3.2. Anatomical anonymisation

In head and brain imaging modalities such as MRI, CT and PET, detailed craniofacial anatomy can be reconstructed from 2D slices into realistic 3D facial renderings and potentially matched to photographs using facial recognition technologies (Schwarz C.G. & Kremers W.K., 2019) (Mazura J.C., 2012) (Schwarz C.G. & Kremers W.K., 2023).

Under HIPAA's Safe Harbor, "full-face photographs and any comparable images" are direct identifiers (PHI), and the GDPR explicitly defines facial images as biometric data and treats their processing as a special-category activity (*U.S. Department of Health and Human Services, 2025*) (*European Data Protection Board, 2020*). Therefore, removing or obscuring recognisable facial features is the most standards-aligned way of releasing head images as anonymous data: it directly addresses HIPAA's "comparable images" risk and implements DICOM PS3.15's Clean Recognizable Visual Features option to reduce identifiability under GDPR.

A common technique within anatomical anonymisation is defacing, which involves algorithmically removing facial features from imaging data while preserving the integrity of the brain and other regions of clinical interest. Other methods include face removal, skull stripping, template masking, face replacement and face distortion. Newer methods leverage deep learning for more targeted facial feature removal.

Several tools support automated anatomical anonymisation, varying in robustness and data quality preservation. We systematically reviewed the currently available defacing tools, summarising their methodologies, supported modalities, release information, licensing status, and reported advantages and limitations. A comprehensive overview of this assessment is provided in Appendix B.

General limitations of defacing approaches:

- **Modality-specific:** Most tools only support T1-weighted MRI; limited or poor performance on CT, PET or other MR sequences.
- **Over-defacing risk:** Can remove parts of the brain or skull base, affecting downstream analyses.
- **Under-defacing risk:** May leave parts of the face visible, especially with atypical anatomy or low-quality scans.
- **Lack of standardisation:** Different tools vary in aggressiveness and accuracy.
- **Need for manual QC:** Visual inspection is often required for adequate anonymisation.

Defacing tool recommendation: mriface (Mayo Clinic)

is a high-performing facial de-identification tool used by the Alzheimer's Disease Neuroimaging Initiative (ADNI). mriface replaces the face and ears with an average face via non-linear registration, which tends to preserve skull and extra-cranial information better than pure masking. It is recommended as a strong candidate tool because published evaluations have shown favourable performance in reducing facial recognisability while generally preserving brain measurements across several MRI, PET and CT research use cases (Schwarz C.G. & Kremers W.K., 2021) (Schwarz C.G. & Choe M., 2024). However, its impact is not universally negligible across all modalities, sequences,

biomarkers or downstream analysis pipelines. Therefore, mriface or any alternative anatomical anonymisation method should be validated for the specific imaging modality, cohort and intended analysis, particularly when quantitative biomarkers, radiomics, segmentation or AI model development are planned.

Defacing is evolving rapidly alongside re-identification techniques. Implementers are responsible for reviewing current methods and re-evaluating the chosen approach to ensure the chosen method balances reducing re-identification risk with data utility.

4.3.2.1 Defacing data format (NIfTI)

Most of the defacing tools take as an input and produce outputs in Neuroimaging Informatics Technology Initiative (NIfTI) format.

The NIfTI format is a widely used standard for storing medical imaging data. It supports both single-file (.nii) and two-file (.hdr/.img) storage structures, where voxel data is accompanied by a compact, fixed-length header (see Figure 2) containing essential metadata such as image dimensions, voxel sizes, orientation and data type.

```
<class 'nibabel.nifti1.Nifti1Header'> object, endian='<'
sizeof_hdr      : 348
data_type       : b''
db_name         : b''
extents         : 0
session_error   : 0
regular         : b'r'
dim_info        : 57
dim             : [ 4 128 96 24 2 1 1 1]
intent_p1       : 0.0
intent_p2       : 0.0
intent_p3       : 0.0
intent_code     : none
datatype        : int16
bitpix         : 16
slice_start     : 0
pixdim          : [ -1.      2.      2.      2.2 2000.      1.      1.      1. ]
vox_offset      : 0.0
scl_slope       : nan
scl_inter       : nan
slice_end       : 23
slice_code      : unknown
xyzt_units      : 10
cal_max         : 1162.0
cal_min         : 0.0
slice_duration  : 0.0
toffset        : 0.0
glmax           : 0
glmin           : 0
descrip         : b'FSL3.3\x00 v2.25 NIfTI-1 Single file format'
aux_file        : b''
qform_code      : scanner
sform_code      : scanner
quatern_b       : -1.94510681403e-26
quatern_c       : -0.996708512306
quatern_d       : -0.081068739295
qoffset_x       : 117.855102539
qoffset_y       : -35.7229423523
qoffset_z       : -7.24879837036
srow_x          : [ -2.      0.      0.      117.86]
srow_y          : [ -0.      1.97  -0.36 -35.72]
srow_z          : [ 0.      0.32  2.17 -7.25]
intent_name     : b''
magic           : b'n+1'
```

Figure 2. Example of an NIfTI file header

Compared to DICOM, NIfTI has a very limited and fixed set of metadata fields that are not dedicated to storing personal data. However, certain free-text fields – notably descrip (80 characters), auxfile (24 characters) and intentname (16 characters) – can inadvertently contain personal data if values are copied directly from DICOM headers during format conversion. Therefore, **NIfTI headers should be reviewed to ensure no inadvertent identifiers are present.**

Where retention of specific DICOM fields is necessary, this can be addressed by:

- 1) Keeping the required metadata fields in separate JSON/XML files
- 2) Maintaining a documented mapping between retained metadata and anonymised image files, where required for downstream analysis
- 3) Exporting the defaced data back to DICOM format.

4.4 Folder and file names

In addition to metadata and pixel-level anonymisation, it is critical to inspect directory structures and file names, which may inadvertently include personal data such as patient IDs and patient names.

All directory and file names should be scanned as part of the anonymisation process to identify potential personal data. If any identifiers are present within directory names or file names, they should be removed or hashed using a secure method to preserve linkability without exposing sensitive information.

Our recommendation is to take a cleaner and safer approach by:

- **Standardising** directory structures and file naming conventions (e.g. Site > Patient > Visit > Exams > Files), and then
- Using **anonymised identifiers** for naming directories and files wherever applicable, rather than relying on source-system naming.

This ensures both removal of any residual personal data, and consistent structure for downstream analysis, transfer and automation.

4.5 Anonymisation summary table

Note: This table summarises recommended actions for common sources of identifying information in clinical imaging data. **It is not an exhaustive list** of all possible personal data-containing elements. Even for standard DICOM tags, additional identifiers may exist depending on acquisition protocols, vendor implementations, etc. We strongly recommend applying a comprehensive automated personal data detection approach (e.g. rule-based DICOM tag scanning, NLP for free-text fields, or AI-/ML-based personal data identification) in addition to these actions, followed by manual or semi-automated quality control to validate results.

Category	Field/Item	Action to be taken
Consistency Across Clinical and Imaging Data	N/A	Apply the same anonymisation rules where applicable to preserve analytical linkages while protecting privacy.
DICOM – Standard Tags	Patient ID	Replace with a salted, irreversible SHA-256 hash that is stable per participant and not recoverable; discard hashing key to ensure irreversibility.
	VR 'PN' Tags	Remove or clear all non-patient Person Name fields.
	Patient Name	Replace with the anonymised Patient ID value.
	Patient Birth Date	Remove entirely or set to "00000000"; do not shift or surrogate.
	Patient Age	Retain only in generalised form (e.g. 090Y, or a value from banded ranges determined by risk assessment).
	Demographic Fields	Retain sex/weight/size if needed; remove race, ethnicity, nationality and similar high-risk demographics.
	Patient Information (group 0010)	Remove remaining sensitive patient information.
	Site ID	Replace with a salted, irreversible SHA-256 hash.
	UIDs	Regenerate all UIDs deterministically under a valid UID root.
	Dates (VR 'DA' and 'DT')	Apply a subject-specific, deterministic date shift.
	Time Fields (VR 'TM')	Remove or zero workflow times; retain only non-identifying technical timing.
	Accession Number and MRNs	Remove or replace with non-identifiable placeholders.
	Institution & Device Information	Remove or replace identifying sites and hardware information with generic placeholders.
	Free-Text Fields (LO, LT, SH, ST, UT)	Remove, replace or retain only after a detailed review.
	De-identification Disclosure Tags	Populate required de-identification methods and identity-removed indicators for transparency.
	Other attributes	See Appendix C for details.
DICOM – Private Tags	Private Tags	Remove all private tags by default. Consider exceptions only via formal review using NEMA "Retain Safe Private Option" guidance and manufacturer's conformance statements; perform targeted personal data review before retention.
DICOM – Other Elements	Sequence Elements	Apply full-depth anonymisation rules consistently at all sequence levels.
	Encapsulated Documents	Parse and redact personal data within content, similar to burned-in text handling.
Pixel Data	Burned-in Annotations	Detect and redact/obfuscate personal data in pixel data; exclude images if redaction is not feasible.
	Anatomical Features (Facial Structures)	Apply automated defacing or facial feature removal; perform QC to ensure adequacy.
	Defacing Output (NIfTI format)	Minimal metadata risk; still verify that metadata, filenames, directory structures, and image content contain no personal data.
File System	Folder & File Names	Scan for personal data; remove or secure hash as needed. The recommendation is to standardise directory structures and use anonymised identifiers for naming files and folders.

4.6 Re-identification risk score assessment

Re-identification refers to the ability to associate anonymised data with a specific individual, either through remaining identifiers, combinations of quasi-identifiers, or linkage to external information sources. Imaging data presents distinct re-identification risks due to both metadata identifiers and the recognisability of facial anatomy or other unique anatomical features.

To mitigate these risks, our anonymisation framework removes or neutralises direct identifiers across DICOM standard and private tags, eliminates burnt-in text from pixel data, applies robust anatomical defacing to remove facial structure, and generalises or suppresses quasi-identifiers such as age, clinical attributes, protocol details and acquisition-site information where necessary. These measures significantly reduce the potential for linking an image or metadata record back to an individual and prevent reconstruction of obvious identifying features.

Because some demographic, clinical and anatomical information must remain to preserve scientific validity, re-identification risk cannot be reduced to zero. Residual risk may arise from combinations of quasi-identifiers, rare conditions, uncommon anatomical features, or patterns of longitudinal imaging that cannot be fully masked without degrading data utility. Aligned with GDPR Recital 26 and HIPAA Expert Determination, our objective is therefore to reduce risk to a very small level that is not reasonably likely to occur, particularly given that data is shared only through a secure, access-controlled platform with contractual, technical and organisational safeguards that further limit the possibility of external linkage.

5. Supporting files

Readme file with anonymisation details:

A short document summarising the anonymisation approach applied to the dataset. It includes a description of key transformations (e.g. hashing rules, date-shifting, removed metadata categories) and any study-specific considerations needed for correct interpretation of the anonymised imaging data.

Dataset inventory file:

A structured listing of all imaging files included in the transfer, along with essential metadata such as anonymised subject identifiers, modality, series, and file names/paths. This file enables recipients to understand dataset composition and verify completeness.

Exclusion logs file:

A record of all images or series excluded from the dataset prior to transfer. Each entry documents the file path, the reason for exclusion (e.g. burned-in text that could not be safely redacted) and any relevant notes. This provides transparency, traceability and auditability for all omitted content.

6. Disclaimer

The opinions expressed in this document are those of the authors and should not be construed to represent the opinions of PHUSE members, respective companies/organisations or regulators' views or policies. The content in this document should not be interpreted as a data standard and/or information required by regulatory authorities.

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Appendix A. DICOM value representations and personal information/PHI risk overview

VR Name	VR Full Name	Description	Data Type	PHI Risk
AE	Application Entity	Application titles	string	Low
AS	Age String	Patient age (e.g. 032Y)	string with formatting	Medium
AT	Attribute Tag	Tag references	ordered pair of 16-bit unsigned integers	Low
CS	Code String	Enumerated values (e.g. "M", "YES")	string	Low
DA	Date	Calendar date (YYYYMMDD)	string with formatting	Medium
DS	Decimal String	Floating point numbers in string format	<u>fixed or floating point</u> number	Low
DT	Date Time	Combined date/time	string with formatting	Medium
FL	Floating Point Single	32-bit float	floating point number	Low
FD	Floating Point Double	64-bit float	floating point number	Low
IS	Integer String	Integers in string format	Integer	Low
LO	Long String	Text up to 64 chars	string	High
LT	Long Text	Longer blocks of text	string	High
OB	Other Byte	Binary data	octet-stream	Medium
OD	Other Double	Binary double-precision	A stream of 64-bit floating point words	Low
OF	Other Float	Binary float	A stream of 32-bit floating point words	Low
OL	Other Long	Binary 32-bit integers	A stream of 32-bit words	Medium
OV	Other 64-bit Very Long	Binary 64-bit values	A stream of 64-bit words	Medium
OW	Other Word	Binary 16-bit values	A stream of 16-bit words	High
PN	Person Name	Structured name fields	string	High
SH	Short String	Text up to 16 chars	string	Medium
SL	Signed Long	32-bit signed integer	Integer	Low
SQ	Sequence of Items	Nested datasets	dataset	Medium
SS	Signed Short	16-bit signed integer	Signed binary integer 16 bits long	Low
ST	Short Text	Limited free text	string	High
SV	Signed 64-bit Very Long	64-bit signed integer	Signed binary integer 64 bits long	Low
TM	Time	Time of day	string with formatting	Medium
UC	Unlimited Characters	Unicode text	string	High
UI	Unique Identifier (UID)	UIDs	string UID	Medium
UL	Unsigned Long	32-bit unsigned integer	Unsigned binary integer 32 bits long	Low
UN	Unknown	Unknown encoding	octet-stream with unknown encoding	Medium
UR	URI/URL	Web addresses or resource identifiers	string URI/URL	High
US	Unsigned Short	16-bit unsigned integer	Unsigned binary integer 16 bits long	Low
UT	Unlimited Text	Very long free text	string	High
UV	Unsigned 64-bit Very Long	64-bit unsigned integer	Unsigned binary integer 64 bits long	Low

Appendix B. Defacing tools overview

Tool	Authors	Year (Latest Update)	Modalities	Method	Used by	Free/Open source	Pros	Cons
mri_reface	Christopher G. Schwarz et al. (Mayo Clinic)	2024	MRI (T1/T2/T2*/FLAIR/ASL), CT, PET (amyloid, tau, FDG)	Face warping	ADNI-4	Free for non-commercial use, not fully open-source code	Strong validation; multi-modality; minimises impact; active support	Requires MATLAB Runtime and ANTs; closed-source core
MiDeFace	FreeSurfer/ Martinos Center team	2023	MRI (primarily T1w)	Face warping	Recent papers, growing adoption	Free for non-commercial use (FreeSurfer)	Integrated in FreeSurfer; minimal brain impact; QC renders	Newer tool; validation literature still maturing
PyDeface	Poldrack Lab & contributors	2022	MRI (primarily T1w)	Face removal	Common in BIDS and OpenNeuro; ReMIND (TCIA)	Yes (MIT)	Simple CLI; open source; widely used; easy to install	May remove more neck/nose; ears retained; can affect certain measures
FSL Deface	FMRIB (Oxford)	2025	MRI (primarily T1/T2)	Face removal	UK Biobank	Free for non-commercial use (FSL deface)	Well maintained; widely deployed	Can impact on some morphometry metrics
SPM Deface	SPM team/ community	2014?	MRI (primarily T1w)	Face removal	OpenNeuro datasets	Yes (GPL)	MATLAB/SPM environments; reproducible; scriptable	MATLAB dependency; may preserve ears; variable impact on measures
AFNI refacer	AFNI team (NIMH)	2020	MRI (primarily T1w)	Face removal or warping	AFNI-centric labs	Yes (GPL)	Header anonymisation option	Reference masks may need tuning
PyFaceWipe	Patel et al. (2024)	2024	MRI (primarily T1w)	Face removal	Early adopters, method paper	Unknown	Promising metrics across contrasts	New; ecosystem/ community still small; no public code repository available yet
mrdefacer	Michael Hanke et al.	2018	MRI (primarily T1w)	Face warping	Used via BIDSonym; NeuroDebian packaging	Yes (GPL)	Flexible; can just generate masks for custom workflows	More DIY; fewer turnkey safeguards/ QC
DeepDefacer	Khazane et al. (with Poldrack Lab)	2022	MRI (primarily T1w)	Face removal	Research studies, hackathons	Yes (MIT)	Fast inference; GPU-enabled; potentially adaptable	Reproducibility/ maintenance vary; generalisation risks; fewer production deployments
Face Masking (mask_face)	XNAT Team (WUSTL)	2018	MRI (primarily T1w), CT	Face blurring	XNAT, HCP, GSP	Yes	Integrates with XNAT; minimally invasive	Tied to XNAT; fewer standalone features
mri_deface	Bischoff-Grethe, Fischl et al. (FreeSurfer/ MBIRN)	2016	MRI (primarily T1w)	Face removal	Older pipelines, evaluated widely	Free for non-commercial use (FreeSurfer)	Established; simple CLI; fast	Legacy; can affect downstream metrics more than newer methods
Quickshear	Schimke et al.	2017	MRI (primarily T1w)	Face removal	Older pipelines, educational comparisons	Yes (open source, legacy)	Simple; transparent; fast	Reported higher failure/brain erosion rates; legacy
AnonyMI	Mikulan et al.	2025	MRI (primarily T1w)	Face removal	Replication/ evaluation studies	Yes (3D Slicer)	GUI-friendly; preserves geometry for EEG/MEG co-registration	Workflow inside Slicer
BRAINS Deface	BRAINS/Slicer community	2020	MRI (T1/T2)	Face removal	Institutional pipelines	Yes (Apache-2.0)	Flexible; visible control over region removed	Manual landmarks
Deface	Biolmage Suite Web (Yale)	2020	MRI (primarily T1w)	Face removal	Small lab workflows	Yes (Apache-2.0)	Very easy UI; web-based	Fewer publications; limited batch/QA features
HD-BET	Isensee et al. (DKFZ/Uni Bonn/ Heidelberg)	2024	MRI (T1/T2/ FLAIR)	Brain extraction	Used for skull stripping	Yes (Apache-2.0)	Accurate, fast DL stripper; open source	Not a defacer
BrainSuite BSE	BrainSuite team (USC)	2024	MRI (primarily T1w)	Brain extraction	Legacy pipelines, education	Free for non-commercial use	Interactive control; established	Not a defacer
FSL-BET	Stephen M. Smith/ FMRIB (Oxford)	2002	MRI (primarily T1w)	Brain extraction	UK Biobank pipelines	Free for non-commercial use (FSL deface)	Fast; battle-tested; widely available	Not a defacer

Appendix C. Recommended anonymisation actions for standard tags

This appendix provides recommended actions **for a selected subset** of standard DICOM attributes. It is not an exhaustive list of all DICOM tags. Instead, it focuses on:

1. Attributes that present a higher risk of re-identification, such as those containing direct or quasi-identifiers
2. Attributes that are commonly required for downstream analysis or scientific use.

For each attribute, the recommended Action (e.g. Keep, Remove, Replace, Hash) reflects both its privacy risk and its analytical relevance.

Important note on the “Remove” action:

When the action is listed as Remove, the implementation depends on the DICOM Attribute Type:

- Type 1 or Type 2 attributes:
These attributes must be present in the dataset according to the DICOM standard. In such cases, the value must be cleared (set to an empty value) rather than removing the attribute entirely.
- Type 3 attributes:
These attributes are optional and may be safely deleted from the dataset when flagged for removal.

This distinction ensures anonymisation does not violate DICOM conformance.

For more details on certain attributes, groups and actions, see Section 4.2.1.

Tag	Name	VR	Action
(0000,1000)	Affected SOP Instance UID	UI	Remove
(0000,1001)	Requested SOP Instance UID	UI	Replace UID
(0002,0001)	File Meta Information Version	OB	Keep
(0002,0003)	Media Storage SOP Instance UID	UI	Replace UID
(0004,1511)	Referenced SOP Instance UID in File	UI	Replace UID
(0008,0012)	Instance Creation Date	DA	Date shift
(0008,0013)	Instance Creation Time	TM	Remove
(0008,0014)	Instance Creator UID	UI	Replace UID
(0008,0015)	Instance Coercion DateTime	DT	Remove
(0008,0017)	Acquisition UID	UI	Replace UID
(0008,0018)	SOP Instance UID	UI	Replace UID
(0008,0019)	Pyramid UID	UI	Replace UID
(0008,0020)	Study Date	DA	Date shift
(0008,0021)	Series Date	DA	Date shift
(0008,0022)	Acquisition Date	DA	Date shift
(0008,0023)	Content Date	DA	Date shift
(0008,0024)	Overlay Date	DA	Remove
(0008,0025)	Curve Date	DA	Remove
(0008,002A)	Acquisition DateTime	DT	Remove
(0008,0030)	Study Time	TM	Keep
(0008,0031)	Series Time	TM	Remove
(0008,0032)	Acquisition Time	TM	Remove
(0008,0033)	Content Time	TM	Keep
(0008,0034)	Overlay Time	TM	Remove

(0008,0035)	Curve Time	TM	Remove
(0008,0050)	Accession Number	SH	Remove
(0008,0051)	Issuer of Accession Number Sequence	SQ	Remove
(0008,0054)	Retrieve AE Title	AE	Remove
(0008,0055)	Station AE Title	AE	Remove
(0008,0058)	Failed SOP Instance UID List	UI	Replace UID
(0008,0070)	Manufacturer	LO	Keep
(0008,0080)	Institution Name	LO	Remove
(0008,0081)	Institution Address	ST	Remove
(0008,0082)	Institution Code Sequence	SQ	Remove
(0008,0090)	Referring Physician's Name	PN	Remove
(0008,0092)	Referring Physician's Address	ST	Remove
(0008,0094)	Referring Physician's Telephone Numbers	SH	Remove
(0008,0096)	Referring Physician's Identification Sequence	SQ	Remove
(0008,009C)	Consulting Physician's Name	PN	Remove
(0008,009D)	Consulting Physician's Identification Sequence	SQ	Remove
(0008,0104)	Code Meaning	LO	Check
(0008,0106)	Context Group Version	DT	Date shift
(0008,0107)	Context Group Local Version	DT	Date shift
(0008,010D)	Context Group Extension Creator UID	UI	Replace UID
(0008,0110)	Coding Scheme Identification Sequence	SQ	Keep
(0008,0112)	Coding Scheme Registry	LO	Keep
(0008,0114)	Coding Scheme External ID	ST	Keep
(0008,0115)	Coding Scheme Name	ST	Keep
(0008,0116)	Coding Scheme Responsible Organization	ST	Keep
(0008,0201)	Timezone Offset From UTC	SH	Remove
(0008,1000)	Network ID	AE	Remove
(0008,1010)	Station Name	SH	Remove
(0008,1030)	Study Description	LO	Check or replace with visit label
(0008,1032)	Procedure Code Sequence	SQ	Remove
(0008,103E)	Series Description	LO	Check
(0008,1040)	Institutional Department Name	LO	Remove
(0008,1041)	Institutional Department Type Code Sequence	SQ	Remove
(0008,1048)	Physician(s) of Record	PN	Remove
(0008,1049)	Physician(s) of Record Identification Sequence	SQ	Remove
(0008,1050)	Performing Physician's Name	PN	Remove
(0008,1052)	Performing Physician's Identification Sequence	SQ	Remove
(0008,1060)	Name of Physician's Reading Study	PN	Remove
(0008,1062)	Physician's Reading Study Identification Sequence	SQ	Remove
(0008,1070)	Operator's Name	PN	Remove
(0008,1072)	Operator's Identification Sequence	SQ	Remove

(0008,1080)	Admitting Diagnoses Description	LO	Remove
(0008,1084)	Admitting Diagnoses Code Sequence	SQ	Remove
(0008,1088)	Pyramid Description	LO	Remove
(0008,1090)	Manufacturer's Model Name	LO	Keep
(0008,1110)	Referenced Study Sequence	SQ	Remove
(0008,1111)	Referenced Performed Procedure Step Sequence	SQ	Remove
(0008,1115)	Referenced Series Sequence	SQ	Keep
(0008,1120)	Referenced Patient Sequence	SQ	Remove
(0008,1125)	Referenced Visit Sequence	SQ	Keep
(0008,1130)	Referenced Overlay Sequence	SQ	Keep
(0008,113A)	Referenced Waveform Sequence	SQ	Keep
(0008,1140)	Referenced Image Sequence	SQ	Remove
(0008,114A)	Referenced Instance Sequence	SQ	Keep
(0008,114B)	Referenced Real World Value Mapping Instance Sequence	SQ	Keep
(0008,1150)	Referenced SOP Class UID	UI	Replace UID
(0008,1155)	Referenced SOP Instance UID	UI	Replace UID
(0008,1164)	Frame Extraction Sequence	SQ	Keep
(0008,1195)	Transaction UID	UI	Replace UID
(0008,1198)	Failed SOP Sequence	SQ	Keep
(0008,1199)	Referenced SOP Sequence	SQ	Keep
(0008,1200)	Studies Containing Other Referenced Instances Sequence	SQ	Keep
(0008,1250)	Related Series Sequence	SQ	Keep
(0008,1301)	Principal Diagnosis Code Sequence	SQ	Remove
(0008,1302)	Primary Diagnosis Code Sequence	SQ	Remove
(0008,1303)	Secondary Diagnoses Code Sequence	SQ	Remove
(0008,1304)	Histological Diagnoses Code Sequence	SQ	Remove
(0008,2111)	Derivation Description	ST	Remove
(0008,2112)	Source Image Sequence	SQ	Remove
(0008,2135)	Event Code Sequence	SQ	Remove
(0008,3010)	Irradiation Event UID	UI	Replace UID
(0008,4000)	Identifying Comments	LT	Remove
(0008,9123)	Creator Version UID	UI	Replace UID
(0010,0010)	Patient's Name	PN	Replace with anonymised Patient ID value
(0010,0011)	Person Names to Use Sequence	SQ	Remove
(0010,0012)	Name to Use	LT	Remove
(0010,0013)	Name to Use Comment	UT	Remove
(0010,0014)	Third Person Pronouns Sequence	SQ	Remove
(0010,0015)	Pronoun Code Sequence	SQ	Remove
(0010,0016)	Pronoun Comment	UT	Remove
(0010,0020)	Patient ID	LO	Hash

(0010,0021)	Issuer of Patient ID	LO	Remove
(0010,0030)	Patient's Birth Date	DA	Remove
(0010,0032)	Patient's Birth Time	TM	Remove
(0010,0040)	Patient's Sex	CS	Keep
(0010,0041)	Gender Identity Sequence	SQ	Remove
(0010,0042)	Sex Parameters for Clinical Use Category Comment	UT	Remove
(0010,0043)	Sex Parameters for Clinical Use Category Sequence	SQ	Remove
(0010,0044)	Gender Identity Code Sequence	SQ	Remove
(0010,0045)	Gender Identity Comment	UT	Remove
(0010,0046)	Sex Parameters for Clinical Use Category Code Sequence	SQ	Remove
(0010,0047)	Sex Parameters for Clinical Use Category Reference	UR	Remove
(0010,0050)	Patient Insurance Plan Code Sequence	SQ	Remove
(0010,0101)	Patient's Primary Language Code Sequence	SQ	Remove
(0010,0102)	Patient's Primary Language Modifier Code Sequence	SQ	Remove
(0010,1000)	Other Patient IDs	LO	Remove
(0010,1001)	Other Patient Names	PN	Remove
(0010,1002)	Other Patient IDs Sequence	SQ	Remove
(0010,1005)	Patient's Birth Name	PN	Remove
(0010,1010)	Patient's Age	AS	Replace with generalised value
(0010,1020)	Patient's Size	DS	Keep
(0010,1030)	Patient's Weight	DS	Keep
(0010,1040)	Patient's Address	LO	Remove
(0010,1050)	Insurance Plan Identification	LO	Remove
(0010,1060)	Patient's Mother's Birth Name	PN	Remove
(0010,1080)	Military Rank	LO	Remove
(0010,1081)	Branch of Service	LO	Remove
(0010,1090)	Medical Record Locator	LO	Remove
(0010,1100)	Referenced Patient Photo Sequence	SQ	Remove
(0010,2000)	Medical Alerts	LO	Remove
(0010,2110)	Allergies	LO	Remove
(0010,2150)	Country of Residence	LO	Remove
(0010,2152)	Region of Residence	LO	Remove
(0010,2154)	Patient's Telephone Numbers	SH	Remove
(0010,2155)	Patient's Telecom Information	LT	Remove
(0010,2160)	Ethnic Group	SH	Remove
(0010,2161)	Ethnic Group Code Sequence	SQ	Remove
(0010,2162)	Ethnic Groups	UC	Remove
(0010,2180)	Occupation	SH	Remove
(0010,21A0)	Smoking Status	CS	Remove
(0010,21B0)	Additional Patient History	LT	Remove
(0010,21C0)	Pregnancy Status	US	Remove

(0010,21D0)	Last Menstrual Date	DA	Remove
(0010,21F0)	Patient's Religious Preference	LO	Remove
(0010,2203)	Patient Sex Neutered	CS	Remove
(0010,2297)	Responsible Person	PN	Remove
(0010,2299)	Responsible Organisation	LO	Remove
(0010,4000)	Patient Comments	LT	Remove
(0012,0010)	Clinical Trial Sponsor Name	LO	Replace with sponsor name
(0012,0020)	Clinical Trial Protocol ID	LO	Replace with study ID
(0012,0021)	Clinical Trial Protocol Name	LO	Replace with study name
(0012,0022)	Clinical Trial Protocol Ethics Committee Name	LO	Remove
(0012,0023)	Other Clinical Trial Protocol IDs Sequence	SQ	Remove
(0012,0030)	Clinical Trial Site ID	LO	Replace with anonymised site ID
(0012,0031)	Clinical Trial Site Name	LO	Replace with anonymised site ID
(0012,0032)	Issuer of Clinical Trial Site ID	LO	Remove
(0012,0040)	Clinical Trial Subject ID	LO	Replace with anonymised patient id
(0012,0041)	Issuer of Clinical Trial Subject ID	LO	Remove
(0012,0042)	Clinical Trial Site Subject Reading ID	LO	Remove
(0012,0043)	Issuer of Clinical Trial Subject Reading ID	LO	Remove
(0012,0050)	Clinical Trial Time Point ID	LO	Check or replace with visit label
(0012,0051)	Clinical Trial Time Point Description	ST	Remove
(0012,0055)	Issuer of Clinical Trial Time Point ID	LO	Remove
(0012,0060)	Clinical Trial Coordinating Center Name	LO	Remove
(0012,0062)	Patient Identity Removed	CS	Redact
(0012,0063)	De-identification Method	LO	Redact
(0012,0064)	De-identification Code Sequence	SQ	Redact
(0012,0071)	Clinical Trial Series ID	LO	Remove
(0012,0072)	Clinical Trial Series Description	LO	Check
(0012,0073)	Issuer of Clinical Trial Series ID	LO	Remove
(0012,0081)	Clinical Trial Protocol Ethics Committee Name	LO	Remove
(0012,0082)	Clinical Trial Protocol Ethics Committee Approval Number	LO	Remove
(0012,0086)	Ethics Committee Approval Effectiveness Start Date	DA	Remove
(0012,0087)	Ethics Committee Approval Effectiveness End Date	DA	Remove
(0014,407C)	Calibration Time	TM	Remove
(0014,407E)	Calibration Date	DA	Remove
(0016,002B)	Maker Note	OB	Remove
(0016,004B)	Device Setting Description	OB	Remove
(0016,004D)	Camera Owner Name	UT	Remove
(0016,004E)	Lens Specification	DS	Remove
(0016,004F)	Lens Make	UT	Remove
(0016,0050)	Lens Model	UT	Remove
(0016,0051)	Lens Serial Number	UT	Remove

(0016,0070)	GPS Version ID	OB	Remove
(0016,0071)	GPS Latitude Ref	CS	Remove
(0016,0072)	GPS Latitude	DS	Remove
(0016,0073)	GPS Longitude Ref	CS	Remove
(0016,0074)	GPS Longitude	DS	Remove
(0016,0075)	GPS Altitude Ref	US	Remove
(0016,0076)	GPS Altitude	DS	Remove
(0016,0077)	GPS Time Stamp	DT	Remove
(0016,0078)	GPS Satellites	UT	Remove
(0016,0079)	GPS Status	CS	Remove
(0016,007A)	GPS Measure Mode	CS	Remove
(0016,007B)	GPS DOP	DS	Remove
(0016,007C)	GPS Speed Ref	CS	Remove
(0016,007D)	GPS Speed	DS	Remove
(0016,007E)	GPS Track Ref	CS	Remove
(0016,007F)	GPS Track	DS	Remove
(0016,0080)	GPS Img Direction Ref	CS	Remove
(0016,0081)	GPS Img Direction	DS	Remove
(0016,0082)	GPS Map Datum	UT	Remove
(0016,0083)	GPS Dest Latitude Ref	CS	Remove
(0016,0084)	GPS Dest Latitude	DS	Remove
(0016,0085)	GPS Dest Longitude Ref	CS	Remove
(0016,0086)	GPS Dest Longitude	DS	Remove
(0016,0087)	GPS Dest Bearing Ref	CS	Remove
(0016,0088)	GPS Dest Bearing	DS	Remove
(0016,0089)	GPS Dest Distance Ref	CS	Remove
(0016,008A)	GPS Dest Distance	DS	Remove
(0016,008B)	GPS Processing Method	OB	Remove
(0016,008C)	GPS Area Information	OB	Remove
(0016,008D)	GPS Date Stamp	DT	Remove
(0016,008E)	GPS Differential	IS	Remove
(0018,0010)	Contrast/Bolus Agent	LO	Check
(0018,0027)	Intervention Drug Stop Time	TM	Remove
(0018,0035)	Intervention Drug Start Time	TM	Remove
(0018,1000)	Device Serial Number	LO	Remove
(0018,1002)	Device UID	UI	Replace UID
(0018,1004)	Plate ID	LO	Remove
(0018,1005)	Generator ID	LO	Remove
(0018,1007)	Cassette ID	LO	Remove
(0018,1008)	Gantry ID	LO	Remove
(0018,1009)	Unique Device Identifier	UT	Remove

(0018,100A)	UDI Sequence	SQ	Remove
(0018,100B)	Manufacturer's Device Class UID	UI	Replace UID
(0018,1012)	Date of Secondary Capture	DA	Remove
(0018,1014)	Time of Secondary Capture	TM	Remove
(0018,1020)	Software Versions	LO	Keep
(0018,1030)	Protocol Name	LO	Remove
(0018,1042)	Contrast/Bolus Start Time	TM	Remove
(0018,1043)	Contrast/Bolus Stop Time	TM	Remove
(0018,1072)	Radiopharmaceutical Start Time	TM	Remove
(0018,1073)	Radiopharmaceutical Stop Time	TM	Remove
(0018,1078)	Radiopharmaceutical Start DateTime	DT	Remove
(0018,1079)	Radiopharmaceutical Stop DateTime	DT	Remove
(0018,11BB)	Acquisition Field of View Label	LO	Remove
(0018,1200)	Date of Last Calibration	DA	Remove
(0018,1201)	Time of Last Calibration	TM	Remove
(0018,1202)	DateTime of Last Calibration	DT	Remove
(0018,1203)	Calibration DateTime	DT	Date shift
(0018,1204)	Date of Manufacture	DA	Remove
(0018,1205)	Date of Installation	DA	Remove
(0018,1400)	Acquisition Device Processing Description	LO	Remove
(0018,2042)	Target UID	UI	Replace UID
(0018,4000)	Acquisition Comments	LT	Remove
(0018,5011)	Transducer Identification Sequence	SQ	Remove
(0018,700A)	Detector ID	SH	Remove
(0018,700C)	Date of Last Detector Calibration	DA	Remove
(0018,700E)	Time of Last Detector Calibration	TM	Remove
(0018,9074)	Frame Acquisition DateTime	DT	Date shift
(0018,9151)	Frame Reference DateTime	DT	Date shift
(0018,9185)	Respiratory Motion Compensation Technique Description	ST	Remove
(0018,9367)	X-Ray Source ID	UC	Remove
(0018,9369)	Source Start DateTime	DT	Date shift
(0018,936A)	Source End DateTime	DT	Date shift
(0018,9371)	X-Ray Detector ID	UC	Remove
(0018,9373)	X-Ray Detector Label	ST	Remove
(0018,937B)	Multi-energy Acquisition Description	UT	Remove
(0018,937F)	Decomposition Description	UT	Remove
(0018,9424)	Acquisition Protocol Description	LT	Remove
(0018,9516)	Start Acquisition DateTime	DT	Remove
(0018,9517)	End Acquisition DateTime	DT	Remove
(0018,9623)	Functional Sync Pulse	DT	Date shift
(0018,9701)	Decay Correction DateTime	DT	Date shift

(0018,9804)	Exclusion Start DateTime	DT	Date shift
(0018,9919)	Instruction Performed DateTime	DT	Date shift
(0018,9937)	Requested Series Description	LO	Remove
(0018,A002)	Contribution DateTime	DT	Remove
(0018,A003)	Contribution Description	ST	Remove
(0020,000D)	Study Instance UID	UI	Replace UID
(0020,000E)	Series Instance UID	UI	Replace UID
(0020,0010)	Study ID	SH	Replace with Study ID
(0020,0027)	Pyramid Label	LO	Remove
(0020,0052)	Frame of Reference UID	UI	Replace UID
(0020,0200)	Synchronization Frame of Reference UID	UI	Replace UID
(0020,3401)	Modifying Device ID	CS	Remove
(0020,3403)	Modified Image Date	DA	Remove
(0020,3404)	Modifying Device Manufacturer	LO	Remove
(0020,3405)	Modified Image Time	TM	Remove
(0020,3406)	Modified Image Description	LO	Remove
(0020,4000)	Image Comments	LT	Remove
(0020,9158)	Frame Comments	LT	Remove
(0020,9161)	Concatenation UID	UI	Replace UID
(0020,9164)	Dimension Organization UID	UI	Replace UID
(0028,1199)	Palette Color Lookup Table UID	UI	Replace UID
(0028,1214)	Large Palette Color Lookup Table UID	UI	Replace UID
(0028,4000)	Image Presentation Comments	LT	Remove
(0032,0012)	Study ID Issuer	LO	Remove
(0032,0032)	Study Verified Date	DA	Remove
(0032,0033)	Study Verified Time	TM	Remove
(0032,0034)	Study Read Date	DA	Remove
(0032,0035)	Study Read Time	TM	Remove
(0032,1000)	Scheduled Study Start Date	DA	Remove
(0032,1001)	Scheduled Study Start Time	TM	Remove
(0032,1010)	Scheduled Study Stop Date	DA	Remove
(0032,1011)	Scheduled Study Stop Time	TM	Remove
(0032,1020)	Scheduled Study Location	LO	Remove
(0032,1021)	Scheduled Study Location AE Title	AE	Remove
(0032,1030)	Reason for Study	LO	Remove
(0032,1032)	Requesting Physician	PN	Remove
(0032,1033)	Requesting Service	LO	Remove
(0032,1040)	Study Arrival Date	DA	Remove
(0032,1041)	Study Arrival Time	TM	Remove
(0032,1050)	Study Completion Date	DA	Remove
(0032,1051)	Study Completion Time	TM	Remove

(0032,1060)	Requested Procedure Description	LO	Remove
(0032,1066)	Reason for Visit	UT	Remove
(0032,1067)	Reason for Visit Code Sequence	SQ	Remove
(0032,1070)	Requested Contrast Agent	LO	Remove
(0032,4000)	Study Comments	LT	Remove
(0034,0001)	Flow Identifier Sequence	SQ	Remove
(0034,0002)	Flow Identifier	OB	Remove
(0034,0005)	Source Identifier	OB	Remove
(0034,0007)	Frame Origin Timestamp	OB	Replace
(0038,0004)	Referenced Patient Alias Sequence	SQ	Remove
(0038,0010)	Admission ID	LO	Remove
(0038,0011)	Issuer of Admission ID	LO	Remove
(0038,0014)	Issuer of Admission ID Sequence	SQ	Remove
(0038,001A)	Scheduled Admission Date	DA	Remove
(0038,001B)	Scheduled Admission Time	TM	Remove
(0038,001C)	Scheduled Discharge Date	DA	Remove
(0038,001D)	Scheduled Discharge Time	TM	Remove
(0038,001E)	Scheduled Patient Institution Residence	LO	Remove
(0038,0020)	Admitting Date	DA	Remove
(0038,0021)	Admitting Time	TM	Remove
(0038,0030)	Discharge Date	DA	Remove
(0038,0032)	Discharge Time	TM	Remove
(0038,0040)	Discharge Diagnosis Description	LO	Remove
(0038,0050)	Special Needs	LO	Remove
(0038,0060)	Service Episode ID	LO	Remove
(0038,0061)	Issuer of Service Episode ID	LO	Remove
(0038,0062)	Service Episode Description	LO	Remove
(0038,0064)	Issuer of Service Episode ID Sequence	SQ	Remove
(0038,0300)	Current Patient Location	LO	Remove
(0038,0400)	Patient's Institution Residence	LO	Remove
(0038,0500)	Patient State	LO	Remove
(0038,0502)	Patient Clinical Trial Participation Sequence	SQ	Remove
(0038,4000)	Visit Comments	LT	Remove
(003A,0310)	Multiplex Group UID	UI	Replace UID
(003A,0314)	Impedance Measurement DateTime	DT	Date shift
(003A,0329)	Waveform Filter Description	ST	Remove
(003A,032B)	Filter Lookup Table Description	ST	Remove
(0040,0001)	Scheduled Station AE Title	AE	Remove
(0040,0002)	Scheduled Procedure Step Start Date	DA	Remove
(0040,0003)	Scheduled Procedure Step Start Time	TM	Remove
(0040,0004)	Scheduled Procedure Step End Date	DA	Remove

(0040,0005)	Scheduled Procedure Step End Time	TM	Remove
(0040,0006)	Scheduled Performing Physician's Name	PN	Remove
(0040,0007)	Scheduled Procedure Step Description	LO	Remove
(0040,0009)	Scheduled Procedure Step ID	SH	Remove
(0040,000B)	Scheduled Performing Physician Identification Sequence	SQ	Remove
(0040,0010)	Scheduled Station Name	SH	Remove
(0040,0011)	Scheduled Procedure Step Location	SH	Remove
(0040,0012)	Pre-Medication	LO	Remove
(0040,0241)	Performed Station AE Title	AE	Remove
(0040,0242)	Performed Station Name	SH	Remove
(0040,0243)	Performed Location	SH	Remove
(0040,0244)	Performed Procedure Step Start Date	DA	Remove
(0040,0245)	Performed Procedure Step Start Time	TM	Remove
(0040,0250)	Performed Procedure Step End Date	DA	Remove
(0040,0251)	Performed Procedure Step End Time	TM	Remove
(0040,0253)	Performed Procedure Step ID	SH	Remove
(0040,0254)	Performed Procedure Step Description	LO	Remove
(0040,0255)	Performed Procedure Type Description	LO	Remove
(0040,0275)	Request Attributes Sequence	SQ	Remove
(0040,0280)	Comments on the Performed Procedure Step	ST	Remove
(0040,0310)	Comments on Radiation Dose	ST	Remove
(0040,050A)	Specimen Accession Number	LO	Remove
(0040,0512)	Container Identifier	LO	Remove
(0040,0513)	Issuer of the Container Identifier Sequence	SQ	Remove
(0040,051A)	Container Description	LO	Remove
(0040,0551)	Specimen Identifier	LO	Remove
(0040,0554)	Specimen UID	UI	Replace UID
(0040,0555)	Acquisition Context Sequence	SQ	Remove
(0040,0562)	Issuer of the Specimen Identifier Sequence	SQ	Remove
(0040,0600)	Specimen Short Description	LO	Remove
(0040,0602)	Specimen Detailed Description	UT	Remove
(0040,0610)	Specimen Preparation Sequence	SQ	Remove
(0040,06FA)	Slide Identifier	LO	Remove
(0040,1001)	Requested Procedure ID	SH	Remove
(0040,1002)	Reason for the Requested Procedure	LO	Remove
(0040,1004)	Patient Transport Arrangements	LO	Remove
(0040,1005)	Requested Procedure Location	LO	Remove
(0040,100A)	Reason for Requested Procedure Code Sequence	SQ	Remove
(0040,1010)	Names of Intended Recipients of Results	PN	Remove
(0040,1011)	Intended Recipients of Results Identification Sequence	SQ	Remove
(0040,1086)	Default Frame of Reference UID	UI	Replace UID

(0040,1101)	Person's Identification Code Sequence	SQ	Remove
(0040,1102)	Person's Address	ST	Remove
(0040,1103)	Person's Telephone Numbers	LO	Remove
(0040,1104)	Person's Telecom Information	LT	Remove
(0040,1400)	Requested Procedure Comments	LT	Remove
(0040,2001)	Reason for the Imaging Service Request	LO	Remove
(0040,2004)	Issue Date of Imaging Service Request	DA	Remove
(0040,2005)	Issue Time of Imaging Service Request	TM	Remove
(0040,2008)	Order Entered By	PN	Remove
(0040,2009)	Order Enterer's Location	SH	Remove
(0040,2010)	Order Callback Phone Number	SH	Remove
(0040,2011)	Order Callback Telecom Information	LT	Remove
(0040,2016)	Placer Order Number of Imaging Service Request	LO	Remove
(0040,2017)	Filler Order Number of Imaging Service Request	LO	Remove
(0040,2400)	Imaging Service Request Comments	LT	Remove
(0040,3001)	Confidentiality Constraint on Patient Data Description	LO	Remove
(0040,4005)	Scheduled Procedure Step Start DateTime	DT	Remove
(0040,4008)	Scheduled Procedure Step Expiration DateTime	DT	Remove
(0040,4010)	Scheduled Procedure Step Modification DateTime	DT	Remove
(0040,4011)	Expected Completion DateTime	DT	Remove
(0040,4023)	Referenced General Purpose Scheduled Procedure Step Transaction UID	UI	Replace UID
(0040,4025)	Scheduled Station Name Code Sequence	SQ	Remove
(0040,4027)	Scheduled Station Geographic Location Code Sequence	SQ	Remove
(0040,4028)	Performed Station Name Code Sequence	SQ	Remove
(0040,4030)	Performed Station Geographic Location Code Sequence	SQ	Remove
(0040,4034)	Scheduled Human Performer's Sequence	SQ	Remove
(0040,4035)	Actual Human Performer's Sequence	SQ	Remove
(0040,4036)	Human Performer's Organization	LO	Remove
(0040,4037)	Human Performer's Name	PN	Remove
(0040,4050)	Performed Procedure Step Start DateTime	DT	Remove
(0040,4051)	Performed Procedure Step End DateTime	DT	Remove
(0040,4052)	Procedure Step Cancellation DateTime	DT	Remove
(0040,A023)	Findings Group Recording Date (Trial)	DA	Remove
(0040,A024)	Findings Group Recording Time (Trial)	TM	Remove
(0040,A027)	Verifying Organization	LO	Remove
(0040,A030)	Verification DateTime	DT	Date Shift
(0040,A032)	Observation DateTime	DT	Remove
(0040,A033)	Observation Start DateTime	DT	Remove
(0040,A034)	Effective Start DateTime	DT	Remove
(0040,A035)	Effective Stop DateTime	DT	Remove
(0040,A073)	Verifying Observer Sequence	SQ	Remove

(0040,A075)	Verifying Observer Name	PN	Remove
(0040,A078)	Author Observer Sequence	SQ	Remove
(0040,A07A)	Participant Sequence	SQ	Remove
(0040,A07C)	Custodial Organization Sequence	SQ	Remove
(0040,A082)	Participation DateTime	DT	Date Shift
(0040,A088)	Verifying Observer Identification Code Sequence	SQ	Remove
(0040,A110)	Date of Document or Verbal Transaction (Trial)	DA	Remove
(0040,A112)	Time of Document Creation or Verbal Transaction (Trial)	TM	Remove
(0040,A120)	DateTime	DT	Date Shift
(0040,A121)	Date	DA	Date Shift
(0040,A122)	Time	TM	Keep
(0040,A123)	Person Name	PN	Remove
(0040,A124)	UID	UI	Replace UID
(0040,A13A)	Referenced DateTime	DT	Date Shift
(0040,A171)	Observation UID	UI	Replace UID
(0040,A172)	Referenced Observation UID (Trial)	UI	Replace UID
(0040,A192)	Observation Date (Trial)	DA	Remove
(0040,A193)	Observation Time (Trial)	TM	Remove
(0040,A307)	Current Observer (Trial)	PN	Remove
(0040,A352)	Verbal Source (Trial)	PN	Remove
(0040,A353)	Address (Trial)	ST	Remove
(0040,A354)	Telephone Number (Trial)	LO	Remove
(0040,A358)	Verbal Source Identifier Code Sequence (Trial)	SQ	Remove
(0040,A402)	Observation Subject UID (Trial)	UI	Replace UID
(0040,A730)	Content Sequence	SQ	Remove
(0040,B034)	Annotation DateTime	DT	Remove
(0040,B036)	Segment Definition DateTime	DT	Remove
(0040,B03B)	Montage Name	LT	Remove
(0040,B03F)	Montage Channel Label	LO	Remove
(0040,DB06)	Template Version	DT	Remove
(0040,DB07)	Template Local Version	DT	Remove
(0040,DB0C)	Template Extension Organization UID	UI	Replace UID
(0040,DB0D)	Template Extension Creator UID	UI	Replace UID
(0040,E004)	HL7 Document Effective Time	DT	Remove
(0042,0011)	Encapsulated Document	OB	Check
(0044,0004)	Approval Status DateTime	DT	Remove
(0044,000B)	Product Expiration DateTime	DT	Remove
(0044,0010)	Substance Administration DateTime	DT	Remove
(0044,0104)	Assertion DateTime	DT	Date Shift
(0044,0105)	Assertion Expiration DateTime	DT	Remove
(0050,001B)	Container Component ID	LO	Remove

(0050,0020)	Device Description	LO	Remove
(0050,0021)	Long Device Description	ST	Remove
(0054,0410)	Patient Orientation Code Sequence	SQ	Keep
(0054,0412)	Patient Orientation Modifier Code Sequence	SQ	Keep
(0062,0021)	Tracking UID	UI	Replace UID
(0064,0003)	Source Frame of Reference UID	UI	Replace UID
(0068,6226)	Effective DateTime	DT	Date Shift
(0068,6270)	Information Issue DateTime	DT	Date Shift
(006A,0003)	Annotation Group UID	UI	Replace UID
(006A,0005)	Annotation Group Label	LO	Check
(006A,0006)	Annotation Group Description	UT	Remove
(0070,0001)	Graphic Annotation Sequence	SQ	Keep
(0070,0082)	Presentation Creation Date	DA	Remove
(0070,0083)	Presentation Creation Time	TM	Remove
(0070,0084)	Content Creator's Name	PN	Remove
(0070,0086)	Content Creator's Identification Code Sequence	SQ	Remove
(0070,031A)	Fiducial UID	UI	Replace UID
(0070,1101)	Presentation Display Collection UID	UI	Replace UID
(0070,1102)	Presentation Sequence Collection UID	UI	Replace UID
(0072,000A)	Hanging Protocol Creation DateTime	DT	Date Shift
(0072,005E)	Selector AE Value	AE	Check
(0072,005F)	Selector AS Value	AS	Replace
(0072,0061)	Selector DA Value	DA	Date Shift
(0072,0063)	Selector DT Value	DT	Date Shift
(0072,0065)	Selector OB Value	OB	Check
(0072,0066)	Selector LO Value	LO	Check
(0072,0068)	Selector LT Value	LT	Check
(0072,006A)	Selector PN Value	PN	Remove
(0072,006B)	Selector TM Value	TM	Keep
(0072,006C)	Selector SH Value	SH	Check
(0072,006D)	Selector UN Value	UN	Check
(0072,006E)	Selector ST Value	ST	Check
(0072,0070)	Selector UT Value	UT	Check
(0072,0071)	Selector UR Value	UR	Check
(0074,1234)	Receiving AE	AE	Remove
(0074,1236)	Requesting AE	AE	Remove
(0088,0140)	Storage Media File-set UID	UI	Replace UID
(0088,0200)	Icon Image Sequence	SQ	Remove
(0088,0904)	Topic Title	LO	Remove
(0088,0906)	Topic Subject	ST	Remove
(0088,0910)	Topic Author	LO	Remove

(0088,0912)	Topic Keywords	LO	Remove
(0100,0420)	SOP Authorization DateTime	DT	Remove
(0400,0100)	Digital Signature UID	UI	Replace UID
(0400,0105)	Digital Signature DateTime	DT	Date Shift
(0400,0115)	Certificate of Signer	OB	Remove
(0400,0120)	Signature	OB	Remove
(0400,0310)	Certified Timestamp	OB	Remove
(0400,0402)	Referenced Digital Signature Sequence	SQ	Remove
(0400,0403)	Referenced SOP Instance MAC Sequence	SQ	Remove
(0400,0404)	MAC	OB	Remove
(0400,0550)	Modified Attributes Sequence	SQ	Remove
(0400,0551)	Nonconforming Modified Attributes Sequence	SQ	Remove
(0400,0552)	Nonconforming Data Element Value	OB	Remove
(0400,0561)	Original Attributes Sequence	SQ	Remove
(0400,0562)	Attribute Modification DateTime	DT	Date Shift
(0400,0563)	Modifying System	LO	Remove
(0400,0564)	Source of Previous Values	LO	Remove
(0400,0565)	Reason for the Attribute Modification	CS	Check
(0400,0600)	Instance Origin Status	CS	Remove
(2030,0020)	Text String	LO	Remove
(2100,0040)	Creation Date	DA	Remove
(2100,0050)	Creation Time	TM	Remove
(2100,0070)	Originator	AE	Remove
(2100,0140)	Destination AE	AE	Check
(2200,0002)	Label Text	UT	Remove
(2200,0005)	Barcode Value	LT	Remove
(3002,0121)	Position Acquisition Template Name	LO	Remove
(3002,0123)	Position Acquisition Template Description	LT	Remove
(3006,0002)	Structure Set Label	SH	Check
(3006,0004)	Structure Set Name	LO	Remove
(3006,0006)	Structure Set Description	ST	Remove
(3006,0008)	Structure Set Date	DA	Date Shift
(3006,0009)	Structure Set Time	TM	Keep
(3006,0024)	Referenced Frame of Reference UID	UI	Replace UID
(3006,0026)	ROI Name	LO	Check
(3006,0028)	ROI Description	ST	Remove
(3006,002D)	ROI DateTime	DT	Remove
(3006,002E)	ROI Observation DateTime	DT	Remove
(3006,0038)	ROI Generation Description	LO	Remove
(3006,004D)	ROI Creator Sequence	SQ	Remove
(3006,004E)	ROI Interpreter Sequence	SQ	Remove

(3006,0085)	ROI Observation Label	SH	Remove
(3006,0088)	ROI Observation Description	ST	Remove
(3006,00A6)	ROI Interpreter	PN	Remove
(3006,00C2)	Related Frame of Reference UID	UI	Replace UID
(3008,0024)	Treatment Control Point Date	DA	Date Shift
(3008,0025)	Treatment Control Point Time	TM	Keep
(3008,0054)	First Treatment Date	DA	Remove
(3008,0056)	Most Recent Treatment Date	DA	Remove
(3008,0105)	Source Serial Number	LO	Remove
(3008,0162)	Safe Position Exit Date	DA	Date Shift
(3008,0164)	Safe Position Exit Time	TM	Keep
(3008,0166)	Safe Position Return Date	DA	Date Shift
(3008,0168)	Safe Position Return Time	TM	Keep
(3008,0250)	Treatment Date	DA	Remove
(3008,0251)	Treatment Time	TM	Remove
(300A,0002)	RT Plan Label	SH	Remove
(300A,0003)	RT Plan Name	LO	Remove
(300A,0004)	RT Plan Description	ST	Remove
(300A,0006)	RT Plan Date	DA	Remove
(300A,0007)	RT Plan Time	TM	Remove
(300A,000B)	Treatment Sites	LO	Remove
(300A,000E)	Prescription Description	ST	Remove
(300A,0013)	Dose Reference UID	UI	Replace UID
(300A,0016)	Dose Reference Description	LO	Remove
(300A,0054)	Table Top Position Alignment UID	UI	Replace UID
(300A,0072)	Fraction Group Description	LO	Remove
(300A,0083)	Referenced Dose Reference UID	UI	Replace UID
(300A,00B2)	Treatment Machine Name	SH	Remove
(300A,00C3)	Beam Description	ST	Remove
(300A,00DD)	Bolus Description	ST	Remove
(300A,0196)	Fixation Device Description	ST	Remove
(300A,01A6)	Shielding Device Description	ST	Remove
(300A,01B2)	Setup Technique Description	ST	Remove
(300A,0216)	Source Manufacturer	LO	Remove
(300A,022C)	Source Strength Reference Date	DA	Date Shift
(300A,022E)	Source Strength Reference Time	TM	Keep
(300A,02EB)	Compensator Description	LT	Remove
(300A,0608)	Treatment Position Group Label	LO	Check
(300A,0609)	Treatment Position Group UID	UI	Replace UID
(300A,0611)	RT Accessory Holder Slot ID	LO	Remove
(300A,0615)	RT Accessory Device Slot ID	LO	Remove

(300A,0619)	Radiation Dose Identification Label	LO	Check
(300A,0623)	Radiation Dose In-Vivo Measurement Label	LO	Check
(300A,062A)	RT Tolerance Set Label	LO	Remove
(300A,0650)	Patient Setup UID	UI	Replace UID
(300A,0676)	Equipment Frame of Reference Description	ST	Remove
(300A,067C)	Radiation Generation Mode Label	SH	Check
(300A,067D)	Radiation Generation Mode Description	ST	Check
(300A,0700)	Treatment Session UID	UI	Replace UID
(300A,0734)	Treatment Tolerance Violation Description	ST	Check
(300A,0736)	Treatment Tolerance Violation DateTime	DT	Date Shift
(300A,073A)	Recorded RT Control Point DateTime	DT	Date Shift
(300A,0741)	Interlock DateTime	DT	Date Shift
(300A,0742)	Interlock Description	ST	Check
(300A,0760)	Override DateTime	DT	Date Shift
(300A,0783)	Interlock Origin Description	ST	Check
(300A,0785)	Referenced Treatment Position Group UID	UI	Replace UID
(300A,078E)	Patient Treatment Preparation Procedure Parameter Description	LT	Remove
(300A,0792)	Patient Treatment Preparation Method Description	LT	Remove
(300A,0794)	Patient Setup Photo Description	LT	Remove
(300A,079A)	Displacement Reference Label	LO	Remove
(300C,0113)	Reason for Omission Description	LO	Remove
(300C,0127)	Beam Hold Transition DateTime	DT	Date Shift
(300E,0004)	Review Date	DA	Date Shift
(300E,0005)	Review Time	TM	Keep
(300E,0008)	Reviewer Name	PN	Remove
(3010,0006)	Conceptual Volume UID	UI	Replace UID
(3010,000B)	Referenced Conceptual Volume UID	UI	Replace UID
(3010,000F)	Conceptual Volume Combination Description	ST	Check
(3010,0013)	Constituent Conceptual Volume UID	UI	Replace UID
(3010,0015)	Source Conceptual Volume UID	UI	Replace UID
(3010,0017)	Conceptual Volume Description	ST	Check
(3010,001B)	Device Alternate Identifier	UC	Remove
(3010,002D)	Device Label	LO	Remove
(3010,0031)	Referenced Fiducials UID	UI	Replace UID
(3010,0033)	User Content Label	SH	Check
(3010,0034)	User Content Long Label	LO	Check
(3010,0035)	Entity Label	SH	Check
(3010,0036)	Entity Name	LO	Remove
(3010,0037)	Entity Description	ST	Remove
(3010,0038)	Entity Long Label	LO	Check
(3010,003B)	RT Treatment Phase UID	UI	Replace UID

(3010,0043)	Manufacturer's Device Identifier	ST	Check
(3010,004C)	Intended Phase Start Date	DA	Remove
(3010,004D)	Intended Phase End Date	DA	Remove
(3010,0054)	RT Prescription Label	LO	Check
(3010,0056)	RT Treatment Approach Label	LO	Remove
(3010,005A)	RT Physician Intent Narrative	UT	Remove
(3010,005C)	Reason for Superseding	ST	Remove
(3010,0061)	Prior Treatment Dose Description	UT	Remove
(3010,006E)	Dosimetric Objective UID	UI	Replace UID
(3010,006F)	Referenced Dosimetric Objective UID	UI	Replace UID
(3010,0077)	Treatment Site	LO	Remove
(3010,007A)	Treatment Technique Notes	UT	Remove
(3010,007B)	Prescription Notes	UT	Remove
(3010,007F)	Fractionation Notes	UT	Remove
(3010,0081)	Prescription Notes Sequence	SQ	Remove
(3010,0085)	Intended Fraction Start Time	TM	Remove
(4000,0010)	Arbitrary	LT	Remove
(4000,4000)	Text Comments	LT	Remove
(4008,0040)	Results ID	SH	Remove
(4008,0042)	Results ID Issuer	LO	Remove
(4008,0100)	Interpretation Recorded Date	DA	Remove
(4008,0101)	Interpretation Recorded Time	TM	Remove
(4008,0102)	Interpretation Recorder	PN	Remove
(4008,0103)	Reference to Recorded Sound	LO	Remove
(4008,0108)	Interpretation Transcription Date	DA	Remove
(4008,0109)	Interpretation Transcription Time	TM	Remove
(4008,010A)	Interpretation Transcriber	PN	Remove
(4008,010B)	Interpretation Text	ST	Remove
(4008,010C)	Interpretation Author	PN	Remove
(4008,0111)	Interpretation Approver Sequence	SQ	Remove
(4008,0112)	Interpretation Approval Date	DA	Remove
(4008,0113)	Interpretation Approval Time	TM	Remove
(4008,0114)	Physician Approving Interpretation	PN	Remove
(4008,0115)	Interpretation Diagnosis Description	LT	Remove
(4008,0118)	Results Distribution List Sequence	SQ	Remove
(4008,0119)	Distribution Name	PN	Remove
(4008,011A)	Distribution Address	LO	Remove
(4008,0200)	Interpretation ID	SH	Remove
(4008,0202)	Interpretation ID Issuer	LO	Remove
(4008,0300)	Impressions	ST	Remove
(4008,4000)	Results Comments	ST	Remove

(50xx,xxxx)	Curve Data		Remove
(60xx,3000)	Overlay Data	OB or OW	Remove
(60xx,4000)	Overlay Comments	LT	Remove
(FFFA,FFFA)	Digital Signatures Sequence	SQ	Remove
(FFFC,FFFC)	Data Set Trailing Padding	OB	Remove