Introduction

A new area of radiology, population-based imaging is gaining importance both clinically, e.g. for screening, and in dedicated scientific cohort studies. A current example represents the German National Cohort Study, a large-scale longitudinal study in Germany, which has been recruiting since 2014 and will enroll approximately 200,000 participants, roughly 30,000 of whom will undergo whole-body MRI [1, 2]. Population-based imaging is characterized above all by its methodical approach in which the radiological modality is indicated and specified not with regard to an individual or an individual concrete clinical request for ima-
ging, but is rather employed in the sense of an overriding approach relevant for the general public [3]. While routine clinical medicine employs evidence-based use of screening examinations for early detection of diseases such as breast cancer or lung cancer [4], vastly broader goals are pursued in the scientific context. Here, population-based imaging is employed to characterize morphological as well as functional alterations of the human body to thereby reach a deeper understanding of the diseases relevant to the population as well as the associated risk factors and outcomes thereof. The newly coined term, “radiomics”, frequently associated with this comprehensive data collection is a combination of the root word “radio” as an abbreviation for radiology or imaging in a broader sense and the suffix “-omics”. In biology the suffix “-omics” denotes methods concerned with the collective characterization and quantification of the entirety of individual elements such as biological molecules, structures, functions and/or dynamics of an organism. Examples of this are “genomics” or “proteomics” [5]. As it was the case in genetics several years ago, population-based imaging is faced with the challenge of having to sift through the amount of data for relevant contents.

The MRI study of the German National Cohort attempted to respond to these challenges and thereby establish a new scientific field in Germany. This article presents the fundamentals of population-based imaging and the challenges it poses as well as the scientific potential of radiomics, while discussing in this context the MRI study of the German National Cohort and comparing it with different studies.

Study design and research questions for population-based imaging

Population-based cohort studies

Unlike diagnostic feasibility studies in which the diagnostic accuracy of new imaging method is examined among a study population of patients with the aid of a reference standard, population-based imaging concentrates on a cohort of study participants which, following inclusion into the study (through baseline examination), is accompanied over a specific period of time and monitored (through follow-up examinations). As a rule, imaging is used in the baseline study to correlated image-based phenotypic features and remarkable or pathological changes in organs with other clinical parameters (cross-sectional analysis). With the aid of follow-ups, phenotypical features gathered at an earlier point can be correlated with a clinical outcome to thereby determine the prognostic relevance thereof (longitudinal analysis). This allows the examination of not only established findings (such as, for example, a myocardial late gadolinium enhancement [6]), but also new, unestablished image-related changes, which can be summarized in the context of novel “imaging biomarkers” [7]. This process of establishing new, prognostic markers is a fundamental goal of this type of study [5, 8]. Another feature is that these studies generally involve a high number of cases (often \( n > 1000 \)), thereby ensuring sufficiently large power in order to identify and verify the expected associations [8].

Design of the German National Cohort Study

The German National Cohort Study constitutes a classic population-based longitudinal observational study [1]. A total of approximately 200 000 residents of the Federal Republic of Germany ranging between 20 and 69 years of age (age at the start of the study) will be included in the German National Cohort Study. The study participants are recruited at a total of 18 study centers, which cover primarily urban and industrial regions as well as a few rural regions throughout the whole of Germany (Fig. 1). Each center will include a minimum of approximately 10 000 study participants randomly selected from the registry office. All 200 000 study participants have gone through a standardized protocol at the local study centers, which was comprised of interviews, questionnaires and medical examinations. In addition, blood and urine samples are taken and other biomaterials collected. Data collection for the baseline-examination is conducted at two intensity levels (Level 1 and 2). All participants undergo a 2.5-hour examination protocol (Level 1), while a subgroup of 40 000 participants receives an expanded protocol (additional 4 hours, Level 2) involving more in-depth examinations and questions [1].

Fig. 1 Design of the National German Cohort MRI Study The German National Cohort Study is comprised of 18 general study centers, 5 of which have a dedicated 3-Tesla MRI scanner (Magnetom Skyra, Siemens Healthcare; in Augsburg, Berlin, Essen, Mannheim, Neubrandenburg). In addition, 4 central facilities known as Imaging Cores were set up to handle imaging-related tasks applicable for all centers [2]: Munich – coordination & training; Greifswald – quality management; Heidelberg – management of incidental findings; Bremen – data management. (Source: German National Cohort Study)
Roughly 30,000 participants will additionally be examined with whole-body MRI; therefore 3-Tesla MRI scanners (Magnetom Skyra, Siemens Healthcare, Erlangen) have been installed at five MRI study centers (Augsburg, Berlin, Essen, Mannheim and New Brandenburg; Fig. 1) [2]. Each of these participants also undergoes the more intense Level 2 examination protocol. So-called imaging cores (Fig. 1) were established in addition to the MRI study centers. The role of the imaging cores is to provide superordinate coordination and communication as well as the planning, issuing and execution of tasks applicable for all centers [2], which can be divided as follows:

- Imaging core for coordination & training in Munich
- Imaging core for data management in Bremen
- Imaging core for incidental findings in Heidelberg
- Imaging core for quality management in Greifswald

Following the base-line examination, each study participant is contacted every 2 to 3 years to answer questionnaires concerning lifestyle, risk factors and the appearance of diseases (“active” follow-up) [1]. For “passive” follow-up, established recording systems e.g. the cancer register in cases of newly diagnosed cancers. In the same manner, an effort is made to link the data of the registries and the mandatory health insurance providers to achieve a comprehensive picture of the occurrence of diseases, therapies administered, clinical outcome and mortality.

**Research questions posed in the German National Cohort Study**

Consistent with the goals of population-based studies, the German National Cohort Study attains through random sampling a generalizable picture of the overall German population and its relevant widespread diseases. Specifically, it pursues the following four primary goals [1]:

- Clarifying the causes of chronic diseases and presenting the connection with genetic factors as well as lifestyle and environmental influences
- Identifying new risk factors and presenting existing geographic and socioeconomic disparities in terms of health status and disease risk in Germany
- Developing risk-forecasting models for diseases that can be used to derive personalized prevention strategies
- Identifying possibilities for early detection of chronic diseases (evaluation of markers as effective aids for disease prevention)

In this regard, the MRI study within the German National Cohort offers above all great potential for identifying new risk factors and developing personalized prevention strategies [9]. In addition, whole-body MRI offers the unique possibility of linking image-based risk factors with the degree of subclinical disease appearance or with normal variants as well as with classic risk factors and making a correlation with clinical outcome (Fig. 2). This provides a host of possibilities especially when combined with a radiomics approach.

**Radiomics**

While the new scientific field radiomics initially emerged from CT-based oncological imaging [10, 11], it is applicable to diverse clinical and scientific areas of imaging regardless of modality [12, 13]. The object of radiomics is to translate and compile the entirety of the available image-based information of an organism into potentially disease-relevant information. In this process the classic radiological approach centered around an organ-based description of anatomical deviations is taken further, and abstract information based primarily on a quantitative characteristic of image data is compiled on a meta level. Radiomics is based on the hypothesis that similar information, that can be obtained from e.g. serum, genetic or protein analyses, can also be derived from images. In addition, imaging has the advantage of being able to gather information in either a circumscribed or comprehensive manner, while a blood sample, for example, can frequently be taken only on a global level; or biopsies, for example, can again capture only local characteristics of a heterogeneous tumor tissue (e.g., with regard to protein expression). The procedure for developing a radiomics concept follows a set of sequential steps (Fig. 3):

**Standardized imaging in large populations**

To facilitate detection of recurrent pathological changes by means of statistical methods amid the complex variability of the human body, comprehensive image data with a high case number, quality and reproducibility, as is the case in the German National Cohort Study, are fundamental to the approach.

**(Semi-) automated segmentation of image files**

Because manual segmentation of image aspects (such as, for example, of certain organs or tumors) would entail a substantial effort and would be subject to increased variability through differences between multiple readers, complete or partial automatization of image segmentation is essential. While there are several functioning approaches [14, 15], this aspect currently poses a challenge for radiomics.

**Extraction of image-based features**

Primarily quantitative features such as volumes, intensity, texture, shape or relation to surrounding tissue are abstracted from image data. Moreover, complex, quantitative parameters can also be gathered using new analytical approaches such as, for example, Geographic Object-Based Image Analysis (GEOBIA) [16]. The capturing of image features must accordingly not remain restricted to a clinically indicated request for information.

**Analysis of image-based features in association with clinical outcome**

For ascertaining clinical relevance, it is imperative to identify from the plethora of extracted parameters (often several hundred) those features and combinations of features that have an association with the desired clinical outcome (e.g. the prognostic value for cardiac events or genetic expression in tumors [10]). In addition to classic statistical analytical methods, approaches such as, for example, machine-based learning, are also pursued [14, 17].

The possibilities of this approach within the actively discussed field of personalized medicine are highly promising. However, they present a major challenge to radiology, particularly with regard to disease prevention [9], given that this concept exceeds the traditional field of radiology and no cor-
**Fig. 3** Schematic process of radiomics using lung cancer as an example. The process covers multiple sequential steps, starting with a standardized image-acquisition in a large population, followed by an automated image segmentation and extraction of image-based features and, finally, an analysis of image-based features in association with medical data, such as genetics or clinical outcomes. The figure is modified by Aerts et al., Nature Communications 2014 [11].

**Fig. 2** Population-based imaging as a possible means for establishing the unclear connection between modifiable (e.g., lifestyle) and non-modifiable (e.g., genetic) risk factors and clinical outcome. In this process, the quantification of subclinical degrees of disease or imaging biomarkers, for example, is expedient, and concepts such as radiomics are can be applied.
responding scientific and technical resources have been established for this type of data collection and analysis.

**Imaging modalities in population-based cohort studies**

The use of modern imaging technology in population-based studies is not new [18], as ultrasound methods have been a regular feature of many population studies for several decades. Clinical imaging modalities such as MRI or computed tomography are playing an increasing role among population-based cohorts. While the choice of modality depends primarily on the research focus, other aspects such as, for example, reproducibility, availability, costs, general legal considerations or risks (e.g. from radiation exposure) play also an important role in the decision.

Ultrasound examination is a relatively cheap easy-to-implement method [19–22] and is thus used frequently (Table 1), even if the reproducibility of the data and measurement values can at times constitute a limitation [23]. Moreover, the possibilities of a retrospective evaluation of the ultrasound images beyond the primary request for information are limited. The most common area of application of ultrasound in cohort studies is echocardiography [20–22] and examination of the carotids [19–22] (Table 1), e.g. regarding the intima-media thickness, which has established itself as a marker for global arteriosclerosis burden [24]. Within the German National Cohort Study 3D-echocardiography is also part of the Level 2 examination protocol [1].

MRI examination is being used increasingly in population-based cohort studies, since it achieves a high tissue contrast even without contrast medium and thus has a very low side-effects profile. In particular, cranial MRI examinations performed in previous studies have generated substantial scientific knowledge, such as regarding brain function in the Rotterdam Study or the 3C Study (Table 1) [25, 26]. MRI is also frequently used for examining cardiac structures and function [27–29] as it continues to be the clinical reference standard in this area.

Due to technical enhancements such as parallel acquisition technologies or continuous table feed as well as high magnetic field strengths allow nowadays comprehensive coverage of the entire body within examination times appropriate for study participants [30]. Accordingly, few studies have incorporated whole-body MRI examinations into the study design (Table 1). In addition to the German National Cohort Study, current studies using whole-body MRI protocols would be the Study of Health in Pomerania (SHIP) [31] as well as to a some extent the UK Biobank [32] and the Cooperative Health Research in the Augsburg Region [Kooperative Gesundheitsforschung in der Region Augsburg] (KORA) [29].

The MRI protocol of the German National Cohort Study is comprised of 12 MRI sequences, which can be divided into the focus areas of neurological, cardiovascular, thoracoabdominal and musculoskeletal examination (Table 2). Each of the focus areas involves an examination time of approximately 15 minutes, yielding a total MRI examination time of 60 minutes (Fig. 4).

Because the reproducibility and comparability of the MRI data depend to a certain degree on the selected sequence parameters [33] significant efforts were made in advance to harmonize with other large population-based cohort studies having the same/common sequences (Table 3), even if each study had its own unique sequences. In the Ger-

<table>
<thead>
<tr>
<th>Table 1</th>
<th>Imaging modalities in population-based cohorts in Europe with existing cohort studies cited as examples.</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>ultrasound</td>
</tr>
<tr>
<td>costs</td>
<td>+</td>
</tr>
<tr>
<td>undesirable effects</td>
<td>none known</td>
</tr>
<tr>
<td>possible exclusion criteria</td>
<td>none BRCA-mutation, clausiophobia, pregnancy implants not compatible with MRI, clausiophobia, pregnancy</td>
</tr>
<tr>
<td>reproducibility</td>
<td>+ +++ +++ +++</td>
</tr>
<tr>
<td>whole-body examinations</td>
<td>no yes yes yes</td>
</tr>
<tr>
<td>mobile solutions</td>
<td>yes limited limited limited</td>
</tr>
<tr>
<td>current implementation (examples, EU only, sorted by n)</td>
<td>Cardiovascular Risk in Young Finns Study (n=1800. carotids) [19]</td>
</tr>
<tr>
<td></td>
<td>SHIP (n=2400. carotids, liver, thyroid gland, heart) [20]</td>
</tr>
<tr>
<td></td>
<td>Heinz Nixdorf Recall (n = 8700. at risk carotids, heart, pancreas, endo-thelial dysfunction) [21]</td>
</tr>
<tr>
<td></td>
<td>Rotterdam Study (n=15000. carotids, heart, cerebral vessels, liver) [22]</td>
</tr>
<tr>
<td></td>
<td>Lung Cancer Screening Intervention Trial (LUSI; n = 2029. Thorax) [36]</td>
</tr>
<tr>
<td></td>
<td>Rotterdam Study (n=2400. carotids, heart) [22]</td>
</tr>
<tr>
<td></td>
<td>Rotterdam Study (n=4800. heart) [34]</td>
</tr>
<tr>
<td></td>
<td>AGES-Reykjavik Study (n=5000. heart, musculoskeletal) [35]</td>
</tr>
<tr>
<td></td>
<td>NELSON Study (n = 7557. thorax) [37]</td>
</tr>
<tr>
<td></td>
<td>Swedish CardioPulmonary Biomage Study (n=30000. thorax, heart) [27]</td>
</tr>
<tr>
<td></td>
<td>Tromso Study (n=5000. hip, spinal column) [39]</td>
</tr>
<tr>
<td></td>
<td>Rotterdam Study (n=15000. hip) [22]</td>
</tr>
<tr>
<td></td>
<td>KORA-study (n = 400. whole body without musculoskeletal) [29]</td>
</tr>
<tr>
<td></td>
<td>SHIP (n = 3300. whole body) [42]</td>
</tr>
<tr>
<td></td>
<td>3C Study (n = 3300. cranial) [25]</td>
</tr>
<tr>
<td></td>
<td>SMART Study (n = 6000. heart) [28]</td>
</tr>
<tr>
<td></td>
<td>Rotterdam Study (n=15000. cranial, carotids, knee) [26]</td>
</tr>
<tr>
<td></td>
<td>Swedish CardioPulmonary Biomage Study (n=30000. carotids) [27]</td>
</tr>
</tbody>
</table>

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### Table 2  German National Cohort Study whole-body MRI protocol Modified by Bamberg et al. [2].

<table>
<thead>
<tr>
<th>MR sequence</th>
<th>Imaging parameters</th>
<th>Anatomical coverage</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>neurological imaging</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>1. T1w 3 D MPRAGE</td>
<td>1.0 × 1.0 × 1.0 mm³ (isotropic) voxel size, sagittal acquisition</td>
<td>neurocranium including upper spinal canal</td>
</tr>
<tr>
<td>2. 2 D FLAIR</td>
<td>0.9 × 0.9 × 4.0 mm³ voxel size, axial acquisition</td>
<td>neurocranium</td>
</tr>
<tr>
<td>3. resting-state EPI BOLD</td>
<td>3.0 × 1.0 × 1.0 mm³ (isotropic) voxel size, axial acquisition</td>
<td>neurocranium</td>
</tr>
<tr>
<td><strong>cardiovascular imaging</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>4. MR-angiography 3 D SPACE STIR</td>
<td>1.9 × 0.9 × 4.0 mm³ voxel size, coronal acquisition</td>
<td>entire thorax</td>
</tr>
<tr>
<td>5. cine SSFP LAX</td>
<td>1.5 × 1.5 × 6.0 mm³ voxel size</td>
<td>two, three, and four-chamber view</td>
</tr>
<tr>
<td>6. cine SSFP SAX</td>
<td>1.5 × 1.5 × 6.0 mm³ voxel size</td>
<td>12 short axis slices from base to apex of heart</td>
</tr>
<tr>
<td>7. MOLLI SAX</td>
<td>1.4 × 1.4 × 8.0 mm³ voxel size</td>
<td>single short axis slice central-ventricular</td>
</tr>
<tr>
<td><strong>thoracic-abdominal imaging</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>8. T2w HASTE</td>
<td>1.4 × 1.4 × 5.0 mm³ voxel size, axial acquisition</td>
<td>shoulders to beneath the kidneys</td>
</tr>
<tr>
<td>9. T1w 3 D VIBE 2-point Dixon</td>
<td>1.4 × 1.4 × 3.0 mm³ voxel size, axial acquisition, CAIPIRINHA</td>
<td>shoulders to beneath the minor trochanter</td>
</tr>
<tr>
<td>10. multi-echo 3 D VIBE</td>
<td>1.6 × 1.6 × 4.0 mm³ voxel size, axial acquisition, 6 echoes</td>
<td>epigastrium</td>
</tr>
<tr>
<td><strong>musculoskeletal imaging</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>11. PD-fs 3 D SPACE</td>
<td>1.0 × 1.0 × 1.0 mm³ (isotropic) voxel size, coronal acquisition</td>
<td>pelvis including the sacroiliac joint and both hips</td>
</tr>
<tr>
<td>12. T2w 2 D Fast Spin-echo</td>
<td>0.9 × 0.9 × 3.0 mm³ voxel size, sagittal acquisition</td>
<td>cervical, thoracic and lumbar spinal column</td>
</tr>
</tbody>
</table>

**Fig. 4**  Examples of the German National Cohort Study divided into 4 research foci: Body with a T1w-GRE, 2-point Dixon of the entire torso, a multi-echo VIBE of the upper abdomen and a T2w-HASTE of the thorax and upper abdomen (no example shown); musculoskeletal system with a T2w-TSE of the spinal column and a PD-fatsat of the pelvis, neurological system with a T1w and FLAIR of the neurocranium as well as a resting state; cardiovascular system with a non-contrast angiogram of the thorax, long- and short-axis CINE of the heart as well as T1-mapping.
man National Cohort Study, specifically the non-contrast thoracic MR-angiography, the T1-mapping of the left ventricle, but also the multi-echo sequences of the upper abdomen and the high-resolution 3D-PD-fs sequence of the pelvis were innovative approaches that set themselves apart from the MRI examination of other studies ([26]).

The use of x-rays must always be restrictedly on possible with CT or MRI. DEXA therefore supplements rather than replaces other modalities. In this context whole-body DEXA allows cost-effective and simple quantification of bone density as it did, for example, in the Tromoso Study or the Rotterdam Study ([22, 39]). Moreover, DEXA allows a quantification of fat and fat-free mass ([40], yet no separation of local fat deposits (e.g. visceral fat) as is possible with CT or MRI. DEXA therefore supplements rather than replaces other modalities. In this context whole-body DEXA examination is used, for example in the UK Biobank ([41]).

### Challenges of population-based imaging

As with other extensive population-based studies, the German National Cohort MRI Study faces multiple major challenges in generating an optimally comprehensive dataset while using a high-resolution imaging modality. Added to this are the high degree of necessary quality assurance, a

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**Table 3** MRI protocol comparison between the German National Cohort Study (NAKO) and other population-based cohort studies using MRI.

<table>
<thead>
<tr>
<th>German National Cohort (NAKO) (3 Tesla)</th>
<th>UK Biobank¹ (3 Tesla &amp; 1.5 Tesla)</th>
<th>SHIP (1.5 Tesla)</th>
<th>Rotterdam Scan Study² (1.5 Tesla)</th>
<th>KORA (3 Tesla)</th>
</tr>
</thead>
<tbody>
<tr>
<td>neurological imaging</td>
<td>T1w 3D MPRAGE</td>
<td>comparable</td>
<td>slice thickness (0.8 mm vs. 1.0 mm in the NAKO)</td>
<td>voxel size (0.3 × 0.3 × 0.3 mm³ vs. 1.0 × 1.0 × 1.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td></td>
<td>slice thickness (0.8 mm vs. 1.0 mm in the NAKO)</td>
<td>voxel size (0.3 × 0.3 × 0.3 mm³ vs. 1.0 × 1.0 × 1.0 mm³ in the NAKO)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>resting-state EPI BOLD</td>
<td>voxel size (2.4 × 2.4 × 2.4 mm³ vs. 3.1 × 3.1 × 3.1 mm³ in the NAKO); measurement time comparable; TR (0.735 s vs. 2.2 s in the NAKO); Multiband (8 × vs. none in the NAKO)</td>
<td>N/A</td>
<td>slice thickness (3.5 mm vs. NAKO 3.1 mm); measurement time (7.73 min vs. NAKO 6 min); TR (2.5 s vs. NAKO 2.2 s)</td>
<td>N/A</td>
</tr>
<tr>
<td>cardiac imaging</td>
<td>CINE LAX/SAX</td>
<td>N/A voxel size (1.8 × 1.8 × 6.0/8.0 mm³ vs. 1.5 × 1.5 × 6.0 mm³ in the NAKO)</td>
<td>voxel size (2.2 × 1.8 × 6.0/7.0 mm³ vs. 1.5 × 1.5 × 6.0 mm³ in the NAKO)</td>
<td>voxel size (1.5 × 1.5 × 8.0 mm³ vs. 1.4 × 1.4 × 8.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td></td>
<td>voxel size (2.2 × 1.8 × 6.0/7.0 mm³ vs. 1.5 × 1.5 × 6.0 mm³ in the NAKO)</td>
<td>voxel size (2.2 × 1.8 × 6.0/7.0 mm³ vs. 1.5 × 1.5 × 6.0 mm³ in the NAKO)</td>
<td>voxel size (1.5 × 1.5 × 8.0 mm³ vs. 1.4 × 1.4 × 8.0 mm³ in the NAKO)</td>
<td>voxel size (1.5 × 1.5 × 8.0 mm³ vs. 1.4 × 1.4 × 8.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td>T1-mapping</td>
<td>sequence (shMOLLI vs. MOLLI in the NAKO)</td>
<td>N/A</td>
<td>voxel size (1.5 × 1.5 × 8.0 mm³ vs. 1.4 × 1.4 × 8.0 mm³ in the NAKO)</td>
<td>voxel size (1.5 × 1.5 × 8.0 mm³ vs. 1.4 × 1.4 × 8.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td>3 D MR-angiography</td>
<td>N/A contrast (with vs. without contrast medium in the NAKO)</td>
<td>N/A voxel size (1.5 × 1.5 × 8.0 mm³ vs. 1.4 × 1.4 × 8.0 mm³ in the NAKO)</td>
<td>voxel size (1.5 × 1.5 × 8.0 mm³ vs. 1.4 × 1.4 × 8.0 mm³ in the NAKO)</td>
<td>voxel size (1.5 × 1.5 × 8.0 mm³ vs. 1.4 × 1.4 × 8.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td>thoracic-abdominal imaging</td>
<td>T1w-3D-GRE</td>
<td>comparable</td>
<td>voxel size (1.8 × 1.8 × 3.0 mm³ vs. 1.4 × 1.4 × 3.0 mm³ in the NAKO)</td>
<td>voxel size (1.7 × 1.7 × 1.7 mm³ vs. 1.4 × 1.4 × 3.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td></td>
<td>voxel size (1.8 × 1.8 × 3.0 mm³ vs. 1.4 × 1.4 × 3.0 mm³ in the NAKO)</td>
<td>voxel size (1.7 × 1.7 × 1.7 mm³ vs. 1.4 × 1.4 × 3.0 mm³ in the NAKO)</td>
<td>voxel size (1.7 × 1.7 × 1.7 mm³ vs. 1.4 × 1.4 × 3.0 mm³ in the NAKO)</td>
<td>voxel size (1.7 × 1.7 × 1.7 mm³ vs. 1.4 × 1.4 × 3.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td>T2w-HASTE</td>
<td>N/A</td>
<td>voxel size (2.3 × 1.8 × 5.0 mm³ vs. 1.4 × 1.4 × 5.0 mm³ in the NAKO)</td>
<td>voxel size (1.2 × 1.2 × 5.0 mm³ vs. 1.4 × 1.4 × 5.0 mm³ in the NAKO)</td>
<td>voxel size (1.2 × 1.2 × 5.0 mm³ vs. 1.4 × 1.4 × 5.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td>multi-echo 3 D</td>
<td>anatomical coverage (single-slice vs. entire liver in the NAKO) Number of echoes (1 – 8 vs. 6 in the NAKO)</td>
<td>N/A voxel size (1.8 × 1.8 × 4.0 mm³ vs. 1.6 × 1.6 × 4.0 mm³ in the NAKO)</td>
<td>voxel size (1.8 × 1.8 × 4.0 mm³ vs. 1.6 × 1.6 × 4.0 mm³ in the NAKO)</td>
<td>voxel size (1.8 × 1.8 × 4.0 mm³ vs. 1.6 × 1.6 × 4.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td>musculoskeletal imaging</td>
<td>3 D PD-fs Hip</td>
<td>N/A voxel size (1.1 × 1.1 × 4.0 mm³ vs. 0.9 × 0.9 × 3.0 mm³ in the NAKO)</td>
<td>voxel size (1.1 × 1.1 × 4.0 mm³ vs. 0.9 × 0.9 × 3.0 mm³ in the NAKO)</td>
<td>voxel size (1.1 × 1.1 × 4.0 mm³ vs. 0.9 × 0.9 × 3.0 mm³ in the NAKO)</td>
</tr>
<tr>
<td></td>
<td>T2w Spine</td>
<td>N/A voxel size (1.1 × 1.1 × 4.0 mm³ vs. 0.9 × 0.9 × 3.0 mm³ in the NAKO)</td>
<td>voxel size (1.1 × 1.1 × 4.0 mm³ vs. 0.9 × 0.9 × 3.0 mm³ in the NAKO)</td>
<td>voxel size (1.1 × 1.1 × 4.0 mm³ vs. 0.9 × 0.9 × 3.0 mm³ in the NAKO)</td>
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N/A (not acquired) denotes sequences used in the NAKO but not in the other study.

¹ Compared with the MRI protocol of the pilot study of the UK Biobank (N=5000 test subjects). UK Biobank uses a 3 Tesla MRI-scanner for neuro imaging and a 1.5 Tesla MRI scanner for all other imaging.

² For the Rotterdam Scan Study only slice thicknesses without X/Y-resolution are published ([26]). “N/A” denotes "not acquired" in the corresponding study.
complex management structure and the evaluation of enormous amounts of data and, in terms of ethical considerations, handling incidental findings.

**Incidental findings in population-based imaging**

Because of the comprehensive, high-resolution MRI data, the prevalence of relevant incidental findings is not insignificant even among asymptomatic study participants [42]. However, dealing with these “incidental findings”, as they are called, in the study setting of the National German Cohort without knowledge of the participants’ medical histories or clinical symptoms is highly complex. Communicating the incidental findings may facilitate curative treatment of the diseases (e.g., early detection of lung cancer). On the other hand, communication constitutes intervention in the natural course of the disease and influences the observed and scientific findings gained through the study. This is especially relevant for the population-based longitudinal study in which the observation of potentially “normal”, disease processes is often one of the declared goals in order to observe the connection between exposure and disease occurrence in a manner with as little influence as possible. The theoretical option of not informing the study participants of these incidental findings has been rejected so far by most ethics committees in the context of population-based imaging in Germany and is not in the personal interest of many study participants [43]. Further it is nearly impossible to provide always final diagnosis for the incidental findings particularly because the MRI protocol is often not designed to detect a specific disease, but rather to provide a morphological or functional phenotyping. Non-definitive reports generally prompt additional tests, which in turn may entail certain risks, side-effects and expenses.

In the German National Cohort MRI Study all acquired images are reviewed by a board certified radiologist for the presence of incidental findings. A list of incidental findings was developed in advance jointly with radiologists, various clinical disciplines, ethicists and epidemiologists that classifies incidental findings as “notification urgently required”, “notification required” and “notification not required”. Criteria for the need to notify are (1) clinical consequence of the incidental finding, (2) negative ramifications for the participant if not notified and (3) optimally low false-positive rate of the MRI report for the assumed disease. This list is continuously updated based on new research findings and recommendations of the clinical associations, and can be viewed online (www.national-kohorte.de).

**Quality management**

With population-based imaging, special challenges arise due to the large size of the studies, with test subjects frequently numbering many thousands, and, as the case may be, the multicentric design. To also detect faint associations between imaging and clinical variables (e.g. outcomes, genetics, serum biomarkers), it is necessary to reduce errors and variability in image acquisition to a technologically feasible minimum. Without consistent homogeneity of data, both among the study centers and over the course of time, investigations with regard to geographical disparities, for example, are possible only to a limited extent. Accordingly, the following quality management measures were established in the German National Cohort Study [2]:

**Defining quality standard**

Quality standards coordinated across interdisciplinary lines were documented in “Standard-Operating-Procedures” (SOP) and communicated to all employees through a training and certification process.

**Checking against the quality standard**

All acquired data are automatically checked for completeness (e.g. “all sequences examined?”, “sequences repeated?”) and correctness (sequence parameters, number of slices, etc.). Moreover, image quality parameters are gathered automatically, such as signal-to-noise-ratio (SNR) or ghosting-level [2]. For additional monitoring, MRI phantoms undergo measurement and automated quantitative evaluation on a weekly basis. The visual quality of images are also evaluated at the study center by the local radiologists. Quality assurance likewise includes an independent evaluation of a 10% random sampling at the imaging core in Greifswald (over-reading).

**Measures for deviations from quality standard**

All parameters recorded in the process of quality assurance are presented in a quality report with dynamic threshold values. When the threshold is exceeded, the study center is notified, which then enacts centrally initiated measures such as, for example, retraining or study center visits if a correction has not been made.

**Data management and evaluation**

Current projections alone place the size of the baseline MRI data in the National German Cohort study at over 90 terabytes, which, for security regulations, are stored long-term at two spatially separated locations [1, 44]. To facilitate rapid access to data, an upload approver were established in the German National Cohort Study, which allows. This model was realized as a web-based platform that facilitates decentralized processing in an internet browser without local software installation. For the long term, scientific evaluation shall follow the same procedure, with scientists gaining access to the MRI data via a web-based platform on which the various evaluation tools are implemented and individual measurement algorithms can be introduced via appropriate interfaces. In addition to the automatization of these algorithms, this infrastructure allows rapid, decentralized, multicentric evaluation and thereby maximum usability of the acquired MRI data while at the same time providing quality control [44].

**Public access and future developments**

According to the statutes of the Nationale Kohorte e.V. [Association of the German National Cohorts Study], the generated data are available both to the national and international scientific community. This results in a usage right extending beyond the initially involved institutions with far-reaching scientific potential. For radiology, the German National Cohort MRI study represents the possibility of significantly and proactively responding to future challenges in healthcare. The use of automated approaches in evaluating comprehensive volumes of data poses a challenge of increasing importance in the clinical setting as well. In this
area the German National Cohort Study can serve to establish the interaction from (semi-) automated analyses with manual review first in the scientific context. The situation is similar in quality management: Approaches such as automated image quality control within the German National Cohort Study are gaining importance in the area of clinical treatment. The use of evaluation strategies such as radiomics, for example, allows the development of new and improved image-based risk stratifications, which offer high potential for personalized medicine and prevention in the future [7]. However, this demands the cooperation and explicit involvement of many different scientists both domestically and internationally, far exceeding the scope of the experts already involved. Moreover, this is possible only with the support of known foundations and recognized specialist associations such as, for example, Deutschen Röntgengesellschaft [German X-Ray Association].

**Summary**

Population-based imaging is an increasingly relevant radiological scientific field that will influence the course of clinical radiology in the future. The MRI study of the German National Cohort, which involves the examination of roughly 30,000 participants using 3-Tesla whole-body MRI, constitutes a unique opportunity to gain key knowledge in the understanding of (subclinical) diseases and to achieve the development of an image-based risk stratification for personalized medicine. **Radiomics** is a relatively new approach in which image information, analogous to laboratory or genome data, is translated and compiled via traditional diagnostic information (qualitative findings) into quantitative, disease-relevant information. With the complex procedures established in the German National Cohort for quality assurance and data standardization, the study may provide a basis for the future developments within Radiology and allows domestic and international scientists access to a unique dataset that can facilitate better understanding of the widest array of pathological processes and the socio-economic relevance thereof.

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