

A review of computer-aided diagnosis in thoracic and colonic imaging

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Abstract: Medical imaging has been indispensable in medicine since the discovery of x-rays. Medical imaging offers useful information on patients' medical conditions and on the causes of their symptoms and diseases. As imaging technologies advance, a large number of medical images are produced which physicians/radiologists must interpret. Thus, computer aids are demanded and become indispensable in physicians' decision making based on medical images. Consequently, computer-aided detection and diagnosis (CAD) has been investigated and has been an active research area in medical imaging. CAD is defined as detection and/or diagnosis made by a radiologist/physician who takes into account the computer output as a "second opinion". In CAD research, detection and diagnosis of lung and colorectal cancer in thoracic and colonic imaging constitute major areas, because lung and colorectal cancers are the leading and second leading causes, respectively, of cancer deaths in the U.S. and also in other countries. In this review, CAD of the thorax and colon, including CAD for detection and diagnosis of lung nodules in thoracic CT, and that for detection of polyps in CT colonography, are reviewed.

Key Words: Computer-aided detection; computer-aided diagnosis; screening; lung nodule; colorectal polyps; thoracic CT; CT colonography; classifier; pixel-based machine learning; machine learning in medical imaging



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Introduction

Medical imaging has been indispensable in medicine since the discovery of X-rays by Wilhelm C. Röntgen in 1895. Medical imaging offers useful information on patients' medical conditions and provides clues to causes of their symptoms and diseases. As imaging technologies advance, a large number of medical images are produced which physicians/radiologists must interpret. Thus, computer aids are demanded and become indispensable in physicians' decision making based on medical images. Consequently, computer-aided detection and diagnosis (CAD) (1-4) has been an active research area in medical imaging. CAD is defined as detection and/or diagnosis made by a radiologist/physician who takes into account the computer output as a "second opinion" (2). CAD is often categorized into two major groups: computer-aided detection (CADE)

and computer-aided diagnosis (CADx). CADE focuses on a detection task, i.e., detection (or localization) of lesions in medical images. CADx focuses on a diagnosis (characterization) task, e.g., distinction between benign and malignant lesions, and classification among different lesion types.

A branch of the history of CAD started in 1955. A radiologist, Lee Lusted, mentioned the potential use of digital computers for large-scale data problems in medicine in (5) in 1955. It was only nine years after the first general-purpose computer, ENIAC, was introduced in 1946. In 1963, Lodwick *et al.* digitized chest radiographs for computer analysis (6). In 1964, Becker *et al.* developed automated measurement of the cardiothoracic ratio in chest radiographs (7,8). The first study on CADE of abnormalities in mammograms was published by Winsberg *et al.* (9) in

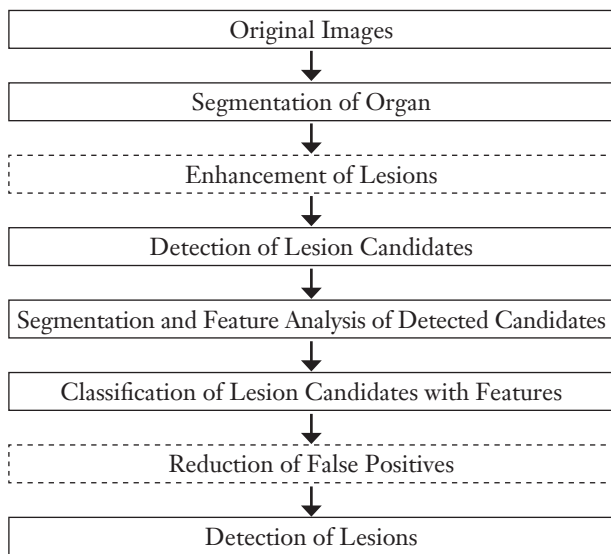


Figure 1 Flowchart for a generic CADe scheme for detection of lesions in medical images. Boxes with solid lines indicate four major steps in the CADe scheme, and those with dashed lines indicate optional, yet important steps

1967. In 1973, Toriwaki *et al.* (10) reported the first study on CADe of a focal abnormality in chest radiographs, and Roellinger *et al.* (11) reported the first study on CADe of a heart abnormality in chest radiographs. In the mid-1980's, investigators in the Kurt Rossmann Laboratories in the Department of Radiology at the University of Chicago began studies on the development and evaluation of CAD. Chan *et al.* (12), Fujita *et al.* (13), Giger *et al.* (14), and Katsuragawa *et al.* (15) published a series of papers on CADe of microcalcifications in mammography, CAD for vessel size measurement in angiography, CADe of lung nodules in chest radiography, and CADe of interstitial lung disease in chest radiography, respectively. In 1988, a venture company, R2 Technology (now Hologic), which obtained licenses for CAD technologies from the University of Chicago, received approval for the first commercial CAD system for mammography from the U.S. Food and Drug Administration (FDA).

Evidence suggests that CAD can help improve the diagnostic performance of radiologists/physicians in their image interpretations (16-22). Consequently, many investigators have participated and developed CAD schemes such as those for detection of lung nodules in chest radiographs (14,23-25) and in thoracic CT (26-29), those for detection of microcalcifications/masses in mammography (12), breast MRI (30), and breast

ultrasound (US) (31,32), and those for detection of polyps in CT colonography (CTC) (33-36).

In this paper, the progress and advancement of the development of CAD schemes of the thorax and colon are reviewed, including CAD schemes for detection and diagnosis of lung nodules in thoracic CT, and those for detection of polyps in CTC.

Generic architectures of CADe and CADx schemes

A flowchart for a generic CADe scheme of lesions in medical images is shown in *Figure 1*. A CADe scheme generally consists of four major steps and two optional steps: (I) segmentation of the organ of interest, (III) detection of lesion candidates from the segmented organ, (IV) segmentation and feature analysis of the detected lesion candidates, (V) classification of the lesion candidates by use of a classifier with features; optionally (II) enhancement of lesions between steps I and III, and (VI) reduction of false-positive (FP) detections. Segmentation of the organ of interest is the first necessary step that aims at making the rest of the steps focus on the organ. The development of the detection of lesion candidates generally aims at obtaining a high sensitivity level, because the sensitivity lost in this step cannot be recovered in the later steps. In the next step, the detected (or localized) lesion candidates are segmented, and connected-component labeling (37,38) is performed for identification of each segmented candidate as an individual isolated object. Pattern features such as gray-level-based features, texture features, and morphologic features are extracted from the segmented candidates. Finally, the detected lesion candidates are classified into lesions or non-lesions by use of a classifier. This final step is very important, because it determines the final performance of a CADe scheme when the additional step of FP reduction is not employed. The development of the classification step aims at removing as many non-lesions (i.e., FPs) as possible while minimizing the removal of lesions (i.e., true-positive detections).

A machine-learning technique is generally used in the step of classification of lesion candidates. The machine-learning technique is trained with sets of input features and correct class labels. This class of machine learning is referred to as feature-based machine learning, or simply as a classifier. The task of machine learning here is to determine "optimal" boundaries for separating classes in the multi-dimensional feature space which is formed by the input features (39). Feature-based machine-learning algorithms

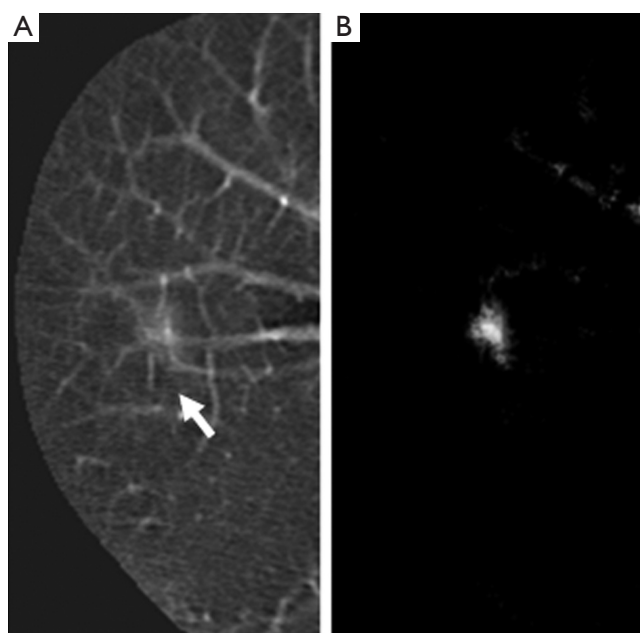


Figure 2 Lesion enhancement by means of a supervised MTANN lesion-enhancement filter. (A) Original axial CT slice with a lung nodule (indicated by an arrow). (B) Output image of the trained MTANN nodule-enhancement filter. In the output image (B), the lung nodule in the original CT image (A) is enhanced, whereas normal structures such as lung vessels are suppressed substantially

include linear discriminant analysis (LDA) (40), quadratic discriminant analysis (QDA) (40), multilayer perceptron [one of the most popular artificial neural network (ANN) models] (41), and support vector machines (SVMs) (42). The structure of an ANN may be designed by use of an automated design method such as sensitivity analysis (43,44). Recently, as available computational power has increased dramatically, pixel/voxel-based machine learning (45) emerged in medical image processing/analysis which uses pixel/voxel values in images directly, instead of features calculated from segmented regions, as input information; thus, feature calculation or segmentation is not required. Pixel-based machine learning has been used in the classification of the detected lesion candidates in CADe and CADx schemes.

To improve the performance of CADe schemes, investigators sometimes employ an additional step that is enhancement of lesions after the step of the segmentation of the organ of interest. This additional step aims at improving the sensitivity for detection of lesion candidates in the subsequent step. It often helps improve the specificity

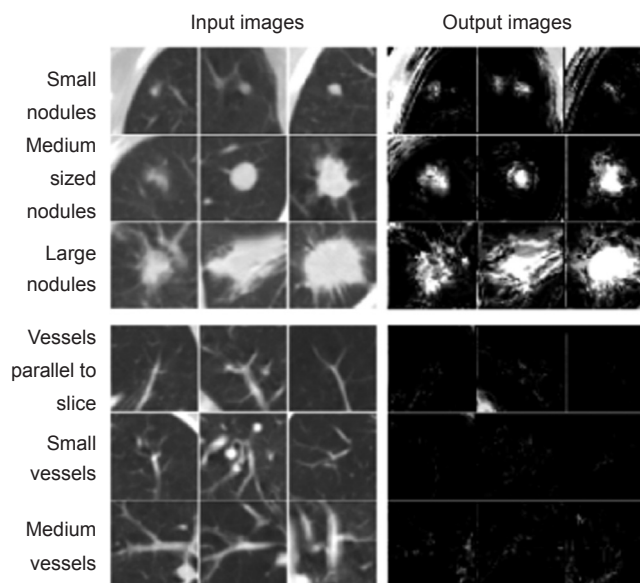


Figure 3 Enhancement of lung nodules and suppression of FPs (i.e., lung vessels) by use of MTANNs for FP reduction. Once lung nodules are enhanced, and FPs are suppressed, FPs can be distinguished from lung nodules by use of scores obtained from the output images

as well. *Figure 2* shows such an example. Suzuki (46) developed a supervised “lesion enhancement” filter based on a massive-training ANN (MTANN) for enhancing lesions and suppressing non-lesions in medical images. *Figure 3* illustrates the enhancement of a lung nodule in a CT image by means of a trained MTANN lesion enhancement filter. In the output image, the lung nodule in the original CT image is enhanced, while normal structures such as lung vessels are suppressed substantially. By use of the MTANN lesion enhancement filter, the performance of the initial nodule candidate detection step in a CADe scheme was substantially improved from a 96% sensitivity with 19.3 FPs per section to a 97% sensitivity with 6.7 FPs per section.

Investigators also often employ an additional step of reduction of FPs at the end in a CADe scheme. The FP reduction step aims at improving the specificity of the CADe scheme. Reduction of FPs is very important, because a large number of FPs could adversely affect the clinical application of CADe. A large number of FPs is likely to confound the radiologist’s task of image interpretation and thus lower his/her efficiency. In addition, radiologists may lose their confidence in CADe as a useful tool. Suzuki *et al.* developed an FP reduction technique based on

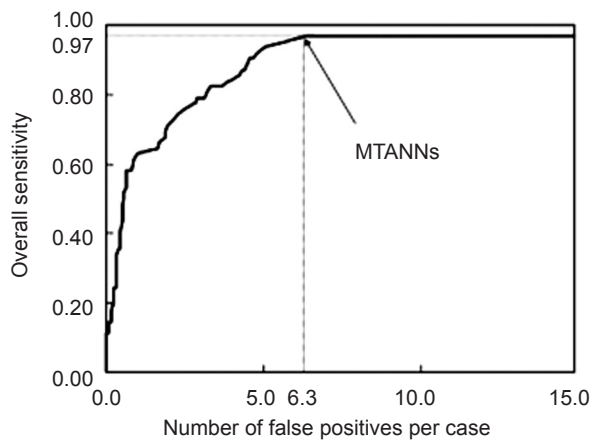


Figure 4 FROC curve indicating the performance of the FP reduction by MTANNs in a CADe scheme for detection of lung nodules in CT. With the trained MTANNs, FPs were removed without any removal of true positives

MTANNs (26) for reduction of FPs in a CADe scheme for lung nodules in CT. The MTANNs were trained to enhance lung nodules and suppress various types of FPs (i.e., non-nodules) such as lung vessels. *Figure 3* shows the results of the enhancement of various-sized lung nodules (a) and those of the suppression of various-sized lung vessels (b). *Figure 4* shows a free-response receiver operating characteristic (FROC) curve (47) indicating the performance of the trained MTANNs in the CADe scheme. With the MTANNs, the specificity of the CADe scheme was improved from 15.0 to 6.3 FPs per case without sacrificing the original sensitivity of 97%.

After the development of a CADe scheme, the evaluation of the stand-alone performance of the developed scheme is the last step in CADe development. The evaluation of radiologists' performance with the use of the developed CAD scheme is the important last step in CAD research.

The flowchart for a generic CADx scheme for diagnosis (or characterization) of detected lesions in medical images is shown in *Figure 5*. CADx schemes start from detected (localized) lesions either automatically or manually. In other words, the location of a lesion of interest is known. CADx schemes include distinction between benign and malignant status of detected lesions, and classification among different types of detected lesions. First, the detected lesions are segmented, and pattern features are extracted from the segmented lesions. With the extracted features, the lesions are classified into benign or malignant, or different lesion

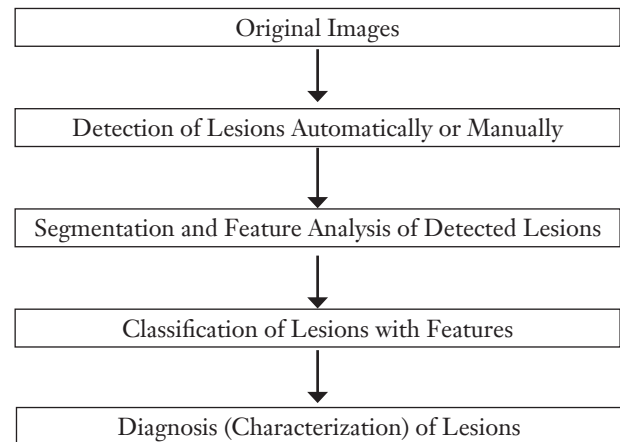


Figure 5 Flowchart for a generic CADx scheme for diagnosis of (detected) lesions in medical images

types. CADx schemes may provide the likelihood of malignancy or the likelihood of being a certain type of lesion, as opposed to categorical classification results (or classes).

CADe and CADx in thoracic imaging

Thoracic imaging for lung cancer detection and diagnosis

Lung cancer continues to rank as the leading cause of cancer deaths in the United States and in other countries such as Japan. Because CT is more sensitive than chest radiography in the detection of small nodules and of lung carcinoma at an early stage (48-51), lung cancer screening programs are being investigated in the United States (52,53), Japan (48,49), and other countries with low-dose (LD) helical CT as the screening modality. Evidence suggests that early detection of lung cancer may allow more timely therapeutic intervention and thus a more favorable prognosis for the patient (49,54). Helical CT, however, generates a large number of images that must be read by radiologists/physicians. This may lead to “information overload” for the radiologists/physicians. Furthermore, they may miss some cancers during their interpretation of CT images (55,56). Therefore, a CADe scheme for detection of lung nodules in CT images has been investigated as a tool for lung cancer screening. *Figure 6* shows an example of CADe outputs on a CT image of the lungs. A CADe



Figure 6 CADe outputs (indicated by circles) on an axial CT slice of the lungs. A lung nodule (indicated by an arrow) was detected correctly by a CADe scheme with one FP detection (branch of lung vessels) on the right

scheme detected a lung nodule correctly with an FP which was a branch of the lung vessels.

CADe of lung nodules in thoracic CT

In 1994, Giger *et al.* (57) developed a CADe scheme for detection of lung nodules in CT based on comparison of geometric features. They applied their CADe scheme to a database of thick-slice diagnostic CT scans of 8 patients with 47 nodules. They achieved a sensitivity of 94% with 1.25 FPs per case. In 1999, Armato *et al.* (28,58) extended the method to include 3D feature analysis, a rule-based scheme, and LDA for classification. They tested their CADe scheme with a database of thick-slice (10 mm) diagnostic CT scans of 43 patients with 171 nodules. They achieved a sensitivity of 70% with 42.2 FPs per case in a leave-one-out cross-validation test. Gurcan *et al.* (59) employed a similar approach, i.e., a rule-based scheme based on 2D and 3D features, followed by LDA for classification. They achieved a sensitivity of 84% with 74.4 FPs per case for a database of thick-slice (2.5-5 mm, mostly 5 mm) diagnostic CT scans of 34 patients with 63 nodules in a leave-one-out test. Lee *et al.* (60) employed a simpler approach which is a rule-based scheme based on 13 features for classification. They achieved a sensitivity of 72% with 30.6 FPs per case for a database of thick-slice (10 mm) diagnostic CT scans of 20 patients with 98 nodules.

Suzuki *et al.* (26) developed a pixel-based machine-learning technique called an MTANN for reduction of a single source of FPs and a multiple MTANN scheme for reduction of multiple sources of FPs that had not been removed by LDA. They achieved a sensitivity of 80.3% with 4.8 FPs per case for a database of thick-slice (10 mm) screening LDCT scans of 63 patients with 71 nodules with solid, part-solid, and non-solid patterns, including 66 cancers in a validation test. This MTANN approach did not require a large number of training cases: the MTANN was able to be trained with 10 positive and 10 negative cases (61-63), whereas feature-based classifiers generally require 400-800 training cases (61-63). Arimura *et al.* (27) employed a rule-based scheme followed by LDA or by the MTANN (26) for classification. They tested their scheme with a database of 106 thick-slice (10 mm) screening LDCT scans of 73 patients with 109 cancers that had solid, part-solid and non-solid patterns, and they achieved a sensitivity of 83% with 5.8 FPs per case in a validation test (or a leave-one-patient-out test for LDA). Farag *et al.* (64) developed a template-modeling approach that uses level sets for classification. They achieved a sensitivity of 93.3% with an FP rate of 3.4% for a database of thin-slice (2.5 mm) screening LDCT scans of 16 patients with 119 nodules and 34 normal patients. Ge *et al.* (65) incorporated 3D gradient field descriptors and ellipsoid features in LDA for classification. They employed Wilks' lambda stepwise feature selection for selecting features before the LDA classification. They achieved a sensitivity of 80% with 14.7 FPs per case for a database of 82 thin-slice (1.0-2.5 mm) CT scans of 56 patients with 116 solid nodules in a leave-one-patient-out test. Matsumoto *et al.* (66) employed LDA with 8 features for classification. They achieved a sensitivity of 90% with 64.1 FPs per case for a database of thick-slice (5 or 7 mm) diagnostic CT scans of 5 patients (4 of which used contrast media) with 50 nodules in a leave-one-out test.

Yuan *et al.* (67) tested a commercially available CADe system (ImageChecker CT, LN-1000, by R2 Technology, Sunnyvale, CA; Hologic now). They achieved a sensitivity of 73% with 3.2 FPs per case for a database of thin-slice (1.25 mm) CT scans of 150 patients with 628 nodules in an independent test. Pu *et al.* (68) developed a scoring method based on the similarity distance of medial axis-like shapes for classification. They achieved a sensitivity of 81.5% with 6.5 FPs per case for a database of thin-slice (2.5 mm) screening CT scans of 52 patients with 184 nodules, including 16 non-solid nodules. Retico *et al.* (69) used a voxel-based neural approach (i.e., a class of the MTANN approach) with pixel

values in a subvolume as input for classification. They obtained sensitivities of 80-85% with 10-13 FPs per case for a database of thin-slice (1 mm) screening CT scans of 39 patients with 102 nodules. Ye *et al.* (70) used a rule-based scheme followed by a weighted SVM for classification. They achieved a sensitivity of 90.2% with 8.2 FPs per case for a database of thin-slice (1 mm) screening CT scans of 54 patients with 118 nodules including 17 non-solid nodules in an independent test. Golosio *et al.* (71) used a fixed-topology ANN for classification, and they evaluated their CADe scheme with a publicly available database from the Lung Image Database Consortium (LIDC) (72). They achieved a sensitivity of 79% with 4 FPs per case for a database of thin-slice (1.5-3.0 mm) CT scans of 83 patients with 148 nodules that one radiologist detected from an LIDC database in an independent test.

Murphy *et al.* (73) used a k-nearest-neighbor classifier with features selected from 135 features for classification. They achieved a sensitivity of 80 with 4.2 FPs per case for a large database of thin-slice screening CT scans of 813 patients with 1,525 nodules in an independent test. Tan *et al.* (74) developed a feature-selective classifier based on a genetic algorithm and ANNs for classification. They achieved a sensitivity of 87.5% with 4 FPs per case for a database of thin-slice CT scans of 125 patients with 80 nodules that 4 radiologists agreed from the LIDC database in an independent test. Messay *et al.* (75) developed a sequential forward selection process for selecting the optimum features for LDA and quadratic discriminant analysis (QDA). They obtained a sensitivity of 83% with 3 FPs per case for a database of thin-slice CT scans of 84 patients with 143 nodules from the LIDC database in a 7-fold cross-validation test. Riccard *et al.* (76) used a heuristic approach based on geometric features, followed by an SVM for classification. They achieved a sensitivity of 71% with 6.5 FPs per case for a database of thin-slice CT scans of 154 patients with 117 nodules that 4 radiologists agreed on from the LIDC database in a 2-fold cross-validation test.

Thus, various approaches have been proposed for CADe schemes for lung nodules in CT. Sensitivities for detection of lung nodules in CT range from 70% to 95%, with from a few to 70 FPs per case. Major sources of FPs are various-sized lung vessels. Major sources of false negatives are ground glass nodules, nodules attached to vessels, and nodules attached to the lung wall (i.e., juxtapleural nodules). Ground glass nodules are difficult to detect, because they are subtle, of low-contrast, and have ill-defined boundaries.

The MTANN approach was able to enhance and thus detect ground-glass nodules (26). The cause of false negatives due to vessel-attached nodules and juxtapleural nodules is mis-segmentation and thus inaccurate feature calculation. Because the MTANN approach does not require segmentation or feature calculation, it was able to detect such nodules (26).

CADx of lung nodules in thoracic CT

Although CT has been shown to be sensitive to the detection lung nodules, it may be difficult for radiologists to distinguish between benign and malignant nodules on LDCT images. In a screening program with LDCT in New York, 88% (206/233) of suspicious lesions were found to be benign on follow-up examinations (50). In a screening program in Japan, only 83 (10%) among 819 scans with suspicious lesions were diagnosed to be cancer cases (56). According to recent findings at the Mayo Clinic, 2,792 (98.6%) of 2,832 nodules detected by a multidetector CT were benign, and 40 (1.4%) nodules were malignant (52). Thus, a large number of benign nodules were found with CT; follow-up examinations such as high-resolution CT (HRCT) and/or biopsy were performed on these patients. Therefore, CADx schemes for distinction between benign and malignant nodules in LDCT would be useful for reducing the number of "unnecessary" follow-up examinations.

A number of researchers developed CADx schemes distinguishing malignant nodules from benign nodules automatically and/or determining the likelihood of malignancy for the detected nodules. The performance of the schemes was generally evaluated by means of receiver-operating-characteristic (ROC) analysis (77), because this task is a two-class classification. The area under the ROC curve (AUC) (78) was often used as a performance index.

In 1999, McNitt-Gray *et al.* (79) developed a classification scheme based on LDA for distinction between malignant and benign nodules in HRCT. They achieved a correct classification rate of 90.3% for a database of 17 malignant and 14 benign nodules. Matsuki *et al.* (80) used an ANN with subjective features determined by radiologists for classification between 99 malignant and 56 benign nodules in HRCT and achieved an AUC value of 0.951. Aoyama *et al.* (81) used LDA for distinction between malignant and benign nodules in thick-slice (10 mm) screening LDCT. They achieved an AUC value of 0.846 for a database of 73 patients with 76 primary cancers and 342 patients with 413 benign nodules. Mori *et al.* (82) developed

a classification scheme for distinction between malignant and benign nodules in contrast-enhanced (CE) CT by using LDA with 3 features (i.e., attenuation, shape index, and curvedness value). They used a database of thin-slice (2 mm) CE-CT scans of 35 malignant and 27 benign nodules for testing their CADx scheme. They achieved AUC values of 0.91 and 1.0 with non-CE CT and CE-CT, respectively, in a leave-one-out test.

Shah *et al.* (83) employed different classifiers such as logistic regression and QDA with features selected from a group of 31 by using stepwise feature selection based on the Akaike information criterion. Their scheme with logistic regression achieved an AUC value of 0.92 in the distinction between 19 malignant and 16 benign nodules in thin-slice CE-CT. Suzuki *et al.* (84) developed a pixel-based machine-learning technique called a multiple MTANN scheme for the classification task. They achieved an AUC value of 0.88 for thick-slice screening LDCT scans of 73 patients with 76 primary cancers and 342 patients with 413 benign nodules. Iwano *et al.* (85) achieved a sensitivity of 76.9% and a specificity of 80% with their scheme based on LDA with 2 features in their evaluation of HRCT images of 52 malignant and 55 benign nodules. Way *et al.* (86) incorporated nodule surface features into their classification based on LDA or an SVM, and they achieved an AUC value of 0.857 in the classification of 124 malignant and 132 benign nodules in 152 patients. Chen *et al.* (87) employed an ANN ensemble to classify 19 malignant and 13 benign nodules, and they achieved an AUC value of 0.915. Lee *et al.* (88) developed a two-step supervised learning scheme combining a genetic algorithm with a random subspace method, and they achieved an AUC value of 0.889 in the classification between 62 malignant and 63 benign nodules.

Thus, various approaches to CADx schemes have been proposed. The database size varied in different studies; CT scans in the databases included screening LDCT, standard diagnostic CT, and HRCT.

CADe in colonic imaging

Colonic imaging for colorectal cancer detection

Colorectal cancer is the second leading cause of cancer deaths in the United States (89). Evidence suggests that early detection and removal of polyps (which are precursors of colorectal cancer) can reduce the incidence of colorectal cancer (90,91). Consequently, the American Cancer

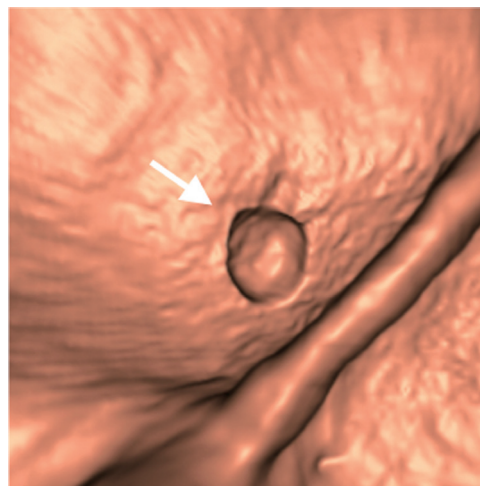


Figure 7 CADe output (indicated by an arrow) for detection of polyps in an endoluminal view in CTC. A polyp (indicated by an arrow) was detected correctly by a CADe scheme

Society recommends that a person who is at average risk for developing colorectal cancer, beginning at age 50, should have colorectal cancer screening with examinations including optical colonoscopy and CTC. CTC, also known as virtual colonoscopy, is a technique for detecting colorectal neoplasms by use of CT scans of the colon (92). The diagnostic performance of CTC in detecting polyps, however, remains uncertain due to a propensity for perceptual errors in the detection of polyps (93). CADe of polyps has been investigated in an effort to address this issue with CTC (94-96). CADe has the potential to (I) increase radiologists' diagnostic accuracy in the detection of polyps, (II) decrease reader variability, and (III) reduce radiologists' interpretation time when CAD is used during the primary read (94,95). A number of investigators have developed automated or semi-automated CADe schemes for the detection of polyps in CTC (33,97-102). *Figure 7* shows an example of a CADe output for detection of polyps in an endoluminal view in CTC. A CADe scheme detected a polyp correctly.

CADe of polyps in CTC

In 2000, Summers *et al.* (36) developed a CADe scheme for detection of polyps in CTC based on curvature analysis. They tested the feasibility of it with simulated polyps. In 2001, they tested their CADe scheme on 50 actual polyps. In 2001, Yoshida *et al.* (33) developed a CADe scheme

based on curvature analysis called a shape index. They tested their CADE scheme on a database of 43 patients with 12 polyps. They achieved a by-patient sensitivity of 100% with 2.0 FPs per case. In 2001, Gokturk *et al.* (103) employed an SVM with histogram input that is used as a shape signature for classification. They achieved a by-patient (by-polyp) sensitivity of 100% (95%) with a specificity of 0.69 (0.74) [14.3 FPs/patient (12.0 FPs/patient)] for CTC data (2.5-3.0 mm collimation) of 48 patients in either supine or prone position, containing 40 polyps (2-15 mm). Näppi *et al.* (104) developed a CADE scheme based on LDA or QDA with 54 volumetric features (9 statistics of 6 features). They achieved a by-patient (by-polyp) sensitivity of 100% (95%) with 2.4 FPs/patient for CTC data (5 mm collimation) of 40 patients in both supine and prone positions, including 12 polyps in 11 patients.

Acar *et al.* (105) used edge-displacement fields to model the changes in consecutive cross-sectional views of CTC data and QDA for classification. They achieved a sensitivity of 100% (95%) with a specificity of 0.47 (0.56) for CTC data (2.5-3.0 mm collimation) of 48 patients in either supine or prone position, containing 40 polyps (2-15 mm). Jerebko *et al.* (106) used a multilayer perceptron to classify polyp candidates in their CADE scheme and improved the performance by incorporating a committee of multilayer perceptrons (107) and a committee of SVMs (108). They used CTC data (5 mm collimation) of 40 patients in both supine and prone positions, including 21 polyps (5-25 mm) for evaluating their CADE scheme. They finally achieved a sensitivity of 86.7% for larger polyps (≥ 10 mm) and 75% for other polyps with 3 FPs/patient in an independent test. Wang *et al.* (109) developed a classification method based on LDA with internal features (geometric, morphologic, and textural) of polyps. They achieved a sensitivity of 100% and 100% for larger polyps (≥ 10 mm) and other polyps with 4 and 6.9 FPs/patient, respectively, for CTC data (5 mm collimation) of 153 patients in both supine and prone positions, including 61 polyps (4-30 mm) in 45 patients. Näppi *et al.* (110) developed a method for false-positive reduction (i.e., classification between lesions and non-lesions) based on supine-prone correspondence of CTC scans where patients are scanned in the supine and prone positions. Näppi *et al.* (111) also developed feature-guided analysis in colon segmentation and feature analysis of polyp candidates and compared them with their previously developed fuzzy clustering. They were able to reduce 70-75% of FPs with their new method, which combines both a non-machine-learning-based method and

a feature-based classifier.

Suzuki *et al.* (112) developed a pixel-based machine-learning technique called a 3D MTANN by extending the structure of a 2D MTANN (26) to process 3D volume data in CTC. Their CADE scheme was based on a Bayesian ANN with texture and geometric features, followed by 3D MTANNs. They removed FPs due to rectal tubes by using a single 3D MTANN (112) and multiple sources of FPs by developing and using a mixture of expert 3D MTANNs (35). They achieved a by-polyp (by-patient) sensitivity of 96.4% (100%) with 2.1 FPs/patient in a leave-one-lesion-out cross-validation test of the classification part for CTC data (1.25-5 mm collimation) of 73 patients in both supine and prone positions, including 28 polyps (5-25 mm) in 15 patients. Li *et al.* (113) developed a classification method based on an SVM classifier with wavelet-based features. They obtained a sensitivity of 71% with 5.4 FPs/patient in a 4-fold cross-validation test of the classification part for CTC data of 44 patients containing 45 polyps (6-9 mm). Wang *et al.* (114) improved the SVM performance by using nonlinear dimensionality reduction (i.e., a diffusion map and locally linear embedding). They used CTC data (1.25-2.5 mm collimation) of 792 patients in both supine and prone positions, including 226 polyps (>6 mm) for evaluating their CADE scheme. They achieved a sensitivity of 83% for polyps (6-9 mm) with 9 FPs/patient.

Yao *et al.* (115) employed a topographic height map for calculating features for an SVM classifier. They used a large database containing CTC data (1.25-2.5 mm collimation) of 792 patients in both supine and prone positions, including 226 polyps (>6 mm) for evaluating their CADE scheme. They obtained a sensitivity of 93% and 76% for larger polyps (≥ 10 mm) and other polyps with 1.2 and 3.1 FPs/patient, respectively, in a 10-fold cross-validation test of the classification part. Suzuki *et al.* (116) tested a CADE scheme based on a Bayesian ANN with texture and geometric features together with MTANNs (i.e., a pixel-based machine-learning technique). They used CTC data (1.25-5 mm collimation) of 24 patients in both supine and prone positions, including 23 polyps (6-25 mm) and a mass (35 mm), that had been "missed" by radiologists (117) in a multicenter clinical trial (118). They achieved a by-polyp (by-patient) sensitivity of 96.4% (100%) with 1.1 FPs/patient in a leave-one-lesion-out cross-validation test of the classification part.

Suzuki *et al.* (119,120) also improved the efficiency of the MTANN approach by incorporating principal-component analysis-based and Laplacian eigenmap-based dimension

reduction techniques. Xu and Suzuki (121) showed that other nonlinear regression models such as support vector and nonlinear Gaussian process regression models instead of the ANN regression model could be used as the core model in the MTANN framework. Zhou *et al.* (122) developed projection features for an SVM classifier. They achieved by-polyp sensitivities of 93.1% and 80.6% for larger polyps (≥ 10 mm) and other polyps with 1.9 and 5.2 FPs/patient, respectively, for CTC data (1.25-5.0 mm collimation) of 325 patients in the supine and/or prone positions, including 347 polyps and masses (5-60 mm). Wang *et al.* (123) improved the performance of a CAD scheme by adding statistical curvature features in multiple-kernel learning. They obtained a sensitivity of 83% with 5 FPs/patient in a leave-one-out cross-validation test of the classification part for a database containing CTC data (1.25-2.5 mm collimation) of 66 patients in supine and/or prone positions, including 96 polyps.

Thus, various approaches have been proposed for CADE schemes for polyps in CTC. Most of the methods for the initial detection of polyp candidates employed morphologic analysis, including curvature analysis such as the shape index. Curvatures are often calculated by using the Hessian matrix. Classifiers used in CADE schemes include LDA, QDA, an SVM, and a Bayesian ANN. Sensitivities for detection of polyps in CTC range from 70% to 100%, with 1 to 30 FPs per case. Major sources of FPs include haustral folds, residual stool, rectal tubes, the ileocecal valve, and extra-colonic structures such as the small bowel and stomach. Existing CADE schemes tend to miss superficially elevated neoplasms (often called flat lesions) (124,125). Recently, Suzuki *et al.* developed a CADE scheme for detection of superficially elevated neoplasms (126). Detection of superficially elevated neoplasms is very important, because they are histologically aggressive, and because they are often missed by radiologists in CTC as well as by gastroenterologists in optical colonoscopy.

Summary

In this paper, CADE and CADx schemes for detection and diagnosis of lung nodules in thoracic CT and those for detection of polyps in CTC have been reviewed. A CADE scheme (i.e., detection of lesions) generally consists of four major steps and two optional steps: segmentation of the organ of interest, detection of lesion candidates from the segmented organ, segmentation and feature analysis of the detected lesion candidates, and classification of the lesion candidates by use of a classifier with features; and

optionally, enhancement of lesions and reduction of FPs. Feature-based machine learning (or classifier), including LDA, QDA, an ANN, a Bayesian ANN, and an SVM, was used in the step of the classification of the lesion candidates. To improve the performance of CADE schemes further, investigators sometimes employ additional steps of enhancement of lesions and FP reduction. CADE schemes that employ those addition steps tend to yield a high performance. Sensitivities for detection of lung nodules in CT range from 70% to 95%, with a few to 70 FPs per case. On the other hand, sensitivities for detection of polyps in CTC range from 70% to 100%, with 1 to 30 FPs per case. Thus, the sensitivities of CADE schemes are relatively high, but the number of FPs is high-compared to radiologists' performance. Therefore, further improvement in specificity is necessary in future research.

A CADx scheme (i.e., diagnosis or characterization of detected lesions) generally consists of two major steps, which are segmentation and feature analysis of the detected lesions and classification of the lesions. CADx schemes start from detected (localized) lesions either automatically or manually. The major component of CADx schemes is classification of lesions by use of a classifier with features. The performance of CADx schemes for distinction between benign and malignant lung nodules in CT range from AUC values of 0.85 to 0.95. Thus, the computer performance is relatively high compared to radiologists' performance.

I hope that this review will be useful for researchers to advance the development of CAD schemes in thoracic and colonic imaging as well as for those who want to start the development of CAD schemes in other imaging areas.

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