Accuracy and Efficiency of Deep-Learning–Based Automation of Dual Stain Cytology in Cervical Cancer Screening

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Abstract

Background: With the advent of primary human papillomavirus testing followed by cytology for cervical cancer screening, visual interpretation of cytology slides remains the last subjective analysis step and suffers from low sensitivity and reproducibility. Methods: We developed a cloud-based whole-slide imaging platform with a deep-learning classifier for p16/Ki-67 dual-stained (DS) slides trained on biopsy-based gold standards. We compared it with conventional Pap and manual DS in 3 epidemiological studies of cervical and anal precancers from Kaiser Permanente Northern California and the University of Oklahoma comprising 4253 patients. All statistical tests were 2-sided. Results: In independent validation at Kaiser Permanente Northern California, artificial intelligence (AI)-based DS had lower positivity than cytology (P < .001) and manual DS (P < .001) with equal sensitivity and substantially higher specificity compared with both Pap (P < .001) and manual DS (P < .001), respectively. Compared with Pap, AI-based DS reduced referral to colposcopy by one-third (41.9% vs 60.1%, P < .001). At a higher cutoff, AI-based DS had similar performance to high-grade squamous intraepithelial lesions cytology, indicating a risk high enough to allow for immediate treatment. The classifier was robust, showing comparable performance in 2 cytology systems and in anal cytology. Conclusions: Automated DS evaluation removes the remaining subjective component from cervical cancer screening and delivers consistent quality for providers and patients. Moving from Pap to automated DS substantially reduces the number of colposcopies and also achieves excellent performance in a simulated fully vaccinated population. Through cloud-based implementation, this approach is globally accessible. Our results demonstrate that AI not only provides automation and objectivity but also delivers a substantial benefit for women by reduction of unnecessary colposcopies.

Advances in digital imaging and machine learning can revolutionize cancer screening, diagnosis, and treatment by improving accuracy and reproducibility of image assessment and streamlining clinical workflow (1–4). With its requirement for high throughput and fast turnaround and its dependence on microscopic and visual technologies, automation can play a major role in improving the efficiency of cervical cancer screening. Many countries are currently switching from Pap cytology to high-risk human papillomavirus (HPV) screening (5–7). Although a negative HPV test provides great reassurance of low cervical cancer risk over the next decade (8–10), only a small subset of women with a positive HPV test require further evaluation. To avoid overburdening the system with HPV-positive women, additional triage is required for colposcopy referral (11, 12). Current triage strategies include partial HPV genotyping and Pap cytology (7, 13). The limited sensitivity and
reproducibility of cytology require laborious quality control procedures and frequent retesting \cite{14,15}. Improving the efficiency of cervical cancer screening is particularly important for vaccinated populations due to lower disease prevalence and higher demands for screening test performance.

A promising triage strategy is concomitant detection of p16 and Ki-67 in the same cell \cite{16,21}. Currently, artificial intelligence (AI) algorithms mostly try to match manual reading accuracy to improve automation but do not offer a substantial improvement for patients. Automated scanning and deep-learning evaluation of DS slides can improve throughput, reproducibility, and accuracy of the assay for better risk stratification and a direct benefit to women \cite{8,22,23}. To achieve this, we developed the CYTOREADER system that combines whole-slide scanning with automated evaluation of DS cytology slides. Cloud-based evaluation provides ample computational capacity and storage space and can provide diagnostic procedures where sufficient personnel, expertise, or infrastructure is lacking. We evaluated the clinical performance of CYTOREADER in 4253 slides from 3 epidemiological studies of HPV-positive cervical and anal precancers.

**Methods**

**General Approach**

CYTOREADER uses whole-slide scanners (Hamamatsu Nanozoomers HT, XR, and S360) for imaging of ThinPrep (Hologic) or SurePath (Becton Dickinson, BD) slides, 2 widely used liquid-based cytology technologies. CYTOREADER is a cloud-based system (Google Cloud Platform) that can also run as a local installation. Training of deep-learning algorithms for automated DS evaluation was performed using small areas (tiles) from whole slides containing individual or small numbers of epithelial cells. For training of the deep-learning algorithms, tiles from training slides were manually evaluated for DS-positive cells by 3 observers (Supplementary Figure 1, available online).

**Deep Learning**

Two deep-learning approaches (Convolutional Neural Network with 4 layers [CNN4] and Inception-v3 with 48 layers [IncV3]) were developed sequentially as shown in Figure 1 and described in Supplementary Methods (available online). The algorithms determine the number of DS-positive cells on a slide by detecting the number of tiles above a certain likelihood threshold. A slide is considered positive if the number of DS-positive cells on a slide exceeds a certain cutoff. Training and validation were conducted on the tile level and the slide level. First, a training set from 450 patients was selected for which the clinical endpoint cervical intraepithelial neoplasia grade 3 or greater (CIN3+) was unblinded. Tiles were selected for initial training (80%) and validation (20%) of the algorithm. The deep-learning network provides a likelihood for each tile above which it is considered positive (0.5 for CNN4 and 0.4 for IncV3). The resulting candidate CNN was applied on the slide level on training slides. A cutoff of positive tiles is used to determine slide positivity (≥3 tiles per cell for CNN4 and ≥2 tiles per cell for IncV3). From misclassified slides, false-positive or false-negative tiles were extracted and fed back into the original CNN training to optimize classification accuracy of the CNN. A final locked CNN was applied on the patient level on the blinded validation set comprising 3803 slides. CNN4 showed good performance in Thinprep slides but not in Surepath slides. Subsequently, a second algorithm (IncV3) was trained specifically for Surepath slides (Supplementary Methods, available online). We published a GitHub repository and created a web page at https://github.com/stcmethub/dual_stain_dl with a source code description of the models and the installation instructions.

**Study Populations**

The Biopsy Study is a population-based study of women aged 18 years or older referred to colposcopy at the University of Oklahoma Health Sciences Center between 2009 and 2011 \cite{24}. We included DS slides from 602 women as previously described \cite{19}. The study population was split into a representative training set (193 slides with 741 DS-positive and 953 DS-negative tiles) and a validation set of 409 slides (Figure 1). This study was approved by the University of Oklahoma and National Cancer Institute (NCI) institutional review boards (IRB); written informed consent was obtained from all participants before study enrollment.

The Anal Cancer Screening Study (ACSS) was based at the San Francisco Kaiser Permanente Northern California (KPNC) Anal Cancer Screening Clinic. HIV-positive men who have sex with men 18 years or older were enrolled at KPNC between 2009 and 2010. DS slides from 318 men were generated as previously described \cite{25}. From 19 training slides, 445 DS-positive and 532 DS-negative tiles were used for training (Figure 1). This study was approved by the KPNC and NCI IRBs; written informed consent was obtained from all participants before study enrollment.

At KPNC, DS was evaluated for triage of HPV-positive women between 2012 and 2015 in a population of women aged 25 years and older who were undergoing routine cervical cancer screening \cite{16}. From a screening population of more than 300 000 women in a year, 3333 slides from HPV-positive women were included. From 238 training slides, 8215 DS-positive and 9739 DS-negative tiles were used for training (Figure 1). The study was approved by the KPNC IRB and was exempted from institutional review at the NCI by the Office of Human Subjects Research. Patient consent was waived because deidentified discarded specimens were used in this study.

**Clinical Endpoints**

All studies followed routine clinical practice at the respective institutions. Cytology was classified by the Bethesda System: negative for intraepithelial lesions or malignancy, atypical squamous cells of undetermined significance, low-grade squamous intraepithelial lesions, and high-grade squamous intraepithelial lesions (HSIL) \cite{26}. Final diagnosis was established by histopathology classified according to the cervical intraepithelial neoplasia (CIN) scale for cervical endpoints, which indicates the extent of dysplastic cells in the cervical epithelium: no indication for biopsy, normal CIN, grade 1 (CIN1), grade 2 (CIN2), grade 3 (CIN3), and cancer. We grouped adenocarcinoma in situ
with CIN3. For anal disease endpoints, the comparable anal intraepithelial neoplasia nomenclature (AIN) was used.

p16/Ki-67 Staining and Evaluation

For the Biopsy Study and ACSS, slides were prepared from residual PreservCyt material using a T2000 processor (Hologic, Bedford, MA). For the KPCN study, slides were prepared from residual SurePath tubes according to the manufacturer’s instructions (BD, Sparks, MD). Immunostaining of cervical cytology slides for p16/Ki-67 was performed using the CINtec Plus Kit (Roche, Tucson, AZ) according to the manufacturer’s instructions. DS-trained cytotechnologists reviewed all slides; a slide was considered positive if 1 or more cervical epithelial cell(s) stained both with a brown cytoplasmic stain (p16) and a red nuclear (Ki-67) irrespective of morphologic abnormalities. Slides from the Biopsy Study and ACSS were stained and evaluated at Roche mtm laboratories AG, Heidelberg, Germany, whereas slides from the Kaiser DS study were stained and evaluated at KPNC. HPV testing with partial genotyping (HPV16 and HPV18) at KPNC was based on the cobas assay (Roche, Pleasanton, CA).

Statistical Analysis

We created boxplots and calculated medians to show the distribution of DS-positive cells in cytology and histology categories. We compared differences in DS cell counts in ordinal cytology and histology categories using 1-way analysis of variance. The primary endpoint for the Biopsy Study and the Kaiser Study was CIN3 or greater (CIN3+). For ACSS, the primary endpoint was AIN2 or AIN3 (AIN2+). Receiver operator characteristics curve analysis was conducted for the number of DS-positive cells against the primary endpoints, and the area under the curve (AUC) was calculated. Sensitivity and specificity coordinates for manual DS evaluation and cytology were plotted on the receiver operator characteristics curve for comparison. We calculated percentage positivity, sensitivity, specificity, and Youden’s index in the Biopsy Study and ACSS for the cutoff determined by CNN4 and for manual DS evaluation. In the Kaiser Study, with a representative population of women who underwent routine screening, we calculated percentage positivity, sensitivity, specificity, and positive and negative predictive values for automated and manual DS. Differences in positivity, sensitivity, specificity, and specificity were evaluated using an exact McNemar’s $\chi^2$, and differences in predictive values were evaluated using the R package DTComPair, using the generalized score statistic (27). To evaluate clinical efficiency of each strategy, we estimated the number of CIN3+ detected for different cutoffs of DS-positive cells and the ratio of the number of tests and colposcopies per case of CIN3+ detected. We also evaluated the theoretical performance of automated DS in a fully vaccinated population by excluding all women who were positive for HPV16 and/or HPV18 from the analysis. Analyses were performed in SPSS, Stata, and R. All statistical tests were 2-sided and $P$ less than .05 was considered statistically significant.

Results

Automated Detection of DS-Positive Cells in Colposcopy and Anoscopy Populations

We developed a deep-learning algorithm for automated detection of DS-positive cells on ThinPrep slides from 2 referral populations (Biopsy Study and ACSS), including 212 training slides with 1186 DS-positive and 1485 DS-negative tiles (Figure 1). We evaluated the algorithm in independent validation slides from...
In both studies (Figure 1), we observed an increase in the number of DS-positive cells by increasing severity of cytology and histology, with higher absolute DS-positive cell numbers in ACSS (P < .001 for all comparison; Supplementary Figure 2, available online).

In the Biopsy Study validation set with 53 CIN3+, the AUC for detecting CIN3+ based on automated DS using CNN4 was 0.74 (Figure 2). At a cutoff of 3 DS-positive cells, the CNN4 algorithm had marginally lower positivity (58% vs 63%, respectively, P = .06) with comparable sensitivity (P = 1.0) and marginally higher specificity compared with manual DS (40.6% vs 45.7%, respectively, P = .07) (Table 1).

In the ACSS validation set with 69 AIN2+, the AUC for detecting AIN2+ based on automated evaluation of DS slides with CNN4 was 0.77 (Figure 2). At a cutoff of 3 DS-positive cells, the positivity of the CNN4 algorithm was lower (63% vs 71%, respectively, P = .001) with comparable sensitivity (P = 1.0) and higher specificity compared with manual DS (36.1% vs 46.1%, respectively, P = .001) (Table 1).

Automated Detection of DS-Positive Cells in an HPV Screening Population

We developed the deep-learning algorithm for SurePath slides using a training set of 238 slides from the Kaiser study with 8215 DS-positive and 9739 DS-negative tiles and applied it in an independent validation set of slides from 3095 women. We observed an increase of DS-positive cells with increasing severity of cytology and histology (Supplementary Figure 2, available online).

In the Kaiser validation study including 218 CIN3+, the AUC for detecting CIN3+ based on automated evaluation of DS slides was 0.82 (Figure 3). At a cutoff of 2 cells, the positivity of the...
algorithm was statistically significantly lower (42% vs 50%, respectively, \(P < .001\)) with equal sensitivity (\(P = 4\)) but statistically significantly higher specificity (61.5% vs 52.6%, respectively, \(P < .001\)) compared with the manual DS. At a cutoff of 100 cells, accuracy approached HSIL cytology that allows for immediate treatment according to current management guidelines (Figure 3). Automated DS provided better risk stratification compared with Pap cytology and manual DS (Figure 4): more women were reassured of a lower risk compared with the other strategies (58% for automated DS vs 50% for manual DS and 40% for cytology), and risk among positives was higher.

Clinical Efficiency of Automated DS Evaluation

We compared the clinical efficiency of Pap cytology, manual DS, and automated DS at 2 cutoffs (2 or more cells and 1 or more cells) for triage of HPV-positive women (Table 2). All DS strategies achieved equal or better sensitivity for detection of CIN3+ compared with Pap cytology while reducing unnecessary colposcopic referrals. Automated DS reduced overall referral to colposcopy by one-third for the primary automated cutoff of 2 cells (41.9% for automated DS vs 60.1% for cytology, \(P < .001\)). Automated DS at a cutoff of 2 or more cells had similar sensitivity but statistically significantly higher specificity compared with manual DS evaluation (61.5% vs 52.6%, \(P < .001\)). Automated DS detection at a cutoff of 1 or more cells achieved the highest sensitivity of all strategies, with statistically significantly higher specificity and lower colposcopy referral compared with Pap cytology. Automated DS at a cutoff of 2 or more cells had the most favorable ratio of colposcopies per CIN3+ detected compared with the least favorable ratio for the current standard, Pap cytology (6.8 vs 9.9, respectively). Extrapolating this to the full Kaiser screening population, out of 300,000 women screened annually, approximately 30,000 would test HPV-positive and more than 18,000 would be referred to colposcopy using the current approach with Pap cytology, while only 12,570 would be referred to colposcopy using automated DS. We also estimated the performance of automated DS in a fully vaccinated population. Similar to the overall evaluation, automated

Figure 3. Receiver operating curve characteristics analysis of number of dual stain (DS)-positive cells detected by CYTOREADER for detection of cervical precancer in a human papillomavirus screening population in Kaiser Permanente Northern California. ASC-US = Atypical Squamous Cells of Undetermined Significance; AUC = area under the curve; HSIL = high-grade squamous intraepithelial lesions.

Figure 4. Absolute risk of precancer for Pap cytology, manual dual stain (DS), and automated DS. ASCUS+ = positive for Atypical Squamous Cells of Undetermined Significance or greater cytology results. The dotted lines show clinical action risk thresholds for colposcopy referral (4% risk) and immediate treatment (50% risk).
**Table 2. Accuracy for cervical precancer based on Pap cytology and manual and automated detection of DS-positive cells on SurePath slides in the Kaiser Validation Study (N = 3095)**

| Evaluation          | Pap cytology (1 cell) (201 CIN3+) | Pap cytology (2 cell) (197 CIN3+) | Pap cytology (3 cell) (197 CIN3+) |
|---------------------|-----------------------------------|-----------------------------------|-----------------------------------|
|                     | Manual DS (197 CIN3+)              | Automated DS (1741)               | Automated DS (1298)               |
| Colposcopy referral, No. (%) | 1850 (60.1)                        | 1536 (48.6)                       | 1298 (41.7)                       |
| Sensitivity, % (95% CI) | 0.90 (0.81 to 0.99)               | 0.89 (0.82 to 0.96)              | 0.89 (0.82 to 0.95)              |
| Specitivity, % (95% CI) | 0.49 (0.43 to 0.56)               | 0.54 (0.50 to 0.59)              | 0.56 (0.52 to 0.60)              |
| PPV, % (95% CI)      | 0.99 (0.96 to 1.00)               | 0.99 (0.96 to 1.00)              | 0.99 (0.96 to 1.00)              |
| NPV, % (95% CI)      | 0.46 (0.42 to 0.49)               | 0.46 (0.42 to 0.49)              | 0.46 (0.42 to 0.49)              |

*Two-sided McNemar’s test. CIN3 = cervical intraepithelial neoplasia grade 3 or worse; DS = dual stain; NPV = negative predictive value; PPV = positive predictive value; Ref = referent.

**Discussion**

Using a rigorous study design, we developed a novel deep-learning-based image analysis platform for automated evaluation of DS cytology. In a large population of women undergoing HPV-based cervical cancer screening, we show that automated evaluation of DS slides dramatically increases the efficiency of cervical cancer screening by substantially reducing unnecessary colposcopies compared with current standards and similarly achieves excellent performance in a simulated fully vaccinated population. Thus, CYTOREADER exceeds human diagnostic accuracy and serves as an example of AI achieving improvements beyond the automation of a human standard.

Our results demonstrate how automation and machine learning can transform cervical cancer screening that is currently undergoing major changes. HPV testing for cervical cancer screening is an objective and reliable approach directly linked to the carcinogenic process (28). HPV-negative women are at very low risk of developing precancer or cancer over the next decade and screening intervals can be extended (8–10). Yet most HPV infections are transient, and women require additional tests to decide who needs further evaluation or treatment (11, 12). Pap cytology is recommended and approved for triage of HPV-positive women but suffers from subjectivity, lack of reproducibility, and relatively low sensitivity (14). Our previous study comparing manual DS to cytology together with the current results demonstrates that automated DS evaluation can supplant and improve the role of Pap cytology for triage of HPV-positive women and should also be evaluated for postcolposcopy and posttreatment surveillance (16). Compared with Pap cytology, manual DS has higher accuracy and can provide longer reassurance against disease when a test is negative, while the risks to patients do not differ from Pap cytology, because the same sample type is used (17, 21). We previously showed that the few DS-negative CIN3s are more likely to have no HPV16/18 and no high-grade cytology, suggesting that these cases are less likely to progress (16). Automated DS evaluation can provide a completely objective cervical cancer-screening approach, improving efficiency and reducing harms and cost related to false-positive screening results. Furthermore, by demonstrating that AI-based DS detection works for anal cytology, we show the robustness of the imaging and analysis platform. Importantly, our approach is also suited for vaccinated populations, where it may achieve even higher specificity and counterbalance the lower disease prevalence in vaccinated women (29).

Automated DS evaluation immediately quantifies the number of DS-positive cells on a slide, allowing tailoring positivity cutoffs for specific clinical decisions. Current guidelines give an option for immediate treatment in women with HSIL cytology, who have a very high probability of having underlying CIN3+ (30). A higher cutoff of DS-positive cells could be used to guide treatment decisions. Moving forward, additional criteria can be developed to expand slide assessment; for example, the presence of abnormal glandular cells to identify adenocarcinoma precursors, which is a particular challenge for Pap cytology (31).

Digitization of glass slides paired with automated evaluation in the cloud can provide high-throughput triage of HPV-positive women with inherent objectivity. Furthermore, the functionality of CYTOREADER can provide an assisted diagnostics mode
for evaluating DS slides. The automatic algorithm can be used for presenting all DS-positive cells found on a slide ranked by the likelihood that a cell is DS-positive to accelerate slide evaluation. Similarly, CYTOREADER can be used for quality control of a program that is based on manual DS evaluation.

Successful implementation of CYTOREADER requires an infrastructure for high-quality staining, full-slide scanning, and running the machine-learning algorithm. However, slide preparation, scanning, and slide evaluation can be geographically separated, providing high-quality cervical cancer screening and triage in locations that currently do not have infrastructure and training to achieve reliable DS evaluation given a reliable courier system is available. Compared with manual evaluation of DS slides, the automated evaluation requires access to scanning infrastructure but may require a smaller cytotechnology workforce. Scanners are increasingly available in pathology laboratories and can process large batches of slides with limited need for a skilled operator (22, 23). Studies are warranted to evaluate if DS is amenable to self-collected specimens, a sampling strategy that is important for low-resource settings. Future efforts also need to evaluate how long a negative automated DS result provides reassurance against precancer and how automated DS can be used in women undergoing surveillance.

We conducted a large, well-powered study to evaluate performance of automated DS for triage of HPV-positive women. However, some limitations should be noted. In contrast to the large KPNC study on HPV triage based on SurePath slides, 2 studies using ThinPrep slides were comparably small, and they were conducted in colposcopy/anoscopy populations. Future studies need to evaluate automated DS in larger HPV screening populations using ThinPrep slides. Also, the positivity and sensitivity of cytology at KPNC is much higher compared with other settings, which may affect the comparison of clinical efficiency estimates.

Our approach to train and validate both on the tile level and the slide level with ground truth disease endpoints sets our work apart from other deep-learning approaches in digital pathology that focus on replicating a subjective evaluation. We recognize that there is substantial subjectivity underlying histologic endpoints of cervical disease (15). In our study, we minimized the impact by relying on the most reproducible correlate of cervical precancer, CIN3, as our primary endpoint for evaluation of triage of HPV-positive women. Our work also emphasizes the importance of integrating epidemiology and AI with the availability of population bases studies to improve medical diagnostics beyond automation. It has been proposed for a long time that “digital pathology” will become an important cornerstone of future health care. Despite this vision, image analysis currently does not contribute substantially to routine clinical practice and to the benefit of the patient. The automated evaluation of DS cytology slides has substantially improved accuracy and efficiency compared with Pap cytology and serves as an important example for introducing digital pathology and deep learning into clinical practice. This approach has the potential to substantially improve screening program performance, potentially affecting millions of women testing HPV-positive in cervical cancer screening each year.

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