

The Normal Epithelium of Crypts Accruing Below Nonpolypoid Adenomas Thrives With Relocated Proliferating Cell-domains and p53-Up-regulated Cells

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Abstract. *Background/Aim:* Colonic crypts with normal epithelium albeit with corrupted shapes (CCS) were previously found beneath nonpolypoid adenomas (NPA). This study aimed to analyze the distribution of proliferating cells (PC) and p53-up-regulated cells in CCS. *Materials and Methods:* Sections from 48 NPA were immunostained with the proliferating-marker Ki67 and against the tumor-suppressor protein p53. *Results:* Asymmetric-haphazardly distributed PC were found in 87.5% of the NPA, continuous PC-domains in 8.3%, asymmetric-haphazardly distributed single PC in 4.2% and p53-up-regulated cells in 29.2%. In 12 controls, the normal-shaped crypts revealed symmetrically-distributed PC-domains in their lower thirds, and no p53-up-regulated cells. *Conclusion:* The normal epithelium that lines the CCS below NPA, thrives with relocated PC-domains, and with occasional p53-up-regulated cells. These findings strongly suggest that the normal epithelium of CCS beneath NPA might harbor somatic mutations. The accretion of putative mutated CCS might play an important role in the evolution of nonpolypoid adenomas in the human colon.

In 1985 Muto *et al.*, detected at colonoscopy, small nonpolypoid mucosal lesions measuring up to one cm in diameter (1). At histologic examination, those lesions were found to be adenomas. Flat adenomas (FA), as they were called, were subsequently found associated with a more aggressive clinical behavior than their polypoid counterpart (2).

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The further improvement of optical devices together with the use of mucosal dyes, permitted endoscopists to more easily visualize previously undetected FA. In 1994, Jaramillo *et al.* (3) identified by the aid of high-resolution video colonoscopy and chromoscopy, a colon carcinoma arising in a FA. In 1995, the same group reported a collection of 109 flat neoplastic lesions: 97% were FA and the remaining 3%, FA with submucosal invasion (4). In 1999, one of us (CAR), trained in Western and Japanese pathology (5), reviewed 781 FA collected at the four main Hospitals in Tokyo, Japan and at the Karolinska Hospital, Stockholm, Sweden (6). The results showed that high-grade dysplasia was present in 42% out of 420 FA in Japanese patients, but only in 14% of the 361 FA in Swedish patients. Intramucosal or submucosal carcinomas were found in 15% of the FA in Japan, but only in 3% of the FA excised in Sweden (6). Thus, Japanese patients had more advanced FA (in terms of high-grade dysplasia) and more aggressive FA (in terms of intramucosal and submucosal invasion) than Swedish patients (6, 7). Those findings confirmed the claims of histological differences between flat colonic neoplasias in Japan and in Western countries (8, 9). According to Gotoda, the differences in the histological classification of gastrointestinal epithelial neoplasias between Western and Japanese pathologists, have been largely resolved by adopting the Vienna classification (10, 11).

In 1997, Kudo *et al.* reported non-polypoid (flat) colon neoplasms with a lateral diameter greater than 10 mm; they were called laterally-spreading tumors (LST) (12). In subsequent surveys, Kim *et al.* (13) found that 85% of the 497 LSTs were nonpolypoid adenomas, Zhao *et al.* (14) that 88% of the 239 LST were nonpolypoid adenomas and Tang *et al.* (15) that all 33 (100%) LST investigated were nonpolypoid adenomas. From the above, it is evident that nonpolypoid adenomas account for the majority of the FA and LST lesions in the human colon. Of note, Sato *et al.* (16) followed-up 14 patients with flat colonic tumors with colonoscopic examinations for 19 months (range=11-26 months). During that

time interval, none of the flat tumors measuring approximately 5 mm in diameter progressed to LST. Notwithstanding, it remained unclear as to whether LST evolved *in situ* as fully developed neoplastic lesions from the start, or progress through the lateral extension of pre-existing FA.

In previous studies, we found beneath the adenomatous tissue of polypoid conventional adenomas (CoA), both in rats (17) and in humans (18), accruing colonic crypts lined with normal epithelium albeit with corrupted shapes (CCS). More recently, we found abnormally distributed proliferating cells (PC) and p53-up-regulated cells in CCS beneath the neoplastic canopy of polypoid CoA (19).

The crucial question was: Could the CCS found beneath the neoplastic tissue of nonpolypoid adenomas (20) also exhibit a disparate distribution of PC and reveal p53 up-regulated cells? In attempts to answer these questions, we explored the distribution of proliferating cells and the possible occurrence of p53 immuno-reactive cells in CCS found in a cohort of nonpolypoid adenomas.

Patients and Methods

The material consists of sections from 48 endoscopically-removed nonpolypoid colon adenomas (NPA), without submucosal invasion. Histological sections (4 µm-thick) were retrieved from the files of the Gastrointestinal Research Laboratory of this Department, stained with hematoxylin and eosin (H&E), and immuno-stained with the proliferation marker Ki67 (batch MIB1, DAKO Automation, 2600 Glostrup, Denmark), and the primary mouse monoclonal antibody (IgG1, kappa) directed against human tumor suppressor p53 protein (antip53, DO-7; Ventana Medical System, Inc, Roche, Basel, Switzerland).

PC and p53 up-regulated cells in CCS. Using 10x oculars and a 4x Apo objective (aperture number 0.20), the Field of Vision (FOV) was 5 mm in diameter. At this magnification, all Ki67 labelled and p53 up-regulated cells in CCS could easily be identified. p53 up-regulated cells were regarded only those having the same intense immunoreactivity for the p53 tumor-suppressor protein as those in the neoplastic tissue on top.

PC in normal colon crypts from historical controls. Sections from 12 grossly normal colonic segments proximal or distal to the surgically removed colonic adenocarcinoma were immunostained with Ki67 (batch MIB1). The PC distribution in the crypts were assessed using a 5mm FOV20.

Statistical analysis. The non-parametric Kruskal-Wallis test was applied, to compare difference between groups. Statistical significance was defined as $p < 0.05$.

This study was approved by The Regional Ethical Review Board in Stockholm (No. 2018/113-32 and 2018/688-32).

Results

PC phenotypes in CCS. When the Ki67 antibody reacts with its cognate antigen, all transit amplifying daughter cells

Table I. The frequency of predominant proliferating cells (PC) phenotypes in crypts with normal epithelial lining but with corrupted shapes (CCS) found underneath the neoplastic canopy in 48 nonpolypoid adenomas of the colon.

Predominant PC phenotype	No. cases (%)
Haphazardly distributed PC clusters*	42 (87.5%)
Continuous PC-domain	4 (8.3%)
Haphazardly-distributed Single PC	2 (4.2%)
Total	48 (100%)

*≥two consecutive PC.

(TADs) become labelled. The following PC-phenotypes were detected in the 48 NPA: i) haphazardly distributed PC clusters (≥two consecutive PC), ii) continuous PC-domain in one or in both sides of entire crypts, and iii) haphazardly-distributed single PC. Since the PC-distribution in CCS could vary in individual NPA, the predominant PC-phenotype in CCS was selected. The intense PC found in the adenomatous tissue above CCS officiated as internal controls.

The distribution of PC in normal epithelium of CCS below the neoplastic tissue of nonpolypoid adenomas. The frequency of predominant PC-phenotypes in the CCS from the 48 NPA is condensed in Table I; examples of PC-phenotypes are illustrated in Figure 1 (upper panel). The Table shows that CCS with asymmetric, haphazardly distributed PC-clusters were recorded in 87.5% (42/48) of the NPA, with continuous PC in one or both slopes of the crypts, in 8.3% (4/48), and with haphazardly distributed single PC in the remaining 4.2% (2/48). The difference between NPA with CCS exhibiting haphazardly distributed PC-clusters and the other two groups in Table I, was significant ($p < 0.05$).

p53 up-regulated cells in the neoplastic tissue of nonpolypoid adenomas. The tumor-suppressor protein p53 was overexpressed in the adenomatous tissue in 39.6% (19/48) of the NPA. This percentage was somewhat higher than that found in a previous survey at this department, where 25% out of the 433 polypoid conventional adenomas showed p53-overexpression in the neoplastic tissue (21).

p53 up-regulated cells in CCS below the neoplastic tissue of nonpolypoid adenomas. In 29.2% (14/48), the CCS beneath NPA showed p53-up-regulated cells; examples are illustrated in Figure 1 (lower panel). No p53-up-regulated cells could be demonstrated in the CCS in the remaining 70.8% (34/48).

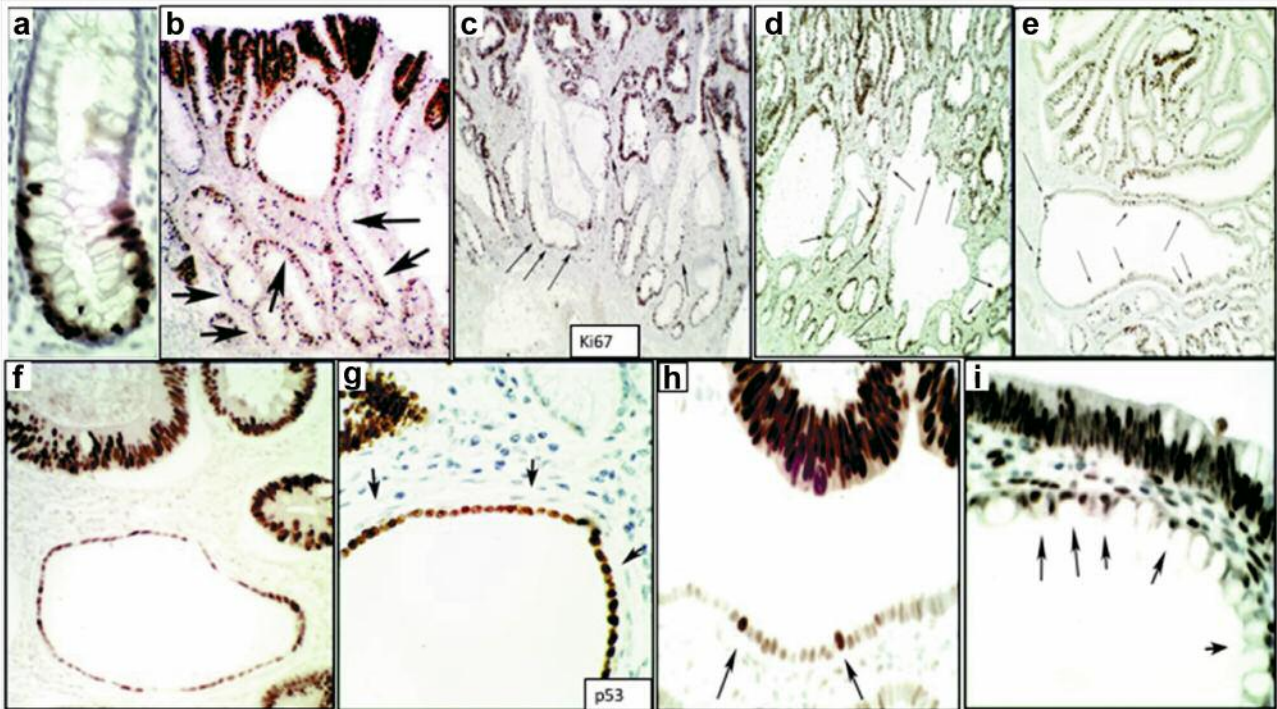


Figure 1. Cell proliferation and p53-up-regulated cells in the normal epithelium of crypts with disparate shapes found beneath nonpolypoid colon adenomas. Upper panel; a: Proliferating cells (PC) in a control crypt (normal colon). Note symmetrically-aligned PC, in both slopes of the lower third of the crypt (control, Ki67, batch MIB1, original $\times 40$), b to e: Cell proliferation in crypts lined with normal epithelium although with corrupted shapes (CCS) found below the neoplastic tissue of nonpolypoid adenomas (NPA), b: Crypts lined with normal epithelium but with CCS found below the neoplastic tissue of a NPA (on top). Note asymmetric-haphazardly distributed PC-domains (Ki67, batch MIB, original $\times 20$), c, d, e: Dilated CCS displaying disparate distributions of PC (Ki67, batch MIB, $\times 10$). Lower panel; f-i: p53-up-regulated cells (arrows) in crypts lined with normal epithelium although with CCS found below the neoplastic tissue of NPA. Note the neoplastic tissue of nonpolypoid adenomas on top (p53 immunostaining, f: original $\times 20$, g: original $\times 40$, h and f: original $\times 40$).

Discussion

This study showed that the normal epithelium of CCS –an integral component of colonic NPA– harbors disparate, haphazardly distributed PC, often in asymmetric PC-clusters or as single PC and unexpectedly, haphazardly distributed p53 up-regulated cells. It became apparent that the CCS underneath NPA thrive with an anomalous operational make-up. In contrast, the normal crypts from control cases displayed symmetrically aligned PC lengthwise their lower thirds, and showed no p53-up-regulated cells.

In the normal colon, the stem cells are located at the base of crypts; stem cells synchronize the repopulation of the crypts by generating progenitor cells called transient amplifying daughter (TAD) cells (22-24). Progenitor cells [120 to 150 TAD cells/crypt according to Testa *et al.* (25)] account for the bulk of the PC in the crypts. Since in normal crypts the PC are generated by stem cells (22-24), the occurrence of multiple PC-domains in the TADs of CCS,

rationality implies that several stem cells exist in these crypts. This deduction is in concert with studies in humans by Baker *et al.* (22), showing that 5 to 6 stem cells are to be found in each normal colon crypt. Thus, at variance with the natural position of stem cells at the base of normal crypts (26-28), the stem cells in CCS seem to have been relocated (rearranged?). The relocation of the normal position of PC in CCS and the presence of up-regulated p53-domains in the normal epithelium of CCS in 29% of the NPA, support the notion that their normal-looking epithelium might had been subjected to somatic mutations.

The crucial question is why some adenomas are nonpolypoid while others are polypoid? In this regard, Voorham *et al.* (29) found that the patterns of DNA copy number changes differed between the two phenotypes. Whereas loss of 5q14.3 and 5q15-q3 were significantly more frequent in flat adenomas, losses of 1p36.32-p35.3, 10q25.3, 17p12, and chromosome 18 were more frequent in polypoid adenomas (29). In addition, APC mutations were more

frequent in polypoid adenomas compared to flat adenomas, suggesting that the disruption of the Wnt-pathway may occur via different mechanisms in these two phenotypes (30). Takahashi *et al.* (31) also found genetic alterations that might play an important role in the development of flat-type advanced adenomas, especially in the distal colon. Epigenetic alterations occurred infrequently in flat-type advanced adenomas, suggesting that they have different genetic and epigenetic alterations than those in protruded-type advanced adenomas (31). From the above it is evident that the molecular make-up in NPA is different to that in polypoid adenomas. Voorham *et al.* (29) found no significant association between the different morphological phenotypes and mutations in key genes of the RAS-RAF-MAPK.

Another pertinent question is: Which are the morphogenetic mechanisms that induce colonic crypts, lined with normal epithelium, to assume corrupted shapes beneath NPA? Morphogenesis stands for the ability of a system to change its form. Jagan *et al.* (32) demonstrated that the formation of colorectal crypts is regulated by phosphatase and tensin homologue deleted on chromosome 10 (PTEN), a protein encoded by the PTEN gene (32). In addition, Georgescu *et al.* (33) in three-dimensional studies of human colon glands, found that NHERF1 protein, a Na⁺/H⁺ exchanger regulatory factor, controls gland morphogenesis. Thus, the CCS-phenotype in NPA might have been generated by alterations in morphogenesis-signals such as NHERF1 and PTEN.

Goodblad *et al.* (34) showed that there are up to 15 times more mitotic figures in a whole colonic crypt than in the 4 µm-thick histological sections that were used in their studies. Since the number of proliferating cells/crypt is much higher than the number of mitotic cells/crypt (34), it is not inconceivable that CCS showing an aberrant PC pattern could contain a higher number of DNA-synthesizing cells or clusters elsewhere, in other areas from the same crypt. It should be stressed that to assess PC in NPA, we also used 4 µm-thick sections.

Boman and Fields (24), stated that normal crypts began to show abnormalities in histology only when they became dysplastic. In contrast, we found that the normal epithelium of CCS in NPA (35-37) had already experienced profound biomolecular alterations.

In sum, the normal epithelium in CCS accruing beneath NPA in the colon disclosed an unprecedented relocation (reorganization?) of PC-domains. Rationally, the relocation of PC might had been preceded by the relocation of the cells that fuel PC, namely the stem cells (22-24). In addition, the CCS of a fraction of NPA disclosed p53-up-regulated cells, strongly suggesting that the normal epithelium in those crypts might harbor somatic mutations. Considering that human colonic crypts typically divide at most once or twice during a lifetime, with an average crypt cycle length of 36

years (22), the accretion of putative mutated CCS emerges as a major event that might play an important role in the evolution of NPA in the human colon.

Conflicts of Interest

The Authors declare no competing interests regarding this study.

Authors' Contributions

CAR: Designed the experiment, performed procedures, data analysis and wrote the manuscript; PTS harvested endoscopically nonpolypoid adenomas, introduced suggestions and approved the final manuscript.

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