

Evaluation of PTGS2 Expression, PIK3CA Mutation, Aspirin Use and Colon Cancer Survival in a Population-Based Cohort Study

Ronan T. Gray, MB, BCh (Hons), MSc, MRCS¹, Marie M. Cantwell, PhD, MPH², Helen G. Coleman, PhD¹, Maurice B. Loughrey, BSc, MRCP, FRCPath, MD^{3,4}, Peter Bankhead, BD, MSc, PhD³, Stephen McQuaid, PhD, FRCPath^{3,5}, Roisin F. O'Neill, PhD¹, Kenneth Arthur, MMedSc, FIBMS, CSci³, Victoria Bingham, MSc³, Claire McGready, BSc³, Anna T. Gavin, MB, BCh, MSc, FFPHM⁶, Chris R. Cardwell, PhD¹, Brian T. Johnston, MD, FRCP⁷, Jacqueline A. James, PhD, FRCPath^{3,4,5}, Peter W. Hamilton, BSc, PhD, FRCPath (Hons)³, Manuel Salto-Tellez, MD-LMS, FRCPath, FRCPI^{3,4} and Liam J. Murray, MD, MRCGP¹

OBJECTIVES: The association between aspirin use and improved survival after colorectal cancer diagnosis may be more pronounced in tumors that have PIK3CA mutations or high PTGS2 expression. However, the evidence of a difference in association by biomarker status lacks consistency. In this population-based colon cancer cohort study the interaction between these biomarkers, aspirin use, and survival was assessed.

METHODS: The cohort consisted of 740 stage II and III colon cancer patients diagnosed between 2004 and 2008. Aspirin use was determined through clinical note review. Tissue blocks were retrieved to determine immunohistochemical assessment of PTGS2 expression and the presence of PIK3CA mutations. Cox proportional hazards models were used to calculate hazard ratios (HR) and 95% confidence intervals (CI) for colorectal cancer-specific and overall survival.

RESULTS: In this cohort aspirin use was associated with a 31% improvement in cancer-specific survival compared to non-use (adjusted HR = 0.69, 95% CI 0.47–0.98). This effect was more pronounced in tumors with high PTGS2 expression (PTGS2-high adjusted HR = 0.55, 95% CI 0.32–0.96) compared to those with low PTGS2 expression (PTGS2-low adjusted HR = 1.19, 95% CI 0.68–2.07, *P* for interaction = 0.09). The aspirin by PTGS2 interaction was significant for overall survival (PTGS2-high adjusted HR = 0.64, 95% CI 0.42–0.98 vs. PTGS2-low adjusted HR = 1.28, 95% CI 0.80–2.03, *P* for interaction = 0.04). However, no interaction was observed between aspirin use and PIK3CA mutation status for colorectal cancer-specific or overall survival.

CONCLUSIONS: Aspirin use was associated with improved survival outcomes in this population-based cohort of colon cancer patients. This association differed according to PTGS2 expression but not PIK3CA mutation status. Limiting adjuvant aspirin trials to PIK3CA-mutant colorectal cancer may be too restrictive.

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INTRODUCTION

Observational studies and long-term follow-up of randomized trials assessing aspirin use in cardiovascular disease indicate that regular aspirin use is associated with a reduced risk of colorectal neoplasia.¹ Further long-term follow-up of these trials suggests aspirin use is associated with a reduction in both the incidence and subsequent development of metastatic disease.² Similarly, a recent population-based cohort study with over 23,000 colorectal cancer patients demonstrated that post-diagnostic aspirin use was associated with improved cancer-specific survival.³ Pre-clinical evidence confirms the biological plausibility of these associations and suggests aspirin may have pleiotropic anti-cancer effects.⁴ Although the

exact mechanisms remain unclear they are likely to involve prostaglandin-endoperoxide synthase 2 (PTGS2, also known as cyclooxygenase-2 or COX-2) dependent and independent pathways.^{4,5} Taken together, this data suggests aspirin could have utility in the adjuvant treatment of colorectal cancer.

A number of clinical trials assessing the role of adjuvant aspirin therapy (low- and high-dose) in colorectal cancer have recently started recruitment (NCT00565708, NCT02607072, NCT02301286, ISRCTN74358648). However, the potential side effects associated with aspirin use, including major gastrointestinal bleeding, and the understanding that colorectal cancers are heterogeneous has led to attempts to better

¹Cancer Epidemiology and Health Services Research Group, Centre for Public Health, Queen's University Belfast, Belfast, Northern Ireland; ²Nutrition and Metabolism Group, Institute for Global Food Security, Queen's University Belfast, Belfast, Northern Ireland; ³Northern Ireland Molecular Pathology Laboratory, Centre for Cancer Research and Cell Biology, Queen's University Belfast, Belfast, Northern Ireland; ⁴Tissue Pathology, Belfast Health and Social Care Trust, Belfast, Northern Ireland; ⁵Northern Ireland Biobank, Queen's University Belfast, Belfast, Northern Ireland; ⁶Northern Ireland Cancer Registry, Queen's University Belfast, Belfast, Northern Ireland and ⁷Department of Gastroenterology, Belfast Health and Social Care Trust, Belfast, Northern Ireland

Correspondence: R.T. Gray, MB BCh (Hons), MSc, MRCS, Cancer Epidemiology and Health Services Research Group, Centre for Public Health, Queen's University Belfast, Royal Victoria Hospital, Belfast BT12 6BA, Northern Ireland. E-mail: rgray05@qub.ac.uk

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identify those patients who may derive a survival benefit.^{6,7} Allowing for tumor heterogeneity, molecular pathological epidemiology studies can facilitate assessment of the interaction between molecular biomarkers and lifestyle or other exogenous factors and the subsequent effect on cancer survival.^{5,8} As a result, they can provide mechanistic insight into the anti-cancer effect of the exposure being assessed and contribute to the development of precision medicine.⁵

PTGS2 is the rate limiting enzyme in the conversion of arachidonic acid to prostaglandins.⁹ Prostaglandins have the potential to enhance the neoplastic process by inhibiting apoptosis, stimulating proliferation and promoting angiogenesis.⁴ Phosphatidylinositol 3-kinase (PI3K) signaling pathways also play an important role in colorectal cancer carcinogenesis.⁶ Phosphatidylinositol-4,5-bisphosphate 3-kinase (PIK3CA) encodes the p110 α catalytic subunit of PI3K and mutations in this gene constitutively activate the PI3K pathway.^{7,10} Subsequent downstream activation of protein kinase B (AKT) results in PTGS2 upregulation and the potential to inhibit apoptosis.^{11,12} Overexpression of PTGS2 and the presence of PIK3CA mutations have been shown to differentiate the survival benefit associated with aspirin use in some^{6,7,9} but not all^{7,13,14} colorectal cancer observational studies. As the evidence of a difference in association by biomarker status lacks consistency further validation is required.

Therefore, the aim of this study was to assess the interaction between aspirin use, the biomarkers PTGS2 and PIK3CA, and survival in a population-based cohort study of patients with stage II and III colon cancer.

METHODS

Study cohort. All stage II and III colon cancer patients (International Classification of Disease code C18) diagnosed in Northern Ireland between 2004 and 2008 were identified from the Northern Ireland Cancer Registry ($n=1,862$). Rectal cancers were excluded from this molecular pathological epidemiology study due to the potential for neoadjuvant radiotherapy to alter tumor expression profiles. Following review of pathology reports the cohort was restricted to patients with a single primary colon adenocarcinoma who had undergone a surgical resection ($n=1,539$). A further 113 patients were excluded after clinical note review for reasons listed in **Figure 1**. To facilitate formalin fixed, paraffin-embedded (FFPE) tissue block retrieval the final cohort was restricted to patients from two of the five Northern Ireland Health and Social Care trusts within the remit of the Northern Ireland Biobank (740 of 1,426, 51.9%). These patients were representative of the overall Northern Ireland cohort in terms of age, sex, tumor stage, adjuvant chemotherapy use and proportion of deaths that occurred (see **Supplementary Table S1** online).

Clinicopathological variables and follow-up. Clinical variables including adjuvant chemotherapy use, prescription medication use, family history of colorectal cancer and Eastern Cooperative Oncology Group (ECOG) performance

status were retrieved from the Northern Ireland Clinical Oncology Information System (COIS), a prospective electronic record of cancer patient management. This process was supplemented by a manual chart review when insufficient information was recorded on COIS or when patients were not referred to an oncologist for consideration of adjuvant therapy. Stage and grade of differentiation were extracted from pathology reports. All patients were followed up for occurrence and cause of death via linkage to the Northern Ireland Registrar General's Office up to 31st December 2013. Colorectal cancer-specific deaths were defined as those with an underlying cause of death International Classification of Disease code C18, C19, C20 (anus) and/or C26 (other and ill-defined digestive organs).

Assessment of aspirin use. Prescription aspirin use (user vs. non-user) was assessed early in the post-operative period for all patients. When medication information was available on COIS this time point was the initial post-operative oncology review where current regularly prescribed medication use is recorded. When medication information was not available on COIS aspirin exposure was determined from the post-operative hospital discharge letter. Information on medication dosage was not consistently recorded on COIS. Our research group has previously demonstrated that 98.5% of aspirin prescriptions after colorectal cancer diagnosis in the UK are for low-dose (75 mg) aspirin.¹⁵ Exposure in this study is therefore considered representative of low-dose aspirin.

Tumor molecular analysis

FFPE block retrieval and tissue microarray construction. The Northern Ireland Biobank retrieved slides from all of the resection specimen blocks for each case. These original hematoxylin and eosin-stained slides were reviewed by an expert gastrointestinal histopathologist to select the two most representative FFPE tumor-rich blocks. Following block retrieval, a new section was cut for hematoxylin and eosin staining and the slides were annotated for tissue microarray (TMA) construction. Blocks were retrieved for 89.3% of the cohort (661 out of 740). Three representative areas within the donor blocks of each case were annotated for targeted coring. Three 1.0mm diameter tissue cores were then extracted from donor blocks and inserted into recipient blocks using a manual tissue arrayer (Estigen, Tartu, Estonia) as described previously.^{16,17}

PTGS2 immunohistochemistry, scoring and assessment. All immunohistochemistry was performed in the Northern Ireland Molecular Pathology Laboratory, a research laboratory which has UK Clinical Pathology Accreditation. In brief, all immunohistochemistry was performed on the fully automated Ventana BenchMark XT platform using an OptiView antibody detection system (Ventana Medical Systems Inc., Tucson, USA) and a Cell Marque ready to use rabbit monoclonal antibody to PTGS2 (Ventana, catalog number 760-4254). *QuPath* (Queen's University Belfast, Northern Ireland) image analysis software facilitated digital immunoscore. More detailed information is provided in the Supplementary Methods (online). A H-score was calculated based on the extent and intensity of cytoplasmic staining ($3 \times \%$ of

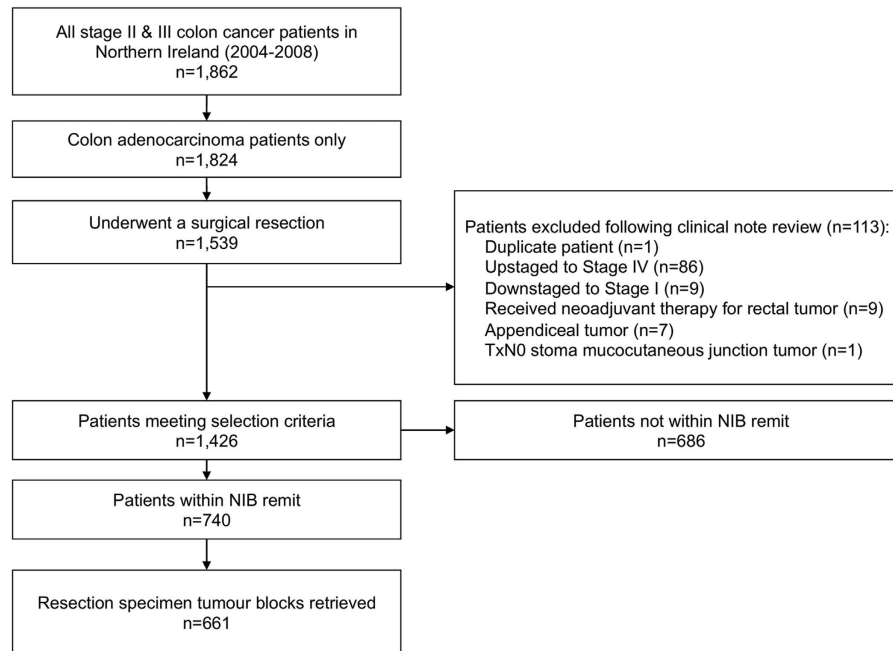


Figure 1 Selection of stage II and III colon cancer (adenocarcinoma) patients and samples. NIB, Northern Ireland Biobank.

strongly staining cytoplasm+2×% of moderately staining cytoplasm+1×% of weakly staining cytoplasm, giving a range of 0–300).¹⁸

Independently, an expert gastrointestinal histopathologist assessed PTGS2 expression in a random 10% sample (207 of the triplicate TMA cores) using the standardized grading system (absent, weak, moderate or strong immunostaining) described by Chan *et al.*⁹ Tumors with low PTGS2 expression (PTGS2-low) were categorized as those demonstrating weak or absent immunostaining while PTGS2-high tumors had moderate or strong levels of immunostaining. Receiver-operating characteristic analysis was subsequently used to determine the optimal H-score cutoff point that predicted PTGS2 grading with maximal sensitivity and specificity (according to Youden's J statistic using Cutoff Finder¹⁹). Using this methodology, a H-score of 145.5 differentiated PTGS2-low from PTGS2-high tumors with good interobserver agreement between the digital H-score and the histopathologist's grading (agreement 86.5%, $\kappa=0.70$). In the final data set, PTGS2 expression grading was determined using the median H-score from the three cores available for each case. Values above and below 145.5 were considered PTGS2-high and PTGS2-low respectively. Representative images are shown in **Figure 2**.

DNA extraction. Tumor-rich areas of the representative blocks from each case were annotated for macrodissection. DNA was extracted according to the manufacturer's instructions from five 5 μ m sections using the Maxwell 16 Instrument (Promega, Southampton, UK) and Promega DNA extraction kit. Quantification of DNA was performed by an absorbance method using the NanoDrop 2000c spectrophotometer (Thermo Scientific, Wilmington, USA).

PIK3CA mutation and microsatellite instability status. Mass spectrometry incorporating the pre-validated ColoCarta Panel (Agena Bioscience, Hamburg, Germany) was used to detect the presence of PIK3CA mutations. Samples were processed at the Genomics Core Technology Unit (Queen's University Belfast, Northern Ireland) and the Assays by Agena Custom Services Laboratory (Hamburg, Germany) using previously described methods.²⁰ Detection of somatic mutations at a frequency of >10% for any of eight alleles (C420R, E542K, E545K, H701P, H1047R, H1047L, Q546K, or R88Q) was considered evidence of a mutation in the PIK3CA gene. A failed reaction at a single position resulted in missing data for PIK3CA status only if the reactions at other positions were wild-type.²¹ This panel includes assays that capture 79% of known PIK3CA mutations including the 'hot spot' mutations in exons 9 and 20 (Q546K, E545K, E542K, H1047R, and H1047L), which confer constitutive kinase activity.¹⁰

Microsatellite instability (MSI) analysis was performed within the Northern Ireland Molecular Pathology Laboratory according to manufacturer's instructions using the MSI Analysis System, version 1.2 kit (Promega, Southampton, UK) for five mononucleotide repeat markers (BAT-25, BAT-26, NR-21, NR-24 and MONO-27). PCR products were separated by capillary electrophoresis using an ABI 3500 Genetic Analyzer (Fisher Scientific—UK Ltd, Loughborough, UK). The output data were analyzed using GeneMapper v4.1 (Fisher Scientific—UK Ltd, Loughborough, UK) to determine MSI-high (MSI-H) status (MSI-H vs. non MSI-H).

Statistical analysis. All statistical analysis was performed using Stata 14 (StataCorp, College Station, TX, USA). The χ^2 test was used to compare characteristics between aspirin

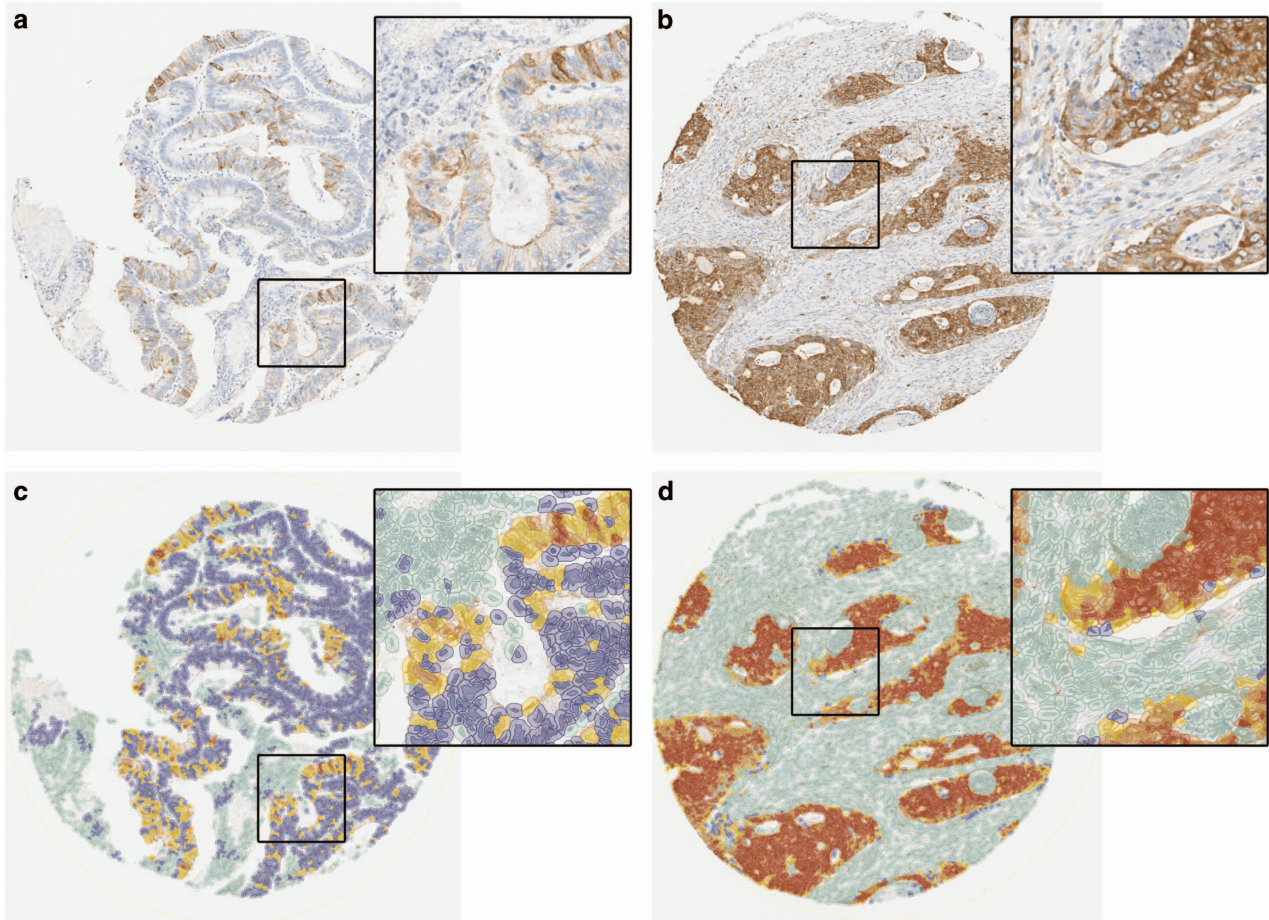


Figure 2 PTGS2 immunohistochemistry in colon cancer tissue microarrays (TMAs) and associated mark-up for digital immunoscore using QuPath image analysis software. Detected cells are color-coded according to their classification: green, non-tumor; blue, negatively staining tumor; yellow, weakly staining tumor; orange, moderately staining tumor; red, strongly staining tumor. (a) Original core from a tumor weakly expressing PTGS2 (PTGS2-low). (b) Original core from a tumor strongly expressing PTGS2 (PTGS2-high). (c) QuPath cellular mark-up in the PTGS2-low core (H-score 30.3). (d) QuPath cellular mark-up in the PTGS2-high core (H-score 243.4).

users and non-users. The primary outcome of this study was colorectal cancer-specific survival and the secondary outcome was overall survival. The association between aspirin use and survival was assessed in the whole cohort and then in analyses stratified by biomarker status. Only cases with information on aspirin exposure (known user vs. known non-user) were included in the former analysis. Only cases with available exposure information and tissue for biomarker assessment were included in the subsequent stratified analyses. Other missing categorical data were coded as unknown.

Survival analysis was performed using the Cox proportional hazards model to calculate hazard ratios (HRs) and associated 95% confidence intervals (CI). To control for confounding the cancer-specific survival multivariable model adjusted for age category at diagnosis (<50/50–<60/60–<70/70–<80/>80), gender, year of diagnosis (continuous variable), stage (II/III), grade (well or moderate/poor/unknown), MSI status (MSI-H/non MSI-H/unknown), adjuvant chemotherapy use within three months of surgery (yes/no), ECOG performance status (0–1/2/3–4/unknown) and family history of colorectal cancer (yes/no/unknown). In addition to the above

variables, the overall survival multivariable model also adjusted for the Charlson Comorbidity Index score (continuous variable).²² Analyses were stratified by biomarker status. Interaction terms for aspirin and PIK3CA or PTGS2 were then included in the Cox model and the Wald test was used to assess for statistical interaction. Sensitivity analyses adjusted for statin use and limited PIK3CA mutations to exons 9 and 20 mutations (confer constitutive kinase activity). Sensitivity analyses were also conducted based on altering the PTGS2 H-score cutoff point. All *P* values were two-sided and a value <0.05 was considered statistically significant.

Ethics. Ethical approval was obtained and tissue was acquired through the Northern Ireland Biobank (NIB ref. 13–0088) under the remit of the biobank’s ethical approval from the Office of Research Ethics Committees Northern Ireland (ORECNI ref. 16/NI/0030).

RESULTS

Patients. Of the 740 patients identified in this population-based cohort study, information on prescription medication use

Table 1 Clinical and demographic characteristics according to aspirin use

Characteristic	Aspirin non-user (n = 534) No. (%)	Aspirin user (n = 146) No. (%)	P value
<i>Age category</i>			
< 50	43 (8.0)	4 (2.7)	0.003
50–<60	63 (11.8)	8 (5.5)	
60–<70	159 (29.9)	38 (26.0)	
70–<80	180 (33.7)	59 (40.4)	
> 80	80 (16.7)	37 (25.3)	
<i>Gender</i>			
Male	286 (53.6)	84 (57.5)	0.39
Female	248 (46.4)	62 (42.5)	
<i>Year of diagnosis</i>			
2004	79 (14.8)	22 (15.1)	0.35
2005	108 (20.2)	26 (17.8)	
2006	110 (20.6)	21 (14.4)	
2007	111 (20.8)	35 (24.0)	
2008	126 (23.6)	42 (28.8)	
<i>Stage</i>			
II	313 (58.6)	76 (52.1)	0.16
III	221 (41.4)	70 (48.0)	
<i>Grade</i>			
Well-moderate	458 (85.8)	119 (81.5)	0.19
Poor	72 (13.5)	27 (18.5)	
Unknown	4 (0.8)	0 (0.0)	
<i>MSI status</i>			
Non MSI-high	337 (63.1)	100 (68.5)	0.29
MSI-high	103 (19.3)	20 (13.7)	
Unknown	94 (17.6)	26 (17.8)	
<i>Adjuvant chemotherapy</i>			
No	346 (64.8)	109 (74.7)	0.03
Yes	188 (35.2)	37 (25.3)	
<i>ECOG performance status</i>			
0–1	296 (55.4)	76 (52.1)	0.41
2	33 (6.2)	14 (9.6)	
3–4	27 (5.1)	5 (3.4)	
Unknown	178 (33.3)	51 (34.9)	
<i>Family history of colorectal cancer</i>			
Yes	289 (54.1)	78 (53.4)	0.44
No	85 (15.9)	18 (12.3)	
Unknown	160 (30.0)	50 (34.3)	
<i>Comorbidity</i>			
Cerebrovascular disease	26 (4.9)	7 (4.8)	0.97
Chronic pulmonary disease	55 (10.3)	10 (6.9)	0.21
Congestive heart failure	22 (4.1)	12 (8.2)	0.04
Diabetes mellitus	54 (10.1)	35 (24.0)	< 0.001
Myocardial infarction	28 (5.2)	33 (22.6)	< 0.001
Peptic ulcer disease	22 (4.1)	3 (2.1)	0.24
Peripheral vascular disease	13 (2.43)	11 (7.5)	0.003
Renal disease	8 (1.5)	5 (3.4)	0.13

ECOG, Eastern Cooperative Oncology Group; MSI, microsatellite instability.

in the early post-operative period was available for 680 (91.9%). Compared to those with available information on medication use, patients with no information on medication use were older and more likely to be diagnosed in the earlier years

Table 2 Colorectal cancer-specific and overall survival according to aspirin use, PIK3CA mutation status and PTGS2 expression

Characteristic	Colorectal cancer-specific survival			Overall survival		
	No. of CRC deaths/patients	Unadjusted HR (95% CI)	P value	No. of deaths/patients	Unadjusted HR (95% CI)	P value
<i>All participants</i>						
Aspirin non-user	172/534	1 (Reference)	1 (Reference)	235/534	1 (Reference)	1 (Reference)
Aspirin user	40/146	0.86 (0.61–1.22)	0.40	64/146	1.02 (0.77–1.34)	0.76 (0.57–1.03)
<i>PIK3CA mutation status</i>						
PIK3CA wild-type	151/422	1 (Reference)	1 (Reference)	209/480	1 (Reference)	1 (Reference)
PIK3CA mutation	33/87	0.99 (0.68–1.44)	0.96	55/109	1.21 (0.90–1.63)	1.32 (0.97–1.79)
<i>PTGS2 expression</i>						
PTGS2 low	69/191	1 (Reference)	1 (Reference)	65/217	1 (Reference)	1 (Reference)
PTGS2 high	119/330	0.92 (0.68–1.23)	0.59	176/387	0.98 (0.77–1.26)	0.98 (0.76–1.27)

CI, confidence interval; CRC, colorectal cancer; HR, hazard ratio; MSI, microsatellite instability; PIK3CA, phosphatidylinositol-4,5-bisphosphate 3-kinase; PTGS2, prostaglandin-endoperoxide synthase 2.

^aMultivariable model adjusted for age, gender, year of diagnosis, grade, MSI status, Eastern Cooperative Oncology Group performance status, family history of colorectal cancer, adjuvant chemotherapy use and stage.

^bMultivariable model adjusted for all variables in footnote (a), and also adjusted for Charlson comorbidity score.

of the study. However, there was no difference in stage, grade of tumor differentiation, or MSI status (see **Supplementary Table S2**). In patients with known information on medication use, there were 146 (21.5%) aspirin users. After a mean follow-up of 5.7 (range 0–10) years there were 299 all-cause and 212 colorectal cancer-specific deaths among these patients.

Aspirin use and survival. **Table 1** summarizes the baseline characteristics between aspirin users and non-users. Compared to aspirin non-users, aspirin users were older and less likely to receive adjuvant chemotherapy. As expected aspirin users also had significantly more cardiovascular disease and diabetes. However, tumor grade, stage, and MSI status were similar between users and non-users. Aspirin use at the time of diagnosis was associated with a 31% reduction in colorectal cancer-specific mortality (adjusted HR = 0.69, 95% CI 0.47–0.98) compared to non-use (**Table 2**). The association between aspirin use and overall mortality was slightly attenuated (adjusted HR = 0.76, 95% CI 0.57–1.03).

PIK3CA mutations. DNA was extracted in 599 tumor blocks of the 680 patients in the cohort with information on medication use at the time of diagnosis. PIK3CA mutation status (mutant vs. wild type) was available for 98.3% of these samples (589 out of 599). There was no difference in the proportion of aspirin users in patients with extracted DNA compared to those without extracted DNA (21.5 vs. 21.0%). There was a lower proportion of

PIK3CA-mutant tumors in aspirin users (17 out of 129, 13.2%) compared to aspirin non-users (92 of 460, 20.0%) although the difference was not statistically significant ($P = 0.08$).

Compared to wild-type PIK3CA, the presence of a PIK3CA mutation was not significantly associated with colorectal cancer-specific survival (adjusted HR = 1.15, 95% CI 0.78–1.70, **Table 2**). In stratified analysis there was also no evidence that the association between aspirin use and colorectal cancer-specific survival differed by PIK3CA mutation status (mutant PIK3CA adjusted HR = 0.66, 95% CI 0.22–2.01 vs. wild-type PIK3CA adjusted HR = 0.69, 95% CI 0.46–1.05, P for interaction 0.80). Similar results were observed for overall survival (**Table 3**).

PTGS2 immunohistochemical expression. TMAs were created from 604 tumor blocks where matched information on medication use was available. There were similar proportions of aspirin users in cases with matched TMA data compared to those without matched TMA data (21.5 vs. 21.1%). The proportion of PTGS2-high tumors was 57.7% (75 of 130) among aspirin users and 65.8% (312 of 474) among aspirin non-users ($P = 0.09$). PTGS2 expression was not associated with colorectal cancer-specific survival when PTGS2-high tumors were compared to PTGS2-low tumors (adjusted HR = 0.94, 95% CI 0.69–1.27, **Table 2**).

In analyses stratified by PTGS2 expression, aspirin users had better colorectal cancer-specific survival compared to

Table 3 Aspirin use and colorectal cancer-specific and overall survival stratified by PIK3CA mutation status and PTGS2 immunohistochemical expression

	Colorectal cancer-specific survival ^a		P value	Overall survival ^b		P value
	Aspirin non-user	Aspirin user		Aspirin non-user	Aspirin user	
<i>PIK3CA mutation</i>						
No. of deaths/patients	28/92	5/17		46/92	9/17	
Unadjusted HR (95% CI)	1 (Reference)	1.03 (0.40–2.67)	0.96	1 (Reference)	1.22 (0.59–2.50)	0.59
Adjusted HR (95% CI)	1 (Reference)	0.66 (0.22–2.01)	0.47	1 (Reference)	0.79 (0.35–1.78)	0.57
<i>PIK3CA wild-type</i>						
No. of deaths/patients	120/368	31/112		159/368	50/112	
Unadjusted HR (95% CI)	1 (Reference)	0.87 (0.58–1.30)	0.50	1 (Reference)	1.07 (0.78–1.48)	0.67
Adjusted HR (95% CI)	1 (Reference)	0.69 (0.46–1.05)	0.09	1 (Reference)	0.80 (0.57–1.13)	0.20
Aspirin by PIK3CA interaction	—	—	0.80	—	—	—
<i>PTGS2-high</i>						
No. of deaths/patients	102/312	17/75		146/312	30/75	
Unadjusted HR (95% CI)	1 (Reference)	0.66 (0.39–1.12)	0.12	1 (Reference)	0.83 (0.56–1.24)	0.37
Adjusted HR (95% CI)	1 (Reference)	0.55 (0.32–0.96)	0.04	1 (Reference)	0.64 (0.42–0.98)	0.04
<i>PTGS2-low</i>						
No. of deaths/patients	50/162	19/55		66/162	29/55	
Unadjusted HR (95% CI)	1 (Reference)	1.25 (0.73–2.12)	0.41	1 (Reference)	1.45 (0.94–2.25)	0.09
Adjusted HR (95% CI)	1 (Reference)	1.19 (0.68–2.07)	0.55	1 (Reference)	1.28 (0.80–2.03)	0.30
Aspirin by PTGS2 interaction	—	—	0.09	—	—	0.04

CI, confidence interval; HR, hazard ratio; MSI, microsatellite instability; PIK3CA, phosphatidylinositol-4,5-bisphosphate 3-kinase; PTGS2, prostaglandin-endoperoxide synthase 2.

^aMultivariable model adjusted for age, gender, year of diagnosis, grade, MSI status, Eastern Cooperative Oncology Group performance status, family history of colorectal cancer, adjuvant chemotherapy use and stage.

^bMultivariable model adjusted for all variables in footnote (a), and also adjusted for Charlson comorbidity score.

Table 4 Sensitivity analyses for aspirin use and survival in colorectal cancer adjusting for statin use

	Colorectal cancer-specific survival				All-cause survival					
	Aspirin non-user	Aspirin user	HR (95% CI) non-user ^a	P value	P for interaction	Aspirin non-user	Aspirin user	HR (95% CI) non-user ^b	P value	P for interaction
Overall (n = 680)	No. of deaths/patients 172/534	No. of deaths/patients 40/146	0.71 (0.49–1.03)	0.07	—	235/534	64/146	0.80 (0.59–1.08)	0.15	—
PIK3CA mutation	28/92	5/17	0.74 (0.24–2.28)	0.60	0.80	46/92	9/17	0.90 (0.39–2.05)	0.80	0.93
PIK3CA wild-type	120/368	31/112	0.70 (0.46–1.09)	0.12		159/368	50/112	0.82 (0.58–1.18)	0.29	
PTGS2 high	102/312	17/75	0.56 (0.31–0.98)	0.04	0.09	146/312	30/75	0.65 (0.43–1.02)	0.06	0.04
PTGS2 low	50/162	19/55	1.27 (0.70–2.30)	0.43		66/162	29/55	1.39 (0.86–2.25)	0.18	

CI, confidence interval; HR, hazard ratio; MSI, microsatellite instability; PIK3CA, phosphatidylinositol-4,5-bisphosphate 3-kinase; PTGS2, prostaglandin-endoperoxide synthase 2.
^aMultivariable model adjusted for age, gender, year of diagnosis, grade, MSI status, Eastern Cooperative Oncology Group performance status, family history of colorectal cancer, adjuvant chemotherapy use, stage and statin use.
^bMultivariable model adjusted for all variables in footnote (a), and also adjusted for Charlson comorbidity score.

non-users in PTGS2-high tumors (adjusted HR = 0.55, 95% CI 0.32–0.96), whereas no improvement was observed in PTGS2-low tumors (adjusted HR = 1.19, 95% CI 0.68–2.07, *P* for interaction 0.09, **Table 3**). Similar results were observed for overall survival with evidence of a significant interaction (PTGS2-high adjusted HR = 0.64, 95% CI 0.42–0.98 vs. PTGS2-low adjusted HR = 1.28, 95% CI 0.80–2.03, *P* for interaction = 0.04, **Table 3**).

Sensitivity/subgroup analyses. The associations described above were not markedly altered when statin use at the time of diagnosis was included in the multivariable model (**Table 4**), or when PIK3CA mutations were limited to exons 9 and 20 (see **Supplementary Table S3**). However, the association between improved cancer-specific survival and aspirin use in PTGS2-high tumors was more pronounced when the PTGS2 H-score cutoff point was increased. This association was attenuated when the cutoff point was decreased (see **Supplementary Table S4**).

DISCUSSION

Exposure to aspirin was associated with improved survival in this population-based cohort study of stage II and III colon cancer patients. In stratified analyses there was no evidence of an interaction between aspirin use and tumor PIK3CA mutation status. However, the association between aspirin use and improved survival was more pronounced in tumors with higher levels of PTGS2 expression.

Previous studies investigating the association between these biomarkers (PTGS2 and PIK3CA), aspirin use and survival in colorectal cancer have reported mixed results. Using data from two US cohort studies, Chan *et al.*⁹ demonstrated that aspirin use after colorectal cancer diagnosis was associated with marked survival benefits in tumors that overexpressed PTGS2 compared to those that did not. Reimers *et al.*¹⁴ were unable to replicate this finding in a large Dutch population-based cohort study. Similarly, Domingo *et al.*⁷ did not observe a differential survival effect for aspirin use according to PTGS2 status following molecular analysis of the cohort of patients enrolled in the VICTOR trial.

The first notable difference between these studies is that Chan *et al.*⁹ considered both high- and low-dose aspirin users whereas the other studies were limited to low-dose aspirin only. The proportion of PTGS2-positive tumors in these cohorts also varied considerably (68.4%⁹ vs. 56.3%¹⁴ vs. 21.0%⁷ respectively). In addition to the choice of primary antibody, differences in the definition of PTGS2 overexpression were also present. In particular, while all studies have assessed the intensity of PTGS2 immunostaining using the methods described by Chan *et al.*⁹ only tumors with the strongest staining intensity were considered overexpressors by Domingo *et al.*⁷ Subsequently there was a marked discrepancy between the proportions of PTGS2-positive tumors (21.0 vs. 68.4%) in the VICTOR trial⁷ cohort and the seminal US cohorts.⁹

The current population-based cohort study is the first to corroborate the findings described by Chan *et al.*⁹

regarding an interaction between aspirin exposure, PTGS2 overexpression and better survival in colorectal cancer. Similar to the Dutch¹⁴ and VICTOR trial⁷ cohorts, the current study is representative of low-dose aspirin exposure only. Importantly though, 64.1% of tumors were defined as having higher levels of PTGS2 expression in our study, which more closely matches the proportions observed using the methods described by Chan *et al.*⁹ This association also became more pronounced in sensitivity analyses that increased the PTGS2 H-score threshold, which increases the robustness and biological plausibility of this finding.

Given the problems associated with standardizing immunohistochemical techniques across pathology laboratories, mutated PIK3CA has been proposed as a more reliable biomarker to identify colorectal tumors sensitive to aspirin.⁶ Liao *et al.*⁶ were the first to report that the presence of PIK3CA mutations differentiated the improved survival benefit observed with aspirin use in colorectal cancer. This finding was subsequently validated using data from the VICTOR trial cohort.⁷ Similar to the findings in the Dutch colorectal cancer cohort¹⁴ however, we did not observe evidence of an interaction between low-dose aspirin use and the presence of PIK3CA mutations. Aspirin use at the time of diagnosis was also not associated with a significant improvement in colorectal cancer-specific survival in a retrospective cohort of 185 patients with mutant PIK3CA tumors.¹³ However, stage IV tumors were included in that cohort and there was no comparative group to assess for an interaction.

Two meta-analyses have assessed the interaction between aspirin use after diagnosis and PIK3CA mutation in observational studies of colorectal cancer patients.^{23,24} Paleari *et al.*²³ found that aspirin use in patients with PIK3CA mutated tumors was associated with a 29% significant reduction in overall mortality.²³ In the meta-regression analysis, however, there was no evidence of a significant interaction between PIK3CA mutation status and aspirin efficacy ($P=0.40$) for overall survival. The authors of this review did, however, recognize the potential for immortal time bias in two of the included studies assessing overall survival,^{6,7} where follow-up commenced at the time of colorectal cancer diagnosis but exposure to aspirin treatment started at a non-specified time during subsequent follow-up.^{23,25} In the present study aspirin use was assessed early in the post-operative period therefore the risk of immortal time bias is minimized.

A more recent meta-analysis by Elwood *et al.*²⁴ has also noted a marked protective association for aspirin and colorectal cancer-specific survival in PIK3CA-mutant tumors. However, the magnitude of their risk estimate (HR = 0.45, 95% CI 0.28–0.71) may be exaggerated due to the inclusion of an estimate for recurrence-free survival from one study that did not report cancer-specific survival⁷ and erroneous inclusion of an estimate relating to PTGS2-high tumors, rather than PIK3CA-mutant tumors, from another study.⁹

Although there appears to be evidence for a benefit from previous work,^{6,7} accurately determining the magnitude of a differential association by PIK3CA status has implications for clinical trial design. The failure to demonstrate a significant

interaction in both the meta-regression of previous observational studies²³ and the present cohort would suggest that caution is required before considering additional randomized clinical trials that only recruit patients with mutated PIK3CA tumors (NCT02647099 and NCT02467582). On this basis, planned PTGS2 and PIK3CA subgroup analysis of trials evaluating adjuvant aspirin use in colorectal cancer patients, irrespective of tumor molecular profile, is perhaps more appropriate.

A major strength of this study is the inclusion of population-representative colon cancer patients. Furthermore, application of a precise, automated and validated digital immunoscore system ensures reproducible immunoscore data. However, there are a number of limitations that warrant further discussion. Most importantly, information on aspirin prescription was only assessed in the early post-operative period and no information was available regarding the duration of previous exposure. Assessment at this single time point may not reflect changes in post-diagnostic use and could lead to misclassification bias.^{23,25} However, the optimal duration and timing of aspirin exposure is not clear from observational data to date. It is possible that the important intervention period for aspirin use is in the early post-operative period, as seen for chemotherapy,^{26,27} in which case our data would capture the optimal aspirin time point. Over the counter medication use is another source of bias in pharmacoepidemiology studies, however medication records in COIS and the medical charts relied on a doctor patient interaction, therefore significant over the counter use is likely to have been captured. Also, previous pharmacoepidemiology studies have confirmed that between 75 and 98% of chronic low-dose aspirin use in the age-group of patients included in this cohort in the UK was by prescription.^{28,29}

The ColoCarta panel also only identifies 79% of known PIK3CA mutations. However, these hotspot mutations are representative of 84% of the exon 9 and 20 mutations detected in the seminal paper by Liao *et al.*⁶ It is therefore unlikely that detection of less common mutations would alter the PIK3CA stratified results as the number of additional aspirin users assigned as having a PIK3CA-mutant tumor would be small. Finally, we were unable to adjust for a number of confounding factors such as physical inactivity and obesity.^{30–32} However, to the best of the authors' knowledge, there is no evidence to suggest these factors would impact analyses stratified by PTGS2 or PIK3CA status.

Caution should be exercised when considering the association between better survival and aspirin use in the PTGS2-high subgroup given the borderline significance and multiple hypotheses tested. However, in sensitivity analyses the relationship appeared to be independent of statin use, a further medication with potential anti-cancer effects,³³ and became more significant when the threshold PTGS2 H-score cutpoint was increased. Statistical power to detect an interaction by PIK3CA mutation status was also limited. However, the numbers were similar to the study by Domingo *et al.*⁷ who only had 14 aspirin users in the PIK3CA-mutant group; the only difference being they had no events in this group whereas five events were observed among 17 patients in our cohort.

In summary, aspirin use was associated with better survival in this population-based colon cancer cohort study and the association was more pronounced in tumors with higher levels of PTGS2 expression. In contrast, no significant interaction was observed with PIK3CA mutation status therefore caution should be exercised when considering trials assessing aspirin use in PIK3CA mutated tumors only. Planned subgroup analysis of trials evaluating adjuvant aspirin therapy in colorectal cancer patients, irrespective of tumor molecular profile, may therefore be more appropriate.

CONFLICT OF INTEREST

Guarantor of the article: Ronan T. Gray, MB BCh (Hons), MSc, MRCS.

Specific author contributions: Study conception and design: Marie M. Cantwell, Helen G. Coleman, Maurice B. Loughrey, Stephen McQuaid, Chris R. Cardwell, Jacqueline A. James, Brian T. Johnston, Manuel Salto-Tellez, and Liam J. Murray; Data acquisition: Ronan T. Gray, Helen G. Coleman, Maurice B. Loughrey, Peter Bankhead, Stephen McQuaid, Roisin F O'Neill, Kenneth Arthur, Victoria Bingham, Claire McGready, and Jacqueline A. James; Data analysis and interpretation: Ronan T. Gray, Helen G. Coleman, Chris R. Cardwell, and Liam J. Murray; Drafting manuscript: Ronan T. Gray, Helen G. Coleman, Stephen McQuaid, Chris R. Cardwell and Liam J. Murray. Manuscript revision: All. Final approval: All.

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Potential competing interests: Peter W. Hamilton is Founder and Director in PathXL Ltd. Manuel Salto-Tellez is a senior advisor to PathXL. The remaining authors declare no conflict of interest.

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Study Highlights

WHAT IS CURRENT KNOWLEDGE

- ✓ Epidemiological studies suggest aspirin use may be associated with improved survival after colorectal cancer diagnosis.
- ✓ This association may vary according to molecular markers of phosphatidylinositol 3-kinase (PI3K) signaling activity.

WHAT IS NEW HERE

- ✓ Aspirin use was associated with improved colorectal cancer-specific survival.
- ✓ The association with mortality differed according to PTGS2 expression but not PIK3CA mutation status.

TRANSLATIONAL IMPACT

- ✓ Limiting adjuvant aspirin trials to PIK3CA-mutant colorectal cancer may be too restrictive.

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