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Association of insulin-related serum factors with colorectal polyp number and type in adult males

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Abstract

Background—Dysregulated insulin signaling is thought to contribute to cancer risk.

Methods—To determine if insulin-related serum factors are associated with colon polyps, 126 asymptomatic men (48–65yr) were recruited at colonoscopy. Blood was collected. Odds ratios were determined using polytomous logistic regression for polyp number and type.

Results—Males with serum C-peptide concentration >3.3 ng/ml were 3.8 times more likely to have an adenoma relative to no polyp than those with C-peptide 1.8 ng/ml. As C-peptide tertile increased, an individual was 2 times more likely to have an adenoma $(p=0.01)$ than no polyp. There were no associations between insulin-like growth factor or its binding proteins with polyp number or type. Males with soluble receptor for advanced glycation end products (sRAGE) concentration >120.4 pg/ml were 0.25 times less likely to have 3 polyps relative to no polyps compared to males with sRAGE 94.5 pg/ml. For each increase in sRAGE tertile, a man was 0.5 times less likely to have $\overline{3}$ polyps than no polyps (p=0.03). Compared to males with a serum vascular endothelial growth factor (VEGF) concentration 104.7 pg/ml, males with a serum VEGF concentration >184.2 pg/ml were 3.4 times more likely to have $\overline{3}$ polyps relative to no polyps. As the VEGF tertile increased, a man was 1.9 times more likely to have 3 polyps than no polyps (p=0.049).

Conclusions—Serum concentrations of C-peptide, sRAGE, and VEGF may indicate which men could benefit most from colonoscopy.

Impact—Identification of biomarkers could reduce medical costs through the elimination of colonoscopies on low-risk individuals.

Keywords

colon; insulin; polyp; prevention; males

No authors have conflict of interests to declare.

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INTRODUCTION

Obesity increases the risk of colorectal cancer (1). Furthermore, there is an association between obesity and polyp formation (2). However, the mechanism(s) for this link are unknown. In the body, insulin is responsible for regulating glucose metabolism by stimulating glucose uptake by cells. Obese individuals commonly have high concentrations of insulin and its associated factors in their blood. These high insulin levels and resulting changes in glucose metabolism may be a contributing link between obesity and colorectal polyp/cancer formation (3).

The role of insulin in the development of colorectal adenomas is under active investigation. Insulin is responsible for regulating glucose metabolism by facilitating glucose uptake by cells including hepatocytes, adipocytes, and myocytes. An increase in glucose levels in the blood normally leads to an increased secretion of insulin. In healthy individuals, secreted insulin signals cells to remove the glucose from the blood. However, obese individuals have an increased risk for insulin resistance wherein their cells do not efficiently remove excess glucose from the blood. In part, this is because adipose tissue releases large amounts of inflammatory cytokines which further impair insulin's action (4). The hyperglycemia then maintains the inflammatory state through a variety of mechanisms and provides a ready source of energy for rapidly dividing neoplastic cells (5, 6). Normoglycemia in a mouse model of diabetes has been achieved through neutralization of vascular endothelial growth factor (VEGF) linking high VEGF levels with insulin resistance (7). Additionally, pre/post bariatric surgery changes in a novel biomarker, soluble receptor for advanced glycation end products (sRAGE), showed a significant negative correlation with well-known measures of insulin resistance. After the surgery, insulin resistance decreased and sRAGE levels increased (8). Thus, insulin-related serum factors may be indicative of the failure to regulate blood glucose and the resulting increased inflammation in obese individuals at a heightened risk for colorectal cancer.

Because the insulin-signaling pathway is associated with cancer (9, 10), molecules involved in these pathways were selected as a focus for this analysis. Evidence that serum factors involved in glucose metabolism are related to colorectal polyps in otherwise healthy adult males is presented. Colonoscopies are costly and undesirable to some adults. Serum markers that could predict which individuals are more likely to have polyps, and thus benefit most from colonoscopy, are desired.

MATERIALS AND METHODS

Study Population

Between August 2009 and February 2011, healthy males ranging from 48–65 years of age were recruited from either Tri-County Gastroenterology Clinic (Macomb, MI) or Michigan State University Clinic (East Lansing, MI) at the time of colonoscopy. These individuals were undergoing routine colonoscopies and were asymptomatic. Exclusion criteria included: 1) cancer within the past two years, 2) surgery within the past two years, 3) inflammatory bowel diseases (i.e. Crohn's, ulcerative colitis), 4) autoimmune disorders (i.e. Rheumatoid

arthritis, HIV/AIDS, Lupus), 5) type I or type II diabetes, 6) chronic liver or kidney disease, 7) history of heart failure, 8) current immunosuppressant usage (i.e. Prednisone), 9) asthma, chronic obstructive pulmonary disease or other lung problems, 10) familial adenomatous polyposis, and 11) Lynch syndrome or hereditary non-polyposis colorectal cancer. 126 men (> 96% Caucasian) participated in the study (Table 1). All enrolled individuals completed the study, and no records were maintained as to the number of individuals approached but not enrolled. At the time of enrollment, immediately prior to routine colonoscopy, written informed consent was obtained and clinical metadata on subject co-morbidities, current medications, and family history were collected. Anthropometric measures were taken to calculate BMI and to record waist circumference (11). Also at the time of enrollment, venous blood was drawn, and serum was isolated by standard procedures and stored at −80°C. The study was approved by the institutional review board of Michigan State University (IRB# 08-786).

Colonoscopy Interpretation

Full colonoscopy was performed on each participant. A gastroenterologist (MSU-affiliated clinics, MI) categorized the location of each polyp during the colonoscopy. Board-certified pathologists assigned a polyp type to each specimen collected during colonoscopy (regional medical center pathology departments).

Serum Biomarker Analysis

C-peptide concentrations (ng/ml) were measured using an ELISA (Calbiotech, Spring Valley, CA, REF: CP1795) following the manufacturer's instructions. IGF-1 was measured by ELISA following the manufacturer's protocol (R&D Systems, DG100, Minneapolis, MN). Both ELISAs were read using the absorbance function on a Bio-Tek (Winooski, VT) plate reader. Insulin growth factor binding proteins (IGFBP) 1 through 7 were measured by multiplex kit per the manufacturer's protocol (HIGFBP-53K, Millipore, Billerica, MA). Serum concentrations of the soluble vascular endothelial growth factor receptors (sVEGFR) including sVEGFR-1, sVEGFR-2, and sVEGFR-3, as well as the soluble receptor for advanced glycation end products (sRAGE), and the soluble endothelial growth factor receptor (sEGFR) were determined using a multiplex kit for soluble cytokine receptors (HSCR-32K, Millipore) as per the manufacturer's instructions. Further, fibroblast growth factor (FGF)-2, endothelial growth factor (EGF), and vascular endothelial growth factor (VEGF) serum concentrations were measured using a multiplex human cytokine/chemokine kit (MPXHCYTO-60K, Millipore). Multiplex assays were analyzed on a Bio-Plex 100 using Bio-Plex 4.1 software (Bio-Rad, Hercules, CA).

Statistical Analyses

Frequencies, means, standard deviations and ranges were calculated for descriptive analysis. Both Pearson and Spearman correlations were calculated. Because some variables were nonnormally distributed, the results from the Spearman correlation are presented (Table 2). All of the biological markers (C-peptide, sRAGE, IGF-1, IGFBP-1, IGFBP-2, IGFBP-3, IGFBP-4, IGFBP-5, IGFBP-6, IGFBP-7, EGF, sEGFR, FGF-2, VEGF, sVEGFR-1, sVEGFR-2, sVEGFR-3) were categorized into tertiles for polytomous logistic regression analysis. Biological cut off points were used for the outcome variables: polyp number and

polyp type. Polyp number was set to four categories: $0, 1, 2$, or $\bar{2}$ 3 polyps. Polyp type was set to three categories: no polyps, hyperplastic polyp(s) or tubular adenoma(s). For polyp type, each individual was assigned a single score based on the colorectal polyp with the worst pathology.

Each individual was assigned a smoking status of "never smoked" or "ever smoked". An individual was classified as "never smoked" if he had smoked fewer than 100 cigarettes over the course of his lifetime. Smoking status was missing for 22 individuals (17.5% missingness). All missing data were considered missing at random.

Multiple imputation (seed $= 20121119$, imputations $= 7$) was used to impute all missing smoking data and the one participant's value missing for EGF, FGF-2, and VEGF (12). The factors: age, smoking, polyp type, polyp number, BMI, C-peptide, IGFBP-1, IGFBP-3, IGFBP-4, IGFBP-5, IGFBP-6, IGFBP-7, EGF, FGF-2, sEGFR, sRAGE, sVEGFR-1, sVEGFR-2, sVEGFR-3, VEGF, and IGF-1 were used in the imputation algorithm of missing values. Sensitivity analyses demonstrated that these imputations did not bias the results.

Odds ratios (OR) were determined using polytomous logistic regression models for categorical outcome data with more than two levels. A generalized logit model (also known as a, "baseline category logit model") formed the basis of the polytomous logistic regression model for our analyses. In this model, men having 3 polyps, 2 polyps or 1 polyp were compared to men having no polyps. Results were obtained for each serum factor categorized into tertiles (with lowest tertile as reference) for each category of polyp number or polyp type relative to no polyp. For clarity, results from the categories featuring the most polyps (≥3) and the polyp type most likely to progress to colorectal cancer (adenoma). Otherwise odds ratios were determined using logistic regression. All models were adjusted for age and smoking status. Test for trend was carried out across tertiles for the factors of interest. Because imputation was used, multiple imputation analyze (Proc MIANALYZE) was used to determine the results from analysis of the imputed data sets. SAS version 9.3 (SAS Institute Inc.; Cary, NC) was used for all statistical analysis. $p = 0.05$ indicates significance.

RESULTS

Participant age, BMI, waist circumference, and average values for some serum factors are given in Table 1. As previously published (11), 57 (45%) of the 126 participants had at least one polyp, and 23 (18%) had $\,$ 3 polyps. 37 (29.4%) of the participants had a tubular adenoma. Age did not differ between individuals with no polyps and those with $\frac{3}{2}$ polyps. Both BMI and waist circumference increased with increasing number of polyps and polyp type. In addition, 17 (13.5%) of the participants had ≥ 3 polyps with at least one tubular adenoma.

Several of the serum factors analyzed correlated with subject characteristics as well as with each other (Table 2). IGFBP-1, IGFBP-3, IGF-1, and sRAGE were inversely correlated with BMI. However, IGFBP-7 and C-peptide were positively correlated with BMI. Although it was inversely correlated with BMI, sRAGE was positively correlated with FGF-2, sEGFR, sVEGFR-1, sVEGFR-2, sVEGFR-3 and IGF-1. A significant correlation was observed

between C-peptide and IGFBP-1 (−), IGF-1 (−), as well as IGFBP-7 (+). IGF-1 was also positively correlated with the soluble VEGF receptor (sVEGFR-2).

Although serum insulin-like growth factor (IGF) and the IGF binding proteins were not associated with colorectal polyp number or type, other insulin-related serum factors were correlated with these parameters. Serum concentrations of two insulin-related factors, sRAGE and VEGF, were significantly associated with the presence of $\overline{3}$ polyps. The odds that a participant with a serum sRAGE concentration > 120.35 pg/ml would have $\frac{3}{2}$ polyps relative to no polyp were 0.25 times (CI: $0.07 - 0.92$) lower than those that a participant with serum sRAGE concentration 94.45 pg/ml would have 3 polyps (p=0.04, Figure 1a) relative to no polyp. Additionally, for each category increase in serum sRAGE concentration, a man was 0.5 (CI: $0.27 - 0.94$) times less likely to have $\overline{3}$ polyps than no polyps (p=0.03). Participants with serum VEGF concentrations > 184.16 tended to be 3.4 times (CI: $0.97 - 11.74$) more likely to have $\overline{3}$ polyps relative to no polyps compared to the likelihood that his counterpart with serum VEGF concentrations 104.67 pg/ml would have β polyps (p=0.055; Figure 1b) relative to no polyps. For each category increase in serum VEGF concentration tertile, a man was 1.9 (CI: 1.003 – 3.53) times more likely to have α 3 polyps than no polyps (p=0.049). Non-significant relationships were observed between polyp number and all other factors analyzed, including C-peptide, IGF-1, IGFBP-3, IGFBP-7, and the IGF-1/IGFBP-3 ratio (Table 3) as well as the remaining IGFBPs, sVEGFR-1, sVEGFR-2, sVEGFR-3, sEGFR, FGF-2 and EGF (data not shown).

sRAGE was the only serum factor in this investigation which was negatively associated with the presence of a hyperplastic polyp (Supplementary Table 1). Males with a serum sRAGE concentration in the highest tertile $(>120.35 \text{ pg/ml})$ were 0.12 (CI: $0.02 - 0.63$) times less likely to have a hyperplastic polyp relative to no polyp than males with a serum sRAGE concentration in the lowest tertile (94.45 pg/ml) (p=0.01). For each category increase in serum sRAGE concentration, a man was 0.4 (CI: $0.2 - 0.79$) times less likely to have a hyperplastic polyp (p=0.01) relative to no polyp. No other analyzed factors were associated with the presence of a hyperplastic polyp.

An insulin-related factor associated with polyp type was C-peptide. Participants with a serum C-peptide concentration in the highest tertile (>3.31 ng/ml) were 3.8 (CI: 1.31 – 11.15) times more likely to have a tubular adenoma relative to no polyp than males with a serum C-peptide concentration in the lowest tertile (1.77 ng/ml) (p=0.01, Figure 2). For each category increase in serum C-peptide concentration, a man was 1.97 (CI: 1.16 – 3.37) times more likely to have a tubular adenoma than no polyp $(p=0.01)$. Non-significant associations were found between the presence of a tubular adenoma and all other serum factors analyzed, including sRAGE, VEGF, IGF-1, IGFBP-3, IGFBP-7, and the ratio between IGF-1 and IGFBP-3 (Table 4).

DISCUSSION

Herein we describe associations between insulin-related serum factors with polyp number or polyp type. Serum concentrations of C-peptide were positively associated with the presence of a tubular adenoma. IGF and its binding proteins were not associated with colorectal polyp

number or type. Two other proteins related to insulin and glucose regulation were associated with polyp number and severity. Serum concentrations of sRAGE were negatively associated with the presence of $\,$ 3 polyps while serum VEGF concentrations were positively associated with the presence of $\,$ 3 polyps. Because polyps are considered precursors to colon cancer (13), the associations demonstrated herein indicate that several insulin-related factors are associated with the risk of colorectal cancer in adult, Caucasian males.

This study had several strengths. The design of this study, cross-sectional with consecutive enrollment, reduced enrollment bias. Additionally, only asymptomatic individuals undergoing a complete screening were enrolled. Anthropometric measurements were taken by trained staff instead of being given by self-report. Participants provided information about relevant confounding variables, thus allowing for control of these confounding variables in our analysis. Because our population is well-defined, the results are likely typical for men of an ethnicity and age similar to that of the study population.

The study also had several limitations. Thus, caution must be used when interpreting the results of this study. Because this study was conducted in a modest-sized sample of primarily Caucasian, adult males, the results may not apply to individuals of other ethnicities or females. Additionally, although sensitivity tests did not show an effect of imputation on the conclusions drawn from the analysis, it is important to note that some data was imputed and that these imputations may have minor effects on the results. The associations described in this study were based on cross-sectional data thus cause cannot be assigned to any factors identified as associated with polyp number or type. Additionally, only a single time point is analyzed, but effects of some serum proteins may be related to the duration of exposure. In addition, local production rather than serum concentrations of the analyzed proteins may be relevant to colorectal polyp formation. However, local production is not easily measured, and serum measurements are more useful in a clinical setting, for instance as a potential precolonoscopy screening tool. A larger sample size would decrease the width of the reported 95% confidence intervals, thus, this research should be repeated with a larger sample size.

It was surprising that IGF-1 levels and the IGF-1 to IGFBP-3 ratios were not associated with colorectal polyp number or type in our population. Several other publications report a link between IGF and colorectal cancer or polyps (14–16). However, a lack of such an association has been demonstrated in prostate cancer (17). We also did not observe associations between the IGF binding proteins and colorectal polyps despite evidence that these binding proteins can block IGF-signaling (18). IGF-1 is important for cellular proliferation, differentiation and apoptosis (19). IGF-1 levels in our study were similar to levels in at least one other study (20). In addition, the inverse relationship between BMI and IGF-1 levels that we observed has also been observed by others (21). Given the outcomes of recent clinical trials (10) showing that anti-IGF-1 receptor treatments are ineffective in treating cancer, our results demonstrating the IGF-1 and its binding proteins are not associated with early steps in the progression to colon cancer are less surprising. Additionally, IGF signaling in the colon rather than blood may be more predictive of colon cancer risk (22).

A study population consisting only of men may also explain the lack of association between IGF and polyps. In women, fasting IGFBP-1 was inversely associated with risk of colon cancer with a relative risk of 0.28. The IGF-I to IGFBP-3 molar ratio was associated with colon cancer risk with a relative risk of 2.82. Further, women with low levels of both IGF-I/ IGFBP-3 and C-peptide (or high IGFBP-1) were at low risk; however, elevation of either was sufficient to increase risk (23). Colon cancer risk showed a nonsignificant positive trend with IGF-1 levels and was significantly increased for women with the highest IGFBP-3 levels (24). Thus, the association between IGF-1 and colorectal cancer may be sex-specific.

Alternatively, glucose regulation rather than insulin signaling itself may be most important in cancer. Anti-diabetic drugs can inhibit carcinogenesis in mice and rats. For instance, Metformin decreases the risk of cancer in individuals with diabetes (25). Significant alterations in tumor glucose levels were identified when metabolomics were used to identify metabolic changes resulting from colorectal neoplasia in a mouse model (26). Furthermore, actual changes in insulin receptor signaling may not become important in colorectal cancer pathogenesis until cancer progression or metastases have occurred. Therefore, we would not be able to detect these changes in our study of colorectal polyp formation.

However, in support of the potential importance of insulin and its related signaling pathways in the development of colon cancer, we found that C-peptide serum concentrations were positively associated with the presence of colonic tubular adenomas. Insulin levels in serum can be indirectly assessed by measuring the C-peptide concentrations in samples. The latter protein is formed in equimolar amounts with insulin when proinsulin is secreted by pancreatic beta cells and cleaved into the two proteins. Insulin is known to have a shorter half-life than C-peptide, which makes C-peptide concentration a better indicator of insulin concentration (reviewed in (27)). Furthermore, serum C-peptide levels are more likely to represent long-term insulin levels since they are not as influenced by disease status in nondiabetic individuals (28).

The increased odds of tubular adenoma with high serum C-peptide concentrations may be due to effects of C-peptide or insulin on inflammatory signaling pathways as C-peptide affects cardiovascular disease risk. For instance, when C-peptide levels were analyzed in 5153 non-diabetic adults between 40 and 74 years of age, individuals with the highest Cpeptide measurements (0.984 nmol/L = 2.95 ng/ml, (27)) had a 60% increase in the adjusted hazards of cardiovascular death compared to those with the lowest (≤0.418 nmol/L $= 1.25$ ng/ml). This trend persisted in risk for cerebrovascular disease, ischemic heart disease, myocardial infarction, and overall mortality (29). C-peptide induces proliferation of smooth muscle cells potentially leading to the development of atherosclerotic lesions (30).

Likewise, high C-peptide levels were associated with increased colorectal adenoma risk in some studies. In a 2012 study by Vidal et al., individuals with C-peptide levels in the fourth quartile had a 2-fold increased risk of adenomas compared to individuals with C-peptide levels in the lowest quartile (31). Furthermore, C-peptide levels were associated with increased cancer risk in the colon, but less so in the rectum (32). Additionally, a metaanalysis reported that subjects with the highest levels of circulating C-peptide were at an increased risk for colon but not rectal cancer (33). The relationship between C-peptide and

colon cancer has been reported to be stronger in men than women (33), but the positive association has also been reported in women (23, 24). Other studies have shown no link between C-peptide and colon cancer (34).

Until recently, C-peptide was not thought to have any physiological function. However, in a 2012 study using BLAST searches and comparative biology, C-peptide was suggested to be a receptor ligand which behaves similarly to a cytokine. Specifically, a highly-conserved portion located around leucine-86 and glutamine-87 of the C-peptide gene sequence is associated with an active receptor binding spot (35). The development of colorectal adenomas was attributed to inflammatory cytokines. For example, levels of TNF-α in patients with hyperplastic polyps were significantly increased compared to those in patients with healthy mucosa (36). These results suggest that C-peptide may behave like a cytokine, inducing inflammatory responses in tissues. Thus, this inflammatory action of C-peptide is potentially important in the development of colon polyps, colorectal cancer or other intestinal inflammatory diseases. Alternatively, C-peptide could lead to adenomas through its anti-apoptotic properties (31).

Notably, individuals who consume high levels of meat, fish, and sweetened beverages and low levels of dairy and whole grains have higher serum C-peptide concentrations (37, 38). This type of diet is positively associated with colorectal cancer risk (39). Depending on the methods of preparation, for instance roasting, broiling or frying, this high C-peptide dietary pattern is also likely to be associated with high intake of advanced glycation end products (AGE) (40). AGE are formed by the nonenzymatic glycation of proteins, lipids, and nucleic acids. Two major contributors of AGE are endogenous AGE that form during normal metabolism and exogenous AGE from tobacco smoke or food processing. AGE accumulation is associated with insulin resistance (41).

The inverse association between serum sRAGE concentrations and the presence of three or more polyps may be due to the anti-inflammatory effects of sRAGE. It is the interaction between AGE and RAGE that leads to inflammation since RAGE signaling mediates NF-kB transcription. sRAGE neutralizes the oxidative stress and inflammatory effects mediated by the AGE/RAGE complex because it competes with RAGE to bind AGE (reviewed in (42)). Therefore, when serum concentrations of sRAGE are high, inflammation may be prevented through inhibition of the AGE/RAGE interaction.

In our subjects, as reported by others (43), serum sRAGE concentrations were negatively correlated with BMI. sRAGE was not correlated with C-peptide. We hypothesize that the increased levels of sRAGE in our subjects are an effective response to chronic low-grade inflammation. Rising levels of serum sRAGE, would, in turn, inhibit the inflammatory response that initiates abnormal colonic cell proliferation. Thus, when sRAGE is present and the inflammatory response is inhibited, fewer complications would arise. Since high levels of sRAGE may prevent progression to adenoma or other inflammatory complications in individuals with colorectal polyps, future research should address strategies to increase serum concentrations of sRAGE. For instance, physical activity can increase sRAGE concentrations in patients with Type 2 diabetes (44). Thus, increased physical activity may be one potential intervention strategy to prevent polyps. Vitamin D supplementation is

another intervention that could potentially affect the progression of polyps through modulation of serum sRAGE concentrations as increased levels of serum calcitriol have been associated with increased sRAGE (45).

Others have reported inverse associations between sRAGE and cancer. One of the first studies to relate sRAGE levels with a decreased risk of colon cancer found that higher prediagnostic levels of serum sRAGE were even associated with a decreased risk in smokers (46). The same group later reported that individuals with colorectal adenomas had higher levels of serum EGF and VEGF and tended to have lower levels of sRAGE than individuals with no adenomas (47). Thus, we are the second group to associate serum sRAGE with colorectal polyps.

A more expected observation from our study was the positive association between VEGF and the presence of three or more polyps. VEGF is a well-known promoter of angiogenesis and, therefore, cancer. But, VEGF is less well known for its association with insulin resistance, where neutralization of VEGF restores insulin sensitivity and normal glucose levels in a mouse model of diabetes (7). Others have reported positive relationships between serum VEGF and colorectal cancer (48) as well as colonic VEGF and cancer (49). In fact anti-VEGF monoclonal antibodies are being used as a therapy in metastatic colorectal cancer (50).

Our research suggests that high serum concentrations of VEGF (> 184.16 pg/ml), high serum concentrations of C-peptide (>3.31 ng/ml), and low serum concentrations of sRAGE (≤94.45 pg/ml) could be used to identify individuals at increased risk for colorectal polyp formation. Since it is known that removal of polyps decreases the incidence of colon cancer (13), this research provides additional evidence that colonoscopy should be recommended for Caucasian males who have inappropriate concentrations of these serum factors. Colonoscopy screening of individuals with these risk factors could potentially decrease colorectal cancer rates. Prior to any policy changes, studies verifying the usefulness of these proteins as predictors of colorectal polyps are needed.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Figure 1. The OR of having ≥ 3 polyps relative to no polyps is negatively associated with serum sRAGE (A) and positively associated with serum VEGF (B)

A, Compared to males with a serum sRAGE concentration 94.45 pg/ml, males with a serum sRAGE concentration > 120.35pg/ml are 0.25 times less likely to have $\frac{3}{2}$ polyps (p=0.0368) relative to no polyps. In addition for each increase in serum sRAGE tertile, a man is 0.498 times less likely to have \rightarrow 3 polyps than no polyps (p=0.0303). Reference group (Ref) = males with sRAGE concentrations in the lowest tertile. B, Compared to males with a serum VEGF concentration 104.67 pg/ml, males with a serum VEGF concentration > 184.16 pg/ml tend to be 3.4 times more likely to have ≥ 3 polyps (p=0.0550) relative to no polyps. For each category increase in serum VEGF tertile, a man is 1.9 times more likely to have $\overline{3}$ polyps than no polyps (p=0.0488). The models were adjusted for age and smoking status (ever/never). Reference group (Ref) = males with VEGF concentrations in the lowest tertile.

Figure 2. Serum C-peptide concentration is associated with the presence of adenomas

Compared to males with a serum C-peptide concentration 1.77 ng/ml , males with a serum C-peptide concentration > 3.31 ng/ml are 3.8 times more likely to have an adenoma (p=0.0144) relative to no polyps. Further, for each increase in serum C-peptide tertile, an individual is 1.97 times more likely to have an adenoma than no polyp (p=0.0127). The model was adjusted for age and smoking status (ever/never). Reference group (Ref) = males with C-peptide concentrations in the lowest tertile.

Table 1

Participant*†* Characteristics

† All participants (n=126) were male, > 96% Caucasian

‡ Reported as mean (range)

*** Does not include those with missing data

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Table 2

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Spearman correlation coefficients

Spearman correlation coefficients

Note: The top number is the Spearman correlation coefficient. The P values are shown under the correlation coefficient. Note: The top number is the Spearman correlation coefficient. The P values are shown under the correlation coefficient.

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Table 3

Association of insulin-related factors with having ≥ 3 polyps relative to no polyps

Note: Model is adjusted for age and ever/never smoked.

Table 4

Association of insulin-related factors with the presence of a tubular adenoma relative to no polyps

Note: Model is adjusted for age and ever/never smoked.