#### **ORIGINAL ARTICLE**



# Carbonic Anhydrases II, IX, and XII in Reflux Esophagitis

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#### Abstract

**Background** The pathogenesis of gastroesophageal reflux disease (GERD) has not been resolved in detail. Esophageal epithelial cells provide resistance to acidic reflux via several mechanisms, many of which involve buffering acid with bicarbonate and transporting protons. Carbonic anhydrases (CAs) are enzymes that control the acid–base balance by catalyzing the reversible hydration of carbon dioxide to produce bicarbonate and hydrogen ions.

**Aims** We aimed to determine the immunohistochemical expression patterns of CAII, CAIX, and CAXII in the normal esophageal squamous epithelium and in patients with GERD.

**Methods** We evaluated 82 biopsy samples, including 26 with a histologically normal esophagus, 26 with histologically mild esophagitis, and 30 with severe esophagitis. Expression patterns of CAII, CAIX, and CAXII in the esophageal squamous epithelium were determined by immunohistochemical staining.

**Results** Cytoplasmic CAII expression was predominantly detected in the upper luminal part of the squamous epithelium and was significantly (p < 0.01) increased in GERD. Expression of CAIX was essentially membranous. The isozyme was constantly present in the peripapillary cells. In the interpapillary areas, clustered expression was observed to emerge and increase significantly (p < 0.01) in esophagitis. CAXII expression was the most abundant of the isozymes and was mainly membranous. In the normal squamous epithelium, CAXII expression was confined to the basal layer; in severe esophagitis, CAXII expression increased significantly in both basal (p < 0.05) and superficial (p < 0.01) halves of the epithelium. **Conclusions** We demonstrate upregulated expression of CAII, CAIX, and CAXII in GERD. The increase in expression likely

contributes to esophageal epithelial resistance to acidic reflux.

Keywords GERD · Reflux esophagitis · CAII · CAIX · CAXII

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## Introduction

Gastroesophageal reflux disease (GERD) has a significant worldwide prevalence of 10-30%, with the highest popularity in the USA, parts of Europe, Latin America, and Middle East [1, 2]. Due to aging population, the prevalence and burden to economies is still increasing [2]. Of all gastrointestinal diseases, the pharmacotherapy of GERD is currently responsible for the greatest direct costs in the USA [3] and causes a significant economic burden in many European countries [4]. Developing countries in Africa, South America, and Asia offer sparse available data on GERD epidemiology or cost analyses, but it is safe to assume GERD-related costs are increasing in the emerging economies as well, even when the racial differences in the prevalence are accounted for [5]. At the same time, complications of GERD, such as Barrett's esophagus (BE) and esophageal adenocarcinoma [3, 6], appear to be increasing. Understanding the pathogenesis of reflux esophagitis is key for treatment and to prevent complications.

Esophageal epithelial cells (EECs) have been suggested to provide resistance to gastroesophageal reflux via several mechanisms, such as the capacity to buffer intra- and intercellular acidification, the efficient transport and disposal of  $HCO_3^-$  and  $H^+$  and the permeability of the epithelium [7]. All of these mechanisms are potentially managed by carbonic anhydrases (CAs), catalysts of reversible  $CO_2$ hydration ( $H_2O + CO_2 \leftrightarrow HCO_3^- + H^+$ ). At present, 12 catalytically active isoforms of carbonic anhydrases in humans have been identified [8]. Cytosolic CAII and membrane-bound isoforms CAIX and CAXII are considered tumor-associated CAs [9], but the basic function of these highly active enzymes is related to the regulation of pH homeostasis and the transport of  $CO_2$  and  $HCO_3^-$  [8, 10].

To date, only fragmentary information about CAs in esophageal physiology is available. CAII, the most widely expressed CA, is present nearly throughout the gastrointestinal tract [11]. Cytosolic CAII is also expressed in EECs [12], and it has been proposed to play a role as a buffer of intracellular acidification in esophageal epithelial resistance [7, 12, 13]. CAII forms transport metabolons with sodium independent Cl<sup>-</sup>/HCO<sub>3</sub><sup>-</sup> anion exchangers (AE1-3) and Na<sup>+</sup>/HCO<sub>3</sub><sup>-</sup> transporters (NBCs) of the family SLC4A and the Na<sup>+</sup>/H<sup>+</sup> exchanger 1 (NHE-1) (SLC9A1) [14-20], all of which participate in esophageal defense [7]. Bile acids, on the other hand, have been suggested to inhibit CAII and CAIX [21]. Basolaterally located CAIX and CAXII are induced by HIF-1 $\alpha$  and hypoxia [22], but they are also expressed in normal EECs [23]. In the stomach of mice, low extracellular pH is an independent inducer of CAIX expression [24], and the enzyme may contribute to gastric mucosa protection by participating in tight junction maintenance [24]. CAXII has a more diverse regulatory system, which includes induction by estrogen receptor  $\alpha$  (ER $\alpha$ ) [25, 26]. CAIX and CAXII have both been shown to form transport metabolons to facilitate anion transport, mainly with AE1 and AE2, but CAIX also with NBCs and NHE-1 [27–29] [30].

We have previously detailed expression of CAII, IX, and XII in normal esophageal squamous epithelium, Barrett's esophagus, and esophageal adenocarcinoma [23]. As the expression patterns and functional roles of CAs II, IX, and XII in the pathogenesis of GERD are largely unexplored, in this study, we evaluated expression of CAII, CAIX, and CAXII in the esophageal squamous epithelium in a representative series of patients with and without reflux esophagitis. We hypothesized that these enzymes are upregulated, contributing to esophageal epithelial resistance to acidinduced injury.

## **Materials and Methods**

A series of 84 cases with esophageal archival biopsy samples were collected from the archives of the Department of Pathology, Oulu University Hospital, from 2011 to 2015. Consecutive cases were selected to obtain three subsets of patients: patients with histologically normal esophagus, histologically mild esophagitis, and histologically severe esophagitis. Information on the indications for the endoscopy and esophageal biopsies was collected from the endoscopy reports. Among subjects with histological esophagitis most had heartburn, regurgitation or dysphagia. Patients with normal esophageal histology and normal endoscopy had presented with a variety of indications such as preoperative work-up prior to bariatric surgery, or as a part of anemia or a weight-loss investigation. None of these subjects had indicated symptoms referring to GERD. Esophageal biopsies in these patients had been taken based on the consideration of the gastroenterologist or gastrointestinal surgeon performing the endoscopy, and in most cases no specific indication for taking these biopsies was recorded. No esophageal pH studies were performed. Esophageal biopsies were collected at the Z-line and +2 cm above. Information regarding the endoscopic degree of esophagitis according to the Los Angeles Classification [31] and the use of proton pump inhibitors 2 weeks prior to the endoscopy (PPIs) was retrieved from Oulu University Hospital medical records, including the original endoscopy reports and endoscopic footage. However, determining the LA class in sufficient accuracy in this retrospective setting was successful only in 60% of patients with GERD, mainly due to the deficient reporting of LA class in the endoscopy reports and often unrepresentative endoscopic images. Therefore, a more simplified grading was applied: endoscopic esophagitis was classified as mild, when erythema or edema was seen (largely corresponding to LA M), and severe in the presence of clear epithelial breaks (LA A-D). Flowchart of patient selection is presented in Fig. 1. All procedures performed in our study were in accordance with the ethical standards of the Oulu University Hospital Ethics Committee and with the 1964 Declaration of Helsinki.

The hematoxylin-and-eosin-stained sections were reevaluated according to the histological criteria of reflux esophagitis [32, 33] by an experienced gastrointestinal pathologist (TJK). The criteria included evaluation of basal cell layer hyperplasia, papillary elongation, dilatation of intercellular spaces, intraepithelial neutrophilic infiltration, intraepithelial eosinophilic infiltration, and the presence of erosion or necrosis. The features were graded on a 0-2 scale (0 = absent, 1 = mild, 2 = severe; for neutrophils and erosion/necrosis grades were 0 = absent and 2 = present [31]. The presence and severity of histological esophagitis was determined using the Global Severity (GS) score introduced by Mastracci et al. [31]. The GS score equals the mean of all histological criteria graded according to highest grade occurring in the case. In addition, GS score was assigned 2 (severe) in the presence of intraepithelial neutrophils or erosion/necrosis, even if calculated mean was less than 2 since these are considered as features of the most severe grade of histological esophagitis [31]. GS score cutoff of 0.35 has been shown to correlate well with the diagnosis of GERD based on pH monitoring. The group of patients with histological [32] esophagitis (GS score  $\geq 0.35$ ) was divided into subgroups of mild (GS 0.35-1.49) and severe (GS 1.5-2.0) histopathological esophagitis based on the median GS score

Fig. 1 Flowchart of patient

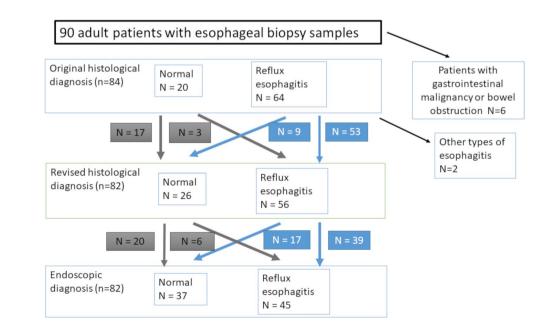
selection

value of 1.5. The presence of lymphocytic [34], infectious and eosinophilic [35] esophagitides was excluded in re-evaluation (TJK).

For immunohistochemistry, sections from formalin-fixed, paraffin-embedded specimens were subjected to high-temperature antigen retrieval in Tris–EDTA buffer for 15 min. CA immunostaining was conducted using polyclonal rabbit anti-human CAII and CAXII sera and monoclonal antihuman CAIX antibody (M75) as described previously for polymer-based detection [36]. The antibodies have been previously characterized and are specific for each isozyme [37–39].

All immunohistochemical stainings were estimated by three independent researchers (TJK, MN, and NV). The region of esophageal squamous epithelium used for evaluation of IHC stainings was selected by an experienced pathologist (TJK) at the time of the light microscopy session for evaluation. The region with most severe histological features of GERD [32, 33] was chosen if features suggesting GERD were present in the specimen. However, necrotic squamous epithelium was not evaluated for CA expression. In other cases, any representative area of the normal squamous epithelium was scored. The other evaluators were blinded for the diagnosis and GS score of the lesion to be evaluated.

For CAII and CAXII assessment, the squamous epithelium was divided into basal and superficial halves. Both were evaluated separately for the intensity of staining (0-3) and the percentage of stained cells (0-100). The histoscore value is the product of the intensity and percentage for the superficial and deep halves, respectively, and the mean of these histoscores is used as the total histoscore. For CAII, a separate assessment was made for nuclear and cytoplasmic staining. For membrane-bound



CAXII, staining was evaluated for membrane-associated and cytoplasmic expression. The CAIX expression profile differed from those of the other two isozymes studied. The majority of the CAIX staining was confined to squamous cells adjacent to subepithelial papillae and was regionally present in interpapillary squamous epithelium. CAIX expression was therefore reported by separately assessing the peripapillary and interpapillary staining. Peripapillary staining was estimated by the intensity (0-3) of the staining and the number of positive cell layers around the papilla. The peripapillary staining score was obtained by multiplying the intensity and the number of stained peripapillary cell layers. Interpapillary regions were estimated in a manner similar to the other CAs; the intensity of staining (0-3) and percentage of cells with positively stained cell membranes in the field of view were evaluated. If the individual estimates of the three observers differed by > 1for the intensity score, > 30% for the percentage, or > 2 for the papillary cell layer count, consensus for the case was reached in a separate consensus meeting.

SPSS Statistics 24.0 (IBM Corp., Armonk, NY, USA) was used for statistical analyses. CAII, CAIX, and CAXII expression levels were dichotomized into two equally sized groups of low and high expression by the median value. Due to the skewed distribution and multiple testing Kruskal–Wallis with Bonferroni correction for multiple tests was used to compare expression levels between groups of histological esophagitis. We applied two-tailed Spearman's rank correlations to evaluate correlations between immunostaining intensities in basal and superficial esophageal epithelium and the degree of histological and endoscopic esophagitis. Cohen's kappa was calculated to analyze interobserver agreement, where values between 0.01–0.20 indicate none

to slight, 0.21–0.40 fair, 0.41–0.60 moderate, 0.61–0.80 substantial, and 0.81–1.00 almost perfect agreement [40].

# Results

Data on the patient demographics are summarized in Table 1. The final study cohort after re-evaluation with GS score included 82 patients; 26 cases with histologically normal esophagus, and 56 patients with reflux esophagitis (26 cases were scored mild and 30 severe). The diagnosis of lymphocytic esophagitis was made in two (2) cases, but zero cases of eosinophilic esophagitis were found. The median age of all patients was 58 years (range 20–94 years), with 52% women (n=44) and 48% men (n=40). The patients with mild reflux esophagitis were mostly women (61%), whereas 63% of the patients with severe reflux esophagitis were men. The presence of histological and endoscopic reflux esophagitis correlated significantly (p < 0.01) (Table 3). Patients with lymphocytic esophagitis were not included in the final cohort of 82 patients or further analyses.

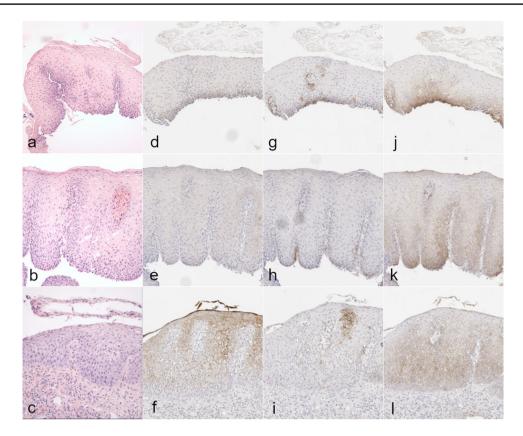
# CAII, CAIX, and CAXII Expression in Normal Esophagus and in Reflux Esophagitis

Carbonic anhydrases CAII, CAIX, and CAXII were all expressed in the normal and inflamed squamous esophageal epithelium (Fig. 2; Table 2). CAII and CAIX assessments showed substantial or almost perfect interobserver agreement between researchers with kappa value ranging from 0.62 to 1.0. For CAIX, the interobserver agreement was slightly weaker with kappa value indicating fair agreement, range 0.3–0.4.

Patient demographics	Characteristic	Histological diagnosis			Endoscopic diagnosis		
		Normal epithelium	Mild esophagitis	Severe esophagitis	Normal endoscopy	Mild esophagitis	Severe esophagi- tis
	Age						
	< 30	13/26	3/28	3/30	6/38	2/16	2/30
	30-60	13/26	14/28	13/30	19/38	5/16	14/30
	>60	10/26	11/28	14/30	13/38	9/16	14/30
	Sex						
	Female	16/26	17/28	11/30	27/38	6/16	10/30
	Male	10/26	11/28	19/30	11/38	10/16	20/30
	LA Class						
	Normal	13/26	0/28	0/30	12/38	1/16	0/30
	LA A-B	4/26	9/28	5/30	1/38	12/16	5/30
	LA C-D	0/26	0/28	21/30	0/38	1/16	20/30
	N/A	9/26	19/28	4/30	25/38	2/16	5/30

N/A, not available

Table



**Fig. 2** Examples of typical expression patterns of CAII (**d**–**f**), CAIX (**g**–**i**), and CAXII (**j**–**l**) in the normal esophageal squamous epithelium (**d**, **g**, **j**), and in squamous epithelium in histologically mild (**e**, **h**, **k**) and severe esophagitis (**f**, **i**, **l**). For comparison, closely corresponding H&E stainings are presented (**a**–**c**). In normal squamous epithelium, CAII expression is mainly cytoplasmic and to lesser extent nuclear, and present in a minority of cells (**d**). In esophagitis (**e**, **f**), CAII expression extends throughout the epithelium with an emphasis in the superficial half and the most luminal cell layers. CAIX expression in

the normal squamous epithelium (g) is mostly detectable in the cell membranes of the peripapillary cells. In mild esophagitis (h), CAIX expression remains mainly constant in the peripapillary cells and shows some increase in severe esophagitis (i). In esophagitis, groups of strongly stained cells emerge in the upper interpapillary region (i). CAXII staining is moderately strong in normal squamous epithelium, localized mainly to the plasma membrane and more evident in the basal half of the epithelium (j). In esophagitis, the expression is more intensive and extends towards the luminal surface (k, l)

Table 2Expression ofCAII, CAIX, and CAXII innormal esophageal squamousepithelium and in mild andsevere esophagitis

Protein/location	Normal epithe- lium (histoscore)	Mild esophagitis (histoscore)	р	Severe esophagi- tis (histoscore)	р
	Median (IQR)	Median (IQR)		Median (IQR)	
CA II					
Basal cytoplasm	17 (6.7–43)	38 (12–67)		43 (20–142)	0.08
Basal nuclei	4.4 (1.1–27)	22 (4.4-42)		20 (5.0-55)	0.61
Superficial cytoplasm	7.7 (0.0–23)	120 (16–236)	0.07	117 (43–223)	< 0.01
Superficial nuclei	0.0 (0.0-8.9)	21 (1.1–49)		23 (0.3-53)	0.12
CA IX					
Peripapillary intensity	1.0 (0.0-2.0)	1.5 (1.0-2.0)		2.5 (1.0-3.0)	0.14
Peripapillary cells, extent	1.0 (0.0–1.3)	1.2 (1.0-2.0)		2.2 (1.0-3.2)	0.08
Non-papillary intensity	1.0 (0.0-1.0)	1.0 (0.8–1.3)	1.0	1.0 (1.0-2.3)	< 0.01
Non-papillary histoscore	10 (0.0–10)	10.0 (5.8–13)	1.0	23 (10-95)	< 0.01
CA XII					
Basal cytoplasm	17 (8.9–20)	17 (13–30)		23 (9.6–34)	0.31
Basal membrane	180 (133–258)	217 (182–245)	0.26	267 (208–292)	< 0.05
Superficial cytoplasm	0.0 (0.0-0.0)	0.6 (0.0-4.4)		0.0 (0.0-1.1)	0.06
Superficial membrane	7.7 (0.0–27)	42 (21–55)	< 0.05	59 (15–119)	< 0.01

# CAII

In the normal squamocellular epithelium, the expression of CAII (Fig. 2d) was located in the cytoplasm and to a lesser extent in the nuclei and was generally faint in intensity. In histologically mild esophagitis, cytoplasmic CAII expression (Fig. 2e) increased in the superficial half of the epithelium, but the change was not statistically significant (p=0.07) (Table 2). Cytoplasmic expression was further increased in severe esophagitis (Fig. 2f), where the CAII immunoreaction was significantly stronger in the superficial half compared with normal esophageal epithelium (p < 0.01)(Table 2, Fig. 3).

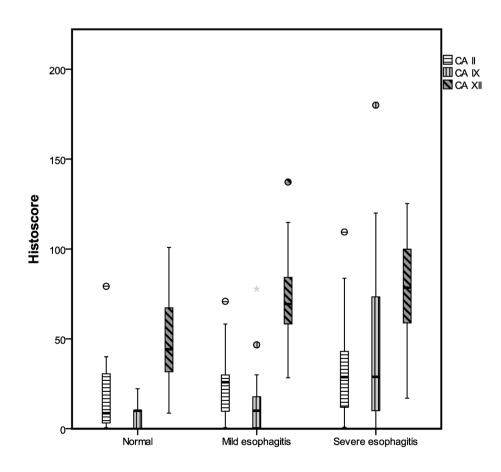
### CAIX

CAIX expression was evaluated in a specific manner as the staining pattern differed distinctly from that of the other isozymes. Expression was mainly associated with cell membranes. Weak expression was occasionally observed in basal cells, mostly in single cells. Interestingly, almost constant expression was observed in the peripapillary region of the squamous epithelium; it was detectable in 1–3 layers of cells and always started at the cells adjacent to the papillary stroma (Fig. 2g–i). The intensity of the staining in peripapillary cells and the number of stained cell layers were not

Fig. 3 Total histoscore of CAII, CAIX, and CAXII in normal esophageal epithelium, mild and severe esophagitis altered in mild esophagitis compared with healthy mucosa (Fig. 2g–h), though the extent of peripapillary staining was non-significantly more abundant in severe esophagitis (p=0.08) (Fig. 2i; Table 2). In addition, positive membranebound staining not related to papillae was observed, mostly in the middle and upper parts of the inflamed squamous epithelium. This interpapillary staining was present as groups of positive cells and increased significantly (p < 0.01) in the histoscore and intensity in severe esophagitis (Figs. 2i, 3 Table 2).

#### CAXII

CAXII membrane-bound expression was the most abundant of the isozymes. In normal squamous epithelium, the staining was most pronounced in the cell membranes of the basal third, with a gradual disappearance in the upper layers (Fig. 2j). Although far lesser than the membranous staining, the cytoplasmic expression showed a similar basal predominant pattern in normal epithelium (Fig. 2j). In esophagitis, expression of CAXII in the basal membranes increased significantly in the presence of severe inflammation (p < 0.05) (Fig. 2l). In the superficial half of the epithelium, a significant increase in the membranous immunoreaction was observed in both mild (p < 0.05) and severe (p < 0.01) esophagitis (Fig. 2k–l; 3 Table 2). The



faint cytoplasmic expression of CAXII showed no changes in expression in mild or severe esophagitis.

# Relationship Between CAII, CAIX, and CAXII Expression and Features of Reflux Esophagitis

The correlations between CAII, CAIX, and CAXII and the presence of histological and endoscopic esophagitis are summarized in Table 3. For CAII, both the total histoscore (p < 0.01) and superficial histoscore (p < 0.01) correlated significantly with histological esophagitis and with endoscopic esophagitis. The CAIX interpapillary histoscore showed a significant positive association (p < 0.01) with histological and endoscopic esophagitis. The superficial (p < 0.01), basal (p < 0.01) and total (p < 0.05) histoscores of CAXII correlated significantly with histological esophagitis but only superficial (p < 0.05) histoscore with endoscopic esophagitis. CAIX interpapillary expression and CAXII superficial expression both correlated with CAII histoscores but not with each other (Table 3).

Expression levels of CAII, CAIX, and CAXII were not related to patient sex in the total set of patients or in subgroups, nor did adjustment for sex-related differences in the severity of GERD reveal any differences (data not shown). Information regarding the use of the PPI within 2 weeks prior to endoscopy was available in 81% of the patients. The use of PPI (n/N=32/82) did not associate with the levels of CA expression or the degree of esophagitis (data not shown).

#### Discussion

Our results show that CAII, IX, and XII are all upregulated in reflux esophagitis. As the essential function of CAs is to control the intra- and extracellular pH, upregulation of expression most likely contributes to epithelial resistance to acid-related injury.

CAII expression in the superficial cytoplasm was significantly upregulated in severe GERD (p < 0.01) and correlated with the histopathological (p < 0.01) and endoscopic (p < 0.01) severity of esophagitis. CAII has potential functional significance in several components of esophageal epithelial resistance against acidic reflux [7]. CAII has been suggested to buffer intracellular acidosis in EECs [12, 13]. In our study, this concept was supported by a significant increase in CAII expression in the superficial, luminal parts of the epithelium in GERD. CAII forms transport metabolons with membrane-bound ion exchangers NHE-1, AEs, and NBCs [14-16, 18, 19]. All of these transporters are located in the basolateral cell walls of EECs [41-45], and they have been proposed to play a protective role in gastroesophageal reflux [46–49]. In vitro studies have shown that the activity of NHE-1, AEs, and NBCs is markedly enhanced in combination with CAII [15, 17, 20]. NHE-1 and NBCs have also been reported to be upregulated in both BE [45] and NHE-1 in GERD without BE [50]. Although the role of acid-loading Cl<sup>-</sup>/HCO<sub>3</sub><sup>-</sup> transporters in GERD is not as well documented, they have been shown in vitro to balance overscaled intracellular alkalization under acidic conditions, such as during GERD [42]. In summary, overexpression of

**Table 3** Correlations between CAII, CAIX, and CAXII expression and esophagitis, calculated using 2-tailed Spearman correlation. Data expressed as correlation coefficient (rho) and significance (p) in

parentheses 0.27 (p=0.027). Histological and endoscopic esophagitis is categorized into normal, mild, and severe

-			-	-				
	CAII basal	CAII superficial	CAII total	CAIX non- papillary	CAXII basal	CAXII super- ficial	CAXII total	Endoscopic esophagitis
Histological esophagitis	0.2 (0.10)	0.35 ( <b>&lt; 0.01</b> )	0.38 ( <b>&lt; 0.01</b> )	0.38 ( <b>&lt; 0.01</b> )	0.33 ( <b>&lt; 0.01</b> )	0.45 (< <b>0.01</b> )	0.30 ( <b>&lt; 0.05</b> )	0.62 ( <b>&lt; 0.01</b> )
CAII basal		0.44 ( <b>&lt; 0.01</b> )	0.56 ( <b>&lt; 0.01</b> )	0.20 (0.14)	0.13 (0.31)	0.27 ( <b>&lt; 0.05</b> )	0.17 (0.20)	0.06 (0.65)
CAII superficial			0.88 ( <b>&lt; 0.01</b> )	0.44 ( <b>&lt;0.01</b> )	0.07 (0.61)	0.34 ( <b>&lt;0.01</b> )	0.10 (0.46)	0.32 ( <b>&lt; 0.01</b> )
CAII total				0.35 ( <b>&lt;0.01</b> )	0.07 (0.61)	0.33 ( <b>&lt;0.01</b> )	0.10 (0.46)	0.31 ( <b>&lt; 0.01</b> )
CAIX non- papillary					0.11 (0.43)	0.23 (0.11)	0.16 (0.25)	0.45 ( <b>&lt; 0.01</b> )
CAXII basal						0.48 ( <b>&lt;0.01</b> )	0.84 ( <b>&lt; 0.01</b> )	0.10 (0.45)
CAXII super- ficial							0.64 ( <b>&lt; 0.01</b> )	0.28 ( <b>&lt; 0.05</b> )
CAXII total								0.14 (0.28)

CAII observed in reflux esophagitis likely has a protective function via enhanced intracellular acid neutralization.

The role of bile acids in regulating CAII responses during GERD is also of interest. In 1982 [51], bile acids were proposed to inhibit CAII, which was later confirmed [21, 52]. However, the antiulcerogenic effect of azetazolamide, a CA inhibitor, appears to not mediate CA inhibition in the stomach of rats [53]. The cumulative effect of bile and acid reflux on CAII expression and activity still requires further study.

Expression of CAIX in esophageal squamous epithelium was present in two main subpopulations of squamous cells, namely in layers 1-3 of peripapillary cells, and clusters of cells within the middle and upper third of the epithelium. CAIX was also only occasionally present in single basal cells outside the peripapillary region. Peripapillary expression and CAIX expression in the odd, positive basal cells, were largely constant and showed no change in esophagitis. Such absence of increase in expression in esophagitis does not support a role of CAIX as a basal cell marker. It is of interest that esophageal stem cells may locate in the peripapillary region [54] which, however, is still a matter of controversy. In the human intestine, CAIX expressing cells in the crypt bases [55, 56], share both the location and morphological features of stem cells [55, 57], suggesting that CAIX might be associated with stemness in at least some epithelial types. Whether the CAIX labeled peripapillary squamous cell population is functionally specific, needs additional studies.

The second subpopulation of CAIX-expressing cells comprised clusters of squamous epithelial cells mainly in the middle and upper layers of epithelium and mostly not in contact with the papillae or basal cell layer. These clusters were rare in normal epithelium and in mild esophagitis, but this expression pattern significantly increased in severe esophagitis (p < 0.01). Mechanisms for the increased expression remain speculative. Main inducers of CAIX are hypoxia via HIF-1α activation, and MAPK activation. Acid-induced MAPK response may occur in GERD [58]. In general, hypoxia worsens inflammation via HIF-1 $\alpha$  downstream effects [59], but the involvement of HIF-1 $\alpha$  in esophagitis is not clear [58, 60]. In GERD, HIF-2 $\alpha$  but not HIF-1 $\alpha$  was induced in the squamous epithelium, although not significantly [60]. One animal study, with an experimental reflux esophagitis model, has found a significant upregulation of HIF-1 $\alpha$  in reflux esophagitis [61]. In our study, the anatomical localization of the cell clusters with increased CAIX expression was in a region distant from mucosal blood vessels in either papillae or beneath the non-papillary region of basal cells. Such specific location might harbor focal epithelial hypoxia and thereby increased CAIX expression. Mechanisms, why the uppermost squamous cells above the CAIX positive clusters remained negative, could be related with the lower oxygen demand in the fully maturated cell layer [62]. Furthermore, heavy epithelial proliferation contributing to both histological severity of inflammation and oxygen demand [62] might explain the abundance of CAIX expression in severe esophagitis.

Considering the functional role of CAIX overexpression, Li et al. [24] demonstrated how CAIX knockout gastric epithelial cells fail to maintain intracellular pH via claudin-18 downregulation, which leads to tight junction failure. Claudin-18 has also been proposed to act as the dominant tight junction protein in BE and to contribute to the more evolved acid resistance in BE compared with healthy esophageal mucosa [63]. The upregulation of CAIX in reflux esophagitis and further in BE [23], in combination with the characteristic disruption of cell-to-cell junctions in GERD [64], supports the suggested role of CAIX in tight junction maintenance. CAIX is also reported to function as a component of membrane-bound transport metabolons with NBCs [29], NHE-1 [30], and AEs 1–3 [27], providing an approximately 30% increase in the function of the transporters. As described in connection with CAII, NHE-1 expression has been shown induced in GERD [50]. NHE-1 and NBCs have been reported upregulated in patients with BE [45]. Similarly, in rat EECs, AEs function in co-operation with acid extruders and counteract excessive alkalinization of pH<sub>i</sub> during acidic reflux [42]. We conclude that the increased interpapillary CAIX expression in reflux esophagitis likely contributes to esophageal defense against acid via several mechanisms.

Expression of CAXII in the squamous esophageal epithelium was most extensive and intensive in the cell membranes of the basal layer, with no change in mild reflux esophagitis, but a significant increase in severe esophagitis (p < 0.05). Expression of CAXII was significantly increased in the membranes of the superficial layer in mild (p < 0.05) and severe esophagitis (p < 0.01). CAXII has been reported to activate AE2 and to coreside at the basolateral membrane to form a transport metabolon in HeLa cells [28]. CAXII upregulation likely leads to more efficient epithelial pH control through collaboration with the ion transporter, as suggested for CAII and CAIX. The CAXII expression pattern also shows an interesting aberration compared with the immunoreactions of other CAs in esophagitis, as shown here and in BE in our previous study [23]. Strong CAXII expression in the squamous epithelium in GERD is almost completely abolished in esophageal metaplastic columnar epithelium [23], implicating selective inhibition of CAXII along with columnar metaplasia; further studies are clearly needed. Finally, as male sex is associated with GERD and evidence exists of ERα-related regulation of CAXII expression [25, 65], we analyzed the relationship of sex and CAXII expression in the esophagus. No association between sex and CAXII or the other CAs was observed.

In addition to correlating with histological and endoscopic esophagitis, the expression levels of CAII, CAIX, and CAXII in esophageal squamous epithelium intercorrelated significantly (Table 3), suggesting shared regulatory mechanisms. However, although statistically significant, several correlation coefficients were rather low, likely related with both pre-analytical issues, such as somewhat inconsistent fixation conditions of the specimens, analytical factors related with IHC, and heterogeneity of the actual disease process. Additionally, histopathological and endoscopic scores may not represent biologically the most relevant grades of severity. Considering alternative explanations, it seems that the variables representing basal half of the epithelium showed the weakest significant correlations (Table 3). We speculate that expression patterns in the basal half could represent more constitutive expression levels, while those in the superficial half may represent direct reactions to GERD and possibly shared regulation in reflux esophagitis. Our results suggest that in reflux esophagitis all three CA isozymes contribute to the response. CAII reaction is dominant in the luminal part of the epithelium, CAXII is strongly expressed in the basal half, but expression in the superficial half is a part of the response. CAXI responds as intense clusters between the elongated papillae. Thus, CAs upregulation seems to synergistically cover the full thickness of the esophageal epithelium and the cumulative contribution to the esophageal defence mechanisms is probably significant. Considering possible shared regulatory mechanisms, we found that expression of the CAs studied correlated with the thickness of basal cell zone, a feature suggesting a link with the regulation of cell proliferation. However, predominant location of the upregulation in the upper half of the squamous epithelium indicates involvement of additional regulatory networks. The identification of the regulatory feedback between human CAs will need considerable research attention in the future.

The limitations of our study include the retrospective design, with no information about pH monitoring or patient symptoms available and difficulties in obtaining reliable endoscopic grading. Regardless, in our study setting, the crucial requirement was not for every patient to meet the full diagnostic criteria designed for clinical decision making; instead, we sought to merely investigate samples that with sufficient accuracy represent epithelial injury caused by acidic reflux and samples without such injury. Diagnosis of the injury was based on a well-characterized histological multifactorial scoring system and a cutoff value able to detect most cases with acid reflux [32], as shown by pH monitoring [66]. In addition, specific types of non-reflux esophagitis were excluded based on a careful histopathological analysis. Secondly, the IHC assessment might include bias. During evaluation of IHC staining, an experienced pathologist selected in each specimen an area of squamous

epithelium in full thickness with either normal tissue, or region showing the most severe histopathological features of reflux esophagitis. Although the severity grade was not indicated to other evaluators, it is obvious that it was not possible to blind the evaluators from the features indicating the presence or severity of esophagitis. This phenomenon is methodologically characteristic to IHC studies in general. The degree of interobserver agreement in IHC assessments in our study, however, ranged from fair to almost perfect. Thirdly, we aimed to study the effect of sex by obtaining similar numbers of male and female patients, but the final distribution of esophagitis with different grades was not even; indeed, males more often had severe esophagitis. Therefore, the detection of any independent role of sex regarding the expression of the studied markers awaits additional studies.

The results of the present study have both research-related and potential clinical implications. Our study was designed as a pilot study and prospective replication studies including pH measurements are needed to confirm the findings and to examine the possible role of CAs as biomarkers of different degrees of reflux esophagitis. CAII has been proposed a function in nociception [67], and therefore, its possible correlation to patient heartburn would be of particular interest. CAII could potentially serve as a biomarker for NERD. Considering the presumed protective effects of increased CA expression, there is clearly a need for experimental evidence of those responses and perhaps studies designed to disclose the regulatory mechanisms of CA expression in EECs. Hypothetically, it could be possible to use various activators of CAs to enhance the potentially protective enzymatic effects without affecting the enzyme levels. There are several compounds that are known to activate CAs, including serotonin reuptake inhibitors (fluoxetine, sertraline, and citalopram) [68], 1-histidine and its derivatives [69], and carnosine derivatives [70]. So far there is no evidence on their role in the treatment of reflux esophagitis.

In conclusion, CAII, CAIX, and CAXII are all upregulated in reflux esophagitis. Functionally, they likely contribute to esophageal defense against acid reflux. Moreover, expression levels of CAII, IX, and XII showed positive correlations, suggesting shared regulatory factors.

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**Author's contribution** TK and MN designed the study; the data were acquired by MN, NV, and TK and analyzed by TK, MN, NV, and HH. MN, TK, SP, HH, and JS contributed to the interpretation of the data. MN wrote the manuscript which was reviewed and edited by all the other authors. All authors have approved the final version to be published and have agreed to be accountable for every aspect of the work.

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**Availability of data and materials** The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

# Declarations

**Conflict of interest** The authors declare that they have no conflicts of interest.

**Ethical approval** All procedures performed in our study were in accordance with the ethical standards of the institutional research committee and with the 1964 Declaration of Helsinki and its later amendments. For this type of study, formal consent is not required. Regional Ethical Committee Approval No: EETTMK: 81/2008. Valvira (National Licensing and Supervising Agency for Social Affairs and Health) Approval No: 10832/06.01.03.01/2014. This article does not contain any studies with animals performed by any of the authors.

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