INTRODUCTION

The process of ureteric bud branching morphogenesis is critical for establishing the location and final number of nephrons that form during kidney development. While much is known about the process of branching morphogenesis including the fact that it is guided by a Turing-type ligand-receptor based model in which the ligand, glial-derived neurotrophic factor, (GDNF) in the mesenchyme binds to and signals to the Ret tyrosine kinase receptor on the ureteric bud (Menshykau et al., 2019) we know little about the cellular events that mediate the process (Costantini, 2012). Ureteric bud cell proliferation is highest in the ureteric bud tips and lowest in the trunks and contributes to the overall growth of the ureteric bud lineage, but does not appear to directly drive the formation of ureteric bud tips (Michael & Davies, 2004). Studies done by Packard et al. demonstrated that ureteric bud tip cells delaminate from the basement membrane and undergo mitosis in the ureteric bud lumen.
such that one daughter cell reinserts in the original position, while the other reinserts randomly in a position 1–3 cells away (Packard et al., 2013). This process of mitosis-associated cell dispersal results in extensive cell rearrangements within the ureteric bud tips, but it is not clear if it results in the formation of new ureteric bud tips. Another cellular event that might be important for ureteric bud tip formation and bifurcation is cell shape change: ureteric bud tip cells tend to be more wedge-shaped with a reduced apical membrane domain when compared to ureteric bud cells within the trunk. Recently, members of the claudin family of tight junction proteins were shown to be important for cell shape changes at the neural plate midline during neural tube closure (Baumholtz et al., 2017).

Claudin mRNA transcripts are detected in a number of expression microarrays and SAGE analyses of the developing rodent kidney (Schmidt-Ott et al., 2005; Siddiqui et al., 2005; Stuart, Bush, & Nigam, 2003) even before the presence of a urinary filtrate. This intriguing observation suggests that claudins may have additional roles in the developing kidney beyond paracellular transport. In previous work, we explored the roles of claudins during kidney development and determined that a number of claudins are expressed in the developing mouse kidney using RT-PCR (Haddad et al., 2011; Khairallah et al., 2014). Here, we took advantage of a protein reagent that selectively removes a subset of claudins, Claudin-3, −4, −6, −7, −8 and −14, from tight junctions without causing any toxic effects. A truncated form of the Clostridium perfringens enterotoxin (Moriwaki, Tsukita, & Furuse, 2007) was cloned into pGEX6P1 and then induced by Isopropyl-1-thio-β-d-galactopyranoside (IPTG). GST fusion proteins were purified from E. coli BL21 strain as described previously (Veshnyakova et al., 2010) and dialyzed against PBS. Protein concentration was determined using the Bio-Rad Protein Assay (BioRad, Mississauga, Canada).

2 | METHODS

2.1 | Mouse explant culture

The Hoxb7/GFP transgene was backcrossed onto an outbred CD1 background (Srinivas, Goldberg, Watanabe, D’Agati, & al-Awqati Costantini, 1999) and Hoxb7/GFP+/− embryos were generated. Kidneys were dissected from Hoxb7/GFP+/− embryos at embryonic day 12 and grown in culture as previously reported (Gupta, Lapointe, & Yu, 2003). All mouse husbandry and breeding was performed in accordance with the regulations of the Canadian Council on Animal Care and approved by the McGill University Animal Care Committee (AUP #4120). Embryonic day 12 mouse kidneys were grown in the presence of GST (200 μg/mL) or GST-C-CPE (200 μg/mL) for 72 hr in culture.

2.2 | Production of C-CPE protein

GST alone and GST fused N-terminal to the C-terminal amino acids 185–319 of Clostridium perfringens enterotoxin (Moriwaki, Tsukita, & Furuse, 2007) was cloned into pGEX6P1 and then induced by Isopropyl-1-thio-β-d-galactopyranoside (IPTG). GST fusion proteins were purified from E. coli BL21 strain as described previously (Veshnyakova et al., 2010) and dialyzed against PBS. Protein concentration was determined using the Bio-Rad Protein Assay (BioRad, Mississauga, Canada).

2.3 | TUNEL and proliferation assays

The TUNEL assay was performed on explants cultured in the presence of C-CPE or GST using the In Situ Cell Death Detection Kit, TMR red (cat No. 12 156 792 910; Roche) using Terminal deoxynucleotidyl transferase (TdT) and by adding tetramethylrhodamine (TMR)-labeled red nucleotides. E12 explants were cultured in the presence of 200 μg/mL of C-CPE or GST for 72 hr and then fixed in formalin for 1 hr at room temperature and processed for cryosectioning. Slides were washed for 30 min in PBS and then permeabilized with 0.1% triton and 0.1% sodium citrate for 2 min on ice. Slides incubated with DNase-1 at a concentration of 3 U/mL for 20 min at room temperature were used as a positive control for the TUNEL assay. Slides were washed two times for 5 min each in PBS and incubated with the labeling reaction mix for 1 hr at 37°C. Slides incubated with the Label Solution that was lacking the terminal transferase enzyme were used as a negative control. Slides were washed for 10 min in PBS three times and mounted with a coverslip.
For the proliferation assay, explants cultured in the presence of C-CPE or GST for 72 hr were fixed in 10% formalin for 1 hr at room temperature and processed for cryosectioning. Ki67 antibody (ab15580; Abcam) was used at a 1:500 dilution. Alexa Fluor 555 goat anti-rabbit (A32732; Invitrogen) was used as the secondary antibody at a concentration of 1:500.

2.4 | Whole mount in situ hybridization

In situ hybridization was performed on E12-13 kidneys as described (Nieto, Patel, & Wilkinson, 1996). All kidneys were fixed overnight in 4% PFA in PBS at 4°C. cDNA clones for Cldn3, Cldn4, Cldn7, Cldn8, and Cldn14 were generated by RT-PCR amplification of the coding sequences using E15 mouse kidney total RNA as a template as indicated (Khairallah et al., 2014). Amplified DNA fragments were sequenced to confirm their identity and then subcloned into the pCRII-TOPO vector (Invitrogen, Camarillo, CA). To generate antisense riboprobes for in situ hybridization analysis, the mouse claudin cDNA sequences were linearized using Sma1 (Cldn4, Cldn6, Cldn8, Cldn14), Xho1 (Cldn3), or BamH1 (Cldn7) restriction enzymes. In vitro gene transcription was performed using 1 µL T3 (Cldn3, Cldn4, Cldn6, Cldn8, Cldn14) or SP6 (Cldn7) RNA polymerase. Anti-DIG antibody conjugated to alkaline phosphatase was used to detect duplexes of DIG-labeled antisense riboprobe hybridized to claudin mRNA sequences. Treated samples were developed using nitro blue tetrazolium chloride, 5-bromo-4-chloro-3-indolyl phosphatase substrate in NTMT.

2.5 | Immunofluorescent detection of claudin proteins

Mouse kidney explants were fixed using 6:3:1 (ethanol: water: 37% PFA). They were then incubated at 4°C overnight with a 1:200 dilution of rabbit anti-Cldn3 (Invitrogen, 34–700), 1:200 dilution of rabbit anti-Cldn4 (Invitrogen, 364,800), 1:25 dilution of rabbit anti-Cldn6 (Abcam, 364,800), 1:100 dilution of rabbit anti-Cldn7 (Invitrogen, 349,100), or a 1:50 dilution of rabbit anti-Cldn8 (Invitrogen, 40-0700Z). R488 and Alexa Fluor 555 goat anti-rabbit (A11034 and A21127; Invitrogen) were used as secondary antibodies at a concentration of 1/500 for 1 hr at room temperature. A Zeiss LSM780 confocal microscope was used to generate images.

2.6 | Quantitative and statistical analysis

TUNEL and proliferation assays quantification: Confocal images were analyzed by ImageJ using the following formula.

\[
\text{Normalized ratio of fluorescence} = \frac{\text{Area of red fluorescent signals within kidney explant}}{\text{Total area of kidney explant} \times \text{Total UB area}}
\]

The cellular area representing apoptotic or proliferating cells was detected by red fluorochromes, Alexa Fluor 555 or TMR, respectively, while the total area of the kidney explant was determined by outlining the DAPI positive cells. The ureteric bud area (Total UB area) was delimited using GFP signal that was expressed by the Hoxb7/GFP+/− embryo (Janke, Ward, & Vogel, 2019).

2.6.1 | Ureteric bud tip counting

Ureteric bud tips were manually counted as single, bifid and trifid ureteric tips. Lumen tip volume was quantified from confocal images by examining six stacks/kidney. The presence of any black space in the lumen was defined as an enlarged lumen. The percentage of enlarged lumens was derived by the number of enlarged UB tips/total number of UB tips X 100.

2.6.2 | Statistical analysis

One-way ANOVA and post hoc Tukey’s multiple comparisons test were used to assess changes in ureteric bud branching. Chi-square analysis was used to assess the proportion of single, bifid, and trifid ureteric bud tips. The Student’s t test was used to determine if there were statistically significant differences between two groups for the TUNEL and proliferation data. Data are presented as mean ± standard deviation for all the experiments.

3 | RESULTS

3.1 | Removal of C-CPE-sensitive claudins decreased ureteric bud branching morphogenesis

Given the large number of claudin family members expressed in the developing mouse kidney (Khairallah et al., 2014), we utilized a protein reagent that binds to multiple claudins and can simultaneously remove them from tight junctions. The full-length enterotoxin of Clostridium perfringens, (CPE) preferentially binds to Claudin-3, −4, −6, −7, −8, and −14 and exerts a cytotoxic effect to create pores within epithelial cells that result in epithelial cell death. In contrast, a modified form of the full-length toxin, containing only the C-terminal domain of the protein, C-CPE, is able to remove the same claudins from tight junctions, but without any cytotoxic effects (Fujita et al., 2000; Gao & McClane, 2012; Katahira,
Inoue, et al., 1997; Katahira, Sugiyama, et al., 1997; Kimura et al., 2010; Lohrberg et al., 2009; Sonoda et al., 1999; Veshnyakova et al., 2010; Winkler et al., 2009).

We treated mouse embryonic day (E) 12 kidney explants expressing the Hoxb7/GFP+/− transgene, which allowed us to easily monitor ureteric bud growth and branching by the green fluorescent protein expressed in this lineage (Srinivas et al., 1999). To minimize the effects of inter-embryo variability, one kidney from each embryo was cultured with a GST-purified form of the C-CPE, and the other kidney was cultured with GST protein alone for 72 hr. We observed that the formation of ureteric bud (UB) tips was inhibited in the presence of C-CPE compared to explants grown in the presence of GST alone (Figure 1a). The inhibitory effect on ureteric bud branching was first observed at 24 hr of culture and continued to be present up to 72 hr after culture (p < .05 at 24, 48 and 72 hr, Figure 1b). The decrease in ureteric bud tip counts correlated with an overall decrease in the perimeter of the ureteric bud tree in C-CPE-treated when compared to GST-treated explants (Figure 1c). The effect of treatment with C-CPE appeared to be limited to the ureteric bud lineage since the perimeter of whole kidney explants measured over time in culture was similar in GST and C-CPE-treated explants (Figure 1d).

A comparison of the complexity of ureteric bud tips revealed significantly more single UB tips (C-CPE: 223/506, 44.1% versus GST: 195/572, 34.1% \( \chi^2 = 11.26, p = .007 \)) and significantly less bifid UB tips in C-CPE-treated explants (C-CPE: 242/506, 47.8% versus GST: 327/572, 57.2%, \( \chi^2 = 9.4, p = .002 \)) compared to GST-treated explants. Similar numbers of trifid UB tips were observed in both groups (C-CPE: 41/506, 8.1% versus 50/572, GST: 8.7%, \( \chi^2 = 0.14, p = .71 \)) (Figure 1e). A total of 20 kidneys in each group were assessed for the analysis.

To determine if there was any histological evidence of necrosis or toxicity, cultured explants were paraffin-embedded, sectioned, and then stained with hematoxylin and eosin. These studies revealed no histological evidence of necrosis or toxicity, cultured explants were paraffin-embedded sectioned, and then stained with hematoxylin and eosin. These studies revealed no histological evidence of necrosis or toxicity with both the ureteric bud and the mesenchymal lineages intact. The sections suggested there was an increase in lumen volume in ureteric bud tips in C-CPE-treated when compared to GST-treated explants (Figure 1f). Taken together, these data show that C-CPE specifically inhibited the formation of ureteric bud tips and this correlated with an overall decrease in the perimeter of the ureteric bud lineage.

To determine if the decrease in ureteric bud tip number was due to either an increase in apoptosis or a decrease in proliferation within the UB lineage, TUNEL and Ki-67 assays were performed on cryosections taken from the cultured explants. These studies revealed that there was no significant difference in the amount of proliferation (mean of total red fluorescent signal expressed as pixels/total UB area expressed as pixels *100 ± standard deviation, GST: 3.07% ± 2.75 versus C-CPE: 3.35% ± 2.33, \( p = .67 \)) or apoptosis (mean of total red fluorescent signal/total UB area *100 ± standard deviation, GST: 0.84% ±0.85 versus C-CPE: 0.87% ±0.92, \( p = .87 \)) in the UB lineage in explants grown in the presence of GST or C-CPE at 72 hr (n = 3 kidneys/group and 10 sections/kidney were quantified).

### 3.2 C-CPE removed Claudins from the Ureteric Bud and Increased the Lumen of Ureteric Bud Tips

To understand which of the C-CPE-sensitive Claudins would be targeted by C-CPE, we characterized their expression patterns during mouse kidney development. By whole mount in situ hybridization, Cldn3, −4, −6, −7, and −8 mRNA transcripts were all detected in the branching ureteric bud lineage in both ureteric trunks and ureteric bud tips at embryonic day 12–13 (Figure 2). Claudin-14 was not detected by whole mount in situ hybridization (data not shown).

To determine if C-CPE-sensitive Claudins were removed from tight junctions in the presence of C-CPE as predicted, immunofluorescent studies were performed on cultured explants. Control mouse explants grown in the presence of GST alone, showed strong expression of Cldn3, −4, −6, and −8 proteins in the apical membrane of the ureteric bud trunks and tips (Figure 3) that colocalized with zona occludens (ZO)-1, another tight junction protein. In contrast, CLDN-7 showed high expression in the basolateral membrane and weak signal in the apical membrane with poor colocalization with ZO-1. CLDN-14 was not expressed in the ureteric bud lineage by immunofluorescence. In the presence of C-CPE, there was much less expression of CLDN-3 and −4 and an even greater loss of CLDN-6 and CLDN-8 in the apical membrane of the ureteric bud lineage. Interestingly, in the presence of C-CPE, CLDN-7 appeared to show relatively more apical compared to basolateral expression, but again there was poor colocalization with ZO-1, suggesting CLDN-7 was not within the tight junction. The immunofluorescent images also suggested that the ureteric bud lumens of C-CPE-treated explants were enlarged compared to GST-treated explants. This was quantified from confocal images and there were significantly more ureteric bud tips with enlarged lumens in C-CPE-treated compared to GST-treated explants (C-CPE: 35/40, 87.5% versus GST: 10/97, 10.3%, \( \chi^2 = 66.77, p = .0001 \), n = 5 kidneys/group). The enlarged lumens were predominantly seen in ureteric bud tips, but some ureteric trunks also showed this phenotype.

### 4 Discussion

In this study, we demonstrate that Claudins are expressed in the ureteric bud lineage during the period of most rapid
branching morphogenesis in vivo (Short et al., 2014). Five claudins that are known to bind to the truncated form of the Clostridium perfringens enterotoxin (C-CPE), Claudin-3, −4, −6, −7, and −8 were expressed in ureteric bud tips and trunks. Embryonic day 12 kidneys that were cultured for 72 hr with C-CPE exhibited impaired branching morphogenesis with a decrease in the formation of ureteric bud tips. The ureteric bud tips that formed in the presence of C-CPE had significantly more single ureteric bud tips and fewer bifid ureteric bud tips in C-CPE-treated kidney explants compared to GST, suggesting a decrease in complexity of ureteric bud branching morphogenesis. Scale bar = 100 μm. The asterisk labels an example of a bifid tip, while the arrow shows an example of a single ureteric bud tip in the C-CPE kidney explant. Representative sections from kidney explants grown for 72 hr that were stained with hematoxylin and eosin are shown. The C-CPE-treated explants exhibited ureteric bud tips with enlarged lumens compared to the GST-treated explants. Red boxes outline the ureteric bud tips shown in the inset images on the right. The inset images show an enlarged ureteric bud tip in the C-CPE-treated explant compared to GST. Scale bar = 100 μm. Inset scale bar = 25 μm Statistical analysis one-way ANOVA and Post hoc Tukey’s multiple comparison test for 1b-d. Chi-square analysis for 1e. * p < .05, ** p < .01, ***p < .001

FIGURE 1  C-CPE Inhibits Branching Morphogenesis in Mouse Embryonic Kidney Explants. (a) For each embryo at E12, one Hoxb7/GFP+/− kidney explant was grown in GST, and the other was grown in C-CPE for 72 hr. Scale bar = 200 μm (b–d) Three graphs are shown from left to right that quantify the total number of ureteric bud tips at 24, 48 and 72 hr (red = GST-treated and blue = C-CPE-treated explants) (b) the change in the number of ureteric bud tips, n = 17 kidneys in each group; (c) the relative change in ureteric bud perimeter (P) (Pfinal(f) − Pinitial(i)/Pinitial(i)) at each time interval, n = 19 kidneys in each group (d), the relative change in the total kidney explant perimeter (Pf − Pi/Pi) at each time interval n = 19 for each group. At 48 and 72 hr, there was a significant decrease in the number of ureteric bud tips observed in C-CPE-treated compared to GST-treated explants. Similarly, at 48 and 72 hr, there was a decrease in the relative change in ureteric bud perimeter in C-CPE-treated compared to GST-treated explants. In contrast, the relative change in perimeter of the total kidney explant was similar in GST and C-CPE-treated explant suggesting C-CPE was specifically affecting the ureteric bud compartment. (e) The total number of single, bifid, and trifid ureteric bud tips was enumerated in GST- and C-CPE-treated kidneys as shown (n = 20/group) after 72 hr of culture. There were significantly more single ureteric bud tips and fewer bifid ureteric bud tips in C-CPE-treated kidney explants compared to GST, suggesting a decrease in complexity of ureteric bud branching morphogenesis. Scale bar = 100 μm. The asterisk labels an example of a bifid tip, while the arrow shows an example of a single ureteric bud tip in the C-CPE kidney explant. (f) Representative sections from kidney explants grown for 72 hr that were stained with hematoxylin and eosin are shown. The C-CPE-treated explants exhibited ureteric bud tips with enlarged lumens compared to the GST-treated explants. Red boxes outline the ureteric bud tips shown in the inset images on the right. The inset images show an enlarged ureteric bud tip in the C-CPE-treated explant compared to GST. Scale bar = 100 μm. Inset scale bar = 25 μm Statistical analysis one-way ANOVA and Post hoc Tukey’s multiple comparison test for 1b-d. Chi-square analysis for 1e. * p < .05, ** p < .01, ***p < .001

branching morphogenesis in vivo (Short et al., 2014). Five claudins that are known to bind to the truncated form of the Clostridium perfringens enterotoxin (C-CPE), Claudin-3, −4, −6, −7, and −8 were expressed in ureteric bud tips and trunks. Embryonic day 12 kidneys that were cultured for 72 hr with C-CPE exhibited impaired branching morphogenesis with a decrease in the formation of ureteric bud tips. The ureteric bud tips that formed in the presence of C-CPE had
larger lumens and were more likely to exist as single rather than bifid tips, suggesting an overall decrease in the complexity of branching. The decrease in ureteric bud tips from treatment with C-CPE correlated with a decrease in expression and co-localization with ZO-1 for all of the C-CPE-sensitive claudins with the exception of CLDN-7, so that they were absent or weakly expressed in the apical membrane. Taken together, claudins are required for ureteric bud branching morphogenesis and regulate the formation of ureteric bud tip lumens.

We removed multiple claudins using the C-CPE reagent because removal of single C-CPE-sensitive claudins has not generated severe renal developmental phenotypes, suggesting there may be functional redundancy amongst members and/or different requirements depending on the model organism. For example, removal of Claudin-3, −4, −6, or −8 has not led to any defect in mouse kidney development based on the fact that the offspring survive and do not succumb to renal failure postnatally (Anderson et al., 2008; Castro Dias et al., 2019; Fujita, Hamazaki, Noda, Oshima, & Minato, 2012; Gong et al., 2015). In the absence of comprehensive counts of nephron number, however, we cannot rule out the possibility that the mice may have mild to moderate defects in nephron number from a deficiency in renal branching morphogenesis. Indeed, we performed nephron number counts on Claudin-7 knockout mice, which die at 2 weeks of age, but we did not observe any difference between mutants and wild-type littermates (Khairallah et al., 2014). In Xenopus, however, knockdown of Claudin-6 did result in decreased apical-basal polarity and cell adhesion within the pronephric tubule (Sun, Wang, Li, & Mao, 2015).

A number of studies have demonstrated that claudins are expressed within the ureteric bud and its derivatives during kidney development at a time when there is no urine filtrate formed by glomerular filtration (Wang et al., 2011). This begs the question, what is their function at this timepoint when there is no significant paracellular transport? In our studies, the decrease in branching morphogenesis following claudin removal was not due to a change in cell proliferation or in apoptosis within the ureteric bud lineage. We did observe larger lumens in the ureteric bud tips of C-CPE-treated kidneys and less complexity of branching morphogenesis compared to GST controls. We speculate that claudins within the ureteric bud lineage maintain tensile forces at the apical membrane through their interactions with the actin cytoskeleton and that these tensile forces are maximally taut where new ureteric buds will arise (Figure 4). Analogous to the purse-string hypothesis, high tensile forces at the apical membrane would alter the shape of individual cells to a wedge as opposed to a columnar shape, and this would confer the emergence of a bud from the ureteric bud epithelium. After binding to C-CPE, C-CPE-sensitive claudins (CLDN3, −4, −6, and −8) are removed from tight junctions and this is predicted to decrease tension along the apical membrane. This would cause a decrease in the number of wedge-shaped cells at the ureteric bud tips that would

**FIGURE 2** C-CPE-Sensitive Claudins Are Detected in Mouse Embryonic Kidneys by Whole Mount In Situ Hybridization. Whole mount in situ hybridization was performed on mouse kidney explants between E12-13 and revealed the presence of Cldn3, −4, −6, −7, and −8 transcripts within the ureteric bud tips and trunks. Cld14 was not detected by whole mount in situ hybridization and is not shown. Whole mount in situ hybridization to c-Ret is shown as a positive control and its pattern reveals that c-Ret is primarily in the ureteric bud tips with much less signal seen in the trunks in contrast to the C-CPE-sensitive claudins. Representative images are shown from a total of three experiments with n = 3 kidneys exposed to each riboprobe.
favor larger ureteric bud tip lumens and less bud formation. Tight junctions are highly dynamic structures based on in vitro cell culture models (Shen, Weber, & Turner, 2008).

In fibroblasts, claudins associate intermittently with ZO-1 and actin and this promotes the formation of tight junction strands between cells. However, claudins can also anneal...
Indeed, it is hypothesized that the ability of tight junction strands to form independently of ZO-1 and actin permits the epithelium to separate its permeability properties from its interactions with the cytoskeleton via tight junction complexes. In summary, we believe that claudins may function at this stage of branching morphogenesis to regulate cell shape through their effects on the tensile forces along the apical membrane.

While we know much about the signaling factors and transcription factors that drive branching morphogenesis, we still have an incomplete understanding of the cellular events that drive this process. Cell proliferation and mitosis-associated dispersal occurs within the ureteric bud lineage during branching, but these do not seem to be the major processes that drive the formation of new ureteric bud tips. Our studies in which we have removed specific claudin members from the tight junctions clearly affects the number and complexity of ureteric bud tips that form, but branching still occurs, albeit less efficiently. To address the role of claudins during renal branching morphogenesis, in vivo models will need to be established in which multiple C-CPE-sensitive claudins are removed. The development of such a model will permit complete removal of claudins as opposed to partial removal as observed in our explant experiments.

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